

### UNIVERSITAT DE BARCELONA

### Polycystic ovary syndrome in adolescent girls: treatment oriented to pathophysiology and new markers of efficacy

Cristina García Beltran

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## POLYCYSTIC OVARY SYNDROME IN ADOLESCENT GIRLS: TREATMENT ORIENTED TO PATHOPHYSIOLOGY AND NEW MARKERS OF EFFICACY

Doctoral thesis dissertation presented by

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## **TABLE OF CONTENTS**

ABBREVIATIONS AND ACRONYMS

- AdipoR: adiponectin receptor
- AE-PCOS: Androgen Excess and PCOS Society
- AGE: advanced glycated end-products
- ALT: alanine aminotransferase
- ASV: amplicon sequence variant
- AMH: anti-Müllerian hormone
- AMPK: AMP-activated protein kinase
- ASRM: American Society for Reproductive Medicine
- AST: aspartate aminotransferase
- AYA: adolescent and young adult
- BAT: brown adipose tissue
- BMI: body mass index
- Brite: brown-in-white
- cIMT: carotid intima-media thickness
- CRP: C-reactive protein
- CXCL14: C-X-C motif chemokine ligand-14
- CYP11A1: cytochrome P450 family 11 subfamily A member 1
- CYP17A1: cytochrome P450 family 17 subfamily A member 1
- CYP19A1: cytochrome P450 family 19 subfamily A member 1
- DBI: diazepam-binding protein-1
- DENN1A: DENN domain-containing protein 1A
- DHEAS: dehydroepiandrosterone sulfate
- DPP4: dipeptidyl peptidase-4
- DXA: dual x-ray absorptiometry
- EDC: endocrine-disrupting chemical
- EMA: European Medicines Agency
- ESHRE: European Society of Human Reproduction and Embryology
- EWAS: epigenome-wide association studies

FAI: free androgen index

FDA: Food and Drug Administration

FFAR: free fatty acid receptor

FGF21: fibroblast growth factor-21

FSH: follicle-stimulating hormone

GDF15: growth-and-differentiation factor-15

GGT: gamma-glutamyl transferase

GLP1: glucagon-like peptide 1

GLUT4: glucose transporter 4

GnRH: gonadotropin-releasing hormone

GSK3: glycogen synthase kinase 3

GWAS: genome-wide association studies

HMW: high-molecular-weight

HPT: hypothalamic-pituitary-axis

HRQoL: health-related quality of life

IL-1  $\beta$ : interleukin-1 $\beta$ 

IL-18: interleukin-18

IL-6: interleukin-6

IR: insulin resistance

KNDy: kisspentin-neurokinin B-dynorphin A

LH: luteinizing hormone

MAPK: mitogen-activated protein kinase

MASLD: metabolic-dysfunction associated steatotic liver disease

MET: metformin

METRNL: meteorin-like protein

miRNA: microRNA

MRI: magnetic resonance imaging

mTOR: mammalian target of rapamycin

- NAFLD: non-alcoholic fatty liver disease
- NIH: National Institutes of Health
- NK3: neurokinin 3
- OC: oral contraceptive
- PCOM: polycystic ovarian morphology
- PCOS: polycystic ovary syndrome
- PI3K-PKB/Akt: phosphoinositide-3-kinase protein kinase B/ Akt
- PIO: pioglitazone
- PPAR: peroxisome proliferator-activated receptor
- ROS: reactive oxygen species
- SCFA: short-chain fatty acids
- SGBS: Simpson Golabi Behmel Syndrome
- SGLT-2: sodium-glucose co-transporter-2
- SHBG: sex hormone binding-globulin
- SOCS-3: suppressor cytokine signaling-3
- SPI: spironolactone
- SPIOMET: spironolactone-pioglitazone-metformin
- T2D: type 2 diabetes
- TH: thyroid hormone
- THRSP: thyroid hormone responsive spot 14
- TNF-α: tumor necrosis factor-alpha
- TRH: thyrotropin-releasing hormone
- TSH: thyroid-stimulating hormone
- TZD: thiazolidinedione
- UCP-1: uncoupling protein-1
- WAT: white adipose tissue

# LIST OF ARTICLES IN THE THESIS

Thesis in compendium of publications format. The thesis consists of five articles, the objectives of which are the same as those of this doctoral thesis, and are detailed in the "Objectives" section.

### **First article**

Ibáñez L, Díaz M, **Garcia-Beltran C**, Malpique R, Garde E, López-Bermejo A, de Zegher F. Toward a treatment normalizing ovulation rate in adolescent girls with polycystic ovary syndrome. Journal of the Endocrine Society. 2020; 4(5), bvaa032. <u>https://doi.org/10.1210/jendso/bvaa032</u>.

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### Second article

**Garcia-Beltran C**, Cereijo R, Quesada-López T, Malpique R, López-Bermejo A, de Zegher F, Ibáñez L, Villarroya F. Reduced circulating levels of chemokine CXCL14 in adolescent girls with polycystic ovary syndrome: normalization after insulin sensitization. BMJ Open Diabetes Research & Care. 2020; 8(1), e001035. https://doi.org/10.1136/bmjdrc-2019-001035.

IF (2020) = 3.388, Q3 (Endocrinology & Metabolism) JCR2020.

### Third article

**Garcia-Beltran C**, Malpique R, Carbonetto B, González-Torres P, Henares D, Brotons P, Muñoz-Almargo C, López-Bermejo A, de Zegher F, Ibáñez L. Gut microbiota in adolescent girls with polycystic ovary syndrome: effects of randomized treatments. Pediatric Obesity. 2021; 16(4), e12734. <u>https://doi.org/10.1111/ijpo.12734</u>.

IF (2021) = 3.910, Q1 (Pediatrics) JCR2021.

### Fourth article

**Garcia-Beltran C**, Bassols J, Carreras-Badosa G, López-Bermejo A, Ibáñez L, de Zegher F. Raised thyroid-stimulating hormone in girls with polycystic ovary syndrome: effects of randomized interventions. Hormone Research in Paediatrics. 2023; 96(5), 458–464. <u>https://doi.org/10.1159/000529183</u>.

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### **Fifth article**

**Garcia-Beltran C**, Peyrou M, Navarro-Gascon A, López-Bermejo A, de Zegher F, Villarroya V, Ibáñez L. Organokines and liver enzymes in adolescent girls with polycystic ovary syndrome during randomized treatments. Frontiers in Endocrinology. 2024; 15:1325230. <u>https://doi.org/10.3389/fendo.2024.1325230</u>.

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## **THESIS SUMMARY IN CATALAN**

### SÍNDROME DE L'OVARI POLIQUÍSTIC EN ADOLESCENTS:

### TERAPÈUTICA ORIENTADA A LA FISIOPATOLOGIA I NOUS MARCADORS D'EFICÀCIA

### INTRODUCCIÓ

La síndrome de l'ovari poliquístic (SOP) podria iniciar-se amb un desajustament entre el creixement prenatal (reduït) i el postnatal (augmentat), provocant un excés de greix hepato-visceral. La disfunció del teixit adipós, l'alteració de la microbiota intestinal, els trastorns de la tiroides i nivells circulants alterats de molècules de senyalització (organocines), poden contribuir a la patogènesi de la SOP.

Actualment no hi ha cap fàrmac aprovat per al tractament d'aquesta síndrome, tot i que habitualment es prescriu un anticonceptiu oral (ACO). Darrerament s'ha proposat un nou tractament basat en la combinació a dosis baixes d'espironolactona, pioglitazona i metformina (SPIOMET). Malgrat que l'evidència és encara limitada, un estudi pilot realitzat en una població d'adolescents amb SOP i sense obesitat (N=34) ha descrit que el tractament amb SPIOMET proporciona efectes més normalitzadors que el tractament amb ACOs.

### HIPÒTESI

En una població d'estudi més àmplia d'adolescents amb SOP, es confirmarà que el tractament amb SPIOMET té efectes més beneficiosos que el tractament amb ACOs.

### OBJECTIUS

 Comparar els efectes del tractament amb SPIOMET o ACOs en una cohort més àmplia d'adolescents amb SOP i sense obesitat

- Caracteritzar els nivells circulants CXCL14 [C-X-C motif chemokine ligand 14, una proteïna secretada pel teixit adipós marró (TAM)] en adolescents amb SOP i avaluar si

es produeixen canvis divergents després d'un any de tractament amb SPIOMET o ACOs. Determinar els efectes dels fàrmacs que composen SPIOMET sobre l'expressió i alliberació de CXCL14 en un model cel·lular d'adipòcits humans.

- Estudiar la composició de la microbiota intestinal en mostres de femta d'adolescents amb SOP i determinar els efectes del tractament durant un any amb SPIOMET o ACOs.

 Determinar si els nivells de tirotropina (TSH) estan elevats en adolescents amb SOP i estudiar si aquests nivells es redueixen després d'un any de tractament amb SPIOMET o ACOs i si es mantenen baixos 1 any després de la suspensió del tractament.

- Avaluar els efectes de 6 mesos de tractament amb SPIOMET o ACOs sobre un conjunt d'organocines [fibroblast growth factor 21 (FGF21), diazepam-binding protein 1 (DBI) i meteorin-like (METRNL)] en adolescents amb SOP i identificar associacions amb biomarcadors circulants de dany hepàtic [aspartat aminotransferasa (AST), alanina aminotransferasa (ALT) i gamma-glutamil transferasa (GGT)] avaluats com a paràmetres de seguretat en els estudis pilot.

### MÈTODES

N=62 adolescents amb SOP i sense obesitat, incloses en dos estudis pilot aleatoritzats per rebre tractament amb SPIOMET (N=31) o ACO (N=31) durant un any i reevaluades després d'un any sense tractament.

Determinació de variables auxològiques, endocrino-metabòliques [insulina, andrògens, adiponectina d'alt pes molecular], de seguretat (TSH, AST, ALT, GGT), de composició corporal (DXA), greix hepato-visceral (ressonància magnètica) durant i post tractament, i la funció ovulatòria (progesterona salival setmanal) durant el segon i el quart trimestre de l'any post-tractament. N=52 adolescents sanes de la mateixa edat s'utilitzen com a controls.

En una selecció de mostres de sèrum de les pacients participants, determinació dels nivells de CXCL14, FGF21, DBI i METRNL mitjançant kits d'ELISA específics.

Pels estudis *in vitro*, utilització d'un model cel·lular humà d'adipogènesis de TAM (cèl·lules SGBS). Quantificació de l'expressió del gen CXCL14 i d'altres gens d'adipogènesis mitjançant PCR a temps real. Determinació dels nivells de CXCL14 en el medi de cultiu mitjançant ELISA.

Per l'estudi de la composició de la microbiota intestinal, seqüenciació de l'amplicó del gen de la subunitat ribosòmica 16S en una selecció de pacients de les que es disposa mostra de femta.

### RESULTATS

Els dos tractaments tenen un efecte similar sobre els andrògens. No obstant, el tractament amb SPIOMET normalitza el greix hepato-visceral i els nivells circulants d'insulina. La funció ovulatòria (segons concentracions de progesterona salival) es normalitza només després del tractament amb SPIOMET.

Les adolescents amb SOP presenten nivells sèrics més baixos de CXCL14 que les controls. Només el tractament amb SPIOMET normalitza les concentracions de CXCL14. La pioglitazona indueix l'expressió de CXCL14 en cultius de pre-adipòcits, en paral·lel amb la inducció de marcadors d'adipogènesis de teixit adipós marró, mentre que l'espironolactona indueix l'expressió i alliberació de CXCL14 en cultius d'adipòcits diferenciats.

Les adolescents amb SOP tenen reduïda la diversitat alfa, alterat el patró de la microbiota i el perfil taxonòmic, amb més abundància de *Family XI* i menys abundància de la família *Prevotellaceae* i dels gèneres *Prevotella* i *Senegalimassilia* en comparació amb les controls. El tractament amb SPIOMET, però no amb ACOs, redueix l'abundància de *Family XI*. L'abundància de *Prevotellaceae*, *Prevotella* i *Senegalimassilia* continua reduïda després del tractament amb SPIOMET i ACOs.

Els nivells de TSH són més elevats en les adolescents amb SOP que en les control. Les concentracions de TSH durant el tractament es mantenen elevades en les pacients tractades amb ACOs i descendeixen ràpidament en les pacients que reben SPIOMET,

assolint valors dins del rang de seguretat. Els nivells de tirotropina després del tractament es mantenen estables en els dos subgrups.

En relació amb les organocines després de 6 mesos de tractament (1) els nivells de FGF21 són significativament més elevats en les adolescents amb SOP que en les controls; (2) els nivells de DBI són més baixos en les pacients tractades amb ACOs comparades amb les controls i amb aquelles pacients tractades amb SPIOMET; (3) no s'observen diferències en les concentracions de METRNL entre les adolescents amb SOP i les controls. Els nivells sèrics d'ALT i GGT correlacionen directament amb els nivells de METRNL només en aquelles pacients tractades amb ACOs.

### CONCLUSIONS

El tractament amb SPIOMET determina un millor estat de salut metabòlica, en comparació amb el tractament amb ACOs, així com una taxa d'ovulació post tractament més normalitzada. Els efectes normalitzadors de la intervenció amb SPIOMET s'amplien per incloure l'abundància de *Famíly XI* a la microbiota intestinal i els nivells circulants de CXCL14 i TSH. L'associació entre les concentracions circulants d'ALT i GGT i els nivells circulants de METRNL, observada només en les pacients tractades amb ACOs suggereix una major síntesi de METRNL reactiva als canvis en els marcadors de danys hepàtic associats al tractament amb ACOs.

# INTRODUCTION

#### **1. POLYCYSTIC OVARY SYNDROME**

Polycystic ovary syndrome (PCOS) is the most common endocrine disorder in women of reproductive age. The prevalence of PCOS is around 5-10% but could be as high as 26% in some populations, and it is estimated that up to 70% of cases are undiagnosed (**1**,**2**). The incidence of PCOS in adolescent girls is around 6-18% (**3**,**4**). This wide range is explained by the different diagnostic criteria used and by the heterogeneity of the studied populations (**5-8**).

The most common features of PCOS include clinical and/ or biochemical androgen excess and menstrual irregularities, as proxy of oligo-anovulation (**6**,**9**). Women with PCOS also have an elevated prevalence of obesity, and a significant proportion also display insulin resistance (IR), even in the absence of obesity (**10**).

PCOS is the most frequent cause of hirsutism and anovulatory subfertility in adolescent girls and young women, and is associated with lifelong co-morbidities such as type 2 diabetes (T2D), premature vascular ageing, premenopausal cancer, metabolic-dysfunction associated steatotic liver disease (MASLD, the new term for non-alcoholic fatty liver disease or NAFLD), gestational complications, higher risk of suicide attempts, and anxiety/depression, with a negative impact on the health-related quality of life (HRQoL) of these subjects and their offspring (**11-18**). Overweight and obesity amplify the metabolic-reproductive abnormalities and the risks for T2D and cardiovascular complications (**10**).

### 2. PATHOPHYSIOLOGY OF PCOS

PCOS is a multifactorial condition; genetic and epigenetic changes, altered ovarian folliculogenesis, neuroendocrine, metabolic and environmental abnormalities, adipose tissue dysfunction and gut microbiota dysbiosis may each contribute to the pathogenesis of this disorder (**Figure 1**). PCOS represents an example of systems biology with multiple interconnected signaling networks, all of which may not be equally operative in every patient (**6**).



Figure 1. Potential factors involved in the pathophysiology of polycystic ovary syndrome (PCOS) (Own elaboration, created with Biorender.com).

### 2.1 Developmental sequence leading to PCOS

Nowadays, PCOS in adolescent girls is thought to be in essence, the result of a mismatch between prenatal adipogenesis and postnatal lipogenesis, which can be inferred from a mismatch between (less) prenatal weight gain and (more) postnatal weight gain (**19-21**). The number of subcutaneous adipocytes, wherein fat can be
stored without complications, is essentially set in early life (age <2 years) (**22,23**). Accordingly, when there is a balance between pre- and postnatal weight gain, the endocrine-metabolic system tends to also remain in balance. However, when there is an upward mismatch between pre- and postnatal weight gain, more fat has to be stored than is safely feasible in the subcutaneous fraction of adipose tissue, and this excess of fat ends up being stored in ectopic depots, notably in the liver and viscera (hepato-visceral or central fat). Indeed, PCOS in adolescent girls is known to be usually preceded by a marked Z-score increment between weight at birth and body mass index (BMI) at PCOS diagnosis (**24**).

The above-mentioned rationale is based on the "adipose tissue expandability" hypothesis (25). This hypothesis suggests that subcutaneous adipose tissue has a limited capacity to increase its mass safely and this capacity varies among individuals and is co-determined by environmental and (epi)genetic factors. According to that, when subcutaneous adipocytes start to be overfilled, then lipids are stored in non-adipose tissue organs, such as muscle, liver or pancreas (Figure 2). This ectopic deposition of lipids is believed to cause IR via lipid-induced toxicity, or lipotoxicity (26).



**Figure 2. Overview of the "adipose tissue expandability" hypothesis.** The ability of the subcutaneous adipose tissue to expand allows the safe storage of the excess of energy. However, when this tissue becomes saturated, lipids are stored in ectopic depots such as muscle, liver or pancreas. Adapted from (27).

As shown in **Figure 3**, the developmental sequence of events leading to PCOS starts with the aforementioned mismatch between prenatal and postnatal weight gain that results in an excess of hepato-visceral fat. As a result, girls tend to develop IR and a trend towards higher blood pressure by early childhood (**28-31**). Actually, a higher Z-score change from weight-at-birth to BMI-in-childhood has been associated with more IR and increased central adiposity (**32**).



**Figure 3.** The developmental sequence of events leading to polycystic ovary syndrome. Abbreviations: DHEAS, dehydroepiandrosterone sulfate; LH, luteinizing hormone; TSH, thyroidstimulating hormone; PCOS, polycystic ovary syndrome. Adapted from (**20**).

During late childhood, as an adaptive/ homeostatic response, mismatch girls accelerate their body growth and maturation to counteract ectopic adiposity. A first step of this response includes a decrease in circulating sex hormone binding-globulin (SHBG) and high-molecular-weight (HMW) adiponectin concentrations, and an exaggerated adrenarche, hallmarked by an increase of dehydroepiandrosterone sulfate (DHEAS) levels, the early appearance of pubic hair (precocious pubarche) and a faster rate of bone age maturation (**33-40**). If the ectopic lipid accumulation continues,

then girls may develop another acceleration of growth and maturation that includes an upregulation of the neuroendocrine axis [luteinizing hormone (LH) hypersecretion and ovarian estrogen secretion] (**41,42**) and thyroid axis [higher thyroid-stimulating hormone (TSH) and free T3] (**43**), more testosterone production (**44**) and an accelerated bone maturation (**45**), potentially resulting in an early menarche (**46**).

In the first years after menarche, when height gain is no longer possible, the homeostatic effect of the accelerated body growth on ectopic fat accumulation is essentially lost (**47**). However, the acceleration-mediating mechanisms (specifically IR, low circulating SHBG and HMW-adiponectin levels, and increased DHEAS, LH TSH and testosterone concentrations) persist, leading to androgen excess and oligo-anovulation (**48,49**).

### 2.2 Ovarian folliculogenesis and the neuroendocrine axis

Ovarian follicle development and the subsequent ovulation process are coordinated by a highly complex interplay between endocrine, paracrine and autocrine signals. Follicle development is regulated mainly by the hypothalamic-pituitary-gonadal axis (**50**). The sequence of events leading to PCOS may be subject in part to alterations in both ovarian folliculogenesis and the hypothalamic-pituitary-gonadal axis.

In a physiological state, the hypothalamus secretes gonadotropin-releasing hormone (GnRH) in a pulsatile fashion, which triggers follicle-stimulating hormone (FSH) and LH release from the anterior pituitary. Importantly, low-frequency GnRH pulses are responsible for FSH secretion, whereas high-frequency pulses are responsible for LH secretion. These, in turn, act on the ovary: FSH stimulates follicle maturation whereas LH is responsible for inducing ovulation (**51**). In addition, LH is the primary stimulus for ovarian androgen production (**52**). Adolescent girls and women with PCOS have downstream disruptions in the regulation of GnRH pulse generator, which increases LH over FSH production (**53-55**). This hormonal imbalance can lead to an excessive ovarian androgen production (due to increased LH levels) and impaired follicular development (caused by insufficient FSH levels), resulting in oligo- or

anovulatory cycles. In turn, hyperadrogenemia may also trigger faster GnRH pulse frequency (**56**). Compelling evidence suggests that elevated androgens disrupt the capacity of ovarian steroids (estrogens and progesterone) to regulate pulsatile GnRH/LH secretion, contributing to the LH hypersecretion characteristic of PCOS (**6**). It has been proposed that oligo-anovulation may result from an adaptive neuroendocrine response to the excess of central fat (**57**).

The aforementioned imbalance between LH, FSH and androgens triggers follicular arrest. Insufficient FSH levels disrupt the selection of a dominant follicle, which leads to accumulation of small pre-antral follicles (**58**). In addition, increased levels of Anti-Müllerian hormone (AMH), which is normally secreted by granulosa cells of pre-antral and early-antral follicles, have been detected in PCOS women (**59**). Increased AMH levels are the consequence of an increased number of small antral follicles (**59**). In turn, AMH blocks FSH action, contributing to hyperandrogenism and inhibiting the recruitment of further primordial follicles (**54**).

## 2.3 Insulin resistance and hyperinsulinemia

IR and compensatory hyperinsulinemia are considered major drives of PCOS imbalances and are involved in the development of hyperandrogenism and reproductive dysfunction through various mechanisms (**60**).

Insulin is a peptide hormone that plays a critical role in the regulation of human metabolism. Under normal circumstances, insulin stimulates nutrient transport into cells, acutely regulates metabolic enzyme activity, controls transcription of metabolic genes, regulates cellular growth and differentiation, and controls its own clearance, all through activation of its receptor (**61**). After being released by pancreatic beta cells, insulin binds to the insulin receptor on the target cell surface. Once activated, the receptor can phosphorylate a number of intracellular substrates that initiate two primary signaling pathways: the phosphoinositide-3-kinase - protein kinase B/ Akt (PI3K-PKB/Akt) pathway and the mitogen-activated protein kinase (MAPK) pathway. Through the PI3K-PKB/Akt pathway, insulin stimulates glucose uptake by promoting

the translocation of glucose transporter 4 (GLUT4) from intracellular vesicles to the cell surface and leads to the inactivation of glycogen synthase kinase 3 (GSK3) increasing glycogen protein syntesis. Akt also increases protein synthesis and blocks autophagy through mammalian target of rapamycin (mTOR). The MAPK pathway ultimately promotes cell division, protein synthesis and cell growth (**60**).

IR and hyperinsulinemia are present in the vast majority of women with PCOS, including in those with overweight and obesity as well as in normal-weight women (**62-64**). Thus, the presence of IR is independent of the degree of adiposity, and also of androgen levels (**65**). IR in PCOS is the consequence of impaired insulin action in target tissues, which results in compensatory hyperinsulinemia and reduced insulin response to a glucose overload (**60**). In addition, IR is tissue-selective; for example, resistance to the metabolic actions of insulin has been reported in skeletal muscle, liver and adipose tissue, whereas steroid-producing tissues (adrenal gland and ovary) remain insulin-sensitive (**66**).

Hyperinsulinemia plays an important role on androgen synthesis. [1] Insulin stimulates GnRH gene transcription through the MAPK pathway and consequently increases LH pulse secretion and ovarian androgen release (**67,68**); [2] insulin can decrease hepatic SHBG release leading to augmented levels of freely circulating androgens (**69**); [3] insulin may also directly stimulate the activity of ovarian cytochrome P450 family 17 subfamily A member 1 (CYP17A1) promoting ovarian androgen steroidogenesis (**70**).

There are other mechanisms identified as contributors of IR in PCOS. For instance, defects downstream of insulin receptor phosphorylation or alterations on GLUT4 translocation through PI3K-PKB/Akt have been detected in PCOS women (**71,72**). In addition, it has been proposed that the accumulation of reactive lipids (such as ceramides) observed in PCOS patients (**73,74**), could impair insulin signaling by blocking insulin stimulation of Akt (**75,76**).

#### 2.4 Hyperandrogenism

The androgen excess observed in PCOS patients is mainly due to enhanced androgen synthesis by follicular ovarian theca cells, which show an increased expression of several genes encoding steroidogenic enzymes (54). For instance, overexpression of DENN\_domain-containing protein 1A (DENN1A) variant 2 has been found to increase androgen biosynthesis in theca cells obtained from women with PCOS (77). Moreover, increased expression of CYP17A1, which encodes the ratelimiting enzyme in androgen biosynthesis, has also been detected in theca cells obtained from women with PCOS (77).

Hyperandrogenism can be biochemical (i.e., increased levels of circulating androgens) or clinical. The most common clinical sign of androgen excess is hirsutism. Androgens (mainly testosterone and dihydrotestosterone), through their effect on the androgen receptor, stimulate ornithine decarboxylase synthesis in the hair follicle, which in turn stimulates polyamine production. Polyamines are multifunctional cationic amines that are indispensable for cellular proliferation, including hair growth in the hair follicle (**54**).

# 2.5 Adipose tissue dysfunction

Adipose tissue is a key endocrine organ in metabolic homeostasis and is thought to underline the pathophysiology of PCOS. Indeed, functional abnormalities in adipose tissue have been reported in patients with PCOS.

There are two main types of adipose tissue: white (WAT) and brown (BAT). WAT is the main depot where metabolic energy is stored (in the form of triglycerides). The major WAT depots are classified according to their anatomic location as either subcutaneous (beneath the skin) or visceral (in the peritoneal cavity) (**78**). In turn, BAT is the main site of non-shivering thermogenesis in mammals by uncoupling protein-1 (UCP-1). BAT activity is associated with protection against obesity and related metabolic alterations, due to its capacity to oxidize metabolites and produce heat (**79**).

BAT is particularly relevant in early human development; in human neonates, BAT accounts for as much as the 5% of the body weight, however, the amount of BAT declines progressively with aging and is rather small in adulthood (**80**). More recently, it has become apparent that adult humans also have functionally relevant BAT and possibly additional capacity to induce the formation of brown-like adipocytes within WAT [named brite ("brown-in-white") or beige adipocytes] under certain conditions. The acquisition of a brown or beige phenotype is considered protective against hyperglycemia and hypertriglyceridemia (**81**).

White, brown, and brite adipocytes secret a wide variety of signaling molecules called adipokines (WAT-secreted molecules), which are bioactive molecules that act on multiple biological process and help maintaining overall homeostasis (**82**), and batokines (released by brown and/or beige adipocytes) that regulate a number of homeostatic systems including food intake, energy expenditure, insulin action and insulin secretion (**79,83**). For instance, HMW-adiponectin is a WAT-secreted molecule with insulin-sensitizing and anti-inflammatory properties (**84**) whereas C-X-C motif chemokine ligand-14 (CXCL14) is a chemokine produced by active brown/beige adipose tissue capable of improving glucose metabolism in insulin-resistant rodent models (**85**).

As explained, PCOS appears to be driven by ectopic fat accumulation coupled with a reduced energy expenditure. Studies have reported that women with PCOS exhibit increased global adiposity (**86**), larger subcutaneous adipocytes (**87**), lower lipoprotein lipase activity (**87**), impaired catecholamine-induced lipolysis (**88**), and altered adipokine release [as judged by decreased levels of circulating HMW-adiponectin (**84**) or increased levels of circulating S100A4, a member of the S100 calcium-binding protein family (**89**)]. Collectively, these results indicate that WAT dysfunction may contribute to the ectopic fat accumulation and the metabolic abnormalities observed in PCOS. Regarding BAT, women with PCOS have lower BAT activity and decreased BAT volume (**90,91**), which results in a deficient thermogenesis that may predispose to central fat accumulation and may explain, at least in part, the tendency for weight gain observed in these patients (**91**).

#### 2.6 Lipotoxicity, low-grade inflammation and oxidative stress

Lipotoxicity may also play a role in the pathophysiology of PCOS. As explained in previous sections, adipose tissue dysfunction may lead to impaired free fatty acid storage and ectopic fat accumulation, resulting in a lipotoxic state. Lipotoxicity is characterized by elevated free fatty acids, hypertriglyceridemia, and an unfavorable adipokine profile [with low levels of HMW-adiponectin, and high levels of interleukin-6 (IL-6) and tumor necrosis factor alpha (TNF- $\alpha$ )], and has adverse effects on metabolism (**26**). HMW-adiponectin promotes glucose uptake, insulin sensitivity and fatty acid oxidation (**92**), whereas TNF- $\alpha$  reduces the expression of GLUT4 (**93**) and IL-6 induces expression of suppressor cytokine signaling-3 (SOCS-3), a potential inhibitor of insulin signaling (**94**). Accordingly, lipotoxicity impairs insulin signaling.

In addition to lipotoxicity, adipocyte dysfunction is also associated with adipose tissue inflammation and low-grade inflammation. Increased levels of inflammatory markers [i.e. C-reactive protein (CRP), interleukin-1 $\beta$  (IL-1 $\beta$ ), IL-6, interleukin-18 (IL-18) or TNF- $\alpha$ ) (**95-98**)] have been detected in women with PCOS. Moreover, inflammation is closely related to oxidative stress as it induces generation of reactive oxygen species (ROS), while oxidative stress in turn promotes and aggravates inflammation (**99**). Women with PCOS have increased levels of oxidative stress (**100-102**).

# 2.7 Organokines and liver enzymes

PCOS in adolescent girls and women is often associated with MASLD (**14,103,104**). The prevalence of MASLD in PCOS has increased significantly in recent years (**105**). IR and hyperandrogenism have been found to be major contributory factors for MASLD, independently of BMI (**14**). MASLD is currently considered a systemic disorder where hepatic steatosis is merely the landmark of a systemic metabolic dysfunction, including IR, increased cardiovascular risk, and low-grade systemic inflammation (**105,106**).

Studies in adults with MASLD indicate a close association between altered hepatic function and systemic dysmetabolism, encompassing a pathogenic re-arrangement of

circulating signaling molecules, the so-called organokines (**107**), which originate in the liver itself (hepatokines), in adipose tissue (adipokines), in the muscle (myokines) or in the heart (cardiokines) (**108**).

The altered circulating levels of these signaling molecules generate multi-organ systemic disturbances, ranging from IR and inflammation to cardiovascular disease and also provide biomarker evidence of existing health risks in patients at distinct stages of disease progression (**109**). Specifically in relation to MASLD, emerging data indicate that an altered secretion of organokines plays an essential role in the pathogenesis of IR and cardiovascular disease. An example is fetuin-A, a hepatokine that elicits low-grade inflammation in MASLD by acting as an endogenous ligand of toll-like receptor-4 and promoting the secretion of proinflammatory cytokines in adipose tissue and other organs (**11**). Fetuin-A also suppresses the expression of adiponectin, overall leading to systemic IR. In MASLD, increased pro-inflammatory cytokines and enhanced levels of hepatokines such as angiopoietin-like proteins and others, also promote endothelial dysfunction, dyslipidemia and atherogenesis (**111**). However, a comprehensive knowledge of the whole set of organokines linking MASLD to systemic dysmetabolism and of the mechanisms underlying these associations is still lacking.

Women with PCOS depict altered levels of organokine signaling molecules (**112,113**); in adolescent PCOS, abnormal organokine concentrations [i.e., HMW-adiponectin (**84**); S100A4 (**89**); growth-and-differentiation factor-15 (GDF15) (**114**); fetuin-A (**115**)] have also been associated to early stages of hepatic and metabolic systemic alterations, even in the absence of overt obesity.

Existing knowledge on the identity and role of the organokines connecting MASLDassociated hepatic disturbances with systemic metabolic alterations and cardiovascular disease is still limited. However, some molecules have been recently recognized to be involved into signal metabolic dysregulation in MASLD in both experimental and clinical studies. For instance, meteorin-like protein (METRNL), has been recently reported to be related to liver injury (**116**); fibroblast growth factor-21 (FGF21), an hepatokine with enhanced expression in liver disease, may potentially have a

protective role against systemic dysmetabolism in MASLD (**117**); in addition, the blockage of diazepam-binding protein-1 (DBI, also named acyl CoA-binding protein), has been reported to improve MASLD in experimental settings (**118**).

Liver enzymes [aspartate aminotransferase (AST), alanine aminotransferase (ALT) and gamma-glutamyl transferase (GGT)] are considered markers of hepatic damage (**119,120**). Indeed, mild elevations of liver enzymes in the upper normal range associate with features of metabolic syndrome and MASLD (**121**) and elevated levels of liver enzymes (ALT, AST and GGT) were detected in children with MASLD (**122**). In addition, increased ALT levels have also been identified in women with PCOS (**123,124**).

# 2.8 Thyroid disorders

Thyroid hormone (TH) is a key regulator of metabolism and energy balance (**125**), and is under the regulation of the hypothalamic-pituitary-thyroid (HPT) axis, a multilevel axis that facilitates the regulation of homeostasis and metabolism. Thyrotropinreleasing hormone (TRH) is released by the hypothalamus in response to the stimuli of energy-sensing molecules, and initiates the cascade of HPT axis function. TRH triggers the secretion of TSH from thyrotropic cells in the anterior pituitary, which stimulates the production and release of TH (**126**).

Thyroid disorders have been associated with menstrual irregularity, infertility, poor pregnancy outcomes and also PCOS (**127**). Women with PCOS have higher concentrations of circulating TSH (**128-131**). However, it is unknown whether TSH levels are already elevated in adolescent girls with PCOS. Raised levels of TSH detected in PCOS women are known to associate with high rates of gestational complications. Recently, a low range of preconception TSH (between 0.91 and 1.82 mIU/L) has been strongly linked to lower rates of gestational complications, specifically to preterm delivery (**132**).

#### 2.9 Dysbiosis of gut microbiota

The gut microbiota provides a set of benefits to the host, including protective, structural and histological functions (**133**). For instance, a relatively increased microbiota gene content and microbial diversity has been associated with enhanced metabolic health (**134**). Gut microbiota can also influence host metabolism and hormone release via the production of several metabolites, such as short-chain fatty acids (SCFA) and bile acids (**135**).

