1   **Phosphorylation at Ser-181 of oncogenic KRAS is required for tumor growth**

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25

26   **Abstract**

27   KRAS phosphorylation has been reported recently to modulate the activity of mutant

28   KRAS protein in vitro. In this study, we defined S181 as a specific phosphorylation site

29   required to license the oncogenic function of mutant KRAS in vivo. The phosphomutant

30   S181A failed to induce tumors in mice, whereas the phosphomimetic mutant S181D

31   exhibited an enhanced tumor formation capacity, compared to the wild-type KRAS

32   protein. Reduced growth of tumors composed of cells expressing the non-

33   phosphorylatable KRAS S181A mutant was correlated with increased apoptosis.

34   Conversely, increased growth of tumors composed of cells expressing the

35   phosphomimetic KRAS S181D mutant was correlated with increased activation of AKT

36   and ERK, two major downstream effectors of KRAS. Pharmacological treatment with

37   PKC inhibitors impaired tumor growth associated with reduced levels of

38   phosphorylated KRAS and reduced effector activation. In a panel of human tumor cell

39   lines expressing various KRAS isoforms, we showed that KRAS phosphorylation was

40   essential for survival and tumorigenic activity. Further, we identified phosphorylated

41   KRAS in a panel of primary human pancreatic tumors. Taken together, our findings

42   establish that KRAS requires S181 phosphorylation to manifest its oncogenic properties,

43   implying that its inhibition represents a relevant target to attack KRAS-driven tumors.

44

Introduction

RAS proteins are well-known small GTPases involved in the regulation of key signal transduction pathways. Cycling from the inactive (GDP-bound) to the active (GTP-bound) state faithfully responds to extracellular signals due to its tight regulation by GTP-exchange factors (GEFs) and GTPase activating proteins (GAPs). Activating point mutations that render RAS proteins insensitive to the extracellular signals are crucial steps in the development of the vast majority of cancers (1-3). Three different genes code for a total of four different Ras isoforms named HRAS, NRAS, KRAS4A and KRAS4B. *RAS* mutations, mainly at the *KRAS4B* (herein after referred to as *KRAS*) genes, occur in pancreatic (95%), colon (40%) and adenocarcinomas of the lung (35%) (1, 4, 5). The most prevalent oncogenic mutations in *RAS* at codons 12, 13 and 61 preserve the GTP-bound, active state by inhibiting intrinsic GTPase activity or interfering with the action of GAPs. In the GTP-bound form, RAS is able to interact with different effector proteins and consequently activates signal transduction pathways. Among those, the best characterized are the RAF1/MEK/ERK and the phosphatidylinositol-3-kinase (PI3K)/AKT (6, 7).

Since oncogenic mutations of *KRAS* give rise to an always GTP-bound protein which constitutively activates the effectors, positive or negative physiological regulation of oncogenic KRAS was not initially expected. Several reversible posttranslational modifications of KRAS have been described that could modulate KRAS oncogenic activity (8). Ubiquitination of oncogenic KRAS at lysine-147 in the guanine nucleotide-binding motif increases its binding to the downstream effectors PI3K and RAF1 thus increasing its tumorigenic activity (9). Furthermore, acetylation at lysine-104 affects interaction with GEFs and inhibits in vitro transforming activity of oncogenic KRAS (10). KRAS has, adjacent of the farnesylated C-terminal cysteine, a

70 stretch of six contiguous lysines in a total of eight lysine residues -known as the

71 polybasic domain- which promotes an electrostatic interaction with the negatively-

72 charged phosphate groups of phospholipids (11, 12). Phosphorylation of KRAS at

73 serine-181 within this domain has been described (13). We previously reported a role of

74 KRAS Ser181 phosphorylation for activation of the wild-type KRAS in vitro and to

75 regulate also in vitro oncogenic KRAS activity (14). By using both a genetic and

76 pharmacological approach we demonstrate here that phosphorylation of oncogenic

77 KRAS is required for tumor growth in vivo and that also this modification can be

78 detected in human tumors. Furthermore, pharmacological inhibition of oncogenic

79 KRAS phosphorylation suppresses KRAS oncogenic activity.

80

81 **Materials and Methods**

82 **Antibodies and reagents**

83 Primary antibodies used for immunoblotting were as follow: Anti-Actin (clone C4)

84 (#691001; 1:1000; MP Biomedicals, Santa Ana, CA, USA), Anti-GAPDH (#MAB374;

85 1:1000; Chemicon, Billerica, MA, USA); Anti-cleaved caspase-3 (Asp175) (#9661;

86 1:1000; Cell Signaling, Danvers, MA, USA); Anti-AKT (#9272; Cell Signaling); Anti-

87 phospho-AKT (Thr308) (#9275; 1:1000, Cell Signaling), Anti-p44/42 MAPK (ERK

88 1/2) (#9102; 1:1000; Cell Signaling); Anti-phospho-p44/42 MAPK (ERK 1/2)

89 (Thr202/Tyr204) (#9101; 1:1000; Cell Signaling), Anti-cyclin B1 (#4138; 1:1000; Cell

90 Signaling); Anti-KRAS (clone Ab-1) mouse (#OP24, 1:400, Calbiochem); Anti-Pan-

91 Ras (clone Ab-3) mouse (#OP40; 1:400; Calbiochem); Anti-HRAS (clone C20) rabbit

92 (#Sc-520, Santa Cruz); Anti-NRAS (clone F155) mouse (#Sc-31, Santa Cruz); Anti-

93 GAP120 (sc-63; 1:100; Santa Cruz, Santa Cruz, CA, USA); Anti-PKCδ (#610398;

94 1:500; BD Transduction Laboratories, San Jose, CA); Anti-phospho-PKCδ

95 (Ser643/676) (#9376; 1:1000; Cell Signaling). For immuhistochemistry we used Anti-

96 Ki-67 (SP6) (#NM-9106S; 1:200; NeoMarkers, Kalamazoo, MI, USA). We used

97 DeadEnd Colorimetric TUNEL System (G7132; Promega) for the TUNEL assays.

98 The reagents used for the detection of phospho-KRAS were: Protein Phosphatase λ

99 (#539514-20KV; Calbiochem); Phos-tag™ (#AAL-107, Wako Chemicals GmbH,

100 Neuss, Germany).

101 The inhibitors of PKC used were: Bryostatin-1 (#BIB0342, Apollo Scientific, Chesire,

102 UK), Edelfosine (1-O-Octadecyl-2-O-methyl-glycero-3-phosphorylcholine) (#BML-

103 L108, Enzo Life Science, Farmingdale, NY, USA), Bisindolylmaleimide I (BIM)

104 (#CAS 176504-36, Millipore), Gö6983 (#G1918, Sigma Aldrich).