Alteration of the composition of the intestinal microbiota, so-called dysbiosis, has been associated with many disorders, such as obesity, IR and T2D (**136,137**). Indeed, dysbiosis of the gut microbiota has also been reported in microbiota-centric studies on PCOS (**138**). Lean adult women with PCOS as well as women and adolescents with obesity and PCOS have gut microbiota communities different from those in healthy controls (**139-143**) suggesting that the dysbiosis of the gut microbiota may contribute to the pathogenesis of the disorder, although the directionality remains so far unclear (**144**). It has been proposed that hyperandrogenism could alter gut microbiota (**142,145**). In turn, altered gut microbiota could lead to abnormal SCFAs production. SCFAs, in turn, can bind to free fatty acid receptor (FFAR)-2 and 3 and suppress the release of appetite-stimulating hormone such as ghrelin (**146**). Ghrelin can block the production of cytochrome P450 family 19 subfamily A member 1 (CYP19A1) in adipostromal cells preventing androgen conversion to estrogens (**147**). However, the profile of the gut microbiota in adolescent girls without obesity has so far not been characterized.

#### 2.10 Genetic, epigenetic and environmental factors

Genetic susceptibility to PCOS is known to be controlled by at least 19 loci identified through genome-wide association studies (GWAS), and mainly linked to genes related to insulin signaling, sexual hormone function and T2D (**148-152**). Moreover, epigenome-wide association studies (EWAS) have demonstrated altered DNA methylation and gene expression patterns in different tissues (**153-156**) or

peripheral blood (**157-159**) from women with PCOS as compared to healthy controls, providing convincing evidences linking epigenetic regulation with disease development. MicroRNAs (miRNAs) have also been described as important players in the onset and progression of PCOS. Altered expression of miRNAs has been reported in serum, adipose tissue, follicular fluid and granulosa theca cells (**160-163**) of women with PCOS and in serum of adolescent girls with PCOS and without obesity (**164**).

Environmental factors are known to have a role in the pathogenesis of PCOS. It has been described that socioeconomic status along with unhealthy behaviors (such as smoking, bad eating habits or sedentary lifestyle) may impact into the development of PCOS (**165**). Among the environmental factors, endocrine-disrupting chemicals (EDCs) or advanced glycated end-products (AGEs) are potential contributors to the pathophysiology of PCOS. For instance, increased levels of bisphenol A, one of the main EDCs, have been reported in serum of women and adolescents with PCOS, and to positively correlate with androgen levels, suggesting a role of bisphenol A in the pathogenesis of PCOS (**166,167**). Elevated serum AEGs levels were also found in women with PCOS (**168,169**).

**Figure 4** summarizes the potential mechanisms underlying the pathophysiology of PCOS.



**Figure 4. Pathophysiology of polycystic ovary syndrome.** [1] Faster gonadotropin-releasing hormone (GnRH) pulse generator increases luteinizing hormone (LH) over follicle-stimulating hormone (FSH) production, leading to an impaired follicular development (due to decreased FSH levels) and to an excessive androgen production (due to increased LH levels); in turn, increased androgen levels also trigger faster GnRH pulsation frequency. [2] Hyperinsulinemia stimulates GnRH gene transcription, decreases hepatic sex hormone binding-globulin (SHBG) release and stimulates the activity of ovarian cytochrome P450-17 alpha, leading to an increase of androgen levels. [3] Ectopic fat accumulation leads to an altered adipokine release that cause insulin resistance. [4] Increased androgen levels alter the composition of the gut microbiota. In turn, gut bacterial metabolites (such as short-chain fatty acids) can favor an increase of androgen levels (Own elaboration, created with Biorender.com).

#### 3. DIAGNOSIS OF PCOS

The wide range of PCOS prevalence among populations can be explained by the use of three different sets of diagnostic criteria issued by: [1] the National Institutes of Health (NIH)'s international conference on PCOS in 1990 (**170**); [2] the European Society of Human Reproduction and Embryology (ESHRE) and the American Society for Reproductive Medicine (ASRM) in 2003 (referred to as the Rotterdam criteria) (**171**); [3] the Androgen Excess and PCOS Society (AE-PCOS) criteria in 2006 (**172**).

The NIH criteria included the presence of clinical and/ or biochemical hyperandrogenism and chronic anovulation (**170**). The Rotterdam criteria are the broadest and use the presence of polycystic ovarian morphology (PCOM) on ultrasound as a new criterion. According to the Rotterdam criteria, diagnosis requires the presence of at least two of three features: hyperandrogenism, anovulation and PCOM on ultrasound (**171**). Finally, the AE-PCOS criteria defined PCOS as hyperandrogenism with ovarian dysfunction or polycystic ovaries (**172**). In all cases, other causes of androgen excess, such as non-classical congenital adrenal hyperplasia, Cushing's syndrome, androgen-secreting tumors, hyperprolactinemia, thyroid dysfunction, and drug-induced androgen excess should be excluded, as well as other causes of oligomenorrhea or anovulation (**173**).

The diagnostic criteria for PCOS in adolescence are controversial because some diagnostic traits used in adult women such as irregular menses or a PCOM on ultrasound may be normal pubertal physiological events in girls (6). This could lead to under-diagnosis, and to delayed and/ or poor diagnosis experiences (174); on the other hand, a timely and not too early diagnosis in girls offers protection against overtreatment (20,175).

Over the last decade, two international consortia have proposed the diagnostic criteria for adolescent PCOS. In 2017, an international collaboration of pediatric endocrinologists suggested a first set of diagnostic criteria for adolescent PCOS (6). In 2020, an international consortium, which focused on the international evidence-based guideline for the assessment and management of PCOS, and recently revised (**7**,**8**), also

phrased the diagnostic criteria for adolescent PCOS (9). The main diagnostic criteria suggested by both consortiums are summarized in **Table 1**.

	Criteria proposed in	Criteria proposed in
	>2 years post menarche	Post-menarche
Menstrual irregularity	Cycles <21 days (dysfunctional bleeding)	1–3 years: cycles <21 or >45 days
	Cycles >45 days (oligomenorrhea)	>3 years: cycles <21 or >35 days, or <8 cycles/year
	Secondary amenorrhea (no menses for >90 days)	>1 year: any cycle >90 days
	Primary amenorrhea after completed growth/puberty	Primary amenorrhea at 15 yrs or >3 yrs after start of Tanner breast stage 2
	Clinical	
Androgen excess	Presence of hirsutism (modified Ferriman-Gallwey score ≥5-6)	Presence of hirsutism (modified Ferriman-Gallwey score ≥4-6)
	And	And/ or
	Biochemical	
	Increased total testosterone or free androgen index (FAI)	Increased total testosterone or free androgen index (FAI)

 Table 1. Diagnostic criteria for adolescent PCOS, adapted from (20):

Despite some differences, both consortiums agree in not using as diagnostic criteria: [1] PCOM on ultrasound, due to the high incidence of physiological multi-follicular ovaries in this life stage; [2] AMH levels, since there is a weak association between AMH levels and adolescent PCOS.

Recently, in 2023 a proposal of merging the two sets of previously recommended criteria into one diagnostic ensemble has emerged (**20**). This proposal defines PCOS as the combination of clinical and/or biochemical androgen excess associated to menstrual irregularity. Irregular menstrual cycles (>2 years post menarche) are defined as:

- cycles <21 days (dysfunctional bleeding)
- cycles >45 days (oligomenorrhea)
- secondary amenorrhea (no menses for >90 days)
- primary amenorrhea after completed growth/puberty

Clinical androgen excess is defined as the presence of hirsutism (modified Ferriman-Gallwey score ≥4-6)]; biochemical androgen excess is defined as the presence of increased total testosterone or free androgen index (FAI)].

#### 4. MANAGEMENT OF PCOS

To date, no treatment for PCOS in adolescent girls or young women (AYAs) has been approved by the European Medicines Agency (EMA) or the US Food and Drug Administration (FDA). Accordingly, this is an unmet need.

Prime recommendation is to reduce body adiposity with lifestyle measures (176,177), such as hypocaloric, ketogenic and anti-inflammatory diets (178), intermittent fasting (179), physical exercise and psychological support (180). The roles of sleep and biorhythm are increasingly recognized (181). However, the long-term effectiveness of lifestyle intervention remains unclear. It has been proposed that lifestyle measures are more efficacious in adolescents than in adults, suggesting that the intervention is more useful when applied early (176,182). Despite this, a recent systematic review of the literature revealed that the evidence is of poor quality with wide variations in trial designs, comparisons, and outcome reporting (183).

When lifestyle measures fail to manage PCOS alterations, a medication can be helpful. Several pharmacological approaches have been used; they are summarized in Table 2. Oral contraceptives (OCs) are frequently prescribed off-label to approximately 98% of young PCOS patients, even to those who do not need contraception (7-9,184). OCs alleviate the clinical symptoms by inducing a combination of anovulatory infertility, regular pseudo-menses and pharmacological elevations of SHBG (177,185). However, they do not revert the underlying pathophysiology so that, upon treatment discontinuation, the entire PCOS phenotype tends to reappear, with subsequent increased risks for subfertility and for potentially lifelong comorbidities that are relevant for public health (186,187). Indeed, OC treatment in AYAs with PCOS fails to normalize IR, the circulating levels of CRP (a marker of low-grade inflammation), carotid intima-media thickness (cIMT, a marker of premature vascular ageing) or ectopic fat (**20,188-191**). Likewise, in the general population, the use of contemporary OCs has been associated with an increased risk for breast cancer, glioma and venous thromboembolism, which may vary with estrogen dose and progestogen type (192-**194**). In women with PCOS, the use of OCs with purportedly low thrombogenic

potential has nevertheless been linked to more risks for inflammatory and coagulation disorders (**195**).

Insulin sensitizers have also been used in PCOS. Metformin (MET), a biguanide approved for the treatment of T2D, is the drug most widely used for the treatment of T2D in adults and in children (aged > 10.0 years) (**40,196,197**). Evidence-based guidelines recommend the use of MET in women with PCOS, obesity and IR, in order to manage endocrine-metabolic abnormalities (**7,8,198**). Although MET has a limited effect on hirsutism, its benefits on other PCOS-associated features and co-morbidities have long been recognized in women; some of these benefits have also been reported in girls, including for ovulation induction (**199,200**). Thiazolidinediones (TZDs) are also insulin sensitizers that have been used in PCOS. TZDs improve IR and dyslipidemia in PCOS more than MET (**201-203**). In addition, the combination of MET and TZDs seems to have a synergistic effect, conferring a greater improvement in SHBG, AMH and postprandial glucose levels (**204**).

Drugs that block androgen action have also been used off-label in the management of PCOS (**205-209**): androgen receptor blockers [such as spironolactone (SPI), flutamide or cyproterone acetate] and 5- $\alpha$ -reductase inhibitors (i.e., finasteride). Anti-androgens significantly reduce hirsutism compared to placebo (**210**) and normalize endocrinemetabolic variables and menstrual cyclicity better than MET in monotherapy (**211**). In addition, the combination of MET and SPI appears to be more effective in reducing BMI and androgen levels than MET alone, and when the treatment course is greater than 6 months, the combination therapy reduces fasting blood glucose and improves IR more effectively than MET alone (**212**). Anti-androgens intake should be accompanied by contraceptive measures due to their teratogenic potential (**206**).

Over the last years, other potential therapies have emerged for the management of PCOS. For instance, treatment with inositol isomers, such as myo-inositol and Dchiro-inositol, appears to improve the regularity of the menstrual cycles, IR and endocrine-metabolic parameters (**213**); however, controversy remains regarding the exact dosing and the extent of the benefit (**214,215**). The evidence supporting the

benefits of dietary supplements like prebiotics, probiotics, vitamin D, polyunsaturated fatty acids or phytochemicals is still limited and inconclusive (**216-219**).

Statins, such as simvastatin or atorvastatin, have also been investigated as a potential therapy for PCOS with obesity, as they have been shown to improve IR and reduce cardiovascular disease risk. However, the level of evidence continues to be low (**220**).

Glucagon-like peptide-1 (GLP-1) agonists, such as semaglutide or liraglutide, have also been studied in patients with PCOS and with obesity. GLP1 agonists decrease body weight and improve androgen excess, metabolic markers and reproductive function in women with PCOS (**221,222**). Currently, there is an ongoing clinical trial comparing the effects of intensive diet versus semaglutide in adolescent girls with PCOS and obesity (**223**). Preliminary results show more weight loss with semaglutide, but similar effects on fasting insulin, androgen excess and menstrual regularity.

Sitagliptin, a dipeptidyl peptidase-4 (DPP4) inhibitor, has also been studied in several clinical trials (**224-226**). Although sitagliptin treatment showed beneficial effects on improving IR and visceral obesity, the evidence is still limited (**227**).

Sodium-glucose co-transporter-2 (SGLT-2) inhibitors, a new class of antidiabetic agents such as canagliflozin, dapagliflozin and empagliflozin, have been approved by the FDA for the treatment of T2D (**228-230**) and currently their potential applications in PCOS are being investigated (**231-236**).

Neurokinin 3 (NK3) receptor antagonism in the kisspentin-neurokinin B-dynorphin A (KNDy) neurons decreases GnRH pulse frequency, reduces basal LH secretion, the LH-to-FSH ratio, suppresses follicle development, and modulates ovarian sex hormone production (**237**). Phase 2 studies examining the effects of the NK3 receptor antagonist Fezolinetant (ESN364) on PCOS are ongoing (**238**).

Artemisinins, which are compounds with well-known antimalarial properties, have been shown to suppress ovarian androgen synthesis by promoting degradation of cytochrome P450 family 11 subfamily A member 1 (CYP11A1), holding thus the

potential for treating PCOS (**239**). However, there is no evidence (yet) on the insulin sensitizing and ectopic-fat reducing effects of these compounds (**239**).

Given the potential key role of ectopic (hepato-visceral) fat in the pathogenesis of PCOS, the focus of the treatment should be to reduce ectopic fat. In turn, such reduction should have normalizing effects on the entire PCOS phenotype. This insight has prompted the exploration of an alternative treatment that consists of a low-dose combination of two insulin sensitizers [pioglitazone (PIO) and metformin (MET)] and one mixed anti-androgen and anti-mineralocorticoid [spironolactone (SPI)] which also increases BAT activity and volume (240). The triple low-dose combination of SPI, PIO and MET—so-called SPIOMET—proved to be superior to an OC in normalizing the PCOS phenotype (199). Ibáñez et al., performed a randomized, open-label, singlecenter, pilot proof-of concept study (12 months on treatment, then 12 months off) in adolescent girls with PCOS and without obesity (N=34; mean age 16 years; BMI 23.5 kg/m<sup>2</sup>). This study aimed to compare the effects of OC versus a low-dose combination of SPI, PIO and MET (SPIOMET). Evidence is still limited but suggests that SPIOMET does indeed hold the potential to revert the PCOS phenotype, particularly on ectopic fat excess, insulin sensitivity and post treatment ovulation rate (199), as shown in Figure 5.



**Figure 5.** Longitudinal changes in hepatic fat and in the mean serum insulin (MSI) Z-score during an oral glucose tolerance test, and post treatment number of ovulations in adolescent girls with polycystic ovary syndrome and without obesity, participating in a randomized pilot study performed by Ibáñez et al. Girls were randomized to receive an oral contraceptive (OC, N=17) or a low dose combination of spironolactone, pioglitazone and metformin (SPIOMET,

N=17) for 12 months (shaded area) and were subsequently followed for another 12 months without treatment. Results are expressed as mean  $\pm$  standard error of the mean. Data from (**199**).

**Table 2.** Medications used off-label or under investigation for polycystic ovarysyndrome (own elaboration):

OFF- LABBEL				
Oral contraceptives	Ethinylestradiol + Levonorgestrel	Inhibition of ovarian androgen secretion and increase in hepatic SHBG production, resulting in less circulating free androgens		
Insulin sensitizers				
Biguanide	Metformin	Reduction of hepatic glucose production and improvement of peripheral insulin sensitivity		
Thiazolidinedione	Pioglitazone	Improvement of IR and dyslipidemia		
Antiandrogens				
Androgen receptor blockers	Spironolactone Flutamide Cyproterone acetate	Reduction of hirsutism and normalization of endocrine-metabolic variables and menstrual cycle		
	Fillasteride			
Inositol isomers	Myo-inositol D-chiro-inositol	regularity and IR		
Statins	Simvastatin Atorvastatin	Improvement of IR and reduction of the risk of cardiovascular disease		
GLP-1 agonists	Semaglutide Liraglutide	Decrease of body weight and improvement of androgen excess		
DPP4 inhibitor	Sitagliptin	Improvement of insulin sensitivity and visceral fat		
SGLT-2 inhibitors	Canagliflozin Dapagliflozin Empagliflozin	Improvement of insulin sensitivity		
NK3 receptor antagonism	Fezolinetant (ESN364)	Reduce LH/FSH ratio and ovarian sex hormone production		
Artemisinins	Artemisinin analog (artemether)	Suppress ovarian androgen synthesis		
SPIOMET	Spironolactone + Pioglitazone + Metformin	Normalization of ectopic fat, insulin sensitivity and ovulation.		

Abbreviations: DPP4, dipeptidyl peptidase-4; FSH, follicle-stimulating hormone; GLP1, glucagon-like peptide 1; IR, insulin resistance; LH, luteinizing hormone; NK3, neurokinin 3; SGLT-2, sodium-glucose co-transporter; SHBG, sex hormone binding-globulin; SPIOMET, spironolactone, pioglitazone, metformin.

SPI, PIO and MET are medications that have long been approved for other indications and used for many years in Europe as well as in the USA.

SPI is a steroid with a structure resembling aldosterone, the main mineralocorticoid hormone, and thus can act as a competitive inhibitor of aldosterone activity. SPI is marketed as diuretic but can serve as an anti-androgen if given in higher doses (up to 200 mg/d). In Europe and in the USA, SPI has for decades been the safe anti-androgen of choice in the treatment of hirsutism with or without hyperandrogenism (**241**). SPI has also been shown to decrease sebum production and to improve acne. In addition, it has recently been identified as a potent activator of BAT which, in turn, may raise energy expenditure, control weight gain, and ultimately result in a lower amount of ectopic fat (**240**). In most countries SPI is marketed within a drug formulation branded as Aldactone<sup>®</sup>.

PIO acts as an agonist of nuclear receptor peroxisome proliferator-activated receptor (PPAR)-gamma and to less extent PPAR-alpha (**242**). It is a TZD acting as an insulin sensitizer in adipose tissue, liver and muscle. It raises circulating HMW-adiponectin (**243**), and also insulin sensitivity via preferentially subcutaneous adipogenesis (**244**). In women with PCOS, PIO improves insulin sensitivity, menstrual cyclicity and ovulation rates (**245**). In most countries PIO is marketed within a drug formulation branded as Actos<sup>®</sup>.

MET has anti-diabetic properties by acting via multiple mechanisms. MET reduces hepatic glucose production (by inhibiting gluconeogenesis and glycogenolysis) and improves peripheral insulin sensitivity. MET is also known to raise AMP-activated protein kinase (AMPK) activity and the circulating concentrations of GDF15, a peptide hormone that may reduce appetite through a specific receptor in the brainstem

(**199,246-250**). In most countries MET is marketed within a drug formulation branded as Metformina<sup>®</sup> or Glucophage<sup>®</sup>.

In the Ibáñez et al. study (**199**), adolescent girls with PCOS and without obesity were randomly assigned to receive once daily, at dinner time, either Loette Diario<sup>®</sup> (Pfizer, Madrid, Spain; 20 µg of ethinylestradiol plus 100 mg levonorgestrel for 21/28 days and placebo for 7/28 days) or SPIOMET, a low dose-combination of 50 mg/d SPI (half of 100 mg tablet of Aldactone<sup>®</sup>, Pfizer, Madrid, Spain), 7.5 mg PIO (half of 15 mg tablet of Actos<sup>®</sup>, Takeda, Madrid, Spain) and 850 mg MET (850 mg tablet of Metformina<sup>®</sup>, Sandoz, Barcelona, Spain).

No safety concerns have been detected with the low doses of SPI, PIO and MET used in the pilot study (**199**). Since its approval in 1960, no safety concerns related to the use of SPI have been raised (**251**), only minor side effects were reported at higher doses (100 mg/d or more) (**210**); thus, there are essentially no safety concerns with only 50 mg/d. The safety and efficacy of PIO in the pediatric population is limited and dosing recommendations are not available; however, PIO appears to be well tolerated by children and adolescents since no side effects have been identified in pediatric patients and young women receiving low-dose PIO (**84,188,243,252,253**). Finally, over the past 60 years, considerable experience has been accumulated regarding the clinical use and safety of MET; the dose of MET used in the pilot study (**850** mg/d) is at the lower limit of the recommended range (**850-2000** mg/d) (**254**).

To sum up, Figure 6 depicts the targeted efficacy of SPIOMET components.



**Figure 6. Targeted efficacy of SPI, PIO, MET and the combination of SPIOMET.** Abbreviations: BAT, brown adipose tissue; GDF15, growth-and-differentiation factor-15; HMW, high-molecular-weight; MET, metformin; PIO, pioglitazone; SPI, spironolactone; SPIOMET, spironolactone, pioglitazone, metformin (Own elaboration).

# **HYPOTHESIS AND OBJECTIVES**

#### **HYPOTHESIS**

Polycystic ovary syndrome is the most common cause of hirsutism and menstrual irregularities in adolescent girls and young women. However, there is no approved treatment for adolescent polycystic ovary syndrome. A pilot study exploring an alternative treatment for polycystic ovary syndrome consisting of the intake of a low-dose combination of spironolactone – pioglitazone – metformin for 1 year proved to have more normalizing effects than the standard oral contraceptive treatment.

The hypotheses of this doctoral thesis are:

- 1- The benefits of spironolactone pioglitazone metformin treatment will be corroborated in a second pilot study with virtually the same design; pooled data will confirm that spironolactone – pioglitazone – metformin intervention will have more normalizing effects than the standard oral contraceptive treatment, especially on ectopic fat, insulin resistance and ovulation.
- 2- Brown adipocytes from patients with polycystic ovary syndrome will secret secrete less C-X-C motif chemokine ligand-14. Adolescent girls with polycystic ovary syndrome will have altered circulating C-X-C motif chemokine ligand-14 concentrations. Such altered levels will be normalized after spironolactone – pioglitazone – metformin treatment.
- 3- Altered gut microbiota might constitute one of the mechanisms underpinning adolescent polycystic ovary syndrome. Girls with polycystic ovary syndrome and without obesity will have a different gut microbiota profile as compared to healthy adolescents. After spironolactone – pioglitazone – metformin or oral contraceptive treatment, divergent changes in the microbial composition will be observed.
- 4- Thyroid disorders might represent a potential factor involved in the pathophysiology of adolescent polycystic ovary syndrome; adult women with polycystic ovary syndrome have higher concentrations of circulating thyroidstimulating hormone. Adolescent girls with polycystic ovary syndrome and without obesity will present increased circulating levels of thyroid-stimulating hormone.

These altered levels will be normalized after spironolactone – pioglitazone – metformin intervention.

5- Altered secretion of organokines may play a role in the pathogenesis of polycystic ovary syndrome. Spironolactone – pioglitazone – metformin and oral contraceptive treatment will have divergent effects on organokine patterns.

#### **OBJECTIVES**

In order expand the knowledge on the pathophysiology of polycystic ovary syndrome as well as on the effects of spironolactone – pioglitazone – metformin treatment, this thesis will formulate the following objectives:

- 1- To compare the effects of a low dose combination of spironolactone pioglitazone – metformin versus those of an oral contraceptive (ethinylestradiol plus levonorgestrel) (1 year on treatment, then 1 year off treatment) on menstrual regularity and ovulation rate, insulin sensitivity and hepato-visceral fat in a wider cohort of adolescents with polycystic ovary syndrome and without obesity.
- 2- To assess C-X-C motif chemokine ligand-14 concentrations in adolescent girls with polycystic ovary syndrome and without obesity as compared to control girls; to evaluate the effects of 1 year of treatment with spironolactone pioglitazone metformin or oral contraceptive on circulating C-X-C motif chemokine ligand-14 concentrations and its relation to metabolic improvement; to determine the individual effects of spironolactone, pioglitazone and metformin on C-X-C motif chemokine ligand-14 in human adipocytes.
- 3- To study the composition of the gut microbiota in adolescent girls with polycystic ovary syndrome and without obesity versus those in age-matched control girls as well as the effects of 1 year of treatment with spironolactone pioglitazone metformin or oral contraceptive on microbiota diversity, pattern and profile.
- 4- To determine whether thyroid-stimulating hormone levels are elevated in adolescent girls with polycystic ovary syndrome and without obesity, and if so whether 1 year of spironolactone – pioglitazone – metformin or oral contraceptive treatment reduces thyroid-stimulating hormone levels, and whether those levels remain low 1 year after treatment discontinuation.
- 5- To assess the effects of 6 months of treatment with spironolactone pioglitazone metformin or oral contraceptive on a set of organokines (fibroblast growth factor-21, diazepam-binding protein-1 and meteorin-like protein) in adolescent girls with polycystic ovary syndrome and without obesity, and to report the associations of

those organokines with circulating biomarkers of liver damage which were assessed longitudinally as safety markers in the aforementioned pilot studies.

# RESULTS

All the data included in this doctoral thesis are part of two randomized, open-label, pilot studies, comparing the effects of an alternative treatment consisting of a low dose-combination of SPI, PIO and MET (SPIOMET) versus those of a standard OC treatment in adolescent girls with PCOS and without obesity. The studies were conducted in the Endocrinology Department of Sant Joan de Déu University Hospital (Barcelona, Spain), between January 2013 and December 2015, after approval by the Institutional Review Board of Sant Joan de Déu University Hospital.

Participating girls were randomized to receive either an OC (20  $\mu$ g of ethinylestradiol plus 100 mg levonorgestrel for 21/28 days and placebo for 7/28 days) or SPIOMET (low dose-combination of 50 mg/d SPI, 7.5 mg/d PIO and 850 mg/d MET) for 1 year and remained subsequently untreated for 1 year. The inclusion/ exclusion criteria were:

- Inclusion criteria:
  - Hirsutism score > 8 on the Ferriman-Gallway scale (255)
  - Oligomenorrhea (menstrual intervals > 45 days)
  - Gynecological age (or time span post-menarche) > 2.0 years
  - No need for contraception
- Exclusion criteria:
  - Evidence of late-onset adrenal hyperplasia due to 21-hidroxylase deficiency (as judged by 17-hydroxy-progesterone levels > 200 ng/dL in the follicular phase or after 2 months of amenorrhea) (256)
  - Glucose intolerance or diabetes mellitus
  - Evidence of thyroid, liver or kidney dysfunction
  - o Hyperprolactinemia
  - Prior use of medication affecting gonadal or adrenal function, or carbohydrate or lipid metabolism

Aged-matched healthy girls recruited in nearby schools served as controls. All had regular menstrual cycles, and none was hirsute or taking medications.

**STUDY 1** 

# TOWARD A TREATMENT NORMALIZING OVULATION RATE IN ADOLESCENT GIRLS WITH POLYCYSTIC OVARY SYNDROME

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#### **STUDY 1**

Toward a treatment normalizing ovulation rate in adolescent girls with polycystic ovary syndrome

**Introduction:** Adolescent polycystic ovary syndrome (PCOS) is characterized by androgen excess and oligomenorrhea, and commonly driven by hepato-visceral fat. There is no approved treatment for adolescent PCOS, however, it is commonly treated with an oral contraceptive (OC). A pilot study tested whether an intervention targeting the reduction of hepato-visceral fat was followed by a higher ovulation rate than OC treatment. This intervention consisted of the intake of a low-dose combination of spironolactone, pioglitazone and metformin (SPIOMET) or OC for 1 year. SPIOMET had more normalizing effects than OC treatment. However, the limited power of the study (N=34) prompted the launch of a second study with virtually identical design.

**Objective:** To compare the effects of a low dose combination of SPIOMET or OC for 1 year on hepato-visceral fat, insulin sensitivity and ovulation rate in a larger population of adolescent girls with PCOS and without obesity.

**Material and methods:** The studied population consisted of 62 adolescent girls with PCOS and without obesity [age, 15.8  $\pm$  0.2 years; BMI, 24.2  $\pm$  0.5 Kg/m<sup>2</sup>] who participated in two randomized, open-label, pilot studies with the same design (1 year on treatment, then 1 year off; N=34 patients from Study 1 and N=28 from Study 2). Compared treatments were OC [20 µg ethinylestradiol plus 100 mg levonorgestrel (21/28 days), and placebo (7/28 days)] versus a low-dose combination of spironolactone 50 mg/d, pioglitazone 7.5 mg/d, and metformin 850 mg/d (SPIOMET). The primary outcome was post treatment ovulation rate inferred from menstrual diaries and weekly salivary progesterone measurements during the second and the fourth quarter of the post treatment year. Secondary outcomes included: auxological and endocrine-metabolic variables [fasting insulin, androgens, high-molecular-weight adiponectin (HMW-adiponectin), and microRNA (miR)-451a], body composition [dual x-ray absorptiometry (DXA)] and hepato-visceral fat [magnetic resonance imaging

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(MRI)] were assessed on and post treatment. Healthy age-matched adolescent girls (N=52; mean age 16.3 years) served as controls.

**Results:** Pooled results of the 2 pilot studies confirmed that OC and SPIOMET treatment reduced the androgen excess comparably and had no differential effects on total-body lean or fat mass. However, SPIOMET was accompanied by more broadly normalizing effects, including on hepato-visceral fat and on circulating insulin, HMW-adiponectin, and miR-451a. On average, there were 3-fold more ovulations post-SPIOMET than post-OC; normovulation was only observed after SPIOMET; anovulation was >10-fold more prevalent post-OC.

**Conclusion:** Pooled results of randomized studies in non-obese adolescent girls with PCOS indicate that SPIOMET treatment leads to an overall healthier, more insulin-sensitive condition-with less ectopic fat than OC treatment, and to a more normal post treatment ovulation rate.

## Toward a Treatment Normalizing Ovulation Rate in Adolescent Girls With Polycystic Ovary Syndrome

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Adolescent polycystic ovary syndrome (PCOS) is characterized by androgen excess and oligomenorrhea, and commonly driven by hepato-visceral fat excess ("central obesity") ensuing from a mismatch between prenatal and postnatal nutrition, on a background of genetic susceptibility. There is no approved treatment for adolescent PCOS.

We report the pooled results of 2 pilot studies in nonobese girls with PCOS (N = 62, age 15.8 years) that compared the effects of randomized treatment for 1 year, either with an oral estro-progestogen contraceptive (OC), or with a low-dose combination of spironolactone-pioglitazone-metformin (SPIOMET, targeting the excess of ectopic fat).

Auxological and endocrine-metabolic variables (including fasting insulin, androgens, highmolecular-weight adiponectin [HMW-adiponectin], and microRNA [miR]-451a), body composition (dual x-ray absorptiometry) and hepato-visceral fat (magnetic resonance imaging) were assessed on- and posttreatment. Data from menstrual diaries were combined with weekly salivary progesterone measurements to infer ovulation rates during the second and fourth quarter of the posttreatment year.

OC and SPIOMET treatment reduced the androgen excess comparably, and had no differential effects on total-body lean or fat mass. However, SPIOMET was accompanied by more broadly normalizing effects, including on hepato-visceral fat and on circulating insulin, HMW-adiponectin, and miR-451a. On average, there were 3-fold more ovulations post-SPIOMET than post-OC; normovulation was only observed after SPIOMET; anovulation was >10-fold more prevalent post-OC.

Pooled results of randomized studies in nonobese adolescent girls with PCOS indicate that SPIOMET treatment leads to an overall healthier, more insulin-sensitive condition—with less ectopic fat—than OC treatment, and to a more normal posttreatment ovulation rate.

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Key Words: PCOS, ovulation, hepatic fat, visceral fat, metformin, spironolactone, pioglitazone

Abbreviations: BMI, body mass index; cIMT, carotid intima-media thickness; CRP, C-reactive protein; HMW, highmolecular-weight; HOMA-IR, homeostatic model assessment-insulin resistance; miR, microRNA; OC, oral contraceptive; MRI, magnetic resonance imaging; PCOS, polycystic ovary syndrome; SEM, standard error of the mean; SPIOMET, spironolactone-pioglitazone-metformin.

There is no approved treatment for polycystic ovary syndrome (PCOS), a prevalent condition in adolescent girls and young women [1, 2]. Many of these patients are guided into a trajectory that starts with oral contraceptive (OC) treatment, leads into oligo-anovulatory subfertility, then into the use of assisted reproductive techniques, and ultimately into pregnancies with a double-to-triple risk for complications (such as gestational diabetes, preeclampsia, and preterm birth) potentially with lifelong sequelae in the offspring [2].

Evidence is converging toward the insight that adolescent PCOS is frequently driven by hepato-visceral fat excess ("central obesity") ensuing from a mismatch between (rather restrictive) prenatal and (rather abundant) postnatal nutrition, on a background of (epi) genetic susceptibility [3, 4, 5]. This insight has prompted the exploration of an alternative treatment for PCOS consisting of the intake of a low-dose combination of spironolactone (a mixed anti-androgen and anti-mineralocorticoid, also activating brown adipose tissue) [6] with pioglitazone and metformin (2 insulin sensitizers acting through different mechanisms) (SPIOMET) for 1 year. This combination proved to have more normalizing effects than OC treatment, in particular, on ectopic fat excess, insulin sensitivity, and posttreatment ovulation rate [7]. The limited power of the first study (N = 34) prompted the launch of a second study with virtually identical design. Here we report the pooled results of both studies in nonobese girls with PCOS (N = 62).

#### 1. Materials and Methods

#### A. Study Population & Design

Both pilot studies (ISRCTN29234515 and ISRCTN11062950) had an open-label, randomized, controlled design, and were conducted in the Adolescent Endocrinology Unit of Sant Joan de Déu University Hospital, Barcelona, Spain. Recruitment was biased against overweight/obesity because, in our setting, overweight/obese adolescent girls are primarily referred to the Adolescent Obesity Unit rather than to the Adolescent Endocrinology Unit. In each study, the on-treatment year was followed by a posttreatment year. Study completion rate was 89% (62/71) (Fig. 1, flow chart).

The inclusion criteria were hirsutism (score > 8 on modified Ferriman-Gallwey scale), oligomenorrhea (menstrual intervals > 45 days), gynecological age > 2.0 years, and absence of sexual activity (no need for contraception). Exclusion criteria were 21-hydroxylase deficiency; glucose intolerance or diabetes; evidence of thyroid, liver, or kidney dysfunction; hyperprolactinemia; and prior use of medications affecting gonadal/adrenal function, or carbohydrate/lipid metabolism [7, 8]. Mediterranean diet and regular exercise were recommended to all participating girls; OC treatment consisted of 20  $\mu$ g ethinylestradiol plus 100 mg levonorgestrel for 21/28 days, and placebo for 7/28 days; SPIOMET treatment consisted of a low-dose combination of spironolactone 50 mg/day, pioglitazone 7.5 mg/day, and metformin 850 mg/day [7].