105

106 **Cell lines**

107 NIH3T3, SW480, A549, MPanc-96 and HPAF-II cells obtained from American Tissue

108 and Cell Collection (ATCC) were grown in Dulbecco's Modified Eagle's Medium

109 (DMEM) containing 10% FCS (Biological Industries), and routinely verified according

110 to the specifications outlined in the ATCC Technical Bulletin. NIH3T3 stable cell lines

111 expressing either HA-KRASG12V, HA-KRASG12V-S181A or HA-KRASG12V-

112 S181D were obtained as previously described (14) .

113 DLD-1 knock-out of mutant KRAS allele DLD1$^{KRASwt/-}$ were obtained from Horizon

114 Discovery Ltd (clone D-WT7; #HD105-002; http://www.horizondiscovery.com;

115 Cambridge, UK). DLD1$^{KRASwt/-}$ cells were generated using the proprietary adeno-

116 associated virus (AAV) gene targeting technology GENESIS$^{®}$. Cells were maintained

117 according to the supplier recommendations in McCoy's modified media containing 10%

118 FBS (Biological Industries). DLD1$^{KRASwt/-}$ stable cell lines expressing either HA-

119 KRASG12V, HA-KRASG12V-S181A or HA-KRASG12V-S181D were obtained from

120 DLD1$^{KRASwt/-}$ after transfecting with the specific HA-KRASV12 plasmids (14) and a

121 puromycine resistance plasmid (pSG5A). After selection with puromycine (4 μg/mL)

122 clones or pools were obtained. .

123

124 **Tumor generation in mice**

125 The day of the injection, one million NIH 3T3 cells stably expressing either HA-

126 KRASG12V, HA-KRASG12V-S181A or HA-KRASG12V-S181D suspended in 0,1

127 mL PBS buffer were subcutaneously injected into both flanks of Swiss nude mice

128 (foxn1$^{-/-}$). Generated tumors were measured over time and at day 18 after injection,

129 mice were euthanized and tumors were harvested, weighed, measured and processed for

130 analysis (each group n=10).

131 For DLD-1 xenografts, one million cells stably expressing either HA-KRASG12V or

132 HA-KRASG12V-S181A suspended in 0,1 mL PBS buffer were subcutaneously injected

133 into both flanks of Swiss nude mice (foxn1$^{-/-}$). Generated tumors were measured over

134 time and at day 28 after injection, mice were euthanized and tumors were harvested,

135 weighed, measured and processed for analysis (each group n=10).

136 For the assays with the PKC inhibitors, one million NIH 3T3 cells stably expressing

137 either HA-KRASG12V or HA-KRASG12V-S181D were subcutaneously injected into

138 both flanks of Swiss nude mice. When tumor reached a designated volume of ~150 mm^3,

139 animals were randomized and divided into vehicle (DMSO), Bryostatin-1 or Edelfosine

140 treatment groups. Mice were weighed daily and received an intraperitoneal injection of

141 either 75 μg/Kg Bryostatin-1 in 5% (v/v) DMSO, 30 mg/Kg Edelfosine in 5% (v/v)

142 DMSO or 5% (v/v) DMSO (vehicle) for 5 days. At day 6 after the beginning of the

143 treatment, mice were euthanized and tumors were harvested, weighed and processed for

144 analysis.

145   All mouse experiments were performed in accordance with protocols approved by the

146   Animal Care and Use Committee of ICO-IDIBELL Hospitalet de Llobregat, Barcelona,

147   Spain.

148

149   **Sample Lysis, Gel electrophoresis, immunoblotting,**

150   Cultured cells were lysed in Ras extraction buffer (20 mM Tris–HCl, pH 7.5, 2 mM

151   EDTA, 100 mM NaCl, 5 mM MgCl$_2$, 1% (v/v) Triton X-100, 5 mM NaF, 10% (v/v)

152   glycerol and 0.5% (v/v) 2-mercaptoethanol) supplemented with a cocktail of protease

153   and phosphatase inhibitors (0.1 mM Na$_3$VO$_4$, 1 mM phenylmethylsulfonyl fluoride, 10

154   mM β-glycerophosphate, 2 μg/ml aprotinin and 10 μg/ml leupeptin)

155   Tumors were lysed using Polytron (Fischer Scientific, Pittsburg, PA, USA) in Ras

156   extraction buffer, and protein resolved using standard SDS-PAGE electrophoresis. After

157   electrotransfer, membranes were incubated using the indicated antibodies and then

158   incubated with peroxidase-coupled secondary antibody. Immunocomplexes were

159   detected by enhanced chemiluminescence reaction ECL western blotting analysis

160   system (Amersham Biosciences, Piscataway, NJ, USA) and imaged by LAS-3000

161   (Fujifilm, Tokyo, Japan). When required, band intensity was determined using the

162   measurement tool of Multigauge 2.0 (FUJIFILM, Tokyo, Japan).

163

164   **Cell viability assay (MTT)**

165   Human cell lines or DLD-1 expressing the same amounts of HA-KRASG12V and HA-

166   KRASG12V-S181D were seeded at 10^4 cells per p96 with DMEM 10% FCS. Next day,

167   were treated with the corresponding concentration of PKC inhibitors for 48h. Then, 10

168   μL of AB solution (MTT Cell Growth Assay Kit, #CT02, Millipore) were added to each

169   well and incubated at 37ºC for 4h. Then, 0.1 mL isopropanol with 0.04 N HCl was

170   added and mixed thoroughly. Absorbance was measured with a test wavelength of 570

171   nm and a reference wavelength of 630 nm according to manufacturer's

172   recommendations.

173

174   **Measurement of Ras isoform activation**

175   RBD (Ras-binding domain of Raf-1) pull-down assays were performed as previously

176   described (14) to determine the amount of active K-, H- and NRAS.

177

178   **Histology**

179   Mice tumors were embedded either in paraffin or frozen in OCT. Paraffin sections were

180   stained following the haematoxylin-eosin standard protocol to study their histological

181   appearance. Mitotic count in 5 consecutive high-power fields (100x) was performed to

182   compare the mitotic index between groups. Frozen section in OCT were used to

183   determine apoptosis by TUNEL assay following manufacturers recommendations

184   (Roche), and to determine the percentage of proliferating cells by

185   immunohistochemistry using Ki-67 antibodies.

186

187   **Human tumors**

188   Five biopsies of Human Pancreatic Ductal Adenorcarcinoma obtained by

189   Doudenopancreatectomy were orthotopically implanted to nude mice and were

190   perpetuated at least four passages. All patients gave informed written consent to

191   participate and to have their biological specimens analyzed. The study was cleared by

192   the Ethical Committee of Hospital de Bellvitge.