Age-matched, healthy girls (N = 52; mean age 16.3 years) recruited from nearby schools served as controls. All had regular menstrual cycles, and none was hirsute or taking medication.

The primary endpoint was posttreatment ovulation rate; secondary outcomes included hirsutism score, fasting insulin, androgens, lipids, high-molecular-weight (HMW) adiponectin, C-reactive protein (CRP), carotid intima-media thickness (cIMT), body composition, and hepato-visceral fat [7]; circulating microRNA (miR)-451a could only be measured in a subset of the participating girls (footnote below Table 1).

Blood sampling in both patients and controls was at all time points performed either in the follicular phase of the menstrual cycle (days 3-7) or after 2 months of amenorrhea; at study start, the ratio of amenorrheic to oligomenorrheic girls was 1 to 7.

#### B. Assessments

Birth weight, birth length, and body mass index (BMI) (and their Z-scores) were retrieved from medical records. Endocrine-metabolic variables and cIMT were assessed as described



Figure 1. Flow chart for Study 1 and Study 2.

[7, 8]. Homeostatic model assessment-insulin resistance (HOMA-IR) was calculated as [fasting insulin in mU/L] x [fasting glucose in mg/dL]/405. Ovulation rates were inferred by combining data from menstrual diaries and from progesterone concentrations assessed in weekly saliva samples, obtained over 12 weeks in the second quarter and then 12 weeks in the fourth quarter of the posttreatment year [7]. Progesterone was measured by enzymelinked immunosorbent assay (ELISA) (Novatec, Inmundiagnostica, cat# DSNOV25, RRID:AB\_2827743) [9]. Circulating miR-451a was measured as described [5], with results expressed in Z-scores, using the data of healthy control girls as reference; circulating miR-451a concentrations are known to be low in adolescent girls with PCOS (average Z-scores between -3 and -4), and to associate negatively with the degree of androgen excess (as judged by circulating testosterone or free androgen index, when the gonadotropic axis is not silenced), with HOMA-IR, and with hepatic and visceral fat; a normalizing rise of circulating miR-451a concentrations in adolescent girls with PCOS can thus point to a normalizing course toward metabolic health, including toward a normal ovulation rate [5]. In a search for a noninvasive, cycle-independent, on-treatment set of markers that allows to anticipate the posttreatment ovulation rate, we tested whether a "metabolic health Z-score" which combines the Z-scores of fasting insulinemia and circulating miR-451a, associated to posttreatment ovulation rate.

Body composition was assessed by dual x-ray absorptiometry with a Lunar Prodigy and Lunar software (version 3.4/3.5, Lunar Corp, Madison, Wisconsin); abdominal fat (subcutaneous and visceral) and hepatic fat were assessed by magnetic resonance imaging (MRI) using a multiple-slice MRI 1.5 Tesla scan (Signa LX Echo Speed Plus Excite, General Electric, Milwaukee, Wisconsin), as described [7, 8].

#### C. Statistics & Ethics

Statistical analyses were performed with SPSS 23.0 (IBM, Armonk, New York). Longitudinal changes in quantitative variables between groups were compared by repeated-measures

	-		Eth	inylestradic	l-Levonorg	estrel (N = ;	31)		SP	IOMET (N=	31)	
	Controls (N = 52)	PCOS (N = 62)	$\operatorname{Start}^{\operatorname{a}}$	12 mo	24 mo	Δ 0–12 mo	A 12–24 mo	${ m Start}^{ m a}$	12 mo	24 mo	Δ 0–12 mo	A 12–24 mo
Birthweight Z-score Age at Menarche	$0.2 \pm 0.1$ 12.4 ± 0.1	$-0.6 \pm 0.1^{***}$ 11.6 $\pm 0.1^{***}$	$-0.6 \pm 0.2$ 11.6 $\pm 0.1$	: :	: :			$-0.6 \pm 0.1$ 11.6 $\pm 0.2$	: :	: :	: :	
(yr) Age $(yr)$	$16.3 \pm 0.2$	$15.8 \pm 0.2$	$15.9 \pm 0.2$			0       		$15.7 \pm 0.2$			4 - - -	
BMI (kg/m²) RMI 7score	$21.3 \pm 0.3$ 0 0 + 0 1	$24.2 \pm 0.5$ 0 8 + 0 1 ***	$24.2 \pm 0.7$ 0 9 + 0 2	$24.9 \pm 0.8^{\circ}$ 1 1 + 0 $2^{\rm b}$	$25.1 \pm 0.8$ 1 2 + 0 2	$0.7 \pm 0.3$ 0.9 + 0.1	$0.2 \pm 0.3$ 0 1 + 0 3	$24.2 \pm 0.7$ 08+02	$23.9 \pm 0.7$ 0 7 + 0 2	$23.9 \pm 0.7$ 0 8 + 0 2	$-0.2 \pm 0.3^{\circ}$ -0.1 + 0.1 <sup>°</sup>	$0.0 \pm 0.2$ 0 1 + 0.3
ΔZ-score Birth-	$-0.2 \pm 0.2$	$1.4 \pm 0.2^{***}$	$1.5 \pm 0.3$	$1.7 \pm 0.3^{b}$	$1.8 \pm 0.3$	$0.2 \pm 0.1$	$0.1 \pm 0.1$	$1.4 \pm 0.3$	$1.4 \pm 0.3$	$1.4 \pm 0.3$	$0.0 \pm 0.1^{e}$	$0.0 \pm 0.1$
weight to BIMI Waist Circumfer-	$74 \pm 1$	$77 \pm 1$	$76 \pm 2$	$78 \pm 2^{\rm b}$	$78 \pm 2$	$2 \pm 1$	$0 \pm 1$	$77 \pm 2$	$74 \pm 1^{\rm d}$	$74 \pm 1$	$-3 \pm 0.8^{g}$	$0 \pm 1$
ence (cm) Hirsutism score		$\begin{array}{c} 17\pm1\\ 22\pm2.** \end{array}$	$17 \pm 1$	$14 \pm 1^{d}$	$14 \pm 1$	$-3 \pm 1$	$0 \pm 1$	$16 \pm 1$	$11 \pm 1^{d}$	$9\pm1^{\circ}$	$-5 \pm 1^{g}$	$-2 \pm 1$
SHBG (nmol/L) Testosterone	$63 \pm 3$ $0.7 \pm 0.1$	$30 \pm 2$ $1.4 \pm 0.1^{***}$	$31 \pm 2$ 1.3 $\pm 0.1$	$61 \pm 5^{\rm u}$ $0.7 \pm 0.1^{\rm d}$	$32 \pm 3^{a}$ 1.6 $\pm 0.2^{d}$	$30 \pm 4$ -0.6 \pm 0.1	$-29 \pm 5$ $0.9 \pm 0.2$	$30 \pm 2$ $1.5 \pm 0.2$	$32 \pm 2$ $0.8 \pm 0.1^{\circ}$	$39 \pm 3^{\circ}$ 1.2 ± 0.2°	$2 \pm 2^{5}$ -0.7 $\pm 0.2$	$7 \pm 2^{8}$ 0.4 ± 0.2
(nmol/L) Androstenedione	$3.5\pm0.2$	$5.3 \pm 0.3^{***}$	$4.8\pm0.3$	$2.5\pm0.2^{ m d}$	$5.7 \pm 0.6^{\mathrm{d}}$	$-2.3 \pm 0.3$	$3.2 \pm 0.5$	$5.7 \pm 0.4$	$3.5\pm0.3^{ m d}$	$5.3 \pm 0.6^{\circ}$	$-2.2 \pm 0.4$	$1.8 \pm 0.6$
(LINULU) Free Testosterone	$0.0 \pm 0.2$	$2.9 \pm 0.5^{***}$	$2.3 \pm 0.5$	$0.3 \pm 0.3^{\circ}$	$3.6\pm0.8^{\rm d}$	$-2.0 \pm 0.6$	$3.3 \pm 0.7$	$3.2 \pm 0.9$	$0.5\pm0.3^{\circ}$	$2.0\pm0.7^{\rm c}$	$-2.7 \pm 0.9$	$1.5 \pm 0.7$
Z-score Free Androstene-	$0.0 \pm 0.2$	$1.8 \pm 0.3^{***}$	$1.1 \pm 0.3$	$-0.9 \pm 0.2^{d}$	$2.2\pm0.5^{\mathrm{d}}$	$-2.0 \pm 0.3$	$3.1 \pm 0.5$	$2.2 \pm 0.4$	$0.1 \pm 0.3^{\rm d}$	$1.8\pm0.6^{\circ}$	$-2.1 \pm 0.4$	$1.7 \pm 0.6$
Tasting Insulin	$49 \pm 7$	$76 \pm 7^{***}$	$83 \pm 7$	$104 \pm 7^{\rm b}$	$76 \pm 7^{c}$	$21 \pm 7$	-28±7	$7 \pm 7$	$42 \pm 7^{\rm d}$	$49 \pm 7$	$-27 \pm 7^{g}$	$7 \pm 7^{\rm f}$
(pmol/l) HOMA-IR OGTT Mean Gly-	$1.5 \pm 0.1$	$2.3 \pm 0.2^{***}$ $0.2 \pm 0.1$	$2.6 \pm 0.3$ $0.1 \pm 0.1$	$3.0 \pm 0.3$ $0.2 \pm 0.1$	$2.2 \pm 0.2^{c}$ $0.1 \pm 0.1$	$0.4 \pm 0.2$ $0.1 \pm 0.1$	$-0.8 \pm 0.3$ $-0.1 \pm 0.1$	$2.1 \pm 0.2$ $0.2 \pm 0.1$	$1.2 \pm 0.1^{\rm d}$ $0.1 \pm 0.1^{\rm c}$	$1.3 \pm 0.2$ $0.1 \pm 0.1$	$-0.9 \pm 0.3^{\rm f}$ $-0.1 \pm 0.1^{\rm f}$	$0.1 \pm 0.2^{\rm f}$ $0.0 \pm 0.1$
cemia z-score Mean Insulinemia	:	$3.2 \pm 0.3$	$3.5 \pm 0.4$	$3.7 \pm 0.5$	$3.1 \pm 0.5$	$0.2 \pm 0.5$	$-0.6 \pm 0.5$	$2.8 \pm 0.4$	$0.6 \pm 0.2^{\mathrm{d}}$	$0.6 \pm 0.2$	$-2.2 \pm 0.3^{g}$	$0.0 \pm 0.2$
Δ-score ALT (μkat/L) AST (μkat/L) GGT (μkat/L)	$\begin{array}{c} 0.30 \pm 0.02 \\ 0.25 \pm 0.02 \\ 0.22 \pm 0.02 \end{array}$	$\begin{array}{c} 0.23 \pm 0.02^{***} \\ 0.27 \pm 0.02 \\ 0.22 \pm 0.02 \end{array}$	$0.23 \pm 0.02$ $0.27 \pm 0.02$ $0.22 \pm 0.02$	$0.32 \pm 0.03^{\circ}$ $0.27 \pm 0.02$ $0.30 \pm 0.02^{d}$	$\begin{array}{c} 0.27 \pm 0.02 \\ 0.27 \pm 0.02 \\ 0.25 \pm 0.02^{\mathrm{b}} \end{array}$	$\begin{array}{c} 0.09 \pm 0.02 \\ 0.00 \pm 0.02 \\ 0.08 \pm 0.02 \end{array}$	$-0.05 \pm 0.03$ $0.00 \pm 0.02$ $-0.05 \pm 0.02$	$0.23 \pm 0.02$ $0.28 \pm 0.02$ $0.22 \pm 0.02$	$\begin{array}{c} 0.23 \pm 0.02 \\ 0.27 \pm 0.02 \\ 0.18 \pm 0.02 \\ \circ \end{array}$	$\begin{array}{c} 0.23 \pm 0.02 \\ 0.27 \pm 0.02 \\ 0.22 \pm 0.02 \end{array}$	$-0.00 \pm 0.02^{\circ}$ $-0.01 \pm 0.02$ $-0.04 \pm 0.02^{\circ}$	$-0.00 \pm 0.02$ $0.00 \pm 0.02$ $0.04 \pm 0.02$
Triacylglycerol (mmol/L)	$0.60 \pm 0.03$	$0.68 \pm 0.03$	$0.66 \pm 0.03$	$0.75 \pm 0.05^{b}$	$0.64 \pm 0.03^{\circ}$	$0.09 \pm 0.03$	$-0.11 \pm 0.03$	$0.70 \pm 0.05$	$0.67 \pm 0.05$	$0.63 \pm 0.05$	$-0.03 \pm 0.05^{\circ}$	$-0.04 \pm 0.03$

Table 1. Continued

			Eth	inylestradic	ol-Levonorg	jestrel (N =	31)		$^{\mathrm{SP}}$	IOMET (N=	31)	
	(N = 52)	$\Gamma = 62$ )	$\operatorname{Start}^{\operatorname{a}}$	12 mo	24 mo	Δ 0–12 mo	A 12–24 mo	$Start^{a}$	12 mo	24 mo	Δ 0–12 mo	Δ 12–24 mo
LDL-cholesterol	$2.2 \pm 0.1$	$2.3 \pm 0.1$	$2.3 \pm 0.1$	$2.7 \pm 0.1^{d}$	$2.2 \pm 0.1^{\mathrm{d}}$	$0.4 \pm 0.1$	$-0.5 \pm 0.1$	$2.2 \pm 0.1$	$2.2 \pm 0.1$	$2.0 \pm 0.1^{d}$	$0.0 \pm 0.1^{\mathrm{f}}$	$-0.2 \pm 0.1^{e}$
HDL-cholesterol	$1.4 \pm 0.1$	$1.3 \pm 0.1$	$1.3 \pm 0.1$	$1.3 \pm 0.1$	$1.4 \pm 0.1$	$0.0 \pm 0.1$	$0.1 \pm 0.1$	$1.3 \pm 0.1$	$1.4 \pm 0.1^{\circ}$	$1.3\pm0.1^{\mathrm{b}}$	$0.1 \pm 0.1^{\mathrm{e}}$	$-0.1 \pm 0.1^{\mathrm{f}}$
(mmol/L) HMW-adiponectin	$9.3 \pm 0.8$	$6.8\pm0.6^*$	$6.5\pm0.6$	$8.9\pm1.3^{ m b}$	$8.6 \pm 0.8$	$2.6 \pm 1.1$	$-0.3 \pm 1.5$	$7.1 \pm 0.9$	$17.1 \pm 2.6^{\mathrm{d}}$	$10.3 \pm 1.5^{\circ}$	$10.0 \pm 2.1^{\mathrm{f}}$	$-7 \pm 2^{\mathrm{e}}$
(mg/L) C-Reactive Protein (nmol/L)	$6.7 \pm 0.9$	$14.3 \pm 1.9^{***}$	$11.4 \pm 1.9$	$24.8\pm3.8^{\circ}$	$18.1 \pm 3.8$	$13.4 \pm 3.8$	-6.7 ± 5.7	$17.1 \pm 3.8$	$6.7\pm0.9^{\circ}$	$6.7 \pm 0.9$	$-10.4 \pm 3.8^{g}$	$0.0 \pm 0.9$
Carotid IMT (mm) Systolic Rlood Pres-	 113+1	$.37 \pm .00$	$.37 \pm .01$ 113 + 2	$.37 \pm .01$	$.36 \pm .01^{b}$ 112 + 2	$00 \pm .00$	$01 \pm .01$ 3 + 2	$.37 \pm 0.01$ 116 + 1	$.35 \pm 0.01^{d}$ 112 + 1 <sup>b</sup>	$.35 \pm 0.01$	$02 \pm 0.01^{e}$ 4 + $2^{e}$	$.00 \pm .01$ 2 + 2
sure (mmHg) Diastolic Blood	70 ± 1	$72 \pm 1$	71 ± 1	$74 \pm 1^{b}$	$73 \pm 1$	00 1 1 1 1 1	- - - - - - - - - - - - - - - - - - -	$73 \pm 1$	71 ± 1	$70 \pm 1$	$-2 \pm 1^{\circ}$	-1+1
Pressure (mmHg) miR-451a Z-score	$0.00 \pm 0.28$	$-3.57 \pm 0.11^{***}$	$-3.75 \pm 0.12$	$-3.31 \pm 0.12$	$-3.59 \pm 0.16$	:	ł	$-3.32 \pm 0.19$	$0.37 \pm 0.31$	$-1.05 \pm 0.43^{\circ}$	ł	1
DXA BMD (g/cm <sup>2</sup> )	: :	$1.19 \pm 0.01$ 35 6 + 0 6	$1.18 \pm 0.02$ $35.7 \pm 0.8$	$1.19 \pm 0.02$ $36.4 \pm 0.0$	$1.20 \pm 0.02^{\rm b}$	$0.01 \pm 0.01$	$0.01 \pm 0.01$	$1.19 \pm 0.02$ $35 5 \pm 0.0$	$1.19 \pm 0.02$ $35.6 \pm 0.8$	$1.21 \pm 0.02^{\rm b}$ $36.1 \pm 0.8$	$0.00 \pm 0.01$	$0.02 \pm 0.01$
Fat Mass (Kg)	1	$22.1 \pm 1.0$	$21.8 \pm 1.4$	$23.2 \pm 1.5^{\circ}$	$23.4 \pm 1.6$	$1.4 \pm 0.5$	$0.2 \pm 0.6$	$22.4 \pm 1.6$	$22.5 \pm 1.4$	$22.1 \pm 1.7$	$0.1 \pm 0.8$	$-0.4 \pm 0.6$
Abd MRI Subc Fat	$94 \pm 9$	$174 \pm 14^{***}$	$169 \pm 18$	$184 \pm 19$	$180 \pm 20$	$15 \pm 9$	$-4 \pm 13$	$179 \pm 21$	$171 \pm 19$	$167 \pm 23$	-8±11	$-4 \pm 9$
(cm <sup>2</sup> ) Visceral Fat (cm <sup>2</sup> ) Liver Fat (%)	$28 \pm 1$ $10 \pm 1$	$43 \pm 2^{***}$ $17 \pm 1^{***}$	$41 \pm 3$ $17 \pm 1$	$45 \pm 4$ $19 \pm 1$	$39 \pm 3$ $17 \pm 1^{b}$	$\begin{array}{c} 4\pm3\\ 2\pm1\end{array}$	-6 ± 3 -2 ± 2	$44 \pm 3$ $18 \pm 1$	$35 \pm 2^{\mathrm{b}}$ $10 \pm 1^{\mathrm{d}}$	$36 \pm 3$ $10 \pm 1$	$-9 \pm 4^{\mathrm{f}}$ -8 $\pm 1^{\mathrm{g}}$	$\begin{array}{c} 1\pm 2\\ 0\pm 1\end{array}$
Values are mean $\pm$ Abbreviations: Abd molecular-weight a tolerance test; SHB "miR-451a (controls "on significant diffe $P < 0.05, {}^{e}P \le 0.01$ $P < 0.05, {}^{*P} \le 0.01$	SEM. MRI, abdomi diponectin; H G, sex hormo G, sex hormo i, $n = 13$ ; OC i rences betwee and <sup>d</sup> $P \le 0.001$ be " $P \le 0.001$ be	nal magnetic r (OMA-IR, hom ne-binding glo at start, n = 12 en randomized 1 within subgrou tween subgrou 001 between a	esonance ims teostasis moc bulin; ; SPIOMET s subgroups a oups for 0-12 n tps for 0-12 n the for st st	uging; BMD, h lel assessmer at start, n = 9 t start ? mo & 12-24 ch no & 12-24 ch cart and conti	one mineral it - insulin r ; OC at 12 m mo change (ι ange (Δ) col group	density; BM esistance; IN io, n = 25; SF	I, body mass AT, intima-m TOMET at 1:	index; DXA, $c$ edia thickne 2 mo, n = 24;	dual x-ray ab ss; miR-4518 OC at 24 mo	sorptiometry , microRNA- , n = 15; SPIG	; HMW adipo 451a; OGTT, OMET at 24 n	aectin, high- oral glucose 10, n = 16)

general linear model. Differences in longitudinal changes between groups were tested by the interaction term among between- and within-subject effects. P < 0.05 was considered significant. Data are presented as mean ± standard error of the mean (SEM).

The studies were conducted after approval by the Institutional Review Board of Sant Joan de Déu Hospital, after written informed consent by the parents, and after assent by each participating girl.

#### 2. Results

Table 1 summarizes the pooled results, which indicate that SPIOMET treatment was accompanied by more broadly normalizing effects than OC, including for waist circumference, circulating insulin, HMW-adiponectin and CRP, cIMT, as well as on visceral and hepatic fat (Fig. 2).

Table 2 shows that there were a mean 3-fold and a median 5-fold more ovulations after SPIOMET than OC; normovulation (as judged by 5 or 6 ovulations over 24 weeks) was only observed after SPIOMET; anovulation (as judged by 0 or 1 ovulation over 24 weeks) was > 10-fold more frequent after OC. Menstrual regularity after SPIOMET (90%) was only 2-fold more prevalent than after OC (42%), thus underestimated the difference in ovulation rates.

Fig. 3 illustrates that the randomized treatments led to marked differences in on-treatment metabolic health (as judged by combined Z-scores of fasting insulin and miR-451a) and in posttreatment ovulation rate, both of which were more normalized after SPIOMET.

#### 3. Discussion

Pooled data corroborated SPIOMET as a combination treatment that is accompanied by more normalization of the endocrine-metabolic status, and is followed by markedly more ovulations than OC in nonobese adolescent girls with PCOS. The consistency of the ovulation rates across the posttreatment year suggests that the lower ovulation rates after OC are attributable to persistence of the underpinning PCOS pathophysiology rather than to residual inhibition of the gonadotropic axis. In healthy young women, ovulatory function is known to recover within 3 months after stopping OC treatment [10, 11].

Ectopic adiposity and insulin resistance failed to improve during standard treatment with OC. In contrast, SPIOMET treatment was accompanied by a loss of hepato-visceral fat excess and by a normalization of insulin sensitivity (as judged by HOMA-IR, and by the insulin response to an oral glucose load), both of which were maintained during the



**Figure 2.** Hepatic fat content (by magnetic resonance imaging) in nonobese adolescent girls with PCOS who were randomized to receive either an oral contraceptive (OC; N = 31; red circles) for 12 months, or a low-dose combination of spironolactone-pioglitazone-metformin (SPIOMET; N = 31; blue circles) for 12 months; subsequently, both subgroups were untreated for 12 months. Body weight did not change in either subgroup. The dotted line indicates the average level in healthy control girls of similar age. Results are expressed as mean  $\pm$  SEM. P < 0.0001 for on-treatment change between subgroups.

Table 2. Posttreatment Ovulation Results in Adolescent Girls With Polycystic Ovary Syndrome Who Were Randomized to Receive an Oral Contraceptive (OC) or Low-Dose Spironolactone + Pioglitazone + Metformin (SPIOMET) for 12 Months, and Were Subsequently Followed for 12 Months Without Treatment. Ovulations Were Assessed Twice Over 12 Weeks, for a Total of 24 Weeks: Between the Study Timepoints of 15 to 18 months (posttreatment months 3-6) and 21 to 24 months (posttreatment months 9-12)

		OC N = 31		S	SPIOMET N = 31	
	15-18 mo (12 wk)	21-24 mo (12 wk)	Total (24 wk)	15-18 mo (12 wk)	21-24 mo (12 wk)	Total (24 wk)
Mean number of ovulations ± SEM	$0.8 \pm 0.1$	$0.8 \pm 0.1$	$1.6 \pm 0.2$	$2.3 \pm 0.2^{\#}$	$2.2 \pm 0.2^{\#}$	$4.5 \pm 0.3^{\#}$
Median number of ovulations (interquartile range)	1 (0-1)	1 (0-1)	1 (1-3)	$3(2-3)^{\#}$	$2(2-3)^{\#}$	$5(3-6)^{\#}$
Normo-ovulatory fraction (%) 5 or 6 ovulations /24 wk			0			62 <sup>#</sup>
Oligo-ovulatory fraction (%) 2. 3. or 4 ovulations /24 wk			47			35
An-ovulatory fraction (%) 0 or 1 ovulation /24 wk			53			3#

 $^{\#}P < 0.0001$  between subgroups



**Figure 3.** Randomized treatment of adolescent girls with PCOS, either with an oral contraceptive (OC) or with a low-dose combination of spironolactone-pioglitazone-metformin (SPIOMET) for 12 months, results in an on-treatment difference of metabolic health (N = 22 *vs* 24) and in a posttreatment difference of ovulation rate (N = 30 *vs* 29), so that this combined metabolic-reproductive outcome is markedly to the advantage of SPIOMET. Metabolic health Z-score was calculated by subtracting the Z-score of fasting insulin from the Z-score of circulating miR-451a after 12 months on treatment. Posttreatment number of ovulations over 6 months was inferred by combining data from menstrual diaries and weekly progesterone measurements in saliva over 12 + 12 weeks, between posttreatment months 3 to 6 and 9 to 12. Body weight did not change in either subgroup. The breadth and height of the boxes represent the ranges from -1 SD to +1 SD, respectively, for metabolic health Z-score and ovulation number. \*\*\* P < 0.0001.

posttreatment year, via mechanisms that remain to be identified. The downward normalization of liver fat on SPIOMET may partly relate to the upward normalization of circulating miR-451a, which reduces the expression of thyroid hormone responsive spot 14 (*THRSP*), the key gene driving liver steatosis [12, 13].

The present findings corroborate the concept that insulin resistance reflects ectopic lipid accumulation, particularly in the liver, and that it precedes the development of disorders such as type 2 diabetes and nonalcoholic fatty liver disease [14]. Increased hepatic fat and insulin resistance are prevalent findings in both nonobese and obese adolescents with PCOS,

and seem to relate to the underpinning PCOS pathophysiology rather than to testosterone concentrations [15, 16]. Targeting a reduction in androgen levels may thus not be the best choice to normalize the entire PCOS phenotype and to address subsequent comorbidities. The diverging effects of OC and SPIOMET on insulin resistance and ectopic fat (Fig. 3) may herald diverging influences on subsequent risk for PCOS-associated disorders such as an-ovulatory subfertility, gestational diabetes, and/or type 2 diabetes.

The present results remain to be further confirmed in larger and more diverse PCOS populations, including in girls with obesity, with different ethnic and developmental backgrounds, and with other environmental exposures. In addition, SPIOMET's capacity to reduce an excess of liver fat while total body weight remains virtually unchanged (Fig. 2), remains to be tested beyond PCOS settings, in older age ranges, and in a cascade of fatty liver diseases, including nonalcoholic steatohepatitis.

In conclusion, pooled results in nonobese adolescent girls with PCOS confirmed SPIOMET as a treatment that attenuates insulin resistance, reduces ectopic adiposity, and is followed by a more normal ovulation rate than OC.

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Author Contributions: L.I. contributed to study design and data interpretation, wrote and reviewed/edited manuscript. M.D. researched data, contributed to data interpretation, and reviewed/edited manuscript. C.G.B. researched data, contributed to data interpretation, and reviewed/edited manuscript. R.M. contributed to data interpretation, wrote the manuscript, and reviewed/edited manuscript. E.G. researched data, and reviewed/edited manuscript. A.L.B. reviewed/edited manuscript. F.d.Z. contributed to study design and data interpretation, wrote and reviewed/edited manuscript.

L.I. had full access to all of the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis.

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**Disclosure Summary:** The authors have nothing to disclose.

**Data Availability:** The datasets generated during and/or analyzed during the current study are not publicly available but are available from the corresponding author on reasonable request.

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**STUDY 2** 

## REDUCED CIRCULATING LEVELS OF CHEMOKINE CXCL14 IN ADOLCESCENT GIRLS WITH POLYCYSTIC OVARY SYNDROME: NORMALIZATION AFTER INSULIN SENSITIZATION

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#### STUDY 2

# Reduced circulating levels of chemokine CXCL14 in adolescent girls with polycystic ovary syndrome: normalization after insulin sensitization

**Introduction:** CXCL14 (C-X-C motif chemokine ligand-14) is a chemokine released by active brown fat, showing protective effects against insulin resistance in experimental models. Polycystic ovary syndrome (PCOS) in adolescent girls is usually related to hepato-visceral fat excess and insulin resistance, and associates with comorbidities such as type 2 diabetes. Treatment with a low-dose combination of one antiandrogen and anti-mineralocorticoid drug (spironolactone) and two insulin sensitizers (pioglitazone/metformin) (SPIOMET) is particularly effective in improving these metabolic derangements. Adipose tissue may be involved in the metabolic alterations of PCOS, and it is a likely target of therapeutic action.

**Objective:** To investigate the alterations in CXCL14 levels in girls with PCOS, the effects of SPIOMET and OC on circulating CXCL14 in relation to metabolic improvement and the effects of the drugs composing SPIOMET treatment on CXCL14 in human adipocytes.

**Material and methods:** We studied 51 adolescents with PCOS and 21 age-matched healthy controls. Thirty-one adolescent patients with PCOS under SPIOMET or oral contraception-based treatment were also studied. For studies *in vitro*, Simpson Golabi Behmel Syndrome (SGBS) adipose cells were used. Gene expression for CXCL14 and other genes was quantified using quantitative real-time PCR. The levels of CXCL14 and adipokines in serum and cell culture media were determined by ELISA.

**Results:** Serum CXCL14 levels are reduced in patients with PCOS. One-year SPIOMET treatment normalized CXCL14 concentrations and improved the metabolic status of patients with PCOS. Pioglitazone induced CXCL14 expression in differentiating human SGBS adipocytes, in parallel with the induction of marker genes of brown adipogenesis. Spironolactone induced CXCL14 expression and release in differentiated human adipocytes.

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**Conclusion:** Insulin sensitization with SPIOMET normalizes the abnormally low levels of CXCL14 in girls with PCOS. This is consistent with the effects of pioglitazone and spironolactone inducing CXCL14 expression and promoting a brown-like phenotype in adipocytes. CXCL14 may be a novel biomarker for PCOS as well as a potential mediator of the beneficial effects of the SPIOMET combination and may hold promise as a therapeutic modulator of the disorder.

### **BMJ Open Diabetes** Research & Care

## **Reduced circulating levels of** chemokine CXCL14 in adolescent girls with polycystic ovary syndrome: normalization after insulin sensitization

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#### ABSTRACT

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Objective CXCL14 (C-X-C motif chemokine ligand-14) is a chemokine released by active brown fat, showing protective effects against insulin resistance in experimental models. Polycystic ovary syndrome (PCOS) in adolescent girls is usually related to hepato-visceral fat excess and insulin resistance, and associates with comorbidities such as type 2 diabetes. Treatment with a low-dose combination of one antiandrogen and antimineralocorticoid drug (spironolactone) and two insulin sensitizers (pioglitazone/ metformin) (SPIOMET) is particularly effective in improving these metabolic derangements. Adipose tissue may be involved in the metabolic alterations of PCOS, and it is a likely target of therapeutic action. We investigated the alterations in CXCL14 levels and the effects of drugs composing SPIOMET treatment on CXCL14 in human adipocytes.

Research design and methods We studied 51 adolescent patients with PCOS and 21 age-matched healthy controls. Thirty-one adolescent patients with PCOS under SPIOMET or oral contraception-based treatment were also studied. For studies in vitro, Simpson Golabi Behmel Syndrome (SGBS) adipose cells were used. Gene expression for CXCL14 and other genes was quantified using quantitative real-time PCR. The levels of CXCL14 and adipokines in serum and cell culture media were determined by ELISA.

**Results** Serum CXCL14 levels are reduced in patients with PCOS. One-year SPIOMET treatment normalized CXCL14 concentrations and improved the metabolic status of patients with PCOS. Pioglitazone induced CXCL14 expression in differentiating human SGBS adipocytes, in parallel with the induction of marker genes of brown adipogenesis. Spironolactone induced CXCL14 expression and release in differentiated human adipocytes. **Conclusion** Insulin sensitization with SPIOMET normalizes the abnormally low levels of CXCL14 in girls with PCOS. This is consistent with the effects of pioglitazone and spironolactone inducing CXCL14 expression and promoting a brown-like phenotype in adipocytes. CXCL14 may be a novel biomarker for PCOS as well as a potential mediator of the beneficial effects of the SPIOMET combination and may hold promise as a therapeutic modulator of the disorder. Trial registration numbers ISRCTN29234515 and ISCRCTN11062950.

#### Significance of this study

#### What is already known about this subject?

- Treatment with a low-dose combination of one mixed antiandrogen and antimineralocorticoid (spironolactone), and two insulin sensitizers (pioglitazone/metformin) (SPIOMET) is particularly effective in improving the endocrine-metabolic derangements in adolescent girls with polycystic ovary syndrome (PCOS).
- CXCL14 (C-X-C motif chemokine ligand-14) is a chemokine released by active brown fat and protective against insulin resistance in experimental models.

#### What are the new findings?

- Serum CXCL14 levels are abnormally reduced in patients with PCOS.
- SPIOMET treatment for 1 year normalized CXCL14 concentrations and improved the endocrine-metabolic status of patients with PCOS.
- Pioglitazone and spironolactone, drug components of SPIOMET, induce CXCL14 expression and release in human adipocytes, in parallel with the induction of marker genes of brown adipogenesis.

#### How might these results change the focus of research or clinical practice?

CXCL14 may become a novel biomarker for PCOS. In addition, CXCL14 appears as a potential mediator of the beneficial effects of the SPIOMET combination and may hold the capacity of serving as therapeutic modulator of the disorder.

#### INTRODUCTION

Adipose tissue plasticity is growingly recognized as a relevant factor for the development of metabolic syndrome, independently of the presence or absence of obesity. The acquisition of a 'brown' or 'beige' phenotype by adipose tissue is considered protective against hyperglycemia and hyperlipidemia and, indeed, the relative protection against these alterations

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in young versus elder individuals is associated with the known prevalence of the brown/beige phenotype in early human development.<sup>1</sup> CXCL14 (C-X-C motif chemokine ligand-14) is a chemokine produced by active brown/beige adipose tissue, capable of improving glucose metabolism in insulin-resistant rodent models.<sup>2</sup>

Central (hepato-visceral) fat excess and insulin resistance are common metabolic comorbidities in girls and young women with polycystic ovary syndrome (PCOS).<sup>3 4</sup> Treatment with a low-dose combination of one mixed antiandrogen and antimineralocorticoid (spironolactone) and two insulin sensitizers (pioglitazone plus metformin) (SPIOMET) has been shown to improve the metabolic condition of these patients to a better extent than oral contraception (OC).<sup>5</sup> However, the specific cellular and tissue targeting, and the relative role of each of the SPIOMET components in the context of PCOS is poorly understood. It has been recently reported that women with PCOS have lower brown adipose tissue (BAT) activity as compared with healthy controls,<sup>67</sup> and experimental data suggest that BAT activation might be a promising therapeutic option for PCOS.<sup>8</sup>

Here, we report abnormally low levels of CXCL14 in girls with PCOS, the differential effects of SPIOMET and OC on circulating CXCL14 in relation to metabolic improvement, and the effects of SPIOMET components on the expression and release of CXCL14 in a cellular model of human adipocytes.

#### **RESEARCH DESIGN AND METHODS** Study population and design

The study population consisted of 52 adolescent girls with PCOS (age, 15.6 years; body mass index (BMI),  $24.3 \text{ kg/m}^2$ ) who were enrolled into two randomized, open-label, controlled trials (with identical design) exploring the effects of OC versus SPIOMET treatment for 1 year, with post-treatment ovulation rate as primary outcome. The results of the first trial (ISRCTN29234515) have been already published,<sup>5</sup> and the second trial (ISCRCTN11062950) will be completed in 2019. Twenty-one out of the 52 girls belonged to the first study and 31 to the second trial; 31 of the 52 had available samples for CXCL14 assessment at 0 and 12 months (online supplementary figure 1). Both trials were performed at Sant Joan de Déu University Hospital, Barcelona, Spain. Inclusion and exclusion criteria have been described previously.<sup>5</sup> OC treatment consisted of 20µg of ethinylestradiol plus 100 mg of levonorgestrel for 21/28 days and placebo for 7/28 days; SPIOMET is a low-dose combination of spironolactone 50 mg/day, pioglitazone 7.5 mg/day, and metformin 850 mg/day. Twenty-one agematched healthy girls recruited in nearby schools served as controls. All had regular menstrual cycles and none was hirsute or taking medications.

#### Clinical, endocrine-metabolic, and imaging assessments

Birth weight and BMI (and their Z-scores) were retrieved, and endocrine-metabolic variables were assessed in the early morning, in the follicular phase (days 3–7) of the cycle, or after 2 months of amenorrhea.<sup>5 9</sup> Serum glucose, insulin, homeostasis model assessment-insulin resistance (HOMA-IR), lipids, ultrasensitive C reactive protein (usCRP), sex hormone-binding globulin, androgens, and high-molecular-weight (HMW) adiponectin were assessed as reported.<sup>5</sup> CXCL14 levels in serum and cell culture medium were determined using a specific ELISA kit (RayBiotech),<sup>2</sup> whose sensitivity was 0.7 ng/mL, and interassay and intra-assay coefficients of variation less than 12%.

Body composition was assessed by dual X-ray absorptiometry with a Lunar Prodigy and Lunar software (version 3.4/3.5; Lunar, Madison, Wisconsin, USA); abdominal (subcutaneous and visceral) and hepatic fat were assessed by MRI using a multiple-slice MRI 1.5 Tesla scan (Signa LX Echo Speed Plus Excite; General Electric, Milwaukee, Wisconsin, USA).<sup>5</sup> Central fat was arbitrarily defined as the sum of visceral fat (in squared centimeter) and hepatic fat (in per cent).

#### Studies in human adipocytes in culture

The effects of pioglitazone, spironolactone, and metformin on adipocytes were studied in human Simpson Golabi Behmel Syndrome (SGBS) cells, a cell model of human beige (brown-like) adipogenesis.<sup>1011</sup> SGBS preadipocytes were maintained in Dulbecco's modified Eagle's (DMEM)/F12 medium, 10% FBS. Adipogenic differentiation was initiated by incubating confluent cell cultures for 4 days in serum-free medium plus 20nM insulin, 0.2 nM triiodothyronine, 100 nM cortisol, 25 nM dexamethasone, 500 µM 3-isobutyl-1-methyl-xanthine, and 2µM rosiglitazone. Subsequently, the cells were switched to DMEM/F12, 20 nM insulin, 0.2 nM triiodothyronine, and 100 nM cortisol and maintained for up to 10 days, when more than 90% cells have acquired differentiated adipocyte morphology. Two experimental designs were followed: (1) cells were treated with the medications across their differentiation process, in the absence of rosiglitazone and maintaining the drugs throughout the 10 days of differentiation and (2) adipocytes were treated with the drugs acutely (24 hours), once the cells have been differentiated. Controls included a 1:1 methanol/ dimethyl sulfoxide mixture (drug solvents) at  $\leq 1/1000$ concentration. Cell culture reagents and drugs were from Sigma. The cell culture medium was collected and concentrated 1:5 prior to measurement of CXCL14 levels. RNA was extracted from cells using an affinity columnbased method (Machery-Nagel). Real-time quantitative reverse transcription PCRs were performed using 0.5 µg RNA and employing TaqMan reagents and probes (Life Technologies), according to supplier indications. PCR was conducted in an ABI/Prism-7700 Sequence Detector System. The following TaqMan probes were used: CXCL14, Hs01557413; uncoupling protein-1 (UCP1), Hs00222453; fatty acid-binding protein-4 (FABP4), Hs00609791, and ribosomal protein lateral stalk subunit 0 (RPLP0) mRNA, Hs99999902. Each sample was run in duplicate and the mean value was used to calculate the

relative amount of individual mRNAs. Each mean value was normalized to that of the RPLP0 mRNA using the comparative  $(2-\Delta CT)$  method.

#### Statistical analysis and ethics

Statistical analyses were performed with SPSS V.23.0 (SPSS, Chicago, Illinois, USA). Baseline differences in CXCL14 concentrations between patients and controls were tested with unpaired t-test; covariance analysis was used to adjust for age and BMI. Longitudinal changes between groups were compared by repeated-measures general linear model. Differences in longitudinal changes between groups were tested by the interaction term among between- and within-subject effects. Associations were sought by Pearson correlation analysis. P value <0.05 was considered statistically significant. Results are expressed as mean±SD.

The study was conducted after approval by the Institutional Review Board of Sant Joan de Déu University Hospital, after written consent by the parents and assent by each of the participants.

#### RESULTS

#### Serum CXCL14 levels are reduced in adolescent girls with **PCOS**

At baseline, PCOS girls showed lower sex hormone-binding protein (SHGB) levels, higher total testosterone concentrations and free androgen index, increased central and hepatic fat, and a trend toward higher insulin and lower HMW adiponectin levels versus controls, as expected (online supplementary table 1). None of the girls were obese (BMI  $\geq$  30 kg/m<sup>2</sup>); however, mean BMI was higher in girls with PCOS as compared with control girls (p=0.01). Fasting glucose levels were within the normal range in both subgroups and were marginally higher in control girls (online supplementary table 1). Serum CXCL14 concentrations were reduced in girls with PCOS close to twothirds of the normal levels (figure 1A), did not correlate with markers of adiposity or glucose homeostasis, and were positively associated with HMW adiponectin levels (online supplementary table 2).

Serum CXCL14 concentrations were reduced in girls with PCOS close to two-thirds of the normal levels (figure 1A) and were positively associated with HMW adiponectin levels (online supplementary table 2). CXCL14 did not correlate with other markers of adiposity or glucose homeostasis; however, there was a trend toward a positive correlation with circulating glucose and a negative correlation with hepatic fat (online supplementary table 2).

#### SPIOMET, but not OC, normalizes serum CXCL14 levels in adolescent girls with PCOS

SPIOMET treatment was associated with more benefits than OC on endocrine-metabolic and imaging markers, including on fasting insulin, HOMA-IR, HMW adiponectin, usCRP, and hepato-visceral (central) fat, consistent with previous observations<sup>5</sup> (online supplementary table 3). Serum CXCL14 levels increased significantly after SPIOMET, reaching levels similar to those in



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Figure 1 (A) Baseline circulating CXCL14 (C-X-C motif chemokine ligand-14) concentrations in girls with polycystic ovary syndrome (PCOS, n=52) and in healthy age-matched controls (n=21). Data are mean±SD. P=0.007, for patients versus controls at baseline. (B) Longitudinal results of CXCL14 serum concentrations in girls with PCOS who received an oral contraceptive (OC, white circles, n=16) or low-dose spironolactone, pioglitazone, and metformin (SPIOMET, black circles, n= 15). The dotted line is the mean in healthy controls (n=21); the shaded area represents the mean±SD in those healthy controls. \*P=0.004, for patients versus controls at baseline; §p=0.02, on treatment differences between controls and the OC subgroup; #p=0.01, for changes 0-1 year in the SPIOMET subgroup; &p=0.04 for changes 0-1 year between the OC subgroup and the SPIOMET subgroup.

controls and remained significantly decreased after OC, as compared with controls and with SPIOMET-treated girls (figure 1B). The increase in CXCL14 levels in the SPIOMET-treated patients correlated significantly with the extent of reduction in fasting insulin and HOMA-IR (online supplementary table 4).

#### Effects of the SPIOMET components on CXCL14 expression and release by human adipocytes

The effects of the three components of SPIOMET on human adipocyte differentiation were assessed individually and in combination (figure 2). Pioglitazone had a doseresponse dramatic effect inducing CXCL14 mRNA expression, whereas neither spironolactone nor metformin did. No concentration higher than 10µM could be tested for spironolactone, because of cell toxicity on preadipocytes. Indeed, pioglitazone at the lowest concentration tested (0.1µM) already caused a close to 10-fold induction of CXCL14 mRNA expression (figure 2A). This effect paralleled a strong positive effect of pioglitazone on adipogenic differentiation, as evidenced by lipid droplet accumulation (figure 2C) and expression of gene markers of general adipogenic (FABP4) and beige adipogenic (UCP1) differentiation (figure 2D).

Addition of metformin to pioglitazone did not modify the extent of pioglitazone induction of CXCL14 mRNA expression (figure 2B). However, the addition of spironolactone reduced significantly the effects of pioglitazone on CXCL14 mRNA expression. The induction of CXCL14 mRNA expression caused by the three drugs in combination was similar to that elicited by pioglitazone plus spironolactone. These data also paralleled the observations on adipogenic differentiation after combined treatments, according to cell morphology and expression of FABP4



Figure 2 Effects of pioglitazone (Pio), spironolactone (Spi), and metformin (Met) on adipogenic differentiation of SGBS human preadipocytes in culture. SGBS human preadipocytes were treated chronically (10 days) with pioglitazone, spironolactone, or metformin alone (A) or in combination (B, C, D) at the indicated doses across the adipogenic differentiation process. (A, B) CXCL14 (C-X-C motif chemokine ligand-14) transcript levels are presented as means±SD from four to five independent experiments and are expressed relative to values from untreated control cells. (C) Representative photomicrographs of adipocyte cell cultures differentiating in the presence of the indicated components: top: control; next to top: pioglitazone (10 µM); middle: pioglitazone (10 µM) and metformin (10 µM); next to bottom: spironolactone (10 µM); bottom: pioglitazone (10  $\mu$ M), spironolactone (10  $\mu$ M), and metformin (10  $\mu$ M). Each column represents duplicate representative pictures per treatment condition. (D) Fatty acid-binding protein-4 (FABP4) and uncoupling protein-1 (UCP1) transcript levels are presented as means±SD from three to four independent experiments and are expressed relative to values from untreated control cells. \*P<0.05, \*\*p<0.01, and \*\*\*p<0.001 for each component versus control; <sup>#</sup>p<0.05 relative to pioglitazone alone. ND, not detected; SGBS, Simpson Golabi Behmel Syndrome.

and UCP1 (figure 2C,D). CXCL14 protein was only detectable in the cell culture medium of differentiating cells under pioglitazone treatment, either alone (0.185 ng/mL) or in combination with spironolactone and metformin (0.179 ng/mL).

Neither pioglitazone nor metformin alone had any effect on CXCL14 mRNA expression on already differentiated human adipocytes (figure 3A). However,



**Figure 3** Effects of pioglitazone (Pio), spironolactone (Spi), and metformin (Met) on *CXCL14* (C-X-C motif chemokine ligand-14) gene expression and CXCL14 protein release in human adipocytes. SGBS cells were treated acutely (24 hours, at the indicated doses) when already differentiated adipocytes. CXCL14 transcript level (A) and CXCL14 protein levels in cell culture medium (B) are presented as means±SD from four to five independent experiments and are expressed relative to values from untreated control cells. \*P<0.05, \*\*p<0.01, and \*\*\*p<0.001 for each component versus control. SGBS, Simpson Golabi Behmel Syndrome.

 $10\,\mu\text{M}$  of spironolactone caused a significant induction of CXCL14 mRNA expression (figure 3A). The combination of three components (with spironolactone dosed at  $10\,\mu\text{M}$ ) induced CXCL14 mRNA to the same extent as treatment with spironolactone alone. Only spironolactone, either alone or in combination with pioglitazone plus metformin, elicited a significant increase (close to eightfold) of the CXCL14 protein levels released by adipocytes to the cell culture medium (figure 3B).

#### DISCUSSION

Our study shows that PCOS associates with reduced levels of CXCL14, a chemokine recently proposed to be secreted preferentially by brown/beige adipose tissue.<sup>2</sup> The restoration of normal levels of CXCL14 with SPIOMET, the capacity of pioglitazone to increase CXCL14 expression in preadipocytes, and the capacity of spironolactone to increase the release of CXCL14 in adipocytes, respectively, suggest that SPIOMET treatment could target adipose tissue to improve the metabolic profile of patients with PCOS. In addition, the normalization of central fat after SPIOMET treatment may indicate that circulating CXCL14 relates to ectopic fat rather than to BMI per se.

High CXCL14 levels have been shown to improve insulin sensitivity in adipocytes in vitro<sup>12</sup> and in rodent models<sup>2</sup>; however, not all reports agree in the antidiabetic actions of CXCL14, and even deleterious prodiabetogenic effects have been proposed in vitro<sup>13</sup> and in vivo.<sup>14</sup> Our study, which is, to our knowledge, the first exploring CXCL14 in human metabolic pathology, indicates that normalization of CXCL14 after SPIOMET associates with decreased insulin resistance. Pioglitazone increases insulin sensitivity as well as adipogenic differentiation and also promotes the

acquisition of a brown/beige phenotype in adipose cells.<sup>15</sup> The induction of CXCL14 expression by pioglitazone is consistent with data in experimental models highlighting increased expression of CXCL14 with brown/beige adipogenic differentiation.<sup>2</sup> The positive effects of spironolactone-inducing CXCL14 in differentiated adipocytes are also consistent with previous reports indicating that spironolactone induces adipose tissue browning in rodents<sup>16</sup> and activates brown fat in humans.<sup>17</sup> Recent reports disclosed impaired BAT activity in PCOS subjects<sup>6</sup><sup>7</sup> and the capacity of brown fat activation to ameliorate PCOS in experimental models.<sup>8</sup> Considering that CXCL14 is preferentially released by brown adipocytes,<sup>2</sup> it may be speculated that SPIOMET treatment drives a shift in adipose tissue plasticity to a more brown/beige phenotype resulting in enhanced CXCL14 release, potentially improving glucose homeostasis and possibly reducing diabetes risk. This sequence may be especially relevant in young girls, where the brown/beige adipose tissue amount is particularly significant.<sup>18</sup>

Our study has obvious limitations. First, the effect of SPIOMET on CXCL14 levels in tissues other than adipose tissue cannot be ruled out. Second, the physiological significance of the drug concentrations used in the in vitro studies is unclear. The concentrations of pioglitazone increasing CXCL14 levels in differentiated adipocytes (from at least  $0.1\,\mu\text{M}$ ) were in the range of those in plasma from SPIOMET-treated healthy women.<sup>19</sup> However, although spironolactone concentrations used here were the same as those employed in previous studies in adipose cells,<sup>16 20</sup> plasma spironolactone levels in SPIOMET-treated controls are lower.<sup>19</sup> Finally, evidence for a direct role of CXCL14 downregulation and reinduction on the metabolic status of patients with PCOS after SPIOMET is obviously lacking, as it would imply intervention experiments beyond the ethical standards of human studies. Nevertheless, our data support a role for CXCL14 as a novel potential biomarker and molecular mediator in the improvement of PCOS-associated metabolic alterations following SPIOMET intervention.

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# **Supplemental Figure 1**



\* Study started in January 2013 and completed in May 2016 § Study started in December 2015, recruitment completed in September 2017

	Controls         PCOS           (N= 21)         (N= 52)		P value
Auxology	· · · ·	· · · · · ·	
Age (years)	$16.0 \pm 1.4$	$15.6 \pm 1.3$	-
Birth weight Z-score	$0.2 \pm 0.8$	$-0.6 \pm 1.0$	0.03
BMI (kg/m <sup>2</sup> )	$21.5 \pm 2.8$	$24.3 \pm 4.0$	0.003
BMI Z-score	$0.1 \pm 1.0$	$0.9 \pm 1.2$	0.02
$\Delta$ Z-score birth weight – BMI	$-0.2 \pm 1.1$	$1.4 \pm 1.5$	0.02
Endocrine-Metabolic variables			
Testosterone (nmol/ L)	$1.0 \pm 0.4$	$1.3 \pm 0.4$	0.04
SHBG (nmol/ L) <sup>†</sup>	61.1 ± 26.5	$30.9 \pm 12.5$	<0.0001
FAI <sup>†</sup>	$1.8 \pm 0.9$	$4.8 \pm 2.5$	0.0004
Glucose (mmol/ L)	$4.9 \pm 0.4$	$4.6 \pm 0.4$	0.004
Fasting insulin (pmol/ L)	$50.2 \pm 28.1$	$74.5 \pm 42.4$	NS
HOMA-IR	$1.6 \pm 0.9$	$2.3 \pm 1.4$	NS
HDL-cholesterol (nmol/ L)	$1.4 \pm 0.2$	$1.3 \pm 0.2$	NS
LDL-cholesterol (nmol/ L)	$2.2 \pm 0.5$	$2.3 \pm 0.5$	NS
Triglycerides (nmol/ L)	$0.6 \pm 0.1$	$0.7 \pm 0.3$	NS
HMW adiponectin (mg/ L)	9.1 ± 4.9	$6.3 \pm 4.7$	NS
usCRP (mg/ L) <sup>†</sup>	$0.6 \pm 0.5$	$1.5 \pm 1.8$	NS
Body composition (DXA) <sup>‡</sup>			
Bone mineral density (g/ cm <sup>2</sup> )	-	$1.18 \pm 0.10$	-
Lean mass (kg)	-	$35.9 \pm 4.9$	-
Fat mass (kg)	-	$22.9 \pm 8.6$	-
Abdominal fat (kg)	-	$6.1 \pm 2.1$	-
Abdominal fat partitioning (MRI)	†		
Subcutaneous fat (cm <sup>2</sup> )	95 ± 52	$184 \pm 119$	NS
Visceral fat (cm <sup>2</sup> )	27 ± 7	$42 \pm 20$	NS
Hepatic fat (%)	$10 \pm 6$	18 ± 6	0.001
Central (hepato-visceral) fat	37 ± 11	$60 \pm 22$	0.01

**Supplemental Table 1.** Study variables in healthy control girls (N= 21) and in girls with Polycystic Ovary Syndrome (PCOS, N= 52).

Values are mean ± standard deviation.

BMI, body mass index; SHBG, sex hormone-binding globulin; FAI, free androgen index; HOMA-IR, homeostasis model assessment insulin resistance; HDL, high-density lipoprotein; LDL, low-density lipoprotein; HMW, high molecular weight; usCRP, ultra-sensitive C-reactive protein; DXA, dual X-ray absorptiometry; MRI, magnetic resonance imaging; NS: not significant.

<sup>†</sup>SHBG, FAI, usCRP and abdominal fat partitioning assessments were performed in 15 out of 21 healthy adolescent girls.

<sup>‡</sup> Indicative DXA values in healthy adolescents, matched for age and height (n=41): lean mass  $35.1 \pm 1.0$  kg; fat mass  $17.6 \pm 1.4$  kg (Ibáñez L, et al. *J Adolesc Health 2017;* 61: 446-453).

P values are adjusted for age and BMI.

All subjects (N=73)							
	CXCL14	(ng/ mL)					
A	R	Р					
Auxological variables							
Age (yr)	-0.130	0.273					
Birth weight Z-score	0.035	0.768					
BMI (kg/m <sup>2</sup> )	-0.157	0.185					
BMI Z-score	-0.137	0.247					
$\Delta$ Z-score birth weight – BMI	-0.124	0.299					
Endocrine-Metabolic variables							
Testosterone (nmol/ L)	-0.023	0.854					
SHBG (nmol/ L) <sup>†</sup>	0.039	0.756					
FAI <sup>†</sup>	-0.099	0.441					
Glucose (mmol/ L)	0.224	0.058					
Fasting insulin (pmol/ L)	-0.102	0.392					
HOMA-IR	-0.083	0.485					
HDL-cholesterol (nmol/ L)	0.039	0.740					
LDL-cholesterol (nmol/ L)	0.058	0.627					
Triglycerides (nmol/ L)	-0.125	0.290					
HMW adiponectin (mg/ L)	0.275	0.023					
usCRP (mg/ L) $^{\dagger}$	-0.195	0.115					
Abdominal fat partitioning $(MRI)^{\dagger}$							
Subcutaneous fat (cm <sup>2</sup> )	-0.158	0.202					
Visceral fat (cm <sup>2</sup> )	0.025	0.844					
Hepatic fat (%)	-0.238	0.052					
Central (hepato-visceral) fat	-0.049	0.692					

**Supplemental Table 2.** Correlations between circulating baseline CXCL14 concentrations and selected variables in the study population [N= 21 controls and N= 52 girls with Polycystic Ovary Syndrome (PCOS].

CXCL14, C-X-C motif chemokine ligand-14, BMI, body mass index; SHBG, sex hormone-binding globulin; FAI, free androgen index; HOMA-IR, homeostasis model assessment insulin resistance; HDL, high-density lipoprotein; LDL, low-density lipoprotein; HMW, high molecular weight; usCRP, ultra-sensitive C-reactive protein; DXA, dual X-ray absorptiometry; MRI, magnetic resonance imaging.

<sup>†</sup> SHBG, FAI, usCRP and abdominal fat partitioning assessments were performed in 67 subjects (15 control girls and 52 girls with PCOS).

Results are Pearson correlation coefficients and P values.

	$\mathbf{OC}  (\mathbf{N} = \mathbf{I0})$		3	FIONET ( $N = 1$	5)	P value	
	Start <sup>‡</sup>	1 year	<b>∆ 0-1</b> year	Start <sup>‡</sup>	1 year	<b>∆ 0-1</b> year	P value
Auxology							
Age (years)	$15.8 \pm 1.4$	$16.9 \pm 1.5$	-	$15.5 \pm 1.4$	$16.6 \pm 1.4$	-	-
Birth weight Z-score	$-0.5 \pm 1.0$	-	-	$-0.2 \pm 1.1$	-	-	-
BMI (kg/m <sup>2</sup> )	$25 \pm 4$	$26 \pm 4$	$1.2 \pm 1.4$	$25 \pm 5$	$25 \pm 5$	$0.0 \pm 2.4$	NS
BMI Z-score	$0.9 \pm 1.2$	$1.3 \pm 1.4$	$0.4 \pm 0.5$	$1.1 \pm 1.4$	$1.1 \pm 1.5$	$-0.1 \pm 0.7$	NS
$\Delta$ Z-score birth weight – BMI	$1.4 \pm 1.2$	-	-	$1.3 \pm 1.8$	-	-	-
<b>Endocrine-Metabolic Variables</b>							
Testosterone (nmol/ L)	$1.3 \pm 0.5$	$0.9 \pm 0.5$	$-0.4 \pm 0.7$	$1.3 \pm 0.4$	$1.0 \pm 0.5$	$-0.3 \pm 0.3$	NS
SHBG (nmol/ L)	$34 \pm 13$	$57 \pm 26$	23 ± 21	$31 \pm 14$	$29 \pm 13$	-2 ± 9	0.0003
FAI	$4.7 \pm 2.4$	$2.6 \pm 4.6$	$-2.1 \pm 5.0$	$5.0 \pm 2.8$	$3.6 \pm 9.2$	$-1.4 \pm 1.0$	NS
Glucose (mmol/ L)	$4.5 \pm 0.3$	$4.5 \pm 0.5$	$0.0 \pm 0.6$	$4.5 \pm 0.4$	$4.2 \pm 0.3$	$-0.3 \pm 0.5$	NS
Fasting insulin (pmol/ L)	$78 \pm 41$	91 ± 45	13 ±33	$69 \pm 28$	$49 \pm 24$	$-20 \pm 35$	0.024
HOMA-IR	$2.3 \pm 1.3$	$2.6 \pm 1.4$	$0.3 \pm 1.1$	$2.0 \pm 0.9$	$1.4 \pm 0.8$	$-0.6 \pm 1.1$	0.035
HDL-cholesterol (nmol/ L)	$1.3 \pm 0.3$	$1.3 \pm 0.3$	$0.0 \pm 0.2$	$1.3 \pm 0.2$	$1.4 \pm 0.2$	$0.1 \pm 0.2$	NS
LDL-cholesterol (nmol/ L)	$2.3 \pm 0.5$	$2.6 \pm 0.6$	$0.3 \pm 0.4$	$2.0 \pm 0.4$	$2.0 \pm 0.5$	$0.0 \pm 0.4$	NS
Triglycerides (nmol/ L)	$0.7 \pm 0.2$	$0.7 \pm 0.3$	$0.0 \pm 0.2$	$0.7 \pm 0.2$	$0.7 \pm 0.2$	$0.0 \pm 0.2$	NS
HMW adiponectin (mg/ L)	5 ± 3	$5 \pm 6$	0 ± 5	$5 \pm 3$	$9 \pm 6$	$4 \pm 6$	0.045
usCRP (mg/ L)	$1.0 \pm 1.0$	$2.1 \pm 2.0$	$1.1 \pm 1.7$	$1.7 \pm 2.8$	$1.1 \pm 1.0$	$-0.6 \pm 2.9$	0.003
Body composition (DXA)							
Bone mineral density (g/cm <sup>2</sup> )	$1.19 \pm 0.1$	$1.19 \pm 0.1$	$0.00 \pm 0.03$	$1.21 \pm 0.1$	$1.20 \pm 0.1$	$-0.01 \pm 0.05$	NS
Lean mass (kg)	$37 \pm 6$	38 ± 7	1 ± 2	$36 \pm 5$	$36 \pm 4$	$0 \pm 2$	NS
Fat mass (kg)	24 ± 8	$26 \pm 9$	$2 \pm 2$	$25 \pm 10$	$25 \pm 9$	$0 \pm 5$	NS
Abdominal fat (kg)	$6.2 \pm 2.2$	$6.7 \pm 2.4$	$0.5 \pm 0.8$	$6.5 \pm 2.3$	$6.2 \pm 2.4$	$-0.3 \pm 1.2$	0.016
Abdominal fat partitioning (MRI)							
Subcutaneous fat (cm <sup>2</sup> )	$192 \pm 112$	$217 \pm 119$	$25 \pm 33$	$188 \pm 135$	$183 \pm 127$	-5 ± 77	0.022
Visceral fat (cm <sup>2</sup> )	41 ± 19	$45 \pm 26$	4 ± 17	$40 \pm 17$	$38 \pm 17$	$-2 \pm 9$	NS
Hepatic fat (%)	17 ± 7	19 ± 4	2 ± 7	$20 \pm 5$	$10 \pm 4$	$-10 \pm 4$	0.00003
Central (hepato-visceral) fat	$58 \pm 23$	$64 \pm 28$	6 ± 18	$60 \pm 17$	$48 \pm 19$	$-12 \pm 10$	0.0051

**Supplemental Table 3.** Auxological, endocrine-metabolic, body composition (by DXA), and abdominal fat partitioning (by MRI) assessments in adolescent girls with PCOS who were randomized to receive oral contraception (OC, N=16) or a low-dose combination of Spironolactone (50mg/d), Pioglitazone (7.5 mg/d) and Metformin (850 mg/d) (SPIOMET, N=15) for 1 year.

SPIOMET (N=15)

**OC**<sup>†</sup> (N=16)

Values are mean  $\pm$  standard deviation.

BMI, body mass index; SHBG, sex hormone-binding globulin; FAI, free androgen index; HOMA-IR, homeostasis model assessment insulin resistance; HDL, high-density lipoprotein; LDL, low-density lipoprotein; HMW, high molecular weight; usCRP, ultra-sensitive C-reactive protein; DXA, dual X-ray absorptiometry; MRI, magnetic resonance imaging. NS: not significant

<sup>†</sup>OC: Ethinylestradiol 20 µg plus Levonorgestrel 100 mg

<sup>‡</sup>Not significant differences between randomized subgroups at start.

*P* values are adjusted for age and BMI.

**Supplemental Table 4.** Correlations between 0-1 year changes ( $\Delta$ ) in serum CXCL14 levels and those in clinical, endocrinemetabolic, body composition, and abdominal fat partitioning variables in girls with Polycystic Ovary Syndrome (PCOS, N=15) who received a low-dose combination of Spironolactone (50 mg/d), Pioglitazone (7.5 mg/d) and Metformin (850 mg/d) (SPIOMET) for 1 year.

SPIOMET (N=15)						
	Δ CXCL1	4 (ng/ mL)				
Auvology	ĸ	P				
	0.100	0.702				
$\Delta$ BMI (kg/m <sup>2</sup> )	-0.108	0.702				
Δ BMI Z-score	-0.108	0.703				
Endocrine-Metabolic variables						
$\Delta$ Testosterone (nmol/ L)	0.293	0.331				
$\Delta$ SHBG (nmol/ L)	-0.144	0.608				
$\Delta$ FAI	0.017	0.955				
$\Delta$ Glucose (mmol/ L)	0.511	0.051				
$\Delta$ Fasting insulin (pmol/ L)	0.588	0.021				
Δ HOMA-IR	0.630	0.012				
$\Delta$ HDL-cholesterol (nmol/ L)	-0.198	0.498				
$\Delta$ LDL-cholesterol (nmol/ L)	-0.446	0.096				
$\Delta$ Triglycerides (nmol/ L)	-0.045	0.874				
$\Delta$ HMW adiponectin (mg/ L)	0.037	0.903				
Δ usCRP (mg/ L)	-0.009	0.980				
Body composition (DXA)						
$\Delta$ Bone mineral density (g/ cm <sup>2</sup> )	-0.205	0.463				
$\Delta$ Lean mass (kg)	-0.176	0.531				
$\Delta$ Fat mass (kg)	-0.036	0.900				
$\Delta$ Abdominal fat (kg)	-0.243	0.384				
Abdominal fat partitioning (MRI)						
$\Delta$ Subcutaneous fat (cm <sup>2</sup> )	-0.111	0.693				
$\Delta$ Visceral fat (cm <sup>2</sup> )	-0.187	0.504				
$\Delta$ Hepatic fat (%)	0.226	0.417				
$\Delta$ Central (hepato-visceral) fat	-0.083	0.769				

CXCL14, C-X-C motif chemokine ligand-14, BMI, body mass index; SHBG, sex hormone-binding globulin; FAI, free androgen index; HOMA-IR, homeostasis model assessment insulin resistance; HDL, high-density lipoprotein; LDL, low-density lipoprotein; HMW, high molecular weight; usCRP, ultra-sensitive C-reactive protein; DXA, dual X-ray absorptiometry; MRI, magnetic resonance imaging.

Results are Pearson correlation coefficients and P values.

**STUDY 3** 

## GUT MICROBIOTA IN ADOLESCENT GIRLS WITH POLYCYSTIC OVARY SYNDROME: EFFECTS OF RANDOMIZED TREATMENTS

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#### STUDY 3

## Gut microbiota in adolescent girls with polycystic ovary syndrome: Effects of randomized treatments

**Introduction:** Girls with obesity and polycystic ovary syndrome (PCOS) and women with PCOS have altered gut microbiota.

**Objective:** To study the gut microbiota composition of girls with PCOS without obesity (age, 15.8 years; body mass index [BMI] 25 kg/m<sup>2</sup>) and the effects of randomized treatments with an oral contraceptive (OC, N=15) or with spironolactone-pioglitazone-metformin (SPIOMET, N=15) for 1 year. Thirty-one age-matched girls served as controls.

**Material and methods:** 16S ribosomal subunit gene amplicon sequencing was performed in stool samples from all subjects; samples from 23 out of 30 girls with PCOS (OC, N=12; SPIOMET, N=11) were available for analysis post treatment. Clinical and endocrine-metabolic variables were measured before and after intervention.

**Results:** Girls with PCOS had decreased diversity alpha, altered microbiota pattern and taxonomic profile with more abundance of *Family XI* (P=0.002), and less abundance of family *Prevotellaceae* (P=0.0006) the genus *Prevotella* (P=0.0001) and *Senegalimassilia* (P<0.0001), as compared to controls. *Family XI* abundance related positively to hepatovisceral fat (R=0.453; P=0.0003). SPIOMET treatment, but not OC, normalized the abundance of *Family XI*. *Prevotellaceae*, *Prevotella* and *Senegalimassilia* abundance remained unchanged after either treatment.

**Conclusion:** SPIOMET's spectrum of normalizing effects in girls with PCOS is herewith broadened as to include *Family XI* abundance in gut microbiota.

#### **ORIGINAL RESEARCH**

#### ediatric WILEY

## Gut microbiota in adolescent girls with polycystic ovary syndrome: Effects of randomized treatments

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#### Summary

Background: Girls with obesity and polycystic ovary syndrome (PCOS) and women with PCOS have altered gut microbiota.

Objective: To study the gut microbiota composition of girls with PCOS without obesity (age, 15.8 years; body mass index [BMI] 25 kg/m<sup>2</sup>) and the effects of randomized treatments with an oral contraceptive (OC, N = 15) or with spironolactonepioglitazone-metformin (SPIOMET, N = 15) for 1 year. Thirty-one age-matched girls served as controls.

Methods: 16S ribosomal subunit gene amplicon sequencing was performed in stool samples from all subjects; samples from 23 out of 30 girls with PCOS (OC, N = 12; SPIOMET, N = 11) were available for analysis post-treatment. Clinical and endocrinemetabolic variables were measured before and after intervention.

**Results:** Girls with PCOS had decreased diversity alpha, altered microbiota pattern and taxonomic profile with more abundance of Family XI (P = .002), and less abundance of family Prevotellaceae (P = .0006) the genus Prevotella (P = .0001) and Senegalimassilia (P < .0001), as compared to controls. Family XI abundance related positively to hepato-visceral fat (R = 0.453; P = .0003). SPIOMET treatment, but not OC, normalized the abundance of Family XI. Prevotellaceae, Prevotella and Senegalimassilia abundance remained unchanged after either treatment.