193

194   **Detection of phospho-KRAS**

195 Phos-tag^{TM} SDS-PAGE. To detect phospho-KRAS from human tumor samples and

196 from nude mice grafts, a fragment of ~ 0,1 g from a tumor biopsy was homogenized in

197 0,4 mL of λ Phosphatase Lysis Buffer (50 mM Tris-HCl pH 8; 150 mM NaCl, 2 mM

198 EDTA, 10% Glycerol, 1% Nonidet P40, 5 mM DTT, 2mM $MnCl_2$) containing either

199 protease inhibitor cocktail (Halt Protease Inhibitor Cocktail, #87786, Thermo Scientific,

200 Rockford, IL USA) alone or plus phosphatase inhibitors (0.2 mM Na_3VO_4, 5 mM NaF).

201 For human cell lines, a 10 cm dish was homogenized in the λ Phosphatase Lysis Buffer

202 as described above. Then, samples homogenized with only protease inhibitors were

203 treated with recombinant Protein Phosphatase λ for 30 minutes at 30ºC according to

204 manufacturer instructions, and finally all tubes were balanced with phosphatase

205 inhibitors in order to equalize both lysis buffers.

206 Protein content was assessed by Lowry method (Lowry et al, 1951) and tubes were

207 balanced. 10 μg protein were loaded into a 12%-polyacrilamide SDS-PAGE gel

208 supplemented with 100 μM Phos-tag^{TM} and 100 μM $MnCl_2$ (according to Phos-tag^{TM}

209 SDS-PAGE protocol indicated by manufacters). The gel was run over night at 5 mA/gel

210 and soaked in a general transfer buffer containing 1 mM EDTA for 20 min followed by

211 10 minutes incubation with a transfer buffer without EDTA. Then, gels were transferred

212 over night at 50 V into a PDVF membrane that was blocked and blotted with anti-

213 KRAS (#OP24, Calbiochem).

214 Two-dimensional gel electrophoresis (2-DE). 100 μg of tumor extract prepared as

215 indicated above, were diluted to a final concentration of 7M urea, 2M thiourea, 4%

216 CHAPS {3-[(3-cholamidopropyl)-dimethylammonio]-1-propanesulfonate}, 65 mM

217 DTE, 0.1% ampholytes (Bio-Lyte 3/10, no. 163-1113; Bio-Rad) and 1.2% DestreakTM

218 Reagent (GE Healthcare, 17-6003-18) to 125 μL volume. Two-dimensional first-

219 dimension electrophoresis was performed as isoelectric focusing (IEF) with precast,

220   immobilized pH gradient (IPG) gel strips (ReadyStripTM IPG Strip, 7 cm, pH 7 to 10;

221   no. 163-2005 [Bio-Rad]) by using a PROTEAN IEF system (Bio-Rad). Sample

222   application and rehydration of the strips were carried out using the active method (50 V

223   contant) according to the manufacturer's instructions (Bio-Rad). Next focusing was

224   performed at 8000 to 20000 V per hour. IEF gels were equilibrated for 10 min in a

225   buffer containing 6 M urea, 0.375 M Tris [pH 8.8], 2% SDS, 20% glycerol, and 2%

226   [wt/vol] DTE, and the second-dimension run was carried out in SDS-polyacrylamide

227   gels. After electrophoresis, gels were transferred to PDVF membranes (Millipore) and

228   immunoblotted with antibodies against KRAS

229

230   **Statistics**

231   All analyses were performed with GraphPad Prism 5.0. Data represent mean ± SEM.

232   Mann-Whitney test was used to analyze significance levels. Specific significance levels

233   are found in figure legends. $P < 0.05$ was considered significant.

234

235

Results and Discussion

Oncogenic KRAS phosphorylation at Ser-181 is required for tumor growth

238   To test the prediction that phosphorylation at Ser181 of oncogenic KRAS was required

239   to support tumor growth, NIH3T3 stable cell lines expressing similar levels of

240   oncogenic HA-tagged G12V KRAS, namely HA-KRAS-G12V-S181 (S181), non-

241   phosphorylatable HA-KRAS-G12V-S181A (S181A) or phosphomimetic HA-KRAS-

242   G12V-S181D (S181D) (Fig. 1A) were subcutaneously injected into nude mice and

243   tumor growth was monitored over time. Tumor formation was nearly abolished in cells

244   expressing non-phosphorylatable S181A (Fig. 1B,C and tables S1 and S2). Furthermore,

245   a dramatic increase in tumor growth was observed for phosphomimetic S181D mutant

246   compared to the phosphorylatable S181. No tumor growth was observed when injecting

247   NIH3T3 cells stably expressing wild-type HA-KRAS (Table S1), which confirmed that

248   both engraftment and growth was driven by our oncogenic KRAS phosphomutants.

249   Interestingly, in spite of the dramatic diminished growth of non phosphorylatable

250   S181A derived tumors, KRAS oncoprotein was overexpressed in those tumors

251   compared to the S181 or S181D derived tumors (Fig. 1D). This suggests that, during the

252   process of tumor development, cells with higher expression of non-phosphorylatable

253   KRAS are positively selected in an attempt to overcome the lower tumorogenic activity

254   exhibited by this mutant. Similar results were obtained when injecting in nude mice

255   two independently immortalized S181A clones with distinct expression levels. Again in

256   S181A clones, tumor growth was highly compromised irrespectively of the KRAS

257   protein expression level (Fig. S1).

258   The impaired tumor growth of the non-phosphorylatable S181 G12V mutant associated

259   with a distinct histological pattern. S181A tumors were composed mostly by cells with

260   an epithelioid appearance and with a significant lymphocytic infiltration (S181A 10.00

261   ± 1.08 lymphocytes per x100 field vs S181 2.25 ± 0.63 vs S181D 1.25 ± 0.25,

262   p<0.0001) (Fig. 3 and Fig.S3). Of note, the same histology has been previously reported

263   for sarcomas harboring the mild *KRAS* codon 13 mutations (15). In contrast, S181 and

264   S181D derived sarcomas were composed of a fusocellular population showing a

265   hemangiopericitoid patterns. The non-phosphorylatable S181A tumors had also a lower

266   mitotic rate (S181A 4.00 ± 1.53 mitotic cells per x100 field vs 27.00 ± 4.12 for S181 vs

267   54.75 ± 4.99 for S181D, p<0.0001) and were the only tumors showing detectable

268   levels of cleaved-caspase 3, a bona fide apoptosis marker (Fig. 1D and Fig S2A),

269   together with a significant increase in TUNEL positive cells (Fig. 2A, B.). This is in

270   accordance with the decreased resistance to apoptosis already reported in vitro for

271   S181A compared to S181 and S181D oncogenic KRAS mutants (14). In agreement with

272   the prediction of a stronger activity of KRAS upon phosphorylation, the

273   phosphomimetic S181D derived tumors exhibited higher ERK and AKT activity (Fig.

274   1D), accompanied by a pronounced increase in the number of positive cells for the

275   proliferative marker Ki-67 (Fig. 2A,B). Although mild increase in TUNEL positive

276   cells was also observed in S181D compared to S181 tumors, S181A tumors were the

277   ones exhibiting the highest degree of apoptosis. Intriguingly, S181A tumors showed

278   higher cyclin B1 expression than the others (Fig. 1D) in line with the requirement of

279   increased Cyclin B1 for apoptosis induction previously reported in several tumor cell

280   lines (16-18). Moreover, Cyclin B1 overexpression has already been related to the mild

281   transforming phenotype of codon 13 *KRAS* mutations in NIH3T3 models (15).

282   Thus, the impossibility of phosphorylating oncogenic KRAS dramatically changes

283   growth pattern rendering activating mutations much less aggressive and demonstrating

284   the relevance of this posttranslational modification in KRAS-driven transformation.

285

286   **PKC inhibitors diminish oncogenic KRAS-mediated tumor growth**

287   The dependence of oncogenic KRAS on S181 phosphorylation makes oncogenic KRAS

288   a putative target for protein kinase inhibitors. Since PKCs are considered to be the

289   putative kinases for KRAS Ser181 phosphorylation (13, 19, 20) we tested whether

290   treatment with two general PKC inhibitors that are clinically relevant (Bryostatin-1 and

291   Edelfosine) (21-25) were able to revert tumor growth in a dephosphorylation-dependent

292   manner.

293   Bryostatin-1 inhibits PKC activity when administrated in vitro at concentrations as low

294   as 0.1 nM (21). In our experiments we used 75 µg/Kg, a dose that was previously used

295   for in vivo PKC inhibition (26). As shown in Fig. 4B, Bryostatin-1 treatment

296   significantly reduced tumor growth of S181 whereas no effect was evident on "non-

297   dephosphorylatable" S181D tumors. Of note, we found that Bryostatin-1 treatment, in

298   accordance to its general PKC inhibitor activity, efficiently downregulated both total

299   and active PKCδ levels as previously described (21) (Fig. 4C). Tumor reduction with

300   Bryostatin-1 treatment was associated with a decreased ERK activity that was specific

301   for S181 phosphorylatable mutant. Moreover, apoptosis was induced as shown by an

302   increase of cleaved caspase-3 levels and TUNEL positive cells (Fig. 4C,D and Fig.

303   S2B). Concomitantly, cell proliferation was inhibited (Fig. 4D.) while cyclin B1

304   expression was increased (Fig. 4C). In this way, Bryostatin-1 treatment showed high

305   specificity for the dephosphorylatable S181 tumors and interestingly, treatment of these

306   tumors efficiently recapitulated the growth and signaling pattern of S181A tumors

307   (21)shown in Fig.1. Concordantly, PKC inhibition did not affect growing and signaling

308   pattern, nor increased apoptosis in the non-dephosphorylatable KRAS S181D tumors.

309   To further confirm the striking results obtained with Bryostatin-1 treatment on mice we

310   treated the same stable transfected NIH3T3 mice grafts with Edelfosine. This is an ether

311   lipid analog to HMG with reported strong PKC inhibitor activity both in vitro (27, 28)

312   and in vivo (29). As shown in Fig S4, we reproduced a significantly reduced tumor

313   growth of S181 and again no significant effect was observed in "non-

314   dephosphorylatable" S181D tumors.

315   Altogether, these results suggested that both Bryostatin-1 and Edelfosine, by blocking

316   PCK activity, impair tumor growth inducing KRAS dephosphorylation and subsequent

317   apoptosis. To formally prove this hypothesis, detection of KRAS phosphorylation was

318   necessary. Since no suitable antibodies are available, we used the Phos-Tag^{TM}-based

319   approach (30, 31) to determine the oncogenic KRAS phosphorylation status in the

320   generated tumors. This method is based on the fact that a complex formation between

321   the phosphate group of a phosphorylated protein and a divalent metal ion in Phos-Tag™

322   reduces the mobility of the phospho-protein during the electrophoresis separation, thus

323   allowing resolution of phosphorylated and non-phosphorylated proteins into different

324   bands. As shown in Fig. 4E, a slow migrating band of HA-KRAS could be observed in

325   the tumors generated from cells expressing the S181 oncogenic KRAS, that was absent

326   in S181D tumors. Disappearance of this band upon λ Phosphatase treatment

327   corroborated it was phosphorylated KRAS. Most interestingly, in Bryostatin-1 and

328   Edelfosine treated animals phosphorylated KRAS was no longer observed (Fig.4E and

329   Fig.S4). Together, these observations reinforce the notion that PKC-dependent Ser181-

330   phosphorylation of oncogenic KRAS is required for tumorigenesis. This effect may

331   account for the previously reported inhibition of different KRAS-driven tumor

332   xenografts by PKC pharmacologic inhibition (29, 32, 33). Interestingly, it has also been

333   shown that PKCδ knock-down prevents apoptosis and promotes tumorigenesis in cells

334   addicted to aberrant KRAS signaling (34-36).

335

336   **Human cell lines require phosphorylation of KRAS for survival and tumor growth**

337   In order to determine whether the requirement for KRAS phosphorylation observed in

338   our NIH3T3 KRAS-transformation model was also involved in human cell lines

339   tumorigenesis, we ectopically expressed the HA-KRAS-G12V phosphomutants

340   described above in the human colorectal cancer cell line DLD-1 but previously

341   knocked-out for the oncogenic endogenous KRAS allele (DLD1$^{KRASwt/-}$).

342   We found that under serum-saturating growth conditions (10%FCS), human colon

343   cancer cells DLD-1 expressing S181A mutant exhibited a significantly reduced growth

344   compared to phosphomimetic S181D expressing cells (Fig. 5A). Trying to recapitulate

345   tumor growth conditions, we evaluated cell growth under serum-limiting conditions

346   (0.1% FCS). After 4 days of starvation, cells stably expressing S181A showed

347   significantly higher reduced growth under serum starvation culture conditions compared

348   to S181 and phosphomimetic S181D (Fig. 5A,). Moreover, S181A exhibited increased

349   sensitivity to apoptosis under serum deprivation or by adriamycin- induced genotoxic

350   damage (Fig 5A,B), thus demonstrating a pro-apoptotic effect of the S181A oncogenic

351   KRAS.

352   In order to evaluate real tumorigenic capacity of these cells, subcutaneous injection of

353   DLD1$^{KRASwt/-}$ expressing either HA-KRASG12V or HA-KRASG12V-S181A

354   phosphomutants was performed. S181A derived tumors were significantly smaller than

355   S181 tumors (Fig. 5C). This confirmed the requirement of KRAS S181 phosphorylation

356   for tumorigenesis of human colon cell lines.