Conclusion: SPIOMET's spectrum of normalizing effects in girls with PCOS is herewith broadened as to include Family XI abundance in gut microbiota.

#### KEYWORDS

gut microbiota, hepatic fat, metformin, PCOS, pioglitazone, spironolactone

Abbreviations: 16S rRNA gene, 16S ribosomal RNA subunit gene; ALT, alanine transaminase; ANOSIM, analysis of similarities; ASV, amplicon sequence variant; BMI, body mass index; dbRDA, distance-based redundancy analysis; FAI, free androgen index; FDR, false discovery rate; HMW, high molecular weight; HOMA-IR, homeostasis model assessment - insulin resistance; LDL, low density lipoprotein; MRI, magnetic resonance imaging; MSG, mean serum glucose; MSI, mean serum insulin; OC, oral contraceptive; PCoA, principal coordinated analysis; PCOS, Polycystic ovary syndrome; SHBG, sex hormone-binding globulin; SPIOMET, spironolactone, pioglitazone and metformin; usCRP, ultrasensitive C reactive protein.

#### 1 | INTRODUCTION

Polycystic ovary syndrome (PCOS) is a highly prevalent, chronic endocrine-metabolic disorder affecting up to 10% of women of reproductive age worldwide.<sup>1</sup> It is the most common cause of hirsutism and menstrual irregularity in adolescent girls and young women, and

appears to result from an excess of fat in subcutaneous adipose tissue -regardless of body weight-, leading to ectopic lipid deposition, especially in the liver and viscera (hepato-visceral or central fat), and subsequently, to insulin resistance and androgen excess.<sup>2,3</sup> The syndrome often associates with comorbidities in adulthood, including subfertility, type 2 diabetes and increased cardiovascular disease risk, among others.<sup>4,5</sup> We have recently shown that in girls with PCOS, a low-dose combination of one mixed anti-androgen and antimineralocorticoid (spironolactone), and two insulin sensitizers (pioglitazone and metformin) (SPIOMET) exerts more benefits on ovulation rate, insulin resistance, ectopic fat deposition, and other cardiometabolic risk markers than does the standard treatment with oral contraceptives (OC), and thus holds the potential for avoiding or delaying PCOS-associated comorbidities.<sup>6</sup>

The gut microbiome provides a set of benefits to the host, including protective, metabolic, structural and histological functions.<sup>7</sup> For example, a relatively increased microbiota gene content and microbial diversity associates with enhanced metabolic health.<sup>8</sup> Alteration of the composition of the intestinal microbiome, so-called dysbiosis, is associated with many disorders, such as obesity, insulin resistance and type 2 diabetes.<sup>9,10</sup> Lean adult women with PCOS as well as women and adolescents with obesity and with PCOS have gut microbiota communities different from those in healthy controls,<sup>11-15</sup> suggesting that the dysbiosis of the gut microbiome may contribute to the pathogenesis of the disorder.<sup>16</sup>

So far, the profile of the gut microbiota as well as its longitudinal changes following prolonged pharmacological intervention have not been characterized in adolescent girls with PCOS and without obesity. Here, we hypothesized that the composition of the gut microbiota in these girls will differ from that in control girls and will have a dissimilar pattern after SPIOMET or OC administration. To test this hypothesis, we used a marker-based approach using the 16S ribosomal RNA subunit gene (16S rRNA gene) to describe and quantify microbial alpha and beta diversity and to study taxonomic profiles in stool samples from healthy control girls and from girls with PCOS before intervention, and after 1 year of treatment with OC or SPIOMET.

#### MATERIALS AND METHODS 2

#### 2.1 Study population and design

The original cohort consisted of 35 Catalan adolescent girls with PCOS (mean age 15.8 years, mean body mass index (BMI) 25 kg/m<sup>2</sup>) enrolled into a randomized, open-label, single-centre controlled trial (ISCRCTN11062950), performed at the Adolescent Endocrinology Unit of Sant Joan de Déu University Hospital, Barcelona, Spain. The study - started in December 2015 and completed in October 2019-, explored the effects of OC vs SPIOMET treatment for 1 year; ovulation rate was the primary endpoint.<sup>6</sup> Gut microbiota composition analysis was a secondary endpoint, and was performed in a subset of 30 girls who provided baseline stool samples (N = 15 on OC; N = 15 on SPIOMET); 23 out of those 30 girls (N = 12 on OC; N = 11 on SPIOMET) collected a stool sample also after 1 year on treatment (Figure S1, flow chart). The characteristics of the girls who did not provide a stool sample after intervention did not differ from those who did (data not shown). Hirsutism score, fasting insulin, androgens, lipids, high molecular weight (HMW) adiponectin, ultrasensitive C reactive protein (usCRP), body composition, and abdominal fat partitioning were also among the secondary outcomes.<sup>6</sup> Mediterranean diet and regular physical exercise were advised to all girls.

The inclusion criteria were hirsutism (score > 8 on modified Ferriman-Gallwey scale), oligomenorrhea (menstrual intervals >45 days), gynaecological age > 2.0 years, and no need for contraception. Exclusion criteria were 21-hydroxylase deficiency; glucose intolerance or diabetes; evidence of thyroid, liver, or kidney dysfunction; hyperprolactinemia; and prior use of medications affecting gonadal/ adrenal function, or carbohydrate/lipid metabolism.<sup>6</sup>

OC treatment consisted of 20  $\mu$ g of ethinylestradiol plus 100 mg of levonorgestrel for 21/28 days and placebo for 7/28 days; SPIOMET is a low dose combination of spironolactone 50 mg/d. pioglitazone 7.5 mg/ d, and metformin 850 mg/d.<sup>6</sup> Thirty-one healthy girls (mean age 15.9 years, mean BMI 22 kg/m<sup>2</sup>) recruited in nearby schools served as controls. All had regular menstrual cycles, a gynaecological age > 2.0 years, and none had chronic diseases, was hirsute or was taking medications. None of the girls was taking antibiotics at the time of collection of the stool samples.

#### 2.2 Clinical, endocrine-metabolic, and imaging assessments

Birth weight and BMI (and their Z-scores) were retrieved from hospital records, and endocrine-metabolic variables were assessed in the early morning, in the follicular phase (days 3-7) of the cycle or after 2 months of amenorrhea.<sup>6</sup> Serum insulin, homeostasis model assessment (HOMA) - insulin resistance (IR), lipids, usCRP, sex hormonebinding globulin (SHBG), androgens, free androgen index (FAI), and HMW-adiponectin were assessed as reported.<sup>6</sup> Standard oral glucose tolerance tests with 75 g glucose were performed after 3 days of consumption of a high carbohydrate diet (300 g/d) and an overnight fast. Blood was sampled before and 30, 60, 90, and 120 minutes after glucose intake for glucose and insulin measurements, from which mean serum glucose (MSG), mean serum insulin (MSI) and their Z-scores were derived. MSG and MSI were calculated as the area under the glucose and insulin response, respectively, from time 0 to time 120.<sup>6</sup>

Ovulatory function was assessed through measurement of weekly salivary progesterone over two periods of 12 consecutive weeks, during the second and fourth trimesters of the post-treatment year, as published.<sup>6</sup>

Body composition was assessed by dual X-ray absorptiometry with a Lunar Prodigy and Lunar software (version 3.4/3.5, Lunar Corp, Madison, Wisconsin); abdominal (subcutaneous and visceral) and hepatic fat were assessed by magnetic resonance imaging (MRI) using a multiple-slice MRI 1.5 Tesla scan (Signa LX Echo Speed Plus Excite, General Electric Milwaukee, Wisconsin).<sup>6</sup> Hepatic fat (%) was

calculated by comparing the intensity of the liver to that of subcutaneous fat and spleen, assuming that the latter organ is fat free. Scans were performed by the same operator, and images were analysed by the same radiologist (both blinded to treatment allocation). Central (hepato-visceral) fat was arbitrarily defined as the sum of visceral fat (in cm<sup>2</sup>) and hepatic fat (in %).

## 2.3 | Stool DNA extraction, library preparation and sequencing

Stool samples were collected at home using a faeces collector (FECOTAINER, Excretas Medical BV, Enschede, The Netherlands) in the morning of the day before the hospital appointment; the samples were frozen in the patient's freezer at  $-20^{\circ}$ C and transported on ice to the hospital, where they were sprayed with liquid nitrogen, aliquoted and stored at -80°C until analysis. The 16S rRNA gene analvsis was performed at Microomics (Barcelona, Spain; http://www. microomics.eu/). DNA was extracted using MagMAX CORE Nucleic Acid Purification Kit 500 RXN (Thermo Fisher, CA, Austin), following the manufacturer's instructions. As a control for downstream procedures. Mock community DNA was included (Zymobiomics Microbial Community DNA, ZymoResearch, Irvine). Samples were amplified using 16S rRNA V3-V4 regions specific primers (V3-V4-Forward 5'-TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGCCTACGGGNGGC WGCAG-3', V3-V4-Reverse 5'GTCTCGTGGGCTCGGAGATGTGTAT AAGAGACAGGACTACHVGGGTATCTAATCC-3'). The PCR was performed in 10-µL final volume with 0.2-µM primer concentration. The PCR cycle included: 3 minutes at 95°C (initial denaturation) followed by 25 cycles: 30 seconds at 95°C 30 s at 55°C, and 30 s at 72°C, and a final elongation step of 5 minutes at 72°C. PCR products were purified using AMPure XP beads (Beckman Coulter, Nyon, Switzerland) with a 0.9x ratio according to manufacturer's instructions. The above described primers contain overhangs allowing the addition of full-length Nextera barcoded adapters for multiplex sequencing in a second PCR step, resulting in sequencing ready libraries with approximately 450 bp insert sizes. In brief, 5 µL of the first PCR purified product were used as template for a second PCR with Nextera XT v2 adaptor primers in a final volume of 30 µL using the same PCR mix and thermal profile as for the first PCR but with only 8 cycles. Twenty-five microliter of the second PCR product were purified with SequalPrep normalization kit (Invitrogen, ThermoFisher Scientific, Waltham, Massachusetts), according to the manufacturer's protocol. Libraries were eluted in 20 µL final volume and pooled for sequencing. The final pool was quantified by qPCR using Kapa library quantification kit for Illumina Platforms (Kapa Biosystems, SigmaAldrich, Saint Louis, Missouri) on an ABI 7900HT real-time cycler (Applied Biosystems, ThermoFisher Scientific, Waltham, Massachusetts). Sequencing was performed using Illumina MiSeq (2 × 300 bp) and v3 chemistry with a loading concentration of 10 PM In all cases, 15% of PhIX control libraries for run quality monitoring were used. Negative controls of the sample collection buffer, DNA extraction, and PCR amplification steps were routinely performed in parallel, using the same conditions and reagents, PCR products after both PCR steps were visualized by electrophoresis gel (1.5% agarose) with SYBR Safe (ThermoFisher Scientific, Waltham, Massachusetts).

## 2.4 | Bioinformatics processing and statistical analysis

Raw demultiplexed forward and reverse reads were processed using the following methods and pipelines as implemented in QIIME2 version 2019.4 with default parameters unless stated.<sup>17</sup> DADA2 was used for quality filtering, denoizing, pair- end merging and amplicon sequence variant calling (ASV) using giime dada2 denoise-paired method.<sup>18</sup> Q20 was used as guality threshold to define read sizes for trimming before merging (parameters: - p-trunc-len-f and -p-trunclen-r). Reads were truncated at the position when the 75th percentile Phred score felt below Q20: 287 bp for forward reads, and 224 bp for reverse reads. ASVs were aligned using the gime alignment mafft method.<sup>19</sup> The alignment was used to create a tree and to calculate phylogenetic relations between ASVs using the gime phylogeny fasttree method.<sup>20</sup> ASV tables were subsampled without replacement in order to even sample sizes for diversity analysis using gime diversity core-metrics-phylogenetic pipeline. ASV tables were used to calculate alpha diversity metrics: community richness (ASVs), Evenness (or Pielou's Evenness; a measure of community evenness) and Shannon's diversity index (quantitative measure of community richness). Alpha diversity comparisons were performed using Kruskal Wallis non-parametric-test. ASV tables and phylogenetic data were used to calculate beta diversity metrics: Bray-Curtis (non-phylogenetic guantitative measure) and Jaccard (non-phylogenetic qualitative measure) dissimilarity matrices, as well as Unweighted Unifrac (phylogenetic qualitative measure) and Weighted Unifrac (phylogenetic quantitative measure) distance matrices.<sup>21</sup> Principal coordinated analysis (PCoA) was the method used for exploring and visualizing similarity of data using the matrices described above. The significance of groups present in community structure was tested using analysis of similarities (ANOSIM) and Permanova tests. Permdisp test was used to identify location vs dispersion effects.<sup>22</sup> Distance-based redundancy analysis (dbRDA) was used to explore which variables constrained PCoA ordinations. Model selection was done stepwise using forward direction and a permutation test using vegan package version 2.5-5. Taxonomic assignment of ASVs was performed using Bayesian Classifier trained with Silva V4 database version 132.23 ASVs were filtered to discard contaminant Eukaryote DNAderived amplicons using Blast against the mentioned database with 90% identity cutoff. Differential abundance of taxa was tested using two methods: ANCOM,<sup>24</sup> and Kruskal Wallis non-parametric test on relative abundance of taxa (total sum of scale). After Kruskal Wallis test, Conover's test was added for pairwise comparison. Benjamini-Hochberg correction was used to control false discovery rate (FDR). The relative abundance of each taxon was calculated as the number of 16S rRNA sequences of a given taxon divided by the total number of 16S rRNA sequence in a participant's sample. Significant threshold was set at 0.05. BiodiversityR version 2.11-1, PMCMR version 4.3, RVAideMemoire version 0.9-7 and vegan version 2.5-5 packages and R software package version 3.6.0 (http://www.R-project.org) were used.

## 2.5 | Statistical analysis of clinical and imaging data and ethics

The statistical analyses of clinical and imaging data were performed with SPSS 23.0 (SPSS, Chicago, Illinois). Baseline differences in clinical, endocrine-metabolic, body composition, and abdominal fat partitioning variables between patients and controls were tested with unpaired *t* test, adjusting for BMI. Longitudinal changes between groups were compared by repeated-measures general linear model. Differences in longitudinal changes between and within- subject effects. The between-group differences in ovulation numbers were sought by the Mann Whitney *U* test. Associations between clinical variables and alpha diversity measures or bacterial taxa were sought by Spearman correlation analysis. *P* < .05 was considered statistically significant. Results are expressed as mean  $\pm$  SEM. The study was conducted after approval by the Institutional Review Board of Sant Joan de Déu University Hospital, after written consent by parents and assent by each of the participants.

#### 3 | RESULTS

## 3.1 | Clinical, endocrine-metabolic and imaging variables

The clinical characteristics of the participants and the effects of OC vs SPIOMET treatment on endocrine-metabolic and imaging parameters are summarized in Table S1. At baseline, girls with PCOS showed lower SHBG and HMW-adiponectin levels, higher MSI, total testosterone concentrations and FAI, and more hepato-visceral (central) fat as compared to control girls, as described.<sup>6</sup> Consistent with previous observations, SPIOMET exerted more benefits than OCs on most studied variables, including on circulating insulin, HMW-adiponectin, usCRP and central fat.<sup>6</sup>

#### 3.2 | Gene sequencing

Samples were available for gene sequencing on the V3 and V4 variable regions of the 16S RNA gene. The average number of reads per sample after quality filtering steps was 71 010 (min: 7048 reads; max: 193 490 reads), and 6869 ASVs were detected. The sample with the smallest size (7048 reads, belonging to a patient with PCOS at baseline) was discarded



**FIGURE 1** Alpha and beta diversity of the gut microbial communities in healthy controls (N = 31) and in girls with polycystic ovary syndrome (PCOS, N = 29) at baseline. A-C, Alpha diversity boxplots (median and interquartile ranges) based on the number of amplicon sequence variants (ASVs), A; Pielou's Evenness, B; and Shannon's diversity Index, C. D and E, Principal coordinate analysis (PCoA) of faecal microbiota based on Jaccard D, (ANOSIM test, P = .01, R = 0.165) and Bray-Curtis dissimilarity matrices E, (ANOSIM test, P = .01, R = 0.202). Each dot represents the bacterial community composition of one individual stool sample; the axis title indicates the percentage of variation explained. One sample from the group with PCOS was excluded from the diversity analysis (alpha and beta) due to the low coverage




**FIGURE 2** Distance-based redundancy analysis (dbRDA) ordination plots illustrating the relationship between clinical variables that best explain the variation of the composition of the gut microbiota in girls with polycystic ovary syndrome (PCOS, N = 30) at baseline and healthy controls (N = 31). A, Central (hepato-visceral) fat (F = 1.06, P = .008) and free androgen index (FAI) (F = 1.05, P = 0.024) for Jaccard B, and hepatic fat (F = 1.32, P = .04) for Bray-Curtis dissimilarity matrices were the majors determinants of the variation of the gut microbiota in girls with PCOS



**FIGURE 3** Baseline composition of gut microbial community in girls with polycystic ovary syndrome (PCOS, N = 30) and healthy controls (N = 31). A, Composition of gut microbial community at phylum level. B-E, Comparison of the relative abundance of the two families, B and C; and the two genera, D and E; that were detected by ANCOM test among groups. Data are mean + SEM. *P* values were obtained using Kruskal Wallis + Conover's test

from the diversity analysis (alpha and beta) in order to take advantage of the sequencing depth of the dataset. The number of reads for the mock control were as expected (Table S2). For the negative control, 17 taxonomic groups at the genus level were detected. When the taxonomic assignments of the reads in the negative control and the samples were compared, the ratio of 15 of these groups was either null or at least one magnitude order lower in the samples (Table S3a). The average abundance of the [Eubacterium] coprostanoligenes group and the

Escherichia-Shigella group was higher in the samples than in the negative control. In order to evaluate potential contaminants, the composition of these two groups was studied at ASV level. Results revealed that the [Eubacterium] coprostanoligenes group was represented by one single ASV which was present in only one sample. Escherichia-Shigella was also represented by a single ASV in the negative control, which was present in three samples (Table S3b). The negative control evaluation was successful to detect environmental contaminants. Results showed that these contaminants were almost absent in samples and did not affect their composition. In addition, our controls systematically provided no visible band or quantifiable DNA amounts by gel visualization or Biolanalyzer, whereas all of our samples provided a clearly visible band (Figure S2).

### The composition of the gut microbiota differs 3.3 between girls with PCOS and healthy controls

Regarding alpha diversity, the number of ASVs was similar at baseline in girls with PCOS and in control girls (Figure 1A). However, Pielou's Evenness and Shannon's diversity Index were both significantly lower in the subgroup with PCOS (P = .03 and P = .04, respectively) (Figures 1B,C).

Beta diversity analysis also unveiled differences in community structure between patients with PCOS at baseline and control girls (Figures 1D,E), showing differences in dispersion for the two nonphylogenetic dissimilarity matrices used: Jaccard (ANOSIM test: P = .001, R = 0.165; Permanova test: P = .01; Permdisp test: P = .19) and Bray-Curtis (ANOSIM test: P = .001, R = 0.202; Permanova test: P = .002; Permdisp test: P = .0001). No differences were detected in microbiota composition overall between girls with PCOS and controls for unweighted and weighted Unifrac distances.

Among the selected clinical variables assessed for the dbRDA (Δ Z-score birth weight - BMI, alanine transaminase [ALT], HOMA-IR, FAI, HMW-adiponectin, visceral fat, hepatic fat and central [hepato-visceral] fat), the greatest amount of variation in the gut microbiota composition for both patients at baseline and controls, was explained by central fat (F = 1.06, P = .008) and FAI (F = 1.05, P = .024) for Jaccard, and hepatic fat (F = 1.32, P = .04) for Bray-Curtis dissimilarity matrices. In addition, the dbRDA diagram for both matrices showed a cluster of control sites away from PCOS sites (Figure 2).

All subjects (N = 61)			
	Bacterial measure	R	Р
Study variables			
Testosterone (nmol/ L)	Segenalimassilia [Relative abundance (%)]	-0.284	.026
SHBG (nmol/ L)	Pielou's Evenness	0.323	.012
	Shannon Index	0.341	.008
	Prevotellaceae [Relative abundance (%)]	0.301	.018
	Prevotella [Relative abundance (%)]	0.298	.020
	Segenalimassilia [Relative abundance (%)]	0.429	.001
FAI	Pielou's Evenness	-0.288	.030
	Segenalimassilia [Relative abundance (%)]	-0.440	.001
ALT (µkat/ L)	ASVs	0.310	.010
	Pielou's Evenness	0.376	.003
	Shannon Index	0.384	.002
LDL-cholesterol (mmol/L)	Prevotellaceae [Relative abundance (%)]	0.297	.020
	Prevotella [Relative abundance (%)]	0.269	.036
HMW-adiponectin (mg/ L)	Family XI [Relative abundance (%)]	-0.345	.011
	Prevotellaceae [Relative abundance (%)]	0.322	.019
	Prevotella [Relative abundance (%)]	0.342	.013
usCRP (nmol/ L)	Prevotella [Relative abundance (%)]	-0.308	.017
	Segenalimassilia [Relative abundance (%)]	-0.319	.013
Visceral fat (cm <sup>2</sup> )	Family XI [Relative abundance (%)]	0.408	.004
Hepatic fat (%)	Segenalimassilia [Relative abundance (%)]	-0.352	.007
Central (hepato-visceral) fat	Family XI [Relative abundance (%)]	0.453	.0003
	Segenalimassilia [Relative abundance (%)]	-0.309	.018

Abbreviations: ALT, alanine transferase; ASVs, amplicon sequence variants; FAI, free androgen index; HMW, high molecular weight; LDL, low density lipoprotein; SHBG, sex hormone-binding globulin; usCRP, ultrasensitive C-reactive protein.

Note: Results are Spearman correlation coefficients and P values. All correlations with P value <.05 are shown.

**TABLE 1** Correlations between clinical, endocrine-metabolic and imaging variables and bacterial measures in the entire study population [control girls (N = 31) and girls with polycystic ovary syndrome (N = 30)] at baseline

Figure 3A shows that the most abundant Phyla in both control girls and in girls with PCOS were *Firmicutes*, *Bacteroidetes* and *Actinobacteria*, representing about 89% of the total community abundance, as expected.<sup>25</sup>

Regarding family composition, 102 different families were identified, but the ANCOM test only detected differences in two: Bacillales Family XI (null hypothesis rejected, W = 8) and Prevotellaceae (null hypothesis rejected, W = 12); belonging to the Firmicutes and Bacteroidetes phylum, respectively. Kruskal Wallis + Conover's test confirmed these results, showing that Family XI was significantly increased in samples from girls with PCOS, being present in 15 out of 30 samples (50%), as compared to only 4 out of 31 samples in controls (13%) (P = .002) (Figure 3B). In contrast, Prevotellaceae was found to be more abundant in control samples than in samples of patients with PCOS (P = .0006) (Figure 3C). At genus level, 337 different types of genera were detected, but the ANCOM test only detected that two genera were statistically different between controls and samples of patients with PCOS: Prevotella (null hypothesis rejected, W = 215) and Senegalimassilia (null hypothesis rejected, W = 247); belonging to Firmicutes and Actinobacteria phylum, respectively. These results were also confirmed by Kruskal Wallis + Conover's test: Prevotella and Senegalimassilia were significantly more abundant in control samples (P = .0001 and P < .0001, respectively) (Figure 3D,E).

# 3.4 | Correlations between microbiota and baseline variables

Table 1 depicts the correlations between baseline clinical, endocrinemetabolic and imaging variables and bacterial measures. Alpha diversity measures (ASVs, Pielou's Evenness and Shannon Index) correlated with markers of hyperandrogenism (SHBG and FAI) and ALT, whereas *Family XI* relative abundance was negatively associated with HMWadiponectin and positively related to hepato-visceral fat. Family *Prevotellaceae* and genus *Prevotella* positively correlated with SHBG, low density lipoprotein (LDL) cholesterol, and HMW-adiponectin. *Prevotella* was negatively associated with usCRP, and *Senegalimassilia* positively associated with markers of hyperandrogenism (testosterone, SHBG and FAI), usCRP and hepato-visceral fat.

# 3.5 | Effects of randomized treatments on the composition of the gut microbiota

No differences between treatments were observed in alpha diversity (richness, evenness or Shannon Index). Within the OC subgroup, no changes 0 to 1 year were detected; in addition, alpha diversity did not differ after 1 year of treatment vs controls. SPIOMET tended to reduce the number of ASVs after 1 year (P = .07) and reduced alpha diversity richness as compared to controls (P = .03).

In relation to beta diversity, no changes in the microbiota composition were observed after OC treatment. In contrast, differences in community structure occurred in girls treated with SPIOMET for Bray-Curtis dissimilarity matrices (ANOSIM test: P = .057, R = 0.102; Permanova test: P = .01; Permdisp test: P = .01). These differences were also detected between controls and SPIOMET-treated patients for Jaccard dissimilarity matrices (ANOSIM test: P = .077, R = 0.095; Permanova test: P = .04; Permdisp test: P < .001). No differences were detected for unweighted and weighted Unifrac distances.

dbRDA analysis revealed that hepatic fat (F = 1.06, P = .029, for Jaccard dissimilarity matrices) was the major determinant of the gut microbiota variation in control girls and in girls with PCOS after treatment with OC or SPIOMET for 1 year. The dbRDA diagram showed a cluster of control and SPIOMET sites away from OC sites (Figure 4).

SPIOMET treatment, but not OC, reduced the abundance of *Family XI*, reaching levels similar to those in controls (Figure 5A). SPIOMET-treated girls also showed a reduction in *Prevotellaceae* levels, becoming less abundant and prevalent than in controls (Figure 5B). Both treatments reduced the abundance of genus *Prevotella* which remained significantly below those in controls (Figure 5C). Neither treatment modified *Senegalimassilia* abundance (Figure 5D).

The unchanged relative abundance of *Bacillales Family XI* in OCtreated patients was positively associated with the increase in fasting insulin (R = 0.633; P = .036), and in visceral (R = 0.686; P = .020) and central fat (R = 0.648; P = .031). No additional associations were detected between changes 0 to 1 year in the study variables and



**FIGURE 4** Distance-based redundancy analysis (dbRDA) ordination plot illustrating that hepatic fat (F = 1.057, P = .029) for Jaccard dissimilarity matrices, was the variable that best explained the variation of the composition of the gut microbiota in girls with polycystic ovary syndrome (PCOS) treated with oral contraceptives (OC, N = 12), or spironolactone, pioglitazone and metformin (SPIOMET, N = 11) for 1 year and in control girls (N = 31)



**FIGURE 5** Longitudinal changes in *Family XI*, A; *Prevotellaceae*, B; *Prevotella*, C; and *Senegalimassilia*, D in girls with PCOS who received oral contraceptives (OC, red circles, N = 12) or spironolactone, pioglitazone and metformin (SPIOMET, blue circles, N = 11). Data are mean  $\pm$  SEM. The dotted line is the mean in healthy controls (N = 31); the shaded area represents the  $\pm$  SEM in those healthy controls. *P* values were obtained using Kruskal Wallis + Conover's test. \**P* < .05; \*\**P* < .001, patients vs controls at baseline. \**P* < .05; \*\**P* < .01, on treatment differences between controls and OC subgroup. \**P* < .01; & *P* < .001, on treatment differences between controls and SPIOMET subgroup. \**P* < .01 for changes 0 to 1 year in the OC and in the SPIOMET subgroup

changes 0 to 1 year of alpha diversity measures or bacterial taxa either in the OC or SPIOMET subgroups.

### 4 | DISCUSSION

The present study appears to be the first assessing the gut microbiota of adolescent girls with PCOS without obesity, and the longitudinal changes after randomized pharmacological treatments. Our study demonstrates the presence of gut dysbiosis in these girls, including decreased evenness and diversity indexes. We confirmed the association between alpha diversity and markers of hyperandrogenism and liver function found in other populations.<sup>13-15</sup> We also showed an altered microbiota pattern in girls with PCOS according to Jaccard and Bray-Curtis dissimilarity matrices. However, the lack of differences in community structure using phylogenetic distances as Unweighted and Weighted Unifrac may indicate that despite detecting different ASVs conforming the microbiota profile of girls with PCOS, these ASVs may be phylogenetically similar to the ASVs

detected in control girls. In line with this observation, no differences were found in the abundance of the most copious phyla between patients with PCOS and healthy girls. However, girls with PCOS had an altered taxonomic profile, with more abundance of *Family XI* and lower abundance of the family *Prevotellaceae* and the genus *Prevotella* and *Senegalimassilia*. The excessive abundance of *Family XI* decreased toward normal after SPIOMET administration but remained unchanged after OCs.

The abundance of *Bacillales Family XI* in patients as compared to controls may have pathophysiological implications. This family has been so far poorly studied and the related literature is thus, scarce; however, the genus *Gemella*, which belongs to *Family XI*, has been linked to liver inflammatory disease.<sup>26</sup> In addition, *Gemella spp* (also belonging to *Family XI*) is a bacterium that produces N-oleoyl serinol, an analogue of ceramides.<sup>27</sup> Serum concentrations of ceramides are elevated in women with PCOS<sup>28,29</sup> and are associated to the degree of hyperandrogenism (as assessed by the FAI) and to central adiposity.<sup>30</sup> Moreover, patients with obesity with increased diabetes risk have elevated circulating ceramide levels that associate to gut

dysbiosis.<sup>31</sup> In addition, ceramides produced by the gut microbiota have the potential to pass the epithelial barrier and interact with the host metabolism.<sup>32</sup> Accordingly, it has been proposed that IR and hyperandrogenemia in patients with PCOS could be at least partly driven by an increase in serine phosphorylation due to the accumulation of reactive lipids, such as ceramides, that activate serine kinase signalling pathways.<sup>33</sup> Moreover, it has also been demonstrated that ceramides induce IR via inhibition of Akt signalling in cell cultures.<sup>34</sup> In our study, SPIOMET treatment, but not OC, decreased Family XI levels toward normal. This reduction could have caused in turn a decrease in N-oleoyl serinol production and subsequently normalize the serine phosphorylation status. Furthermore, the unchanged Family XI abundance in OC-treated girls was found to associate with increased fasting insulin and hepato-visceral fat. Hence, it is tempting to speculate that the metabolic benefits exerted by SPIOMET could be mediated in part by a reduction of Family XI abundance.

Previous studies have disclosed the association between adiponectin and ceramides; for example, adiponectin decreases cellular ceramides by triggering ceramidase activities of adiponectin receptors (AdipoRs)<sup>35</sup> and by stimulating ceramide efflux to the exosome.<sup>36</sup> Other studies suggest that ceramides might be involved in the regulation of adiponectin secretion.<sup>37</sup> In the present study, the increase in circulating HMW-adiponectin concentrations after SPIOMET treatment could have modified N-oleoyl serinol levels, and ultimately explain the negative association between HMW-adiponectin and *Family XI* relative abundance.

In line with previous observations,<sup>12</sup> we disclosed that girls with PCOS had less abundance of the family *Prevotellaceae* and the genus *Prevotella* (belonging to *Prevotellaceae* family). In contrast, other studies have shown an abundance of *Prevotella* in adolescent girls with PCOS and with obesity,<sup>15</sup> in women with irregular menstrual cycles<sup>38</sup> and in animal models of PCOS.<sup>39</sup> In addition, the reports on the involvement of *Prevotella* in other pathological conditions have been inconsistent. For example, increased abundance of *Prevotella* has been associated with IR, non-alcoholic fatty liver disease (NAFLD) and chronic inflammatory disorders.<sup>26,40-43</sup> However, other studies relate the abundance of *Prevotella* with an improved glucose metabolism<sup>44</sup> or fail to demonstrate an association with type 2 diabetes.<sup>45</sup>

The abundance of *Prevotellaceae* and *Prevotella* was significantly reduced after SPIOMET treatment. These results are in line with others demonstrating a non-significant reduction of *Prevotella* abundance after metformin intervention in women with obesity,<sup>46</sup> but are in contrast with others showing that metformin increases its abundance in healthy men<sup>47</sup> and in men with diabetes,<sup>48</sup> but not in groups combining men and women.<sup>48</sup> These discrepancies may be due to the presence of functional diversities within the *Prevotella* genus or to the influence of sex in the composition of gut microbiota.<sup>14</sup> Further analyses are needed to investigate the implications of the different *Prevotella* species within the context of PCOS.

In the present study, we also showed that girls with PCOS had lower abundance of the genus *Senegalimassilia*, and that this abundance associated inversely with markers of androgen excess and inflammation, and with hepato-visceral fat. Along these lines, a higher abundance of *Senegalimassilia* has been associated with healthy traits; Pediatric

for example, a higher occurrence of *Senegalimassilia anaerobia* (a specie belonging to *Senegalimassilia* genus) was observed in faecal samples of children without obesity as compared to children with overweight.<sup>49</sup> In addition, rats treated with a formula that effectively alleviated type 2 diabetes had higher abundance of gut *Senegalimassilia*.<sup>50</sup> Interestingly, although metformin treatment increases *Senegalimassilia* abundance in young healthy men<sup>47</sup>; in our study, the combination of metformin with spironolactone and pioglitazone failed to restore *Senegalimassilia* abundance in girls with PCOS.

Finally, we disclosed that the dysbiosis of the gut microbiota observed in girls with PCOS was explained by the FAI and by hepatovisceral fat, suggesting that both hyperandrogenism and central adiposity may alter the gut microbiota, which in turn, could impact the pathophysiology of PCOS. Moreover, we showed that the variation of gut microbiota in OC-treated girls was explained by hepatic fat accumulation; hepato-visceral fat remained unchanged on OCs and was significantly reduced after SPIOMET. These results corroborate the notion that targeting a reduction of ectopic fat normalizes the PCOS phenotype including a recovery of the gut microbiota, represented here by the normalization of *Family XI* relative abundance. Further studies are needed to determine whether the changes in microbiota composition observed in girls with PCOS persist after therapy discontinuation and may, in turn, impact androgen metabolism.

Study limitations include the relatively low number of participants, the absence of an untreated group of girls with PCOS, the absence of a follow-up period after therapy discontinuation, and the lack of analysis of ceramide serum concentrations due to additional sample unavailability. Study strengths include the strict inclusion and exclusion criteria, the longitudinal design, and the assessment of the impact of two interventions.

In summary, our study demonstrates a dysbiosis of gut microbial community in girls with PCOS without obesity, including a reduction of alpha diversity, differences in community structure, a reduction of *Prevotellaceae, Prevotella* and *Senegalimassilia* and an increase of *Bacillales Family XI*. SPIOMET, a medication targeting hepato-visceral (central) fat normalized the PCOS phenotype and the abundance of *Family XI*.