357   A preferential activation of endogenous wild-type H- and N- RAS alleles induced by the

358   oncogenic KRAS has recently been reported (37). To check whether diminished growth

359   capacity of S181A was due to lack of activation of the endogenous RAS isoforms,

360   RBD pull-down assays were performed to test GTP loading of endogenous RAS

361   isoforms. Lower GTP loading of endogenous Ras in the S181A expressing cells was not

362   observed compared with the other phosphomutants (Fig. S5).

363   Next, we wanted to determine whether BIM and Gö6983, two PKC inhibitors (38, 39)

364   were able to affect DLD-1 cells in a KRAS S181-specific manner. To this aim, dose-

365   response to these PKC inhibitors was evaluated in DLD1$^{KRASwt/-}$ cells expressing HA-

366   KRASG12V and using DLD-1 expressing HA-KRASG12V-S181D as a non-

367   dephosphorylatable control. After 48h of treatment, it was shown that at doses between

368   1µM and 20µM for BIM and 1.5µM and 10µM for Gö6983, cells expressing oncogenic

369   KRAS with S181 exhibited significantly enhanced sensitivity to PKC inhibition

370   compared to the phosphomimetic non-dephosphorylatable mutant (Fig 6A). Most

371   importantly, after PKC inhibition, S181 cells lost its KRAS phosphorylation as shown

372   by Phos-tag^{TM} SDS-PAGE gels (Fig 6A).

373   Finally, we evaluated the ability of a set of PKC inhibitors to reduce proliferation

374   together with KRAS phosphorylation in a panel of human cell lines from different

375   origin harboring oncogenic KRAS. We found that at doses reported to inhibit PKC (21,

376   28, 38, 40), cell growth was compromised. Most importantly, after 12h of treatment,

377   band corresponding to phospho-KRAS was lost, thus reinforcing the idea that PKC

378   inhibition is able to revert growth in a KRAS S181-dependent manner (Fig. 6B).

379

380   **KRAS is phosphorylated in human tumors.**

381   Next, we investigated whether the S181 phosphorylation observed in our model system

382   was also present in human tumors. To do so, a set of orthotopic xenografts derived from

383   carcinomas of the exocrine pancreas were analyzed. Five tumors harboring codon 12

384   KRAS mutations were tested. As shown in Fig. 7, by using Phos-Tag^{TM} SDS-PAGE,

385   several bands were detected in all tumors using the anti-KRAS antibody. Treatment

386   with λ phosphatase (Fig. 7) and two-dimensional electrophoresis analysis corroborated

387   the presence of phosphorylated KRAS in these human tumors (Fig. S6). Thus, the

388   presence of phospho-KRAS in human malignancies emphasizes the alleged requirement

389   of this modification for human KRAS-driven tumorigenesis.

390   Altogether, the results depict a scenario of a novel tight regulation of KRAS

391   oncogenicity by phosphorylation at S181. We have recently shown that although

392   phosphorylated KRAS is mainly found at the plasma membrane (40, 41),

393   phosphorylated K-Ras forms distinct plasma membrane signaling platforms that induce

394   preferential activation of main KRAS effectors involved in oncogenesis . Interestingly,

395   this distinct functionality could be reverted by PKC inhibitors (40).   This would give a

396   rationale for the strikingly different tumorigenic activity of oncogenic KRAS according

397   to its S181 phosphorylation status .

398   The fact that, as we show, this could be efficiently pharmacologically inhibited raises

399   the possibility of novel therapeutic strategies targeting KRAS-driven human

400   malignancies. So far, clinical trials with PKC regulators have been disappointing

401   mostly because of the lack of selectivity and unacceptable toxicity (39). The

402   identification of KRAS as a key PKC target may help in developing specific inhibitors

403   of KRAS phosphorylation.

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521   **Figure Legends**

522   **Fig. 1. Phosphorylation at Ser181 is necessary for tumor growth of oncogenic**

523   **KRAS-G12V.** NIH 3T3 cells stably expressing either, HA-KRAS-G12V (S181), HA-

524   KRAS-G12V-S181A (S181A) or HA-KRAS-G12V-S181D (S181D) were injected into

525   each flank of nude mice (each group n=10). A) Levels of exogenous HA-KRAS from

526   the different NIH 3T3 pools were analyzed by immunoblot the day of injection into

527   mice. Pool S181#2 was injected into mice (herein after referred to as S181); B) Tumor

528   volumes were measured at days 13, 16 and 19 after injection; C) At day 19 mice were

529   euthanized and tumors were dissected and weighed. Graph showing the weight of

530   excised tumors (each dot corresponds to a tumor); D) Total cell lysates of representative

531   excised tumors were immunoblotted to detect the indicated proteins (numbers indicate

532   different tumors). Anti-GAP120 was used as loading control;

533   **Fig. 2. Lack of growth of non-phospohorylatable KRAS tumors correlates with**

534   **decreased proliferation (Ki-67) and increased apoptosis (TUNEL) markers**. A)

535   Tumor sections were stained for Ki-67 or TUNEL. Scale bars represent 50 µm;B)

536   Quantifications of Ki-67 labeling (left) and TUNEL labeling (right) were made from at

537   least 2 different tumors per mutant (each point represents a counted field). (***,

538   $p < 0.0001$, **, $p < 0.001$ and *, $p < 0.01$, p for Student's two-tailed t test; mean and SEM

539   are represented).

540   **Fig. 3. Differential tumor growth according to KRAS phosphorylation is associated**

541   **with a distinct histological patern**. Paraffin sections were stained following the

542   haematoxylin-eosin protocol to study their histological appearance. Arrow caps indicate

543   lymphocyte infiltration. Scale bars represent 50 µm.

544   **Fig. 4. Pharmacologic inhibition of PKC activity inhibits tumor growth and**

545   **KRAS-G12V dependent signaling pathways in a K-RasG12V Ser181-**

546   **phosphorylation dependent manner.** NIH 3T3 cells stably expressing either HA-

547   KRAS-G12V (S181); or HA-KRAS-G12V-S181D (S181D) were injected into each

548   flank of nude mice. When tumor reached a designated volume of ~150 mm^3 (latency

549   time shorter for S181D tumors), animals were divided into two groups (each group

550   n=10) and treated daily either with vehicle (5% DMSO) or Bryostatin-1 (Bryo) (75

551   µg/Kg) for 6 days, and euthanized next day. **A)** Western blot showing HA-K-Ras

552   expression in different pools of NIH 3T3 cells the day of injection. Injected NIH 3T3

553   pools (red arrows) were chosen among the ones with equivalent expression for HA-

554   KRAS-G12V (S181) or HA-KRAS-G12V-S181D (S181D); **B)** Increment in tumor size

555   was obtained by comparing tumor volume at the starting day (day 1) and at day 7 of

556   treatment. Dissected tumors from the nude mice are displayed below the graph. Scale

557   bar represents 25 mm **C)** Total cell lysates of representative excised tumors were

558   immunoblotted to detect the indicated proteins (numbers indicate different tumors).

559   Anti-GAP120 was used as loading control **D)** Quantifications of Ki-67 labeling (left)

560   and TUNEL labeling (right) were made from at least 2 different tumors per mutant

561   (each point represents a counted field). (***, $p<0.0001$, **, $p<0.001$ and *, $p<0.01$, p

562   value for Student's two-tailed t test; ns: non-significant differences; mean and SEM are

563   represented). **E)** Cellular extract from tumors were resolved in Phos-Tag^{TM} SDS-PAGE

564   gels and immunoblot was performed using anti-HA antibody. An aliquot of a S181

565   tumor from an animal treated with DMSO was incubated, prior to electrophoresis, at

566   30ºC with phosphatase λ (λ) or only with buffer (Ctl) to discard unspecific effects due to
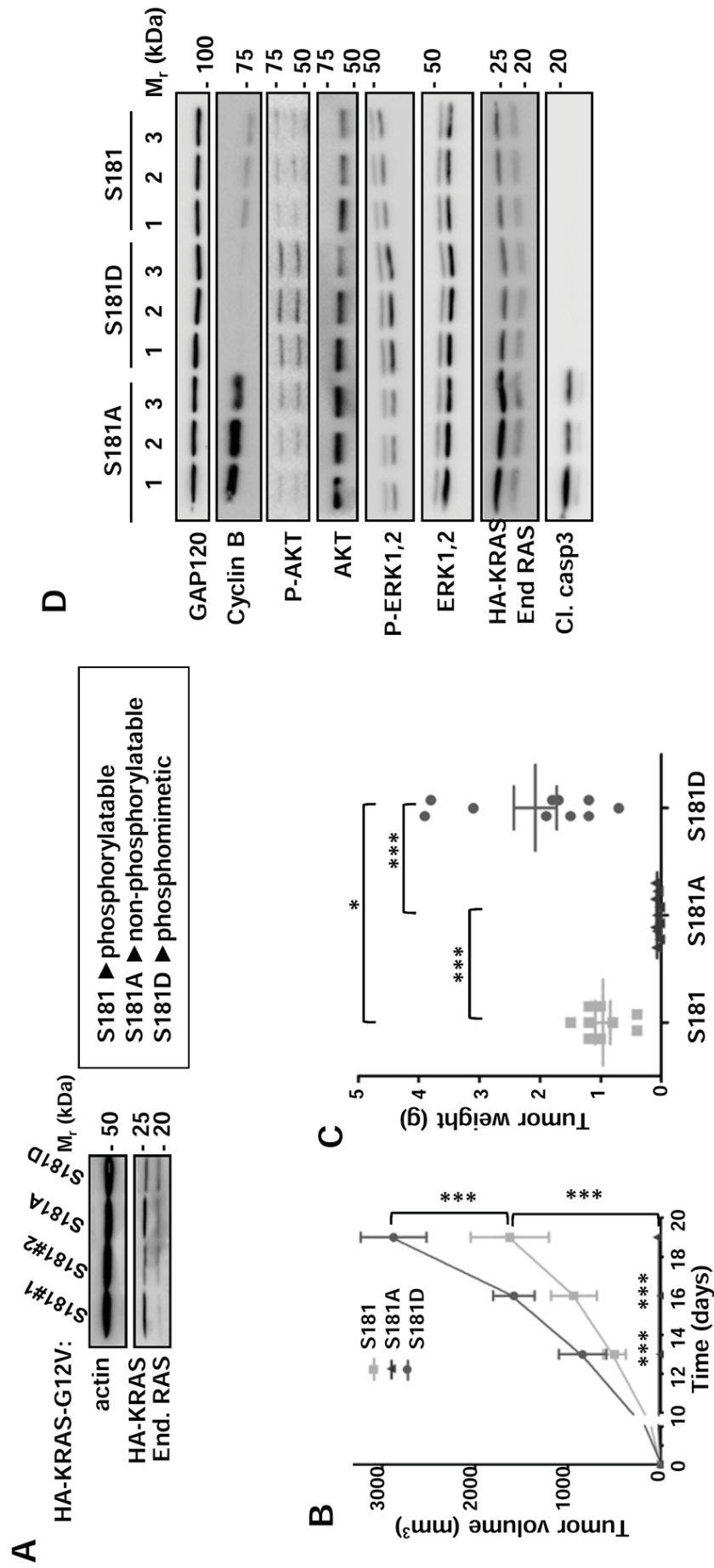
567   heating samples.

568   **Fig. 5**. **Phosphorylation at Ser181 is necessary for cell proliferation, survival and**

569   **tumor growth of DLD1** $^{KRAS\ wt/-}$ **cell expressing KRASG12V phosphomutants.** A)

570   $3·10^4$ DLD1 $^{KRAS\ wt/-}$ cells stably expressing either HA-KRASG12V-S181A, HA-

571   KRASG12V-S181D or HA-KRASG12V-S181 were cultured under serum-saturated

572   (10% FCS) or serum-limiting (0.1% FCS) conditions for 4 days and counted to evaluate

573   the proliferation rate. KRAS expression of the different mutants at the initial day and

574   levels of cleaved caspase 3 (Cl. casp 3) at the fourth day are showed. B) The different

575   mutants were cultured for 2 days with adryamicin (5µM) to induce genotoxic damage.

576   The sensitivity to apoptosis was analyzed by the levels of cleaved caspase 3 (Cl. casp

577   3). C) Pools of DLD1 $^{KRAS\ wt/-}$ expressing HA-KRAS-G12V S181 or HA-KRAS-G12V

578   S181A were injected into each flank of nude mice. Levels of exogenous HA-KRAS

579   from the pools injected were analyzed by immunoblot the day of injection into mice. At

580   day 28 after injection mice were euthanized and tumors were weighed. Graph showing

581   the weight of excised tumors.