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### CONFLICTS OF INTERESTS

The authors declare no conflicts of interest.

### AUTHOR CONTRIBUTIONS

Cristina Garcia-Beltran researched data, contributed to study design and data interpretation, wrote the manuscript, and reviewed/edited 10 of 11 WILEY

the manuscript. Rita Malpique researched data, contributed to study design and data interpretation, and reviewed/edited the manuscript. Belen Carbonetto and Pedro González-Torres performed the bioinformatics and draft part of the manuscript. Desirée Henares, Pedro Brotons and Carmen Muñoz-Almagro contributed to data interpretation and reviewed/edited the manuscript. Abel López-Bermejo contributed to data interpretation and reviewed/edited the manuscript. Francis de Zegher contributed to study design and data interpretation and reviewed/edited the manuscript. Lourdes Ibáñez contributed to study design and data interpretation, wrote the manuscript, and reviewed/edited the manuscript.

### DATA AVAILABILITY STATEMENT

The datasets generated and/or analysed during the current study have been archived in a public repository (BioProject ID: PRJNA659664) and are accessible at: http://www.ncbi.nlm.nih.gov/bioproject/659664.

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### SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section at the end of this article.

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### Gut Microbiota in Non-Obese Adolescent Girls with Polycystic Ovary Syndrome: Effects of Randomized Treatments

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# Figure S2. Results of electrophoresis







**Table S1.** Study variables in control girls (N= 31) and in girls with polycystic ovary syndrome. Girls with PCOS were randomized to receive oral contraception (OC, Ethinylestratiol 20 µg plus Levonorgestrel 100 mg) or a low-dose combination of Spironolactone (50mg/ d), Pioglitazone (7.5 mg/ d) and Metformin (850 mg/ d) (SPIOMET) for 1 year. Baseline data are presented for N= 30 girls with PCOS. Longitudinal data are shown from N= 23 girls with PCOS (N= 12 on OC; N= 11 on SPIOMET) with available stool samples for microbiota analysis at baseline and after 1 year.

	Controls	PCOS		OC (N= 12)			SPIOMET (N= 11)	
	(N= 31)	(N= 30)	Start <sup>a</sup>	1 year	Δ 0-1 year	Start <sup>a</sup>	1 year	Δ 0-1 year
Auxology								
Age (years)	$15.9 \pm 0.2$	$15.8 \pm 0.3$	$15.6 \pm 0.5$			15.6 ± 0.4		ı
Birth weight Z-score	$0.1 \pm 0.2$	-0.6 ± 0.2 **	-0.8 ± 0.3	ı	ı	-0.5 ± 0.3	ı	ı
BMI (kg/ m²)	22 ± 0	$25 \pm 1$ **	$25 \pm 1$	26±1	$1.6 \pm 0.4$	$24 \pm 1$	$24 \pm 1$	-0.2 ± 0.9
BMI Z-score	$0.1 \pm 0.2$	$1.1 \pm 0.2$	$1.0 \pm 0.3$	$1.5 \pm 0.4$	$0.5 \pm 0.1$	$0.9 \pm 0.4$	0.8 ± 0.4 <sup>b</sup>	-0.1 ± 0.5
Δ Z-score birth weight - BMI	-0.1±0.2	$1.6 \pm 0.3$	$1.7 \pm 0.4$	ı	-	$1.4 \pm 0.6$	ı	I
Endocrine – Metabolic Variables								
Testosterone (nmol/L)	$1.0 \pm 0.1$	$1.3 \pm 0.1$	$1.3 \pm 0.3$	0.9 ± 0.1 <sup>h</sup>	-0.4 ± 0.2	1.3 ± 0.2	$1.0 \pm 0.2^{h}$	-0.3 ± 0.2
SHBG (nmol/ L)	$65 \pm 4$	32 ± 3 ***	35 ± 5	58±9 <sup>1</sup>	24±6	33 ± 4	32 ± 4 °	-2 ± 2 <sup>g</sup>
FAI	$1.7 \pm 0.2$	$6.4 \pm 0.8$	$6.8 \pm 1.2$	$3.0 \pm 1.1^{10}$	-3.8±1.1	$6.1 \pm 1.6$	3.9±0.4	-2.2 ± 1.3
Fasting insulin (pmol/ L)	63±5	70 ± 8	67 ± 12	77 ± 9	$11 \pm 10$	55 ± 8	39 ± 6 °	-16 ± 10 <sup>e</sup>
HOMA-IR	$2.0 \pm 0.2$	$2.1 \pm 0.3$	$2.0 \pm 0.4$	2.2 ± 0.2	0.2±0.3	$1.7 \pm 0.2$	$1.1 \pm 0.2^{\circ}$	-0.6±0.3 e
0GTT								
Mean Glycemia Z-score	I	$0.1 \pm 0.0$	$0.1 \pm 0.0$	$0.1 \pm 0.1$	$0.1 \pm 0.1$	$0.1 \pm 0.1$	$0.0 \pm 0.1$	-0.1 ± 0.1
Mean Insulinemia Z-score	I	$3.1 \pm 0.5$	$3.1 \pm 0.8$	2.9 ± 0.7	-0.3±0.9	$2.5 \pm 1.0$	$0.9 \pm 0.6$	-1.6±0.6
ALT (µkat/ L)	$0.27 \pm 0.01$	$0.25 \pm 0.02$	$0.24 \pm 0.02$	0.35 ± 0.05 <sup>h</sup>	0.10 ± 0.05	0.27 ± 0.03	$0.24 \pm 0.03$	-0.03 ± 0.02 °
AST (µkat/ L)	$0.32 \pm 0.01$	$0.29 \pm 0.02$	0.27 ± 0.02	$0.28 \pm 0.02$	$0.01 \pm 0.01$	$0.34 \pm 0.04$	$0.28 \pm 0.02$	-0.05 ± 0.07
GGT (μkat/ L)	$0.22 \pm 0.01$	$0.24 \pm 0.01$	0.23 ± 0.02	0.30 ± 0.04 <sup>h</sup>	0.06±0.03	$0.23 \pm 0.01$	0.20±0.01 <sup>bh</sup>	-0.03 ± 0.01 <sup>f</sup>
HDL-cholesterol (mmol/ L)	$1.4 \pm 0.0$	$1.3 \pm 0.1$	$1.4 \pm 0.1$	$1.4 \pm 0.1$	$0.0 \pm 0.1$	$1.4 \pm 0.1$	$1.4 \pm 0.1$	$0.1 \pm 0.1$
LDL-cholesterol (mmol/ L)	$2.3 \pm 0.1$	$2.2 \pm 0.1$	$2.3 \pm 0.2$	2.8 ± 0.2 <sup>h</sup>	0.5±0.2	2.1 ± 0.2	2.1±0.2 <sup>b</sup>	0.0 ± 0.1 <sup>e</sup>
Triglycerides (mmol/ L)	$0.6 \pm 0.0$	$0.7 \pm 0.0$	$0.7 \pm 0.1$	$0.8 \pm 0.1$	$0.1 \pm 0.1$	$0.7 \pm 0.1$	$0.8 \pm 0.1$	$0.1 \pm 0.1$
HMW-adiponectin (mg/ L)	7.4 ± 0.7	$4.7 \pm 0.6$	$5.3 \pm 1.0$	3.9 ± 0.5	-1.3±0.8	4.7 ± 0.9	7.1 ± 1.0 <sup>b</sup>	$2.4 \pm 1^{f}$
usCRP (nmol/ L)	$6.4 \pm 1.0$	$15.4 \pm 3.5$	$11.0 \pm 3.1$	25.6±6.4 <sup>h</sup>	14.5±5.9	$14.9 \pm 4.4$	8.1±3.5 <sup>b</sup>	-6.8±4.1 <sup>f</sup>
Ovulatory function ${}^{*}$								
Total cycles	ı	ı	ı	3 ± 0.4	ı	I	6 ± 0.2 #	I
Total ovulations			I	2 ± 0.5	·	ı	4 ± 0.5 #	I
Body composition (DXA) <sup>‡</sup>								
Bone mineral density (g/ cm <sup>2</sup> )	I	$1.21 \pm 0.02$	$1.18 \pm 0.04$	$1.18 \pm 0.04$	0.00 ± 0.01	1.21 ± 0.02	$1.19 \pm 0.03$	-0.02 ± 0.01
Lean mass (kg)	ı	$37 \pm 1$	37 ± 2	38±2	$1 \pm 1$	$36 \pm 1$	$37 \pm 1$	$1 \pm 1$
Fat mass (kg)	ı	25 ± 2	23 ± 2	27 ± 2 <sup>j</sup>	$3\pm 1$	24 ± 3	23 ± 3	0±2
Abdominal fat (kg)		$6.5 \pm 0.4$	$6.4 \pm 0.6$	7.2 ± 0.7 <sup>j</sup>	0.8±0.3	$6.1 \pm 0.7$	5.9±0.7	-0.2 ± 0.4 <sup>e</sup>
Abdominal fat partioning (MRI)								
Subcutaneous fat (cm <sup>2</sup> )	$98 \pm 10$	205 ± 24	194 ± 35	220 ± 35 <sup>h</sup>	26±10	194 ± 51	181 ± 42	-13 ± 28 <sup>e</sup>
Visceral fat (cm <sup>2</sup> )	29±1	40 ± 4	37 ± 5 <sup>j</sup>	43 ± 5	6±5	32 ± 3	$34 \pm 4$	2 ± 4
Hepatic fat (%)	$11 \pm 1$	$17 \pm 1$	$16 \pm 1$	18±2	2±3	$19 \pm 1$	$10 \pm 1^{dj}$	-9±1ª
Central (hepato-visceral) fat	39 ± 2	57 ± 4 **	53 ± 5	62 ± 5	9±6	51 ± 3	45 ± 5 °	-7±4°

Values are mean ± SEM.

BMI, body mass index; SHBG, sex hormone-binding globulin; FAI, free androgen index; HOMA-IR, homeostasis model assessment insulin resistance; oGTT, oral glucose tolerance test; ALT, alanine transaminase; AST, aspartate transaminase; GGT, gamma-glutamyl transpeptidase; HDL, high-density lipoprotein; LDL, low-density lipoprotein; HMW, high molecular weight; usCRP, ultrasensitive C-reactive protein; DXA, dual X-ray absorptiometry; MRI, magnetic resonance imaging.

\* Total numbers of cycles and ovulations over two periods of 12 consecutive weeks, during the second and fourth trimesters of the post-treatment year.

<sup>±</sup> Indicative DXA values in healthy adolescents, matched for age and height (N= 41): lean mass 35.1 ± 1.0 kg; fat mass 17.6 ± 1.4 kg (6).

\* P≤ 0.05; \*\* P≤ 0.01 and \*\*\* P≤ 0.001 between controls and girls with PCOS at baseline.

<sup>a</sup> no significant differences between randomized subgroups at start.

<sup>b</sup>  $P \le 0.05$ , <sup>c</sup>  $P \le 0.01$  and <sup>d</sup>  $P \le 0.001$  for differences between subgroups at 1 year. <sup>e</sup>  $P \le 0.05$ , <sup>f</sup>  $P \le 0.01$  and <sup>g</sup>  $P \le 0.001$  for differences between subgroup changes ( $\Delta$ ) 0-1 year.

 $h \ge 0.05$ ,  $h \le 0.01$  and  $h \ge 0.001$  for differences within-subgroup changes from start.

 $^{\#}$  P< 0.0001 between subgropups differences in ovulatory function.

*P* values for differences between controls and girls with PCOS at baseline and between subgroups are adjusted for BMI.

Table S2. Number of reads and relative abundance observed, historical and theoretical of the mock community.

Bacteria; Firmicutes; Bacilli; Lactobacillales; Lactobacillaceae; Lactobacillus32187Bacteria; Firmicutes; Bacilli; Bacillales; Staphylococcaceae; Staphylococcus22727Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales; Enterobacteriaceae21841	20.5 14 5	21.0 14.3	18.4 15.5
Bacteria; Firmicutes; Bacilli; Bacillales; Staphylococcaceae; Staphylococcus Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales; Enterobacteriaceae 21841	14 F	14.3	15.5
Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales; Enterobacteriaceae	0:1-1		
	13.9	13.8	10.4
Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales; Pseudomonadaceae; Pseudomonas 19245	12.1	12.4	17.4
Bacteria; Firmicutes; Bacilli; Bacillales; Bacillaceae; Bacillus 18965	12.2	11.0	4.2
Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales; Enterobacteriaceae; Escherichia-Shigella 16116	10.1	10.5	14.1
Bacteria; Firmicutes; Bacilli; Bacillales; Listeriaceae; Listeria	10.3	10.4	10.1
Bacteria; Firmicutes; Bacilli; Lactobacillales; Enterococcaceae; Enterococcus	6.4	6.6	9.9
Others 256	0.2	0.4	ı

<sup>1</sup> Mean of the historical composition observed according to Microomics (http://www.microomics.eu/)

<sup>2</sup> Theoretical composition according to the manufacturer (Zymobiomics Microbial Community DNA, ZymoResearch, Irvine, USA)

Таха	Number of reads in the NC	Average of the number of reads in the samples	Ratio samples/ NC
Actinobacteria; Actinobacteria; Micrococcales; Microbacteriaceae; Curtobacterium	51	0	0
Bacteroidetes; Bacteroidia; Bacteroidales; Porphyromonadaceae; Porphyromonas	87	25.388	0.292
Firmicutes; Clostridia; Clostridiales; Ruminococcaceae; [Eubacterium] coprostanoligenes group	81	4097.365	50.585
Proteobacteria; Alphaproteobacteria; Caulobacterales; Caulobacteraceae, unassigned genus	1266	3.447	0.003
Proteobacteria; Alphaproteobacteria; Rhizobiales; Rhizobiaceae; Mesorhizobium	745	3.412	0.005
Proteobacteria; Alphaproteobacteria; Rhizobiales; Rhizobiales Incertae Sedis; Phreatobacter	56	0	0
Proteobacteria; Alphaproteobacteria; Rhizobiales; Xanthobacteraceae, unassigned genus	374	0	0
Proteobacteria; Alphaproteobacteria; Rhizobiales; Xanthobacteraceae; Bradyrhizobium	11141	37.047	0.003
Proteobacteria; Alphaproteobacteria; Sphingomonadales; Sphingomonadaceae; Sphingomonas	5578	28.812	0.005
Proteobacteria; Deltaproteobacteria; Bdellovibrionales; Bdellovibrionaceae; Bdellovibrio	33	0	0
Proteobacteria; Gammaproteobacteria; Betaproteobacteriales; Burkholderiaceae; Curvibacter	62	0	0
Proteobacteria; Gammaproteobacteria; Betaproteobacteriales; Burkholderiaceae; Pelomonas	155	0.082	0.001
Proteobacteria; Gammaproteobacteria; Betaproteobacteriales; Burkholderiaceae; Ralstonia	1236	5.624	0.005
Proteobacteria; Gammaproteobacteria; Enterobacteriales; Enterobacteriaceae; Escherichia-Shigella	203	1107.812	5.457
Proteobacteria; Gammaproteobacteria; Enterobacteriales; Enterobacteriaceae; Serratia	206	0.306	0.001
Proteobacteria; Gammaproteobacteria; Pseudomonadales; Pseudomonadaceae; Pseudomonas	905	0.106	0.000
Proteobacteria; Gammaproteobacteria; Xanthomonadales; Xanthomonadaceae; Stenotrophomonas	787	5.588	0.007

Table S3b. Taxonomic assignment at the amplicon sequence variants (ASV) level of abundant groups.

Taur	A CV/ I dow+ifinotion			Sample Ide	entification	
I GYG			R0002	R0026	R0027	R0067
[Eubacterium] coprostanoligenes group	da2535464b1eb41b5c616b4ea1fd8645	81		ı	-	15
Enterobacteriaceae;Escherichia-Shigella	c6f0efea3a269137720ca341ec873169	203	186	30	113	I

Table S3a. Taxonomic assignment of the reads at the genus level of the negative control (NC)

**STUDY 4** 

## RAISED THYROID-STIMULATING HORMONE IN GIRLS WITH POLYCYSTIC OVARY SYNDROME: EFFECTS OF RANDOMIZED INTERVENTIONS

<u>Cristina Garcia-Beltran</u>, Judit Bassols, Gemma Carreras-Badosa, Abel López-Bermejo, Lourdes Ibáñez, Francis de Zegher

Hormone Research in Paediatrics. 2023; 96(5), 458-464

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### STUDY 4

# Raised thyroid-stimulating hormone in girls with polycystic ovary syndrome: effects of randomized interventions

**Introduction:** Polycystic ovary syndrome (PCOS) in women associates with raised levels of circulating thyroid-stimulating hormone (TSH) and with high rates of gestational complications. A low range of preconception TSH is followed by low rates of gestational complications. It is unknown whether TSH levels are elevated in adolescents with PCOS and, if so, whether traditional or exploratory treatments can lower them into safe preconception range.

**Objective**: To investigate TSH in non-obese adolescents with PCOS, including the effects of randomized interventions.

**Material and methods:** Morning TSH was a safety marker in randomized pilot studies comparing the effects of an oral contraceptive (OC) versus those of a low-dose combination of spironolactone-pioglitazone-metformin (SPIOMET) in non-obese adolescents with PCOS. A post hoc analysis compared TSH levels in PCOS (N=62) versus controls, TSH changes on treatment (for 1 year), and TSH levels post treatment (for 1 year).

**Results:** Mean TSH levels were higher in PCOS patients than in control girls (P<0.01). On treatment TSH levels diverged (P<0.001), remaining elevated on OC, and descending swiftly on SPIOMET, well into safe preconception range. Post treatment TSH levels were stable in both subgroups. On treatment changes of circulating TSH associated to those of liver fat (R=0.307, P=0.017).

**Conclusion:** The endocrine signature of early PCOS is herewith extended to include modestly raised levels of circulating TSH; the normalizing effects of SPIOMET intervention in non-obese adolescents with PCOS are herewith extended to include on and post treatment TSH.

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Hormone Research in Paediatrics

### **Research Article**

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# Raised Thyroid-Stimulating Hormone in Girls with Polycystic Ovary Syndrome: Effects of Randomized Interventions

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### Keywords

Polycystic ovary syndrome · Thyroid-stimulating hormone · Spironolactone · Pioglitazone · Metformin

### Abstract

**Introduction:** Polycystic ovary syndrome (PCOS) in women associates with raised levels of circulating thyroidstimulating hormone (TSH) and with high rates of gestational complications. A low range of preconception TSH is followed by low rates of gestational complications. It is unknown whether TSH levels are elevated in adolescents with PCOS and, if so, whether traditional or exploratory treatments can lower them into safe preconception range. We investigated TSH in nonobese adolescents with PCOS, including the effects of randomized interventions. **Methods:** Morning TSH was a safety marker in randomized pilot studies comparing the effects of an oral contraceptive (OC) versus those of a low-dose combination of spironolactonepioglitazone-metformin (SPIOMET) in nonobese adolescents with PCOS. A post hoc analysis compared TSH levels in PCOS (N = 62) versus controls, TSH changes on treatment (for 1 year), and TSH levels posttreatment (for 1 year). **Results:** Mean TSH levels were higher in PCOS patients than in control girls (p < 0.01). On-treatment TSH levels diverged (p < 0.001), remaining elevated on OC, and descending swiftly on SPIOMET, well into safe preconception range. Posttreatment TSH levels were stable in both subgroups. On-treatment changes of circulating TSH associated to those of liver fat (R = 0.307, p = 0.017). **Conclusion:** The endocrine signature of early PCOS is herewith extended to include modestly raised levels of circulating TSH; the normalizing effects of SPIOMET intervention in nonobese adolescents with PCOS are herewith extended to include on- and posttreatment TSH.

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### Introduction

Polycystic ovary syndrome (PCOS) is the most common cause of hirsutism and menstrual irregularity in adolescent girls and young women [1, 2]. Adolescent PCOS is nowadays thought to be essentially elicited by ectopic fat and insulin resistance [2, 3]. The diagnostic criteria of adolescent PCOS include the copresence of androgen excess (clinical/biochemical) and menstrual irregularity (oligo-amenorrhea) >2 years beyond menarche [1]. An early diagnosis of PCOS has implications for a woman's long-term health [4-8]. For example, the incidence of gestational complications is elevated in women with PCOS, and this includes a twofold risk for preterm delivery [9]. Recently, the lowest risk for preterm delivery has been strongly linked to a preconception concentration of circulating thyroid stimulating hormone (TSH) between 0.91 and 1.82 mIU/L [10]; the middle of this range happens to be the same as the recently reported median TSH concentration (1.35 mIU/L) in nonobese adolescent girls aged 16 years [11]. These observations open the perspective of including not only a normal TSH level for chronological age but also a safe level of preconception TSH among the criteria whereby adolescent PCOS treatments could henceforth be gauged.

There is no approved treatment for adolescent PCOS. International guidelines recommend silencing the gonadotropic axis with an oral estro-progestogen contraceptive (OC) [12]. An alternative approach (in nonobese girls who are not sexually active) is to give a low-dose combination of spironolactone, pioglitazone, and metformin (SPIOMET) which appears to be well tolerated, to reduce hirsutism and androgen excess effectively and to exert more normalizing effects than an OC on key outcomes of longer term relevance such as ectopic fat, insulin resistance, and posttreatment ovulation rate [13, 14]. The mechanisms contributing to SPIOMET's normalizing actions include an activation of brown adipose tissue by low-dose spironolactone [15], a doubling of high molecular weight adiponectinemia by low-dose pioglitazone [16], and a tripling of circulating growth and differentiation factor-15 (GDF15) by low-dose metformin [17].

The concentrations of circulating TSH are known to be relatively high in women with PCOS [18–21]. It is unknown whether TSH levels are already elevated in adolescent girls with PCOS and, if so, whether any recommended or exploratory treatments can lower them into the safest range of preconception TSH. In the aforementioned OC-versus-SPIOMET studies [13, 14], TSH was a [hitherto unreported] safety marker; here we report a post hoc analysis of pre-, on-, and posttreatment TSH data in nonobese adolescent girls with PCOS.

### **Materials and Methods**

### Study Population and Design

A total of 62 nonobese girls with PCOS (mean age 15.8 year; mean body mass index [BMI] 24.2 kg/m<sup>2</sup>; and 32% overweight [22]) participated in two randomized (1:1) trials (ISRCTN29234515; ISRCTN11062950) that compared the on-treatment (over 1 year) and posttreatment (over 1 year) effects of OC-versus-SPIOMET (online suppl. Fig. 1; for all online suppl. material, see www.karger.com/doi/ 10.1159/000529183), as reported [13, 14]. In summary, both openlabel pilot studies were performed in the Endocrinology Department at Sant Joan de Déu University Hospital in Barcelona, Spain. Obesity was not an exclusion criterion but, in this hospital, adolescents with obesity are mostly referred to the obesity unit [14]. Inclusion criteria were hirsutism (modified Ferriman-Gallwey score >8), biochemical androgen excess (serum testosterone >45 ng/dL and/or free androgen index >3.5), oligomenorrhea (menstrual intervals >45 days), gynecological age >2.0 years, and the absence of sexual intercourse. Exclusion criteria were 21-hydroxylase deficiency; glucose intolerance or diabetes; evidence of liver or kidney dysfunction; hyperprolactinemia; prior use of medications affecting gonadal/adrenal function or carbohydrate/lipid metabolism; and circulating TSH <0.3 or >4.5 mIU/ L. The latter criterion ended up excluding none of the study candidates.

All study participants received standardized instructions for a Mediterranean diet and regular exercise; OC treatment consisted of 20 µg ethinylestradiol plus 100 mg levonorgestrel (21/28 days) and placebo (7/28 days); SPIOMET treatment consisted of the concomitant intake of spironolactone 50 mg, pioglitazone 7.5 mg, and metformin 850 mg in separate tablets, once daily at dinner time [13, 14]. A total of 29 asymptomatic girls (mean age 15.8 year; mean BMI 21.6 kg/m<sup>2</sup>; and 11% overweight [22]) were recruited in nearby schools to serve as controls; all had regular menses and a gynecological age >2.0 years; none was hirsute, had chronic disease, or took medication. Blood sampling in patients and controls was performed in fasting state in the early morning, thereby reducing the variability related to the circadian rhythm of circulating TSH [23]. The sampling was scheduled in the follicular phase (days 3–7) of the cycle or after 2 months of amenorhea.

### Assessments

Circulating TSH was a pre-, on-, and posttreatment safety marker in the OC-versus-SPIOMET studies [13, 14] and was measured by chemiluminescent microparticle immunoassay (CMIA, ARCHITECT i200, Abbott Diagnostics, IL, USA). Birth weight and BMI (and their Z-scores) were retrieved from hospital records. Serum insulin and sex hormone-binding globulin (SHBG) were measured by immunochemiluminiscence (IMMULITE 2000, Diagnostic Products, Los Angeles, CA). The intra- and inter-assay coefficients of variation were <0.1% and 7.2%, respectively, for insulin and <0.5% and 8%, respectively, for SHBG; the lower detection limit was 1.4 pmol/L for insulin and 1.05 nmol/L for SHBG. Circulating testosterone was measured by liquid chromatography-tandem mass spectrometry, as described [13, 14]; intra- and inter-assay coefficients of variation were 8% and 12%, respectively, and the lower detection limit was 0.002 nmol/L. Free androgen index and ultrasensitive C-reactive protein (usCRP) were assessed as previously reported [13, 14]. Standard oral glucose tolerance tests with 75 g of glucose were performed after an overnight fast; blood was sampled before and 30, 60, 90, and 120 min after glucose intake for glucose, mean serum insulin, and their Z-scores were derived [13, 14]. Body composition was assessed by dual X-ray absorptiometry with a Lunar Prodigy and Lunar software (version 3.4/3.5; Lunar Corp, Madison, WI, USA). Abdominal fat partitioning (subcutaneous and visceral) and liver fat were assessed by magnetic resonance imaging (MRI) using multiple-slice MRI 1.5 Tesla scan (Signa LX Echo Speed Plus Excite, General Electric, Milwaukee, WI, USA), as described [13, 14].

### Statistics and Ethics

Statistical analyses were performed with SPSS version 27.0 (SPSS software, IBM, Armok, NY, USA). Longitudinal changes in quantitative variables between groups were compared by general linear model for repeated measures. Differences in longitudinal changes between subgroups were tested by the interaction among between- and within-subject effects. Associations were sought by Pearson correlation analysis. p < 0.05 was considered significant. Data are presented as mean  $\pm$  standard error of the mean.

The OC-versus-SPIOMET studies were conducted after approval by the Institutional Review Board of Sant Joan de Déu Hospital, after written informed consent by parents, and after assent by each participating girl. All methods followed the relevant guidelines and regulations.

### Results

Key TSH results, including individual end-of-treatment data, are depicted in Figure 1; statistical comparisons are shown in Table 1.

### Controls versus PCOS

As expected, nonobese PCOS patients had a lower birth weight and higher BMI and were centrally more adipose than controls [2]. Mean TSH levels in nonobese girls with PCOS were similar in the two randomized subgroups at baseline and were 27% higher than those in controls (p < 0.01; Table 1).

### OC-versus-SPIOMET Treatment in PCOS Girls

Figure 1 (panel a) shows that on-treatment TSH levels diverged: on OC, they remained relatively high and on SPIOMET, they descended within 6 months into safe preconception range; posttreatment TSH levels remained stable, respectively, above and inside safe preconception range. Figure 1 (panel b) shows that the upward shift of TSH concentrations in individual PCOS patients was still present after 1 year on OC treatment but was essentially absent after SPIOMET treatment. On-treatment changes in TSH did not associate to changes in circulating androgens or in total body weight, lean mass, or fat mass but paralleled those in liver fat (R = 0.307, p = 0.017).

### Discussion

TSH – but not free T3 or free T4 – was a safety marker in OC-versus-SPIOMET pilot studies performed in nonobese adolescent girls with PCOS [13, 14]. A post hoc analysis of these TSH data was prompted by the recent description of relatively low TSH reference intervals in nonobese adolescent girls [11] and by the even more recent delineation of the safest range of preconception TSH [10]. In the present report, novel findings include that (1) TSH levels in nonobese adolescents with PCOS were already shifted upward, on average by 27%; (2) this upward TSH shift did persist on/posttreatment with OC but disappeared on/posttreatment with SPIOMET, so that TSH levels entered swiftly into the safest preconception range and remained stable inside this range; and (3) on-treatment changes in TSH associated most closely to those in liver fat (by MRI).

The present pre-, on-, and posttreatment observations corroborate the hypothesis that a TSH elevation in an adolescent girl with PCOS may in essence be no more than an epiphenomenon that points to the presence of insulin resistance within a context of lipid accumulation in ectopic depots. TSH elevations have previously been described as epiphenomena of insulin resistance and/or ectopic fat in a variety of study populations including overweight children and adolescents and also nonobese and obese women with PCOS [18–21, 24–31].

The on-treatment data challenge the concept that TSH elevations in PCOS relate primarily to androgen excess [18] or body weight [19, 25, 26] given that the tested treatments had virtually parallel effects on androgenemia and body weight while having markedly divergent effects on TSH, insulin resistance, and ectopic fat. The post-treatment data in PCOS patients strengthen the notion that SPIOMET has the capacity to confer broadly normalizing benefits that persist beyond its intake and that are herewith extended to include a lowering of TSH concentrations into the safest preconception range. These surprisingly prolonged effects are incompletely explained by factors such as CXCL14 [15], adiponectin [16], GDF15 [17], miRNA profiles, [32] and gut microbiota [33] and thus remain to be further clarified.