582

583   **Fig.6. Pharmacologic PKC inhibition impairs KRAS phosphorylation and cell**

584   **survival.** A) DLD1 $^{KRAS\ wt/-}$ expressing KRASG12V -S181 or KRASG12V-S181D (as

585   non-dephosphorylatable control) were treated with the PKC inhibitors Gö6938 (Gö)

586   (1.5, 5 and 10 µM) and BIM (1, 10 and 20 µM) for 48 hours. The columns represent

587   the growth rate estimated by the measurement of the absorbance following the MTT

588   assay as a function of the initial cell number (left). Extracts from DLD1 KRAS wt/-

589   expressing KRASG12V -S181 or KRASG12V-S181D treated or no with phosphatase λ

590   (λ) or with PKC inhibitors (Gö6983 or BIM) were resolved in Phos-Tag^{TM} SDS-PAGE

591   gel following by immunoblotting using anti-KRAS antibodies (right).**B)** A pannel of

592   human cell lines from different origin harboring oncogenic KRAS were treated with 5

593   µM BIM, 1 µM Gö6983 (Gö), 1 µM Bryostatin-1(Bryo) and 10 µg/mL Edelfosine

594   (Edelf) for 48h. Cells were harvested and extracts were resolved in Phos-Tag^{TM} SDS-

595   PAGE and immunoblotted using anti-KRAS antibody.

596   **Fig. 7. Detection of the phosphorylated form of oncogenic KRAS in human**

597   **pancreatic ductal adenocarcinomas.** Extract from 5 different human pancreatic ductal

598   adenocarcinomas with oncogenic mutations in codon 12 of KRAS (#1 G12D

599   heterozygous; #2 G12D heterozygous; #3 G12D homozygous; #4 G12V heterozygous;

600   #5 G12V heterozygous), were resolved in Phos-Tag^{TM} SDS-PAGE or SDS-PAGE

601   followed by immunoblotting using anti-KRAS antibodies. An aliquot of each extract

602   was previously incubated with phosphatase λ. Anti-GAP120 was used as loading
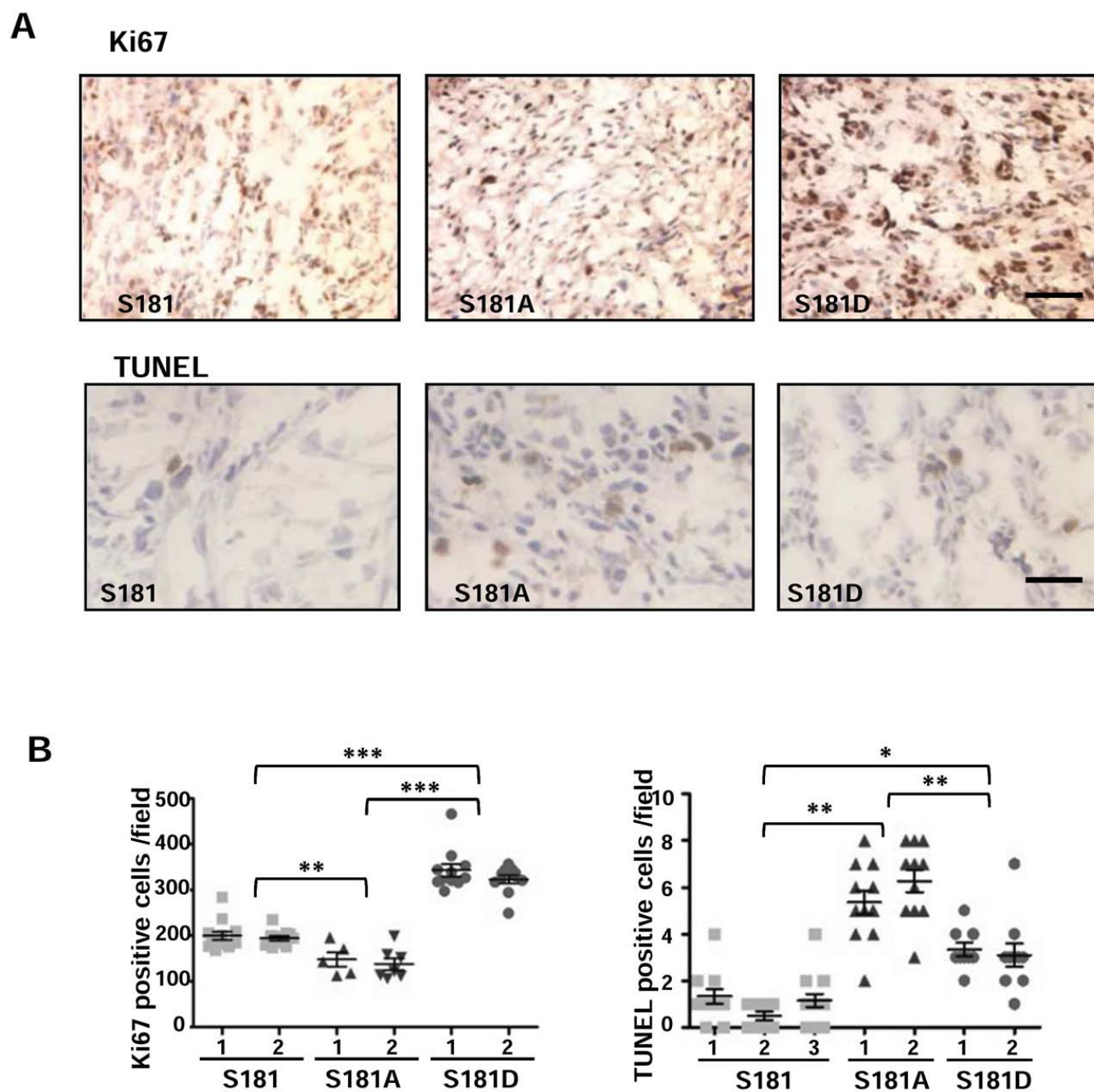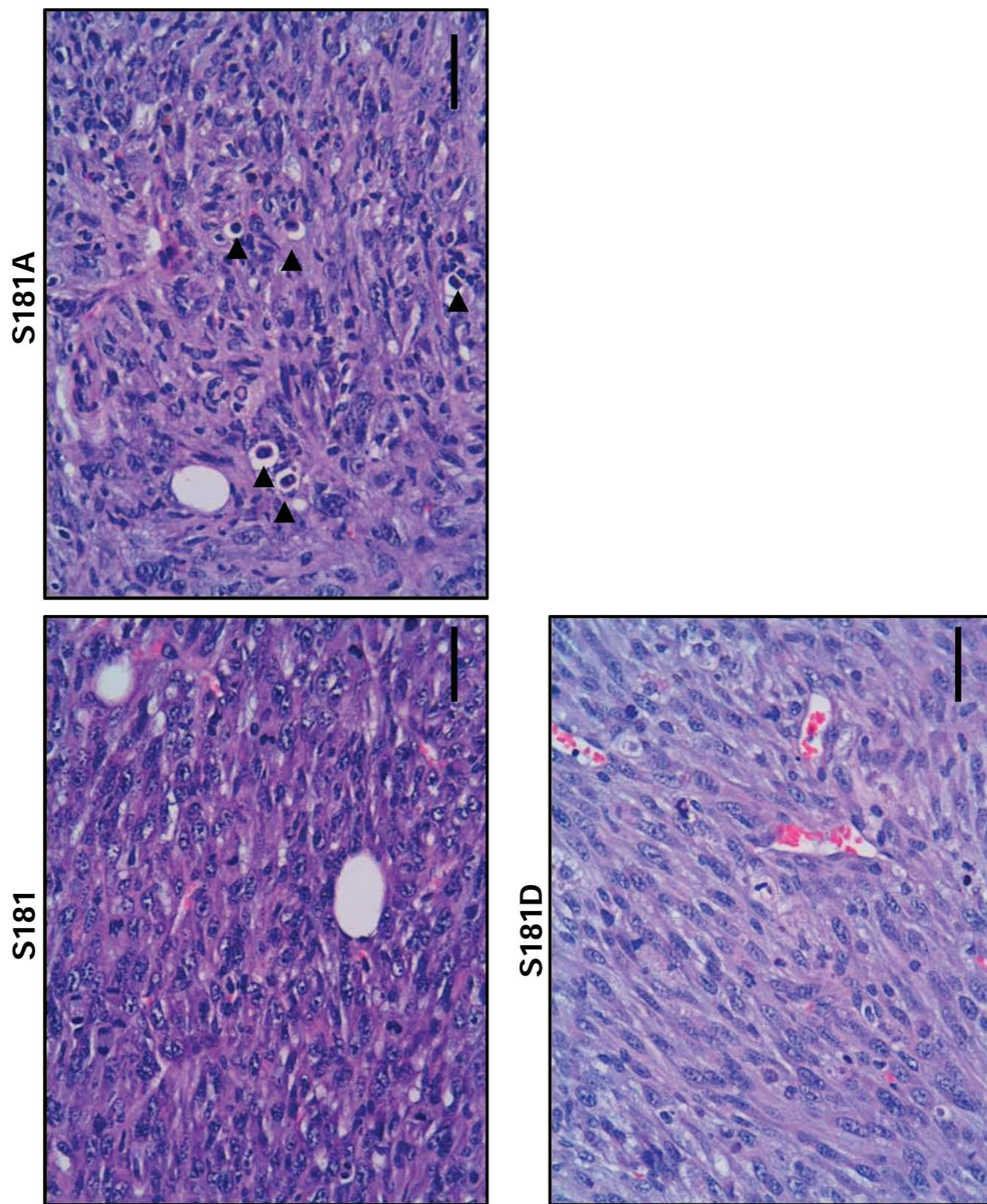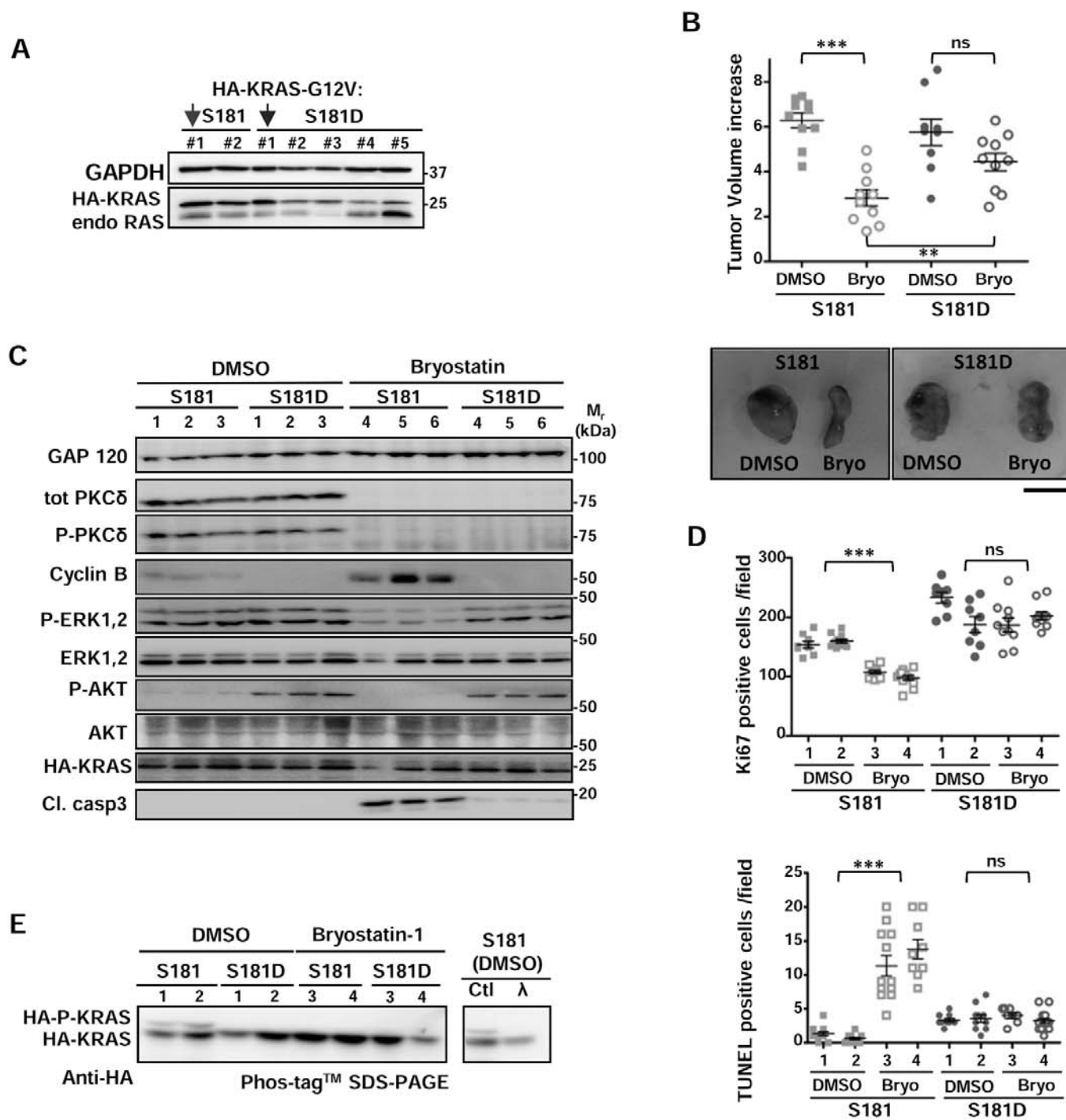