There is still no evidence suggesting that the intake of spironolactone and/or pioglitazone has TSHlowering effects. In contrast, metformin is known to exert **Table 1.** Selected study variables in control girls (*N* = 29) and in girls with PCOS (*N* = 62) who were randomized to receive an OC (*N* = 31) or a low-dose combination of SPIOMET (N = 31) for 12 months and who remained untreated during the subsequent 12 months

	Controls	PCOS	OC (N = 3	(1					SPIOMET (I	V = 31)				
			baseline <sup>a</sup>	6 months	12 months	24 months	$\Delta$ 0–12 months	$\Delta$ 12–24 months	baseline	6 months	12 months	24 months	$\Delta$ 0–12 months	∆ 12–24 months
Birth weight Z-score	0.2±0.2	-0.6±0.1***	-0.6±0.2	I	I	I	1	I	-0.6±0.2	1	1	1	ı	ı
Age at menarcne, year. Age, year	15.8±0.1	11.0±0.1	11.0±0.1	1 1 1	<u>-</u> - - - 1 -	1 1	1 1		11.0±0.2 15.8±0.3	1 1	1 1	1 1	- - - -	
BMI Z-score	0.1±0.2 0.2±0.3	0.8±0.1** 1.4±0.2***	0.9±0.2 1.5±0.3	0.9±0.2 -	1.1±0.2 <sup>19</sup> −	1.2±0.25 -	0.2±0.1 -	0.1±0.1 -	0.8±0.2 1.4±0.3	0.8±0.2 -	0.7±0.2 -	0.8±0.2 -	−0.1±0.1ª	0.1±0.1
TSH, mIU/L Testosterone. nmol/L	$1.58\pm0.12$ $0.9\pm0.1$	2.00±0.09** 1.8+0.1***	1.85±0.12 1.9±0.1	1.95±0.13 1.0±0.1 <sup>f</sup>	2.00±0.16 1.0±0.1 <sup>f</sup>	2.02±0.14 1.6±0.2	0.14±0.13 −0.9±0.1	0.02±0.16 0.7±0.2	2.14±0.12 1.7±0.1	1.49±0.09 <sup>e, f</sup> 1.1±0.1 <sup>f</sup>	1.45±0.11 <sup>e, f</sup> 1.1±0.1 <sup>f</sup>	1.51±0.10 <sup>e, f</sup> 1.3±0.1 <sup>c</sup>	$-0.65\pm0.12^{9}$ $-0.6\pm0.1$	0.05±0.11 0.2+0.1 <sup>d</sup>
SHBG, nmol/L	64±5	30±2***	31.2±2.4	64.2±5.8 <sup>f</sup>	60.6±4.9 <sup>f</sup>	32.5±2.9	29.4±4.2	-28.1±4.5	29.6±2.0	30.4±1.9 <sup>h</sup>	31.9±2.2 <sup>h</sup>	38.6±2.9℃	2.3±2.2 <sup>9</sup>	6.7±2.09
FAI OGTT	1.7±0.2	6.8±0.5***	<b>6.9±0.6</b>	1.8±0.2 <sup>f</sup>	2.3±0.4 <sup>f</sup>	<b>6.4±0.9</b>	-4.6±0.6	<b>4.1±0.8</b>	6.7±0.7	4.2±0.4 <sup>h, f</sup>	3.2±0.3 <sup>f</sup>	4.1±0.5 <sup>c, i</sup>	-3.5±0.7	0.9±0.49
Mean glycemia Z-score	I	0.18±0.03	0.13±0.03	I	0.17±0.04	0.13±0.04	0.04±0.04	-0.04±0.04	0.23±0.05	I	0.09±0.04 <sup>c</sup>	0.13±0.04	−0.14±0.05 <sup>j</sup>	0.05±0.04
Mean insulinemia	I	3.19±0.31	3.54±0.45	I	3.75±0.53	3.01±0.48	0.21±0.46	-0.74±0.54	2.83±0.43	I	0.65±0.25 <sup>h, f</sup>	0.61±0.19 <sup>h, f</sup>	-2.18±0.31 <sup>g</sup>	-0.04±0.23
z-score usCRP, nmol/L	6.0±1.0	13.2±1.6**	11.8±1.8	23.1±3.3°	25.1±3.9°	17.8±3.9	13.2±4.1	-7.3±5.2	14.6±2.7	7.6±2.1 <sup>h, f</sup>	6.8±1.3 <sup>h, c</sup>	6.5±0.8 <sup>c, e</sup>	-7.9±2.49	-0.3±2.4
Body composition (DX/ Bone mineral		1.19±0.01	1.18±0.02	1.19±0.02	1.19±0.02	1.21±0.02	0.00±0.01	0.02±0.01	1.19±0.02	1.18±0.02	1.19±0.02	1.21±0.02	0.00±0.01	0.02±0.01
density, g/cm² Lean mass, kg	I	35.6±0.6	35.7±0.8	35.8±0.8	36.4±1.0	36.5±1.0 <sup>b</sup>	0.7±0.4	0.2±0.3	35.5±0.9	35.6±0.7	35.6±0.8	36.1±0.8	0.1±0.3	0.5±0.3
Fat mass, kg	I	22.1±1.0	21.8±1.4	22.3±1.3	23.2±1.5°	23.4±1.6 <sup>b</sup>	1.4±0.5	0.3±0.6	22.4±1.6	22.5±1.6	22.5±1.4	22.1±1.7	0.1±0.8	-0.3±0.6
Abdominal fat, kg Abdominal fat nartition	- MRI)	5.9±0.2	6.0±0.3	<b>6.0±0.3</b>	6.0±0.4	<b>6.</b> 3±0.4	0.1±0.2	0.4±0.2	5.9±0.4	5.7±0.4	5.7±0.3	5.7±0.4	−0.3±0.2	0.0±0.1
Subcutaneous	98±8	174±14*	169±18	171±18	184±19	180±20	15±9	-5±13	179±21	172±19	171±19	167±23	-7±11	-4±9
fat, cm² Visceral fat, cm²	29±1	43±2***	41±3	<b>4</b> 3±4	45±4	39±3	4±3	-5±3	44±3	37±2 <sup>b</sup>	35±2 <sup>b, i</sup>	36±3 <sup>b</sup>	-9±4i	1±2
Liver fat, %	11±1	17土1***	17±1	20±1	20±1	17±1	3±1	-2±1	18±1	12±1 <sup>h, f</sup>	10±1 <sup>h, f</sup>	10±1 <sup>h, f</sup>	-8±1 <sup>9</sup>	0±1
Values are me BMI. bodv mass i	an ± stand dex: SHB(	ard error of 3. sex hormo	the mea bino	n. PCOS, I ling alobi	polycystik ulin: FAL 1	c ovary sy ree andro	ndrome; O( agen index:	C, oral contraction of the contr	ceptive; S ucose tol	PIOMET, sl erance tes	pironolact t: TSH. thv	one, piogli roid-stimu	itazone, and Ilating horn	I metformin

ultrasensitive C-reactive protein; DXA, dual X-ray absorptiometry; MRI, magnetic resonance imaging. \* $p \le 0.05$ . \*\* $p \le 0.01$ . \*\*\* $p \le 0.001$  between controls and girls with PCOS at baseline. <sup>a</sup>No significant differences between randomized subgroups at start. <sup>b</sup> $p \le 0.05$ . <sup>c</sup> $p \le 0.01$ . <sup>d</sup> $p \le 0.05$ . <sup>e</sup> $p \le 0.01$ . <sup>f</sup> $p \le 0.001$  for differences within subgroup changes from baseline.  $^{9}p \le 0.001$  for differences between subgroup changes ( $\Delta$ ) 0–12 months and 12–24 months.  $^{h}p \le 0.001$  for differences between subgroups at 6, 12, and 24 months.  $^{h}p \le 0.001$  for differences between subgroups at 6, 12, and 24 months.  $^{h}p \le 0.001$  for differences between subgroups at 6, 12, and 24 months.  $^{h}p \le 0.001$  for differences between subgroups at 6, 12, and 24 months.  $^{h}p \le 0.051$   $^{h}p \le 0.001$  for differences between subgroups at 6, 12, and 24 months.  $^{h}p \le 0.001$  for differences between subgroups at 6, 12, and 24 months.  $^{h}p \le 0.051$   $^{h}p \le 0.001$  for differences between subgroups at 6, 12, and 24 months.  $^{h}p \le 0.001$  for differences between subgroups at 6, 12, and 24 months.  $^{h}p \le 0.001$  for differences between the subgroups at 6, 12, and 24 months.  $^{h}p \le 0.001$  for differences between the subgroups at 6, 12, and 24 months.  $^{h}p \le 0.051$   $^{h}p \le 0.001$  for a subgroups at 6, 12, and 24 months.  $^{h}p \le 0.001$   $^{h}p \le 0.001$  for a subgroups at 6, 12, and 24 months.  $^{h}p \le 0.001$   $^{h}p \le$ and fat mass  $17.6 \pm 1.4 \text{ kg}$  [13]. Fig. 1. a Longitudinal results of serum thyroidstimulating hormone (TSH) concentrations in adolescent girls with polycystic ovary syndrome (PCOS) who received an oral contraceptive (OC, red circles, N = 31) or a low-dose combination of spironolactone, pioglitazone, and metformin (SPIOMET, blue circles, N = 31) for 12 months and who remained untreated over the subsequent 12 months. The yellow area represents the active treatment phase. The grayshaded area represents the safest range of preconception TSH [10] whose average is at the same level as the median TSH in adolescent girls aged 16 years [11]. \*p <0.01 for differences between subgroups at 6, 12, and 24 months. #p <0.001 for 0-6-month changes within the SPIOMET subgroup. Data are expressed as mean  $\pm$  standard error of the mean. **b** Circulating thyroid-stimulating hormone (TSH) concentrations in nonobese adolescent girls with polycystic ovary syndrome (PCOS) after 1 year of randomized treatment either with an oral estro-progestogen contraceptive (OC, red dots, N = 31) or with low-dose combination of spironolactone, pioglitazone, and metformin (SPIOMET, blue dots, N = 31). The shaded background represents the reference range of morning TSH levels (centiles 2.5, 50, and 97.5) in nonobese girls aged 16 years, who were studied with the same TSH assay [11].

TSH-lowering effects [34] if given for at least 1 year [35] to older adults with type 2 diabetes and with TSH levels in the upper-normal range for age (above 2.27 mIU/L) [36]. The present findings suggest that metformin's TSH-lowering effects in nonobese adolescents with PCOS occur within 6 months (Fig. 1, panel a) and also mainly in patients with TSH levels in the upper-normal range for age (above 1.35 mIU/L; Fig. 1, panel b). The mechanisms whereby metformin exerts its TSH-lowering effects are largely unknown but may involve an altered set point at the hypothalamo-pituitary level [34] and/or a sensitization of TSH receptors to TSH in the thyroid gland [37].

Strengths of the present report include the randomized study design and the abundance of on- and posttreatment data over relatively long timespans. Weaknesses include the unblinded study design and, in particular, the nonavailability of free T3 and free T4 data. In conclusion, the endocrine signature of early PCOS is herewith extended to include modestly raised concentrations of circulating TSH; in turn, the safety and normalizing efficacy of SPIOMET intervention in nonobese adolescent girls with



PCOS are herewith extended to include on- and posttreatment TSH.

### **Statement of Ethics**

The study protocol was reviewed and approved by the Institutional Review Board of Sant Joan de Déu Hospital and registered at ClinicalTrial.gov (ISRCTN29234515; ISRCTN11062950). Written informed consent was obtained by the parents/legal and assent by each of the participants. Each participant girl assented to participate.

### **Conflict of Interest Statement**

The authors have no conflicts of interest to declare.

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**Author Contributions** 

C.G.-B. contributed to literature research, design of figures and tables, data collection, data analysis and interpretation, and wrote the manuscript. J.B. and A.L.B. contributed to data interpretation

and reviewed/edited the manuscript. G.C.-B. contributed to data interpretation, and reviewed/edited the manuscript. L.I. contributed to study design, data interpretation, and reviewed/edited the manuscript. F.d.Z. contributed to study design, data interpretation, wrote the manuscript, and reviewed/edited the manuscript.

### **Data Availability Statement**

All data generated or analyzed during this study are included in this article and its supplementary material files. Further inquiries can be directed to the corresponding author.

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Supplementary Figure 1. Recruitment of the study population

**STUDY 5** 

# ORGANOKINES AND LIVER ENZYMES IN ADOLESCENT GIRLS WITH POLYCYSTIC OVARY SYNDROME DURING RANDOMIZED TREATMENTS

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### **STUDY 5**

# Organokines and liver enzymes in adolescent girls with polycystic ovary syndrome during randomized treatments

**Introduction:** Polycystic ovary syndrome (PCOS) is often related to metabolicdysfunction associated steatotic liver disease (MASLD). MASLD has been associated with altered hepatic function and systemic dysmetabolism and with abnormal circulating levels of signaling molecules, so-called organokines.

**Objective:** To assess the effects of two randomized treatments on a set of organokines in adolescent girls with PCOS and without obesity, and report the associations with circulating biomarkers of liver damage which were assessed longitudinally in the aforementioned studies as safety markers.

**Materials and methods:** Liver enzymes [aspartate aminotransferase (AST), alanine aminotransferase (ALT) and gamma-glutamyl transferase (GGT)] were assessed as safety markers in previous randomized pilot studies comparing the effects of an oral contraceptive (OC) to those of a low-dose combination of spironolactone-pioglitazone-metformin (SPIOMET) for 1 yr. As a post-hoc endpoint, the organokines fibroblast growth factor-21 (FGF21), diazepam-binding protein-1 (DBI) and meteorin-like protein (METRNL), were assessed by ELISA after 6 months on OC (N=26) or SPIOMET (N= 28). Auxological, endocrine-metabolic, body composition (by DXA) and abdominal fat partitioning (by MRI) were also evaluated. Healthy age-matched adolescent girls (N=17) served as controls.

**Results:** Circulating ALT and GGT levels raised during OC treatment and returned to baseline concentrations in the post treatment phase; in contrast, SPIOMET treatment elicited no detectable changes in ALT and GGT concentrations. In relation to organokines after 6 months on treatment, (1) FGF21 levels were significantly higher in PCOS adolescents than in control girls; (2) DBI levels were lower in OC-treated girls as compared to controls and SPIOMET -treated girls; (3) no differences were observed in METRNL concentrations between PCOS girls and controls. Serum ALT and GGT levels

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directly correlated with circulating METRNL levels only in OC-treated girls (R=0.449; P=0.036 and R=0.552; P=0.004, respectively).

**Conclusion:** The on treatment increase in ALT and GGT levels occurring only in OCtreated girls associates with circulating METRNL levels suggesting an enhanced METRNL synthesis as a reaction to the hepatic changes elicited by OC treatment.

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# Organokines and liver enzymes in adolescent girls with polycystic ovary syndrome during randomized treatments

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**Introduction:** Polycystic ovary syndrome (PCOS) is often associated with metabolic-associated fatty liver disease (MAFLD). MAFLD has been associated with altered hepatic function, systemic dysmetabolism, and abnormal circulating levels of signaling molecules called organokines. Here, we assessed the effects of two randomized treatments on a set of organokines in adolescent girls with PCOS and without obesity, and report the associations with circulating biomarkers of liver damage, which were assessed longitudinally in the aforementioned studies as safety markers.

**Materials and methods:** Liver enzymes [aspartate aminotransferase (AST), alanine aminotransferase (ALT), and gamma-glutamyl transferase (GGT)] were assessed as safety markers in previous randomized pilot studies comparing the effects of an oral contraceptive (OC) with those of a low-dose combination of spironolactone-pioglitazone-metformin (spiomet) for 1 year. As a *post hoc* endpoint, the organokines fibroblast growth factor-21 (FGF21), diazepambinding protein-1 (DBI), and meteorin-like protein (METRNL) were assessed by ELISA after 6 months of OC (N = 26) or spiomet (N = 28). Auxological, endocrine-metabolic, body composition (using DXA), and abdominal fat partitioning (using MRI) were also evaluated. Healthy, age-matched adolescent girls (N = 17) served as controls.

**Results:** Circulating ALT and GGT levels increased during OC treatment and returned to baseline concentrations in the post-treatment phase; in contrast, spiomet treatment elicited no detectable changes in ALT and GGT concentrations. In relation to organokines after 6 months of treatment, (1) FGF21 levels were significantly higher in PCOS adolescents than in control girls; (2) DBI levels were lower in OC-treated girls than in controls and spiomet-treated girls; and (3) no differences were observed in METRNL concentrations between PCOS girls and controls. Serum ALT and GGT levels

were directly correlated with circulating METRNL levels only in OC-treated girls (R = 0.449, P = 0.036 and R = 0.552, P = 0.004, respectively).

**Conclusion:** The on-treatment increase in ALT and GGT levels occurring only in OC-treated girls is associated with circulating METRNL levels, suggesting enhanced METRNL synthesis as a reaction to the hepatic changes elicited by OC treatment.

Clinical Trial Registration: https://doi.org, identifiers 10.1186/ISRCTN29234515, 10.1186/ISRCTN11062950.

KEYWORDS

organokines, PCOS, spironolactone, pioglitazone, metformin, oral contraceptives, METRNL, liver enzymes

### 1 Introduction

Polycystic Ovary Syndrome (PCOS) is a common condition in adolescents and young women (1, 2). Adolescent PCOS is characterized by a combination of clinical and/or biochemical androgen excess and anovulatory oligo-amenorrhea that presents between 2 and 8 years after menarche (3). The entity appears to be driven by ectopic fat accumulation, particularly in the liver, leading to insulin resistance (3), and by reduced energy expenditure, partly due to a lower activity of brown adipose tissue (4), favoring weight gain, and increasing the risk for type 2 diabetes (5).

PCOS in adolescents and women is often associated with nonalcoholic fatty liver disease (NAFLD) (6, 7). Insulin resistance and hyperandrogenemia have been found to be major contributory factors independent of body mass index (BMI) (7). Genetically predicted NAFLD is associated with a higher risk of PCOS, as judged by bidirectional two-sample Mendelian randomization analyses (8). Mitochondrial dysfunction, gut microbiome dysbiosis, and endocannabinoid system overactivation are among the proposed molecular mechanisms linking NAFDL and PCOS (9). NAFLD is currently considered a systemic disorder in which hepatic steatosis is merely the landmark of systemic metabolic dysfunction, including insulin resistance, increased cardiovascular risk, and low-grade systemic inflammation (10, 11). This notion has recently led to the expansion of the designation of NAFLD to "metabolically associated fatty liver disease" or MAFLD (11).

Studies in adults affected by MAFLD indicate a close association between altered hepatic function and systemic dysmetabolism, encompassing a pathogenic rearrangement of circulating signaling molecules, the so-called organokines (12), which originate in the liver (hepatokines) and adipose tissue (adipokines) or other organs and tissues (13). The altered circulating levels of these signaling molecules generate multiorgan systemic disturbances and provide biomarker evidence of existing health risks in patients at distinct stages of disease progression (14). Especifically in relation to MAFLD, emerging data indicate that an altered secretion of organokines plays an essential role in the pathogenesis of insulin resistance and cardiovascular diseases. For example, fetuin-A, a hepatokine that elicits low-grade inflammation in MAFLD by acting as an endogenous ligand of toll-like receptor-4 and promotes the secretion of proinflammatory cytokines in adipose tissue and other organs (15). Fetuin-A also suppresses the expression of the insulin-sensitizing adipokine adiponectin, which leads to systemic insulin resistance. In MAFLD, increased proinflammatory cytokines and enhanced levels of hepatokines such as angiopoietin-like proteins, also promote endothelial dysfunction, dyslipidemia, and atherogenesis (16). However, comprehensive knowledge of the entire set of organokines involved in linking MAFLD to systemic alterations and the associated mechanisms of action is still lacking.

Women with PCOS show altered levels of organokine signaling molecules (17, 18). In adolescent PCOS, abnormal organokine concentrations [i.e., high molecular weight adiponectin (19, 20), growth-and-differentiation factor-15 (21), fetuin-A (22), and chemokine ligand-14 (23)], have also been associated with earlier stages of hepatic and metabolic systemic alterations, even in the absence of overt obesity.

Currently, there is no approved pharmacological treatment for adolescent PCOS. The usually recommended off-label medication is

Abbreviations: ALT, alanine aminotransferase; AST, aspartate aminotransferase; BMI, body mass index CV, coefficients of variation; DBI, diazepam-binding protein-1; ELISA, human enzyme-linked immunosorbent assay; FGF21, fibroblast growth factor-21; GDF15, growth-and-differentiation factor-15; GGT, gamma-glutamyl transferase; HOMA-IR, homeostasis model assessment of insulin resistance; NAFLD, non-alcoholic fatty liver disease; MAFLD, Metabolic associated fatty liver disease; METRNL, meteorin-like protein; OC, oral contraceptive; PCOS, polycystic ovary syndrome; Spiomet, spironolactonepioglitazone-metformin.

an oral estroprogestagen contraceptive (OC) that is primarily used to revert androgen excess and restore menstrual regularity (24). However, this approach has a limited capacity to improve metabolic status (3, 25, 26) and may cause sustained unfavorable changes in hepatic markers (25, 27). Currently, the research in progress focuses on the development of safer medications that reduce ectopic fat and/or increase energy expenditure (3). In adolescents with PCOS and without obesity, a low-dose combination of one mixed antiandrogen and anti-mineralocorticoid (spironolactone) which increases brown adipose tissue activity (23, 28), and two insulin sensitizers (pioglitazone plus metformin) (spiomet) results in a better improvement in the metabolic condition as compared to OCs, including increased insulin sensitivity, reduced inflammation and liver fat accumulation, and more normalization of circulating hepatokines (20-23, 29). However, it is unclear to what extent the targets of spiomet include extrahepatic tissues, as well as the effects of randomized treatments on new bioactive organokines in young girls.

Existing knowledge on the identity and role of organokines connecting MAFLD hepatic disturbances with systemic metabolic and cardiovascular diseases is still limited, and there are a number of recently recognized circulating molecules that potentially play this role. We chose to analyze the effects of OCs vs spiomet on organokines recently related to MAFLD, such as meteorin-like protein (METRNL), recently reported to be related to liver injury (30); fibroblast growth factor-21 (FGF21), a hepatokine with enhanced expression in liver disease, potentially protective against systemic dysmetabolism in MALFD (31), and diazepam-binding protein-1 (DBI, also named acyl CoA-binding protein), whose blockage has been reported to improve MAFLD in recent experimental settings (32). We described their associations with circulating biomarkers of hepatic damage, which were assessed longitudinally in the aforementioned studies as safety markers (19, 20).

### 2 Materials and methods

### 2.1 Study population and design

The study population consisted of 54 adolescent girls with PCOS and without obesity [age,  $16.3 \pm 0.2$  yr; BMI,  $24.1 \pm 0.5$  Kg/m<sup>2</sup>], who participated in two randomized, open-label, pilot studies with the same design (Study 1, ISRCTN29234515 and Study 2, ISRCTN11062950, Supplementary Figure 1), comparing on-treatment (over 1 year) and post-treatment (over 1 year) effects of OC versus spiomet; the primary endpoint was ovulation rate after OC or spiomet intervention (19, 20). The trials were performed at the Endocrinology Department of Sant Joan de Déu University Hospital, Barcelona, Spain and the pooled results have been previously reported in detail, including the primary endpoint and secondary endpoints, namely, hirsutism score, androgens, carotid intima-media thickness, body composition, abdominal fat distribution, and hepatic fat. Serum aspartate aminotransferase (AST), alanine aminotransferase (ALT), and gamma-glutamyl transferase (GGT) levels were assessed as pre-, on-, and post-treatment safety markers (19, 20).

Due to the limited availability of spare serum, the present report focuses on 6-month on-treatment assessments. At this time point, serum was available for METRNL and DBI measurements in 33 out of the 34 randomized girls with complete data (97%) in Study 1 and in 21 out of 28 girls with complete data (75%) in Study 2, while FGF21 measurement could be performed in 30 (88%) and 20 (75%) patients, respectively. All studied patients finalized the treatment and post-treatment phases of the trials and had complete longitudinal data (Supplementary Figure 1).

The inclusion and exclusion criteria have been previously described in detail (19, 20). OC treatment consisted of 20  $\mu$ g ethinylestradiol plus 100 mg levonorgestrel (21/28 days), and placebo (7/28 days); spiomet is a low-dose combination of spironolactone 50 mg, pioglitazone 7.5 mg, and metformin 850 mg, taken together, once daily at dinner time. A total of 17 agematched, healthy girls recruited in nearby schools for the original studies (19, 20) in whom a spare sample was available served as controls; all had regular menses and a gynecological age >2.0 years and none were hirsute or taking medications.

### 2.2 Clinical and endocrinemetabolic assessments

Height, weight, and BMI were retrieved from medical records. Blood sampling for the assessment of endocrine-metabolic and safety parameters was performed in the early morning, after an overnight fast, in the follicular phase (days 3–7) of the cycle or after two months of amenorrhea.

Circulating testosterone was measured by liquid chromatography-tandem mass spectrometry, as described previously (19, 20); sex hormone-binding globulin (SHBG) was assessed by immunochemiluminiscence (Immulite 2000, Diagnostic Products, Los Angeles, USA), and the intra- and inter-assay coefficients of variation (CVs) were <0.5% and <8%, respectively. The free androgen index (FAI) was calculated as the ratio of serum testosterone (nmol/L) to that of SHBG (nmol/L) ×100. Serum glucose was measured by the glucose oxidase method; circulating insulin was assessed by immunochemiluminiscence (Immulite 2000, Diagnostic Products, Los Angeles, USA); intra- and interassay CVs were <0.1% and <7.2%, respectively. Homeostasis model assessment of insulin resistance (HOMA-IR) was calculated as [fasting insulin (mU/L)] × [fasting glucose (mg/dL)]/405. Serum AST, ALT, and GGT were assessed by molecular absorption spectrometry. Ultrasensitive C-reactive protein (us-CRP) was measured using a highly sensitive method (Architect c8000; Abbott, Wiesbaden, Germany); the intra- and inter-assay CVs were <1% and <5%, respectively.

Serum METRNL levels were assessed with a specific human enzyme-linked immunosorbent assay (ELISA) (R&D Systems, Minneapolis, MN, USA), sensitivity: 0.64 ng/mL; intra- and interassay CVs were <10% and <12%, respectively] (33). Serum concentrations of DBI were assessed using ELISA (Abnova, Taipei, Taiwan); intra- and inter-assay CVs were <9% (34). Circulating FGF21 levels were determined using a specific noncross-reactive ELISA kit (Biovendor, Brno, Czech Republic), and intra- and inter-assay CVs were 3.5% and 3.7%, respectively (35).

Body composition was assessed using dual X-ray absorptiometry (DXA) with the Lunar Prodigy and Lunar software (version 3.4/3.5, Lunar Corp, WI) (15, 16). Abdominal fat partitioning (subcutaneous and visceral) and hepatic fat were assessed by magnetic resonance imaging (MRI) using a multiple-slice MRI 1.5 T scan (Signa LX Echo Speed Plus Excite, General Electric, Milwaukee, Wisconsin, USA), as previously described (19, 20).

### 2.3 Statistics and ethics

Statistical analyses were performed using SPSS version 27.0 (SPSS software, IBM Corp., Armonk, NY, USA) and GraphPad Prism 5 (GraphPad Software, CA, USA). Results are shown as the mean ± standard error of the mean. Variables with a normal distribution were compared using two-tailed Student's t-test. When necessary, logarithmic transformation was used to achieve a normal distribution of continuous variables. Correlations and stepwise multi-regression analysis were used to study associations between liver enzymes (AST, ALT, and GGT) and study variables, and between organokines (METRNL, FGF21, and DBI) and study variables. A covariance analysis was used to adjust for BMI. Statistical significance was set at p-value <0.05.

The study was approved by the Institutional Review Board of the Sant Joan de Déu University Hospital. Written informed consent was obtained from the parents and assent of each of the participating girls.

### **3** Results

# 3.1 Key variables in PCOS-treated girls vs. controls

The auxological, endocrine-metabolic, and imaging results in both patients and controls are shown in Table 1. As previously described (19, 20), spiomet intervention was associated with more normalizing effects than OC, as judged by fasting insulin, HOMA-IR, us-CRP, and hepato-visceral fat.

# 3.2 Longitudinal results of liver enzymes in PCOS-treated girls

The longitudinal results of liver enzymes (AST, ALT, and GGT) are shown in Figure 1. The AST levels on- and post-treatment (Figure 1A) were lower than those in the controls in both study subgroups. In contrast, on-treatment ALT (Figure 1B) and GGT (Figure 1C) levels were significantly increased in patients receiving OCs and remained unchanged on spiomet. After treatment, ALT and GGT concentrations decreased in the OC-treated girls, reaching levels similar to those in the control and spiomet-treated girls.

TABLE 1 Study variables in healthy control girls and girls with polycystic ovary syndrome (PCOS) without obesity who were randomized to receive an oral contraceptive (OC) or a low-dose combination of spironolactone plus pioglitazone plus metformin (spiomet) for 6 months.

	Controls	PCOS on-tr	(6 months eatment)
	(N = 17)	OC (N = 26)	spiomet (N = 28)
Auxology			
Age (years)	$16.1 \pm 0.3$	16.3 ± 0.3	$16.1 \pm 0.3$
Weight (kg)	58.1 ± 1.5	62.9 ± 2.3	$60.1 \pm 1.8$
BMI (kg/m <sup>2</sup> )	21.8 ± 0.6	24.5 ± 0.7	23.6 ± 0.6
Endocrine-Metab	oolic Variable	S	
Testosterone (nmol/L)	1.0 ± 0.1	1.0 ± 0.1	$1.1 \pm 0.1$
SHBG (nmol/L)	64 ± 7	63 ± 6	$32 \pm 2$ <sup>c f</sup>
FAI	1.9 ± 0.3	1.8 ± 0.2	$4.2\pm0.4$ $^{\rm b~f}$
Glucose (mmol/L)	4.6 ± 0.3	$4.8\pm0.1$	$4.7 \pm 0.1$
Fasting insulin (pmol/L)	67 ± 8	100 ± 8	$55 \pm 6^{\rm f}$
HOMA-IR	1.9 ± 0.2	3.2 ± 0.3	$1.7\pm0.2$ $^{\rm f}$
AST (µkat/L)	0.32 ± 0.01	$0.26\pm0.01^{\rm \ b}$	0.28 ± 0.01
ALT (µkat/L)	0.26 ± 0.01	0.28 ± 0.02	$0.24 \pm 0.02$
GGT (µkat/L)	0.21 ± 0.01	0.25 ± 0.01	0.18 $\pm$ 0.01 a $^{\rm f}$
usCRP (nmol/L)	6.4 ± 1.2	$22.7\pm3.3$ $^{\rm b}$	$5.7 \pm 1.1$ $^{\rm f}$
METRNL (pg/mL)	369.7 ± 31.5	384.8 ± 26.7	437.4 ± 32.4
log DBI (ng/mL)	$2.43 \pm 0.04$	$2.17$ $\pm$ 0.04 $^{\rm c}$	$2.34\pm0.05$ $^{\rm d}$
log FGF21 (pg/mL) <sup>&amp;</sup>	$1.27\pm0.07$	$1.53 \pm 0.09$ <sup>b</sup>	$1.67\pm0.06$ $^{\rm c}$
Body composition	n (DXA) ‡		
Lean mass (kg)	-	37 ± 1	36 ± 1
Fat mass (kg)	_	23 ± 2	22 ± 2
Abdominal fat (kg)	-	$6.2\pm0.4$	$5.5 \pm 0.4$
Abdominal fat par	titioning (MR	RI)	
Subcutaneous fat (cm <sup>2</sup> )	111 ± 13	175 ± 19	159 ± 19
Visceral fat (cm <sup>2</sup> )	29 ± 2	$45 \pm 4^{a}$	35 ± 2 <sup>d</sup>
Hepatic fat (%)	10 ± 1	$20~{\pm}~1$ $^{\rm c}$	$12 \pm 1$ <sup>f</sup>

Values are mean ± standard error of the mean (SEM).

ALT, alanine transaminase; AST, aspartate transaminase; BMI, body mass index; DBI, diazepam-binding inhibitor; DXA, dual X-ray absorptiometry; FAI, free androgen index; FGF21, fibroblast growth factor 21; GGT, gamma-glutamyl transferase; HOMA-IR, homeostasis model assessment insulin resistance; METRNL, meteorin-like; MRI, magnetic resonance imaging; OC, oral contraceptive; SHBG, sex hormone-binding globulin; spiomet, spironolactone plus pioglitazone plus metformin; usCRP, ultrasensitive C-reactive protein.  $^{+}$ Indicative DXA values in healthy adolescents, matched for age and height (N = 41): lean mass 35.1 ± 1.0 kg; fat mass 17.6 ± 1.4 kg (doi.org/10.1210/jendso/bvaa032).

 $^{\rm d}P$   $^{\rm <0.05;}$   $^{\rm e}P$   $^{\rm <0.01}$  and  $^{\rm f}P$   $^{\rm <0.001}$  for differences between randomized subgroups.

P values are adjusted for BMI. The table depicts the study variables after 6 months of active treatment.

<sup>&</sup>lt;sup>88</sup>FGF21 assessment was performed in N = 24 girls on OCs and in N = 26 girls on spiomet <sup>a</sup>P  $^{\circ}0.05$ ; <sup>b</sup>P  $^{\circ}0.01$  and <sup>c</sup>P  $^{\circ}0.001$  vs controls.



FIGURE 1

Longitudinal results of aspartate aminotransferase (AST, **A**), alanine aminotransferase (ALT, **B**), and gamma-glutamyl transferase (GGT, **C**) concentrations in adolescent girls with polycystic ovary syndrome (PCOS) who received an oral contraceptive (OC, red circles, N = 26) or a low-dose combination of spironolactone-pioglitazone-metformin (spiomet, blue circles, N = 28) for 12 months and remained untreated for 12 months. The yellow area represents the active treatment phase. The dotted line represents the mean value in healthy controls (N = 17), and the shaded area represents the mean  $\pm$  standard error in healthy controls. \*P <0.05; \*\*P <0.01; \*\*\*P <0.001 for differences between subgroups at 6, 12, and 18 months.  $^{+}P$  <0.05;  $^{+#}P$  <0.01, differences between controls and the OC subgroup. \*P <0.05, for differences between controls and spiomet subgroup.

# 3.3 Organokine levels in PCOS-treated girls vs. controls

Regarding organokines (Table 1), on-treatment FGF21 levels were significantly increased in both PCOS subgroups compared with those in control girls. Circulating DBI levels were lower in the OC-treated girls than in the spiomet-treated girls and controls. Lastly, no differences were observed in METRNL levels between the controls and OC- or spiomet-treated girls.

# 3.4 Associations among liver enzymes, organokines, and study variables

The associations between liver enzymes and endocrinemetabolic, body composition, and abdominal fat partitioning variables after 6 months of OC or spiomet treatment are shown in Table 2 and Figure 2. AST levels after 6 months of spiomet treatment correlated negatively with the FAI (R = -0.537; P = 0.02) and FGF21 (R = -0.531; P = 0.042) levels, and positively with HOMA-IR (R = 0.546; P = 0.035). No significant association was observed in the OC subgroup. ALT levels in spiomet-treated girls also associated negatively with the FAI (R = -0.458; P = 0.014) and with HOMA-IR (R = 0.524; P = 0.014); in the OC subgroup, ALT concentrations positively correlated with SHBG (R = 0.496; P = 0.011) and METRNL (R = 0.449; P = 0.036) levels. GGT levels in OC-treated girls were also found to be strongly correlated with METRNL concentrations (R = 0.552; P = 0.004).

The correlations between organokines and the study variables are shown in Table 3. In spiomet-treated girls, FGF21 negatively correlated with visceral fat (R = -0.750; P = 0.001).

TABLE 2 Correlation between liver enzymes [aspartate transaminase (AST), alanine aminotransferase (ALT), and gamma-glutamyl transpeptidase
(GGT)] and study variables in girls with polycystic ovary syndrome (PCOS) after 6 months on an oral contraceptive (OC) or a low-dose combination of
spironolactone plus pioglitazone plus metformin (spiomet).

	AST (µkat/L)				ALT (µkat/L)				GGT (µkat/L)			
	PCOS	(6 mon	ths on-tre	eatment)	PCOS	(6 mor	ths on-tre	eatment)	PCOS	(6 mon	ths on-tre	atment)
	OC (N	= 26)	spiomet	(N = 28)	OC (N	= 26)	spiomet	(N = 28)	OC (N	= 26)	spiomet	(N = 28)
	R	Р	R	Р	R	Р	R	Р	R	Р	R	Р
Weight (kg)	-0.226	0.480	0.167	0.551	0.371	0.235	0.176	0.530	0.097	0.765	0.089	0.754
Testosterone (nmol/L)	0.031	0.924	-0.439	0.102	-0.023	0.943	-0.200	0.474	0.208	0.517	0.076	0.788
SHBG (nmol/L)	0.103	0.750	0.360	0.188	0.496	0.011	0.279	0.315	0.072	0.730	0.540	0.087
FAI	0.168	0.535	-0.537	0.002	-0.346	0.271	-0.458	0.014	0.185	0.364	-0.081	0.685
Glucose (mmol/L)	0.568	0.064	0.239	0.391	-0.185	0.565	0.474	0.074	0.153	0.635	-0.041	0.886
Insulin (pmol/L)	-0.409	0.186	0.496	0.060	-0.045	0.890	0.485	0.067	-0.433	0.160	0.199	0.478
HOMA-IR	0.200	0.532	0.546	0.035	0.014	0.819	0.524	0.028	-0.375	0.229	0.198	0.477
usCRP (nmol/L)	0.456	0.136	0.553	0.073	0.406	0.190	0.461	0.082	0.154	0.756	0.151	0.442
METRNL (pg/mL)	0.255	0.359	-0.210	0.435	0.449	0.036	0.149	0.423	0.552	0.004	0.157	0.424
log DBI (ng/mL)	0.270	0.396	-0.206	0.461	-0.397	0.202	-0.321	0.244	-0.101	0.755	0.069	0.824
log FGF21 (pg/mL) <sup>&amp;</sup>	0.014	0.946	-0.531	0.042	-0.256	0.422	-0.104	0.712	0.025	0.939	0.183	0.513
Lean mass (kg)	-0.211	0.510	0.151	0.591	0.019	0.924	0.178	0.364	0.210	0.512	-0.313	0.256
Fat mass (kg)	-0.063	0.845	0.098	0.727	0.064	0.753	0.244	0.209	-0.074	0.820	0.489	0.075
Abdominal fat (kg)	-0.125	0.699	0.011	0.970	0.012	0.950	0.203	0.299	-0.254	0.425	0.133	0.635
Subcutaneous fat (cm <sup>2</sup> )	-0.161	0.617	-0.042	0.882	0.023	0.932	0.228	0.241	0.169	0.408	0.291	0.132
Visceral fat (cm <sup>2</sup> )	-0.374	0.231	0.281	0.311	0.187	0.560	0.084	0.667	-0.358	0.077	0.154	0.432
Hepatic fat (%)	0.334	0.288	-0.214	0.443	0.187	0.560	0.054	0.848	-0.233	0.251	0.237	0.221

ALT, alanine transaminase; AST, aspartate transaminase; DBI, diazepam-binding inhibitor; FAI, free androgen index; FGF21, fibroblast growth factor 21; GGT, gamma-glutamyl transferase; HOMA-IR, homeostasis model assessment insulin resistance; METRNL, meteorin-like; OC, oral contraceptive; SHBG, sex hormone-binding globulin; spiomet, spironolactone, pioglitazone, metformin; usCRP, ultrasensitive C-reactive protein.

Results are shown as R coefficients and P-values, adjusted for body mass index in multiple regression analysis. Significant values are in bold.

 $^{\&}$ FGF21 assessment was performed in N = 50 girls with PCOS (N = 24 OC, N = 26 spiomet).

### 4 Discussion

The present longitudinal on- and post-treatment observations in adolescent girls with PCOS and without obesity receiving either OCs or spiomet revealed that circulating levels of ALT and GGT increased only under OC intervention, indicating a stressful effect on the liver (36, 37), which nevertheless reverted upon treatment discontinuation.

Our data are in line with those of previous studies reporting the influence of OCs on liver enzymes (38) and the effects of metformin on circulating ALT and GGT (but not AST) levels (39). Indeed, serum ALT activity is considered a highly sensitive biomarker of hepatic damage, ahead of circulating AST levels, and serum GGT activity is considered an additional biomarker of liver function used to extend the information provided by ALT activity (36).

Both treatments upregulated FGF21 levels, an effect previously observed after ethinylestradiol-cyproterone acetate-based OC treatment in PCOS adolescents (35). The inverse correlation between circulating FGF21 levels and visceral fat only in spiomettreated girls could reflect an increase in FGF21 signaling in visceral adipose tissue, followed by a more insulin-sensitive status (40). No significant associations were found between the on-treatment FGF21 changes and ALT and GGT concentrations.

To our knowledge, this is the first study in PCOS adolescents exploring the effects of two randomized treatments on the circulating concentrations of DBI, a multifunctional protein that mediates broad hepatoprotective effects (41), and METRNL, a regulatory protein involved in adipose tissue plasticity, inflammation, and cardiac function (42), recently identified as a potential hepatokine (30). DBI levels were lower in OC-treated patients than in spiomet-treated patients, but showed no correlation with indicators of hepatic damage. In contrast, METRNL levels did not differ between the two randomized subgroups but showed a strong positive association with ALT and GGT only in OC-treated PCOS girls, in whom hepatic enzymes experienced an on-treatment upward change. This finding is somewhat unexpected, as METRNL, previously considered mostly an adipokine and myokine, has been proposed to play a protective role against insulin resistance and inflammation in experimental models of obesity (43), and is usually downregulated in adult patients with obesity (44). However, our


#### FIGURE 2

Correlations between circulating meteorin-like (METRNL) levels and alanine aminotransferase (ALT, **A**) and gamma-glutamyl transferase (GGT, **B**) concentrations in adolescent girls with polycystic ovary syndrome (PCOS) after 6 months of treatment with an oral contraceptive (OC, red circles, N = 26) or a low-dose combination of spironolactone-pioglitazone-metformin (spiomet, blue circles, N = 28). P-values were adjusted for body mass index.

TABLE 3 Correlations between organokines [meteorin-like (METRNL), fibroblast growth factor 21 (FGF21) and diazepam-binding inhibitor (DBI)] and study variables in girls with polycystic ovary syndrome (PCOS) after 6 months on an oral contraceptive (OC) or on a low-dose combination of spironolactone plus pioglitazone plus metformin (spiomet).

	METRNL (pg/mL) PCOS (6 months on-treatment)				log DBI (ng/mL)				log FGF21 (pg/mL)			
					PCOS (6 months on-treatment)				PCOS (6 months on-treatment)			
			intrib-on-treatment)									
	OC (N = 26)		spiomet (N = 28)		OC (N = 26)		spiomet (N = 28)		OC (N = 24)		spiomet (N = 26)	
	R	Р	R	Р	R	Р	R	Р	R	Р	R	Р
Weight (kg)	0.217	0.436	-0.360	0.171	-0.471	0.122	-0.122	0.554	-0.313	0.321	-0.087	0.759
Testosterone (nmol/L)	-0.154	0.584	-0.249	0.352	0.477	0.117	0.083	0.788	-0.327	0.300	0.489	0.064
SHBG (nmol/L)	0.158	0.575	-0.430	0.097	-0.150	0.642	0.107	0.729	-0.363	0.248	0.079	0.778
FAI	-0.334	0.289	0.230	0.408	0.367	0.240	-0.040	0.896	0.167	0.603	0.433	0.107
Glucose (mmol/L)	-0.169	0.548	-0.034	0.900	0.229	0.475	-0.057	0.852	-0.172	0.594	-0.012	0.968
Insulin (pmol/L)	-0.450	0.092	0.247	0.357	-0.337	0.284	-0.003	0.865	-0.216	0.499	-0.307	0.895
HOMA-IR	-0.460	0.085	0.245	0.360	-0.259	0.196	-0.001	0.985	-0.255	0.242	-0.039	0.891
usCRP (nmol/L)	0.366	0.252	-0.259	0.333	-0.045	0.825	-0.249	0.201	-0.115	0.723	-0.177	0.528
Lean mass (kg)	0.219	0.434	-0.280	0.293	-0.245	0.443	-0.049	0.873	-0.321	0.309	-0.385	0.157
Fat mass (kg)	0.034	0.905	-0.134	0.621	-0.427	0.167	-0.445	0.127	-0.029	0.930	0.377	0.166
Abdominal fat (kg)	-0.003	0.993	-0.168	0.533	-0.310	0.326	-0.464	0.111	-0.059	0.855	0.348	0.204
Subcutaneous fat (cm <sup>2</sup> )	-0.470	0.077	-0.066	0.983	-0.014	0.964	-0.187	0.079	-0.208	0.931	0.358	0.190
Visceral fat (cm <sup>2</sup> )	-0.241	0.384	0.059	0.829	-0.085	0.792	-0.341	0.255	0.274	0.389	-0.750	0.001
Hepatic fat (%)	-0.060	0.831	-0.119	0.661	-0.273	0.230	-0.133	0.664	0.096	0.767	0.208	0.458

DBI, diazepam-binding inhibitor; FAI, free androgen index; FGF21, fibroblast growth factor 21; HOMA-IR, homeostasis model assessment insulin resistance; METRNL, meteorin-like; OC, oral contraceptive; SHBG, sex hormone-binding globulin; spiomet, spionolactone, pioglitazone, metformin; usCRP, ultrasensitive C-reactive protein. Results are shown as R coefficients and P-values, adjusted for body mass index in multiple regression analysis. Significant values are in bold.

findings are consistent with a recent report pointing to METRNL as a hepatokine specifically induced by hepatic injury in obese patients (30). Considering the aforementioned positive role of METRNL in systemic metabolism, our results suggest enhanced METRNL synthesis in OC-treated girls as a reaction to the hepatic changes elicited by OC treatment. We cannot unequivocally establish that the association between METRNL levels and markers of hepatic damage corresponds to altered METRNL synthesis, specifically in the liver. However, hepatic stress may lead to increased production of protective agents, and FGF21 is an example of such a response (31). METRNL has anti-inflammatory properties and acts via the c-Kit receptor (45), which is expressed in the liver and peripheral tissues (46). Enhanced METRNL signaling may be speculated to be a compensatory mechanism intended to prevent systemic alterations, including inflammation, in response to hepatic insults. In any case, our findings suggest a potential role of METRNL as a molecular factor related to changes in liver enzymes with systemic metabolism, which warrants further investigation.

This study has several limitations. First, the small sample size and limited availability of samples precluded a longitudinal analysis of organokine concentrations that had to be restricted to a single time point of treatment. Second, access to liver biopsy samples, unfeasible for obvious ethical reasons in this type of study, would have provided particularly relevant information on the hepatic expression of METRNL in relation to OC. The strengths of the present report include the randomized study design, rather homogeneous study population, and assessment of novel organokines under two randomized treatments.

In conclusion, the pattern of circulating DBI, but not of FGF21 and METRNL, differs in adolescent girls with PCOS receiving OCs or spiomet in randomized studies. The on-treatment increase in ALT and GGT levels, occurring only in OC-treated girls, directly associated with the circulating levels of METRNL, suggesting enhanced METRNL synthesis as a reaction to the hepatic changes elicited by OC treatment.

### Data availability statement

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

### **Ethics statement**

The studies involving humans were approved by Institutional Review Board of University of Barcelona, Sant Joan de Déu University Hospital. The studies were conducted in accordance with the local legislation and institutional requirements. Written informed consent for participation in this study was provided by the participants' legal guardians/next of kin. Written informed consent was obtained from the individual(s), and minor(s)' legal guardian/ next of kin, for the publication of any potentially identifiable images or data included in this article.

### Author contributions

CG-B: Writing – original draft, Data curation, Formal analysis, Investigation, Methodology. MP: Data curation, Formal analysis, Writing – review & editing. AN-G: Writing – review & editing, Data curation. AL-B: Conceptualization, Writing – review & editing. FZ: Conceptualization, Writing – review & editing. FV: Writing – original draft, Conceptualization, Supervision, Writing – review & editing. LI: Writing – review & editing, Conceptualization, Funding acquisition, Supervision, Writing – original draft.

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## Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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## Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fendo.2024. 1325230/full#supplementary-material 1. Wolf WM, Wattick RA, Kinkade ON, Olfert MD. Geographical prevalence of polycystic ovary syndrome as determined by region and race/ethnicity. *Int J Environ Res Public Health*. (2018) 15:2589. doi: 10.3390/ijerph15112589

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## Supplementary Material

## Organokines and liver enzymes in adolescent girls with polycystic ovary syndrome during randomized treatments

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Supplementary Figure 1. Recruitment of the study population

PCOS is the most common endocrine disorder affecting 5 to 10% of women of reproductive age worldwide. PCOS is associated with numerous comorbidities, has a negative impact on HRQoL and a huge global yearly cost for the health sector. However, there is no approved therapy for PCOS.

So far, the usual off-label recommended treatment for PCOS is an OC. OCs alleviate the clinical symptoms but they do not revert the underlying pathophysiology of PCOS; i.e., they do not target a reduction of ectopic (hepato-visceral) fat, so that upon discontinuation of such treatment, the PCOS phenotype tends to return. In recent years, alternative potential therapies have emerged for managing PCOS. In patients without obesity, a low-dose combination of SPI, PIO and MET (SPIOMET), targeting ectopic fat, the root cause of the disorder, appears to hold the potential to revert the PCOS phenotype.

The results derived from two-randomized, open-label, pilot studies, comparing the effects of SPIOMET vs the standard OC treatment in girls with PCOS and without obesity are presented in this doctoral thesis.

#### Effects of a low-dose combination of SPI, PIO and MET (SPIOMET) versus OC

The first study of this doctoral thesis presents combined data from two pilot studies (ISRCTN29234515 and ISRCTN11062950, respectively) comparing the effects of SPIOMET versus OC treatment. Pooled data corroborate SPIOMET as a combination treatment that leads to more normalizing effects than OCs, as judged by the reduction of ectopic (hepato-visceral) fat, by the increase in the annualized number of ovulations, and by the outcome of several markers of metabolic and cardiovascular health, including for waist circumference, circulating HMW-adiponectin and cIMT.

Waist circumference is a reliable indicator of visceral fat (**257**). There is a significant association between waist circumference and the risk of developing T2D or coronary heart disease (**258-260**). Moreover, women with PCOS tend to have higher waist circumference (**261**). SPIOMET-treated girls experienced a reduction in waist circumference which is considered critically important for decreasing adverse health risks (**262**).

Adiponectin is an adipocyte-secreted hormone that exists in multiple forms. Among those, HMW-adiponectin is the biologically active form (**263**) and has insulinsensitizing and anti-inflammatory properties (**84**). In concordance with previous studies (**264-266**), our pooled results showed that adolescent girls with PCOS had lower circulating HMW-adiponectin levels. Moreover, the study also confirmed that SPIOMET treatment -but not OCs- raised HMW-adiponectin levels, most likely, resulting from PIO action (**243,267**).

cIMT is an established and non-invasive marker of cardiovascular risk. cIMT is increased in women and adolescent girls with PCOS and directly correlates with components of the metabolic syndrome (**113,268,269**). In line with previous studies, pooled results also confirmed that SPIOMET reduces the progressive increase of cIMT in PCOS patients (**270-272**).

The normalization of the above-mentioned parameters observed in SPIOMETtreated girls, but not in those receiving OC treatment, highlights the potential benefits of SPIOMET in the management of PCOS. Moreover, pooled results of the randomized studies also indicate that SPIOMET treatment led to an overall healthier, more insulinsensitive condition, with less ectopic fat, and to a more normal post treatment ovulation rate. The low ovulation rates observed after OC treatment may be the result of persisting pathophysiological abnormalities rather than the consequence of a residual inhibition of the gonadotropic axis. Indeed, in healthy young women, ovulatory function recovers within three months after discontinuing OC treatment (**273,274**).

Pooled data supports the concept that ectopic (hepato-visceral) fat and/ or hyperinsulinemia are key drivers of PCOS. Whereas ectopic adiposity and IR failed to improve during standard treatment with OC, SPIOMET intervention went along with a reduction of hepato-visceral fat and by a normalization of IR, both of which were maintained during the post treatment year.

The decrease in hepato-visceral fat observed in the SPIOMET subgroup was not accompanied by a reduction of lean mass, total fat mass or abdominal subcutaneous

fat, suggesting a redistribution of ectopic fat towards subcutaneous stores, potentially reducing the comorbidities associated with ectopic fat storage (**191,275**).

The downward normalization of hepato-visceral fat may have been influenced in part by the increase in circulating miR-451a, which reduces the expression of thyroid hormone responsive spot 14 (*THRSP*), a key gene driving hepatic steatosis (**276,277**).

Ectopic lipid accumulation, particularly in the liver, leads to IR that may precede the development of disorders such as T2D and MASLD (**278**). Increased hepatic fat and IR are common in adolescents with PCOS with and without obesity, and appear to be linked to the underlying PCOS pathophysiology rather than to testosterone levels (**279,280**). It can be speculated that ectopic fat accumulation is the cause rather than the consequence of androgen excess. Therefore, targeting a reduction in androgen levels may not be the best choice to normalize the entire PCOS phenotype and to address subsequent comorbidities. The diverging effects of OC and SPIOMET on IR and ectopic fat may herald diverging influences on subsequent risks for PCOS-associated disorders including anovulatory subfertility, gestational diabetes, T2D and/or MASLD.

# CXCL14 levels in adolescent girls with PCOS and the effects of 1 year on treatment of SPIOMET and OC on circulating CXCL14 concentrations

The second study of this doctoral thesis reports CXCL14 levels in adolescent girls with PCOS, the effects of SPIOMET and OC on circulating CXCL14 concentrations as well as the effects of SPIOMET components on the expression and release of CXCL14 in a cellular model of human adipocytes.

The study shows that adolescent girls with PCOS have reduced levels of CXCL14, a chemokine secreted preferentially by brown/ beige adipose tissue previously identified in experimental studies (**85**). In line with our results, another study reported decreased circulating levels of CXCL14 in disorders associated to IR, such as obesity, especially in patients also suffering from T2D (**281**). Moreover, a study conducted in a cohort of children from birth to 1 year of age detected that CXCL14 levels positively correlated with the area of active cervical BAT only in 1-yr-old girls, which concurred with a high CXCL14 gene expression in neonatal BAT (**282**). These data support the usefulness of

CXCL14 as a potential biomarker for BAT activity as well as for metabolic health. In women with PCOS, reduced levels of CXCL14 have also been detected in follicular fluid and human-luteinized granulosa cells, in parallel with decreased progesterone concentrations (**283**). These abnormalities are partially reversed after CXCL14 administration, suggesting a novel role of CXCL14 in progesterone synthesis (**283**).

Our data suggest that CXCL14 could be a target of SPIOMET treatment, as CXL14 levels were restored after SPIOMET but not after OC intervention. Consistent with our results, another study also considered CXCL14 as a potential pharmacological target for PCOS treatment (**284**). In this study, Ye et al., reported that Compound K, a final intestinal metabolite of Ginseng herb, could activate BAT and up-regulate CXCL14 expression leading to an improvement of PCOS phenotype in an animal model (**284**).

High CXCL14 levels have been shown to improve insulin sensitivity in adipocytes *in vitro* (**285**) and in rodent models (**85**); however, not all reports agree in the antidiabetic actions of CXCL14, and even deleterious pro-diabetogenic effects have been proposed *in vitro* (**286**) and *in vivo* (**287**). Nevertheless, our study indicated that normalization of CXCL14 after SPIOMET associated with decreased IR.

The results of our study also support the evidence that PIO increased CXCL14 expression in pre-adipocytes. PIO is known to increase insulin sensitivity as well as adipogenic differentiation and to promote the acquisition of a brown/beige phenotype in adipose cells (**288**). The induction of CXCL14 expression by PIO is consistent with data in experimental models highlighting increased expression of CXCL14 with brown/ beige adipogenic differentiation (**85**). Thus, PIO could elicit a so-called "browning" of WAT. Our data also show that SPI increased the release of CXCL14 in differentiated adipocytes. This is consistent with previous reports indicating that SPI activates brown fat in humans (**240**). So, it could be proposed that SPI raised energy expenditure by BAT activation.

The physiological significance of the drug concentrations used in the *in vitro* studies is unclear. The concentrations of PIO increasing CXCL14 levels in differentiated adipocytes (from at least  $0.1 \mu$ M) were in the range of those in plasma from SPIOMET-

treated healthy women (**289**). However, although the SPI concentrations used here were the same as those employed in previous studies in adipose cells (**290,291**) plasma SPI levels in SPIOMET-treated controls are lower (**289**).

Women with PCOS have lower BAT activity and decreased BAT volume (**90,91**). Moreover, it has recently been described that women with PCOS have reduced resting energy expenditure - when corrected for fat-free mass - which might potentially favor weight gain (**292**). Interestingly, transplantation of BAT was reported to exert a beneficial effect in rodent models with PCOS (**293,294**), reflecting the important role of BAT in PCOS pathogenesis. Considering that CXCL14 is preferentially released by brown adipocytes (**85**) it may be speculated that SPIOMET treatment drives a shift in adipose tissue plasticity to a more brown/beige phenotype resulting in enhanced CXCL14 release, potentially improving glucose homeostasis and possibly reducing diabetes risk. This sequence may be especially relevant in young girls, where the brown/beige adipose tissue amount is particularly significant (**295**).

In conclusion, insulin sensitization with SPIOMET normalizes the abnormally low levels of CXCL14 in girls with PCOS. This is consistent with the effects of pioglitazone and spironolactone inducing CXCL14 expression and promoting a brown-like phenotype in adipocytes.

## Gut microbiota composition in adolescent girls with PCOS and the effects of SPIOMET and OC interventions

The third study of this doctoral thesis assessed the gut microbiota of adolescent girls with PCOS and without obesity and the longitudinal changes after SPIOMET and OC interventions.

The study demonstrated the presence of gut dysbiosis in these girls, including decreased evenness and diversity indexes, and confirmed the association between alpha diversity and markers of hyperandrogenism and liver function found in other populations (**141-143**). Regarding beta diversity, the study also confirmed an altered microbiota pattern in girls with PCOS according to Jaccard and Bray-Curtis dissimilarity matrices. However, the lack of differences in community structure using phylogenetic

distances as Unweighted and Weighted Unifrac may indicate that despite detecting different amplicon sequence variants (ASVs) conforming the microbiota profile of girls with PCOS, these ASVs may be phylogenetically similar to the ASVs detected in control girls. In line with this observation, no differences were found in the abundance of the most copious phyla between patients with PCOS and healthy girls. However, girls with PCOS had an altered taxonomic profile, with more abundance of *Family XI* and lower abundance of the family *Prevotellaceae* and the genus *Prevotella* and *Senegalimassilia*. The excessive abundance of *Family XI* decreased toward normal after SPIOMET administration but remained unchanged after OCs.

The abundance of Bacillales Family XI in patients as compared to controls may have pathophysiological implications. This family has been so far poorly studied and the related literature is thus, scarce; however, the genus *Gemella*, which belongs to *Family* XI, has been linked to liver inflammatory disease, gut inflammation and obesity (296,297). In addition, Gemella spp (also belonging to Family XI) is a bacterium that produces N-oleoyl serinol, an analogue of ceramides (298). Serum concentrations of ceramides are elevated in women with PCOS (73,299) and are associated to the degree of hyperandrogenism (as assessed by the FAI) and to central adiposity (74). Moreover, patients with obesity with increased diabetes risk have elevated circulating ceramide levels that associate to gut dysbiosis (300). In addition, ceramides produced by the gut microbiota have the potential to pass the epithelial barrier and interact with the host metabolism (301). Accordingly, it has been proposed that IR and hyperandrogenemia in patients with PCOS could be at least partly driven by an increase in serine phosphorylation due to the accumulation of reactive lipids, such as ceramides, that activate serine kinase signaling pathways (**302**). Moreover, it has also been demonstrated that ceramides induce IR via inhibition of Akt signaling in cell cultures (75). In our study, SPIOMET treatment, but not OC, decreased Family XI levels toward normal. This reduction could have caused in turn a decrease in N-oleoyl serinol production and subsequently normalize the serine phosphorylation status. Furthermore, the unchanged Family XI abundance in OC-treated girls was found to associate with increased fasting insulin and hepato-visceral fat. Hence, it is tempting to

speculate that the metabolic benefits exerted by SPIOMET could be mediated in part by a reduction of *Family XI* abundance.

Previous studies have disclosed the association between adiponectin and ceramides; for example, adiponectin decreases cellular ceramides by triggering ceramidase activities of adiponectin receptors (AdipoRs) (**303**) and by stimulating ceramide efflux to the exosome (**304**). Other studies suggest that ceramides might be involved in the regulation of adiponectin secretion (**305**). In the present study, the increase in circulating HMW-adiponectin concentrations after SPIOMET treatment could have modified N-oleoyl serinol levels, and ultimately explain the negative association between HMW-adiponectin and *Family XI* relative abundance.

In line with previous observations (140), we disclosed that girls with PCOS had less abundance of the family *Prevotellaceae* and the genus *Prevotella* (belonging to *Prevotellaceae* family). In contrast, other studies have shown an abundance of *Prevotella* in adolescent girls with PCOS and with obesity (143), in women with irregular menstrual cycles (306) and in animal models of PCOS (307). In addition, the reports on the involvement of *Prevotella* in other pathological conditions have been inconsistent. For example, increased abundance of *Prevotella* has been associated with IR, MASLD and chronic inflammatory disorders (293,308-311). However, other studies relate the abundance of *Prevotella* with an improved glucose metabolism (312) or fail to demonstrate an association with T2D (313).

The abundance of *Prevotellaceae* and *Prevotella* was significantly reduced after SPIOMET treatment. These results are in line with others demonstrating a non-significant reduction of *Prevotella* abundance after metformin intervention in women with obesity (**314**), but are in contrast with others showing that metformin increases its abundance in healthy men (**315**) and in men with diabetes (**316**) but not in groups combining men and women (**316**). These discrepancies may be due to the presence of functional diversities within the *Prevotella* genus or to the influence of sex in the composition of gut microbiota (**142**). Further analyses are needed to investigate the implications of the different *Prevotella* species within the context of PCOS.

In the present study, we also showed that girls with PCOS had lower abundance of the genus *Senegalimassilia*, and that this abundance associated inversely with markers of androgen excess and inflammation, and with hepato-visceral fat. Along these lines, a higher abundance of *Senegalimassilia* has been associated with healthy traits; for example, a higher occurrence of *Senegalimassilia* anaerobia (a species belonging to *Senegalimassilia* genus) was observed in faecal samples of children without obesity as compared to children with overweight (**317**). In addition, rats treated with a formula that effectively alleviated T2D had higher abundance of gut *Senegalimassilia* (**318**). Recently, the genus *Senegalimassilia* was proposed to act as a potential biomarker for early metabolic syndrome diagnosis, as it was down-regulated in patients with metabolic syndrome (**319**). Interestingly, although metformin treatment increases *Senegalimassilia* abundance in young healthy men (**315**); in our study, the combination of metformin with spironolactone and pioglitazone failed to restore *Senegalimassilia* abundance in girls with PCOS.

Finally, we disclosed that the dysbiosis of the gut microbiota observed in girls with PCOS was explained by the FAI and by hepato-visceral fat, suggesting that both hyperandrogenism and central adiposity may alter the gut microbiota, which in turn, could impact the pathophysiology of PCOS. Moreover, we showed that the variation of gut microbiota in OC-treated girls was explained by hepatic fat accumulation; hepato-visceral fat remained unchanged on OCs and was significantly reduced after SPIOMET. These results corroborate the notion that targeting a reduction of ectopic fat normalizes the PCOS phenotype including a recovery of the gut microbiota, represented here by the normalization of *Family XI* relative abundance. Further studies are needed to determine whether the changes in microbiota composition observed in girls with PCOS persist after therapy discontinuation and may, in turn, impact androgen metabolism.

In summary, the third study of this doctoral thesis demonstrates a dysbiosis of gut microbial community in girls with PCOS without obesity, including a reduction of alpha diversity, differences in community structure, a reduction of *Prevotellaceae*, *Prevotella* and *Senegalimassilia* and an increase of *Bacillales Family XI*.

## TSH levels in adolescents with PCOS and the effects of SPIOMET and OC interventions

The fourth study of this doctoral thesis reports TSH levels in adolescent girls with PCOS and without obesity and the on and post treatment effects of SPIOMET and OC on TSH concentrations.

TSH (but not free T3 o free T4) was a safety marker in the two-randomized, openlabel, pilot studies, comparing the effects of SPIOMET vs OC treatment in non-obese adolescent girls with PCOS. A post-hoc analysis of these TSH data was prompted by the recent description of relatively low TSH reference intervals in non-obese adolescent girls (**320**) and by the even more recent delineation of the safest range of preconception TSH (**132**). In the present report, novel findings include that [1] TSH levels in non-obese adolescents with PCOS were already shifted upward, on average by 27%; [2] this upward TSH shift did persist on/post treatment with OC but disappeared on/post treatment with SPIOMET, so that TSH levels entered swiftly into the safest preconception range and remained stable inside this range; and [3] on treatment changes in TSH associated most closely to those in liver fat.

The present pre, on, and post treatment observations corroborate the hypothesis that a TSH elevation in an adolescent girl with PCOS may in essence be no more than an epiphenomenon that points to the presence of insulin resistance within a context of lipid accumulation in ectopic depots. TSH elevations have previously been described as epiphenomena of insulin resistance and/ or ectopic fat in a variety of study populations including overweight children and adolescents and also in women with PCOS with or without obesity (**43,128-131,321-327**).

The on treatment data challenge the concept that TSH elevations in PCOS relate primarily to androgen excess (**128**) or body weight (**129,321,322**) given that the tested treatments had virtually parallel effects on androgenemia and body weight while having markedly divergent effects on TSH, insulin resistance, and ectopic fat. The post treatment data in PCOS patients strengthen the notion that SPIOMET has the capacity to confer broadly normalizing benefits that persist beyond its intake and that are herewith extended to include a lowering of TSH concentrations into the safest preconception range. These surprisingly prolonged effects are incompletely explained by factors such as HMW-adiponectin (84), GDF15 (114), miRNA profiles (164), as well as CXCL14 and gut microbiota (both data presented in the second and the third paper of this doctoral thesis), and thus remain to be further clarified.

There is still no evidence suggesting that the intake of spironolactone and/or pioglitazone has TSH-lowering effects. In contrast, metformin is known to exert TSH-lowering effects (**328**) if given for at least 1 year (**329**) to older adults with T2D and with TSH levels in the upper-normal range for age (above 2.27 mIU/L) (**330**). The present findings suggest that metformin's TSH-lowering effects in non-obese adolescents with PCOS occur within 6 months and also mainly in patients with TSH levels in the upper-normal range for age (above 1.35 mIU/L). The mechanisms whereby metformin exerts its TSH-lowering effects are largely unknown but may involve an altered set point at the hypothalamo-pituitary level (**328**) and/or a sensitization of TSH receptors to TSH in the thyroid gland (**331**).

# Organokines and liver enzymes in adolescent girls with PCOS and the effects of SPIOMET and OC treatments

The last study of this doctoral thesis reports the longitudinal on and post treatment observations in adolescent girls with PCOS and without obesity receiving either OCs or SPIOMET and revealed that circulating levels of ALT and GGT increase only under OC intervention, indicating a stressful effect on the liver (**119,120**), which nevertheless reverts upon treatment discontinuation.

Our data are in line with those of previous studies reporting the influence of OCs on liver enzymes (**332**) and the effects of metformin on circulating ALT and GGT (but not AST) levels (**333**). Indeed, serum ALT activity is considered a highly sensitive biomarker of hepatic damage, ahead of circulating AST levels, and serum GGT activity is considered an additional biomarker of liver function used to extend the information provided by ALT activity (**119**).

Both treatments upregulated FGF21 levels, an effect previously observed after ethinylestradiol–cyproterone acetate-based OC treatment in PCOS adolescents (**334**). The inverse correlation between circulating FGF21 levels and visceral fat only in SPIOMET-treated girls could reflect an increase in FGF21 signaling in visceral adipose tissue, followed by a more insulin-sensitive status (**335**). No significant associations were found between the on treatment FGF21 changes and ALT and GGT concentrations.

To our knowledge, this is the first study in PCOS adolescents exploring the effects of two randomized treatments on the circulating concentrations of DBI, a multifunctional protein that mediates broad hepatoprotective effects (336), and METRNL, a regulatory protein involved in adipose tissue plasticity, inflammation, and cardiac function (337), recently identified as a potential hepatokine (116). DBI levels were lower in OC-treated patients than in SPIOMET-treated patients, but showed no correlation with indicators of hepatic damage. In contrast, METRNL levels did not differ between the two randomized subgroups but showed a strong positive association with ALT and GGT only in OC-treated PCOS girls, in whom hepatic enzymes experienced an on treatment upward change. This finding is somewhat unexpected, as METRNL, previously considered mostly an adipokine and myokine, has been proposed to play a protective role against insulin resistance and inflammation in experimental models of obesity (338), and is usually downregulated in adult patients with obesity (339). However, our findings are consistent with a recent report pointing to METRNL as a hepatokine specifically induced by hepatic injury in obese patients (116). Considering the aforementioned positive role of METRNL in systemic metabolism, our results suggest enhanced METRNL synthesis in OC-treated girls as a reaction to the hepatic changes elicited by OC treatment. We cannot unequivocally establish that the association between METRNL levels and markers of hepatic damage corresponds to altered METRNL synthesis, specifically in the liver. However, hepatic stress may lead to increased production of protective agents, and FGF21 is an example of such a response (117). METRNL has anti-inflammatory properties and acts via the cKit receptor (340), which is expressed in the liver and peripheral tissues (341). Enhanced METRNL

signaling may be speculated to be a compensatory mechanism intended to prevent systemic alterations, including inflammation, in response to hepatic insults. In any case, our findings suggest a potential role of METRNL as a molecular factor related to changes in liver enzymes with systemic metabolism, which warrants further investigation.

The main strengths of this doctoral thesis include the strict inclusion and exclusion criteria, the longitudinal design and the assessment of the impact of two interventions. The limitations include the relatively low number of participants, the unblinded design and the absence of a control (untreated) group. Additional specific limitations include:

- The impossibility of studying the effect of SPIOMET on CXCL14 levels in tissues besides the adipose tissue.
- The lack of awareness of the physiological significance of the drug concentrations used in the *in vitro* studies.
- The lack of evidence for a direct role of CXCL14 downregulation and reinduction on the metabolic status of patients with PCOS after SPIOMET, as it would imply intervention experiments beyond the ethical standards for human studies.
- The absence of a follow-up period to ascertain whether the changes in the gut microbiota do persist after therapy discontinuation.
- The lack of additional serum samples to analyze ceramide serum concentrations.
- The non-availability of free T3 and free T4 data.
- The lack of a longitudinal analysis of organokine concentrations (being restricted to a single time point of treatment) due to the limited availability of samples.
- The unfeasibility to access to liver biopsy samples (for ethical reasons), which would have provided particularly relevant information on the hepatic expression of METRNL in relation to OC.

The results presented in this doctoral thesis remain to be further confirmed in larger and more diverse PCOS populations, including in girls with PCOS and obesity,

with different ethnic and developmental backgrounds, with other environmental exposures and with older ages. Currently, the potential of SPIOMET intervention is studied further by a European consortium working on adolescent PCOS (**342**). The efficacy and safety of SPIOMET components plus lifestyle intervention are being tested in a multi-centre, randomized, double-blind, placebo-controlled, phase II clinical trial in AYAs with PCOS (**342**). Indeed, patients are being recruited from 7 clinical centres across Europe, the BMI of the participants is >18 kg/m<sup>3</sup> and <35 kg/m<sup>3</sup> and the age is >12.0 years and ≤23.9 years at study start, which ensures a large an ethnically diverse cohort as well as the participation of older girls with and without obesity.

The potential of SPIOMET to prevent the development of adolescent PCOS is being currently explored in girls with a mismatch between (less) prenatal weight gain and (more) postnatal weight gain. Practically, the effects of half-dose SPIOMET are being tested in a randomized, placebo-controlled, phase II clinical trial performed in mismatch girls with advanced puberty (breast development before age 9 years) (**343**).

# CONCLUSIONS

- 1- Pooled results in adolescent girls with polycystic ovary syndrome and without obesity confirms spironolactone – pioglitazone – metformin as a treatment that attenuates insulin resistance, reduces ectopic adiposity, and is followed by a more normal ovulation rate and by an overall normalization of the polycystic ovary syndrome phenotype.
- 2- C-X-C motif chemokine ligand-14 may be a novel potential biomarker and molecular mediator in the improvement of polycystic ovary syndrome-associated metabolic alterations following spironolactone – pioglitazone – metformin intervention.
- 3- Spironolactone pioglitazone metformin's spectrum of normalizing effects in girls with polycystic ovary syndrome is herewith broadened as to include *Family XI* abundance in gut microbiota.
- 4- The endocrine signature of early polycystic ovary syndrome additionally incorporates modestly raised levels of circulating thyroid-stimulating hormone; the normalizing effects of spironolactone – pioglitazone – metformin intervention in adolescents with polycystic ovary syndrome and without obesity include the normalization of on and post treatment thyroid-stimulating hormone concentrations within a safe pre-gestational range.
- 5- The on treatment increase in alanine aminotransferase and gamma-glutamyl transferase levels occurring only in oral contraceptive-treated girls is associated with circulating meteorin-like protein levels, suggesting enhanced meteorin-like protein synthesis as a reaction to the hepatic changes elicited by oral contraceptive treatment.

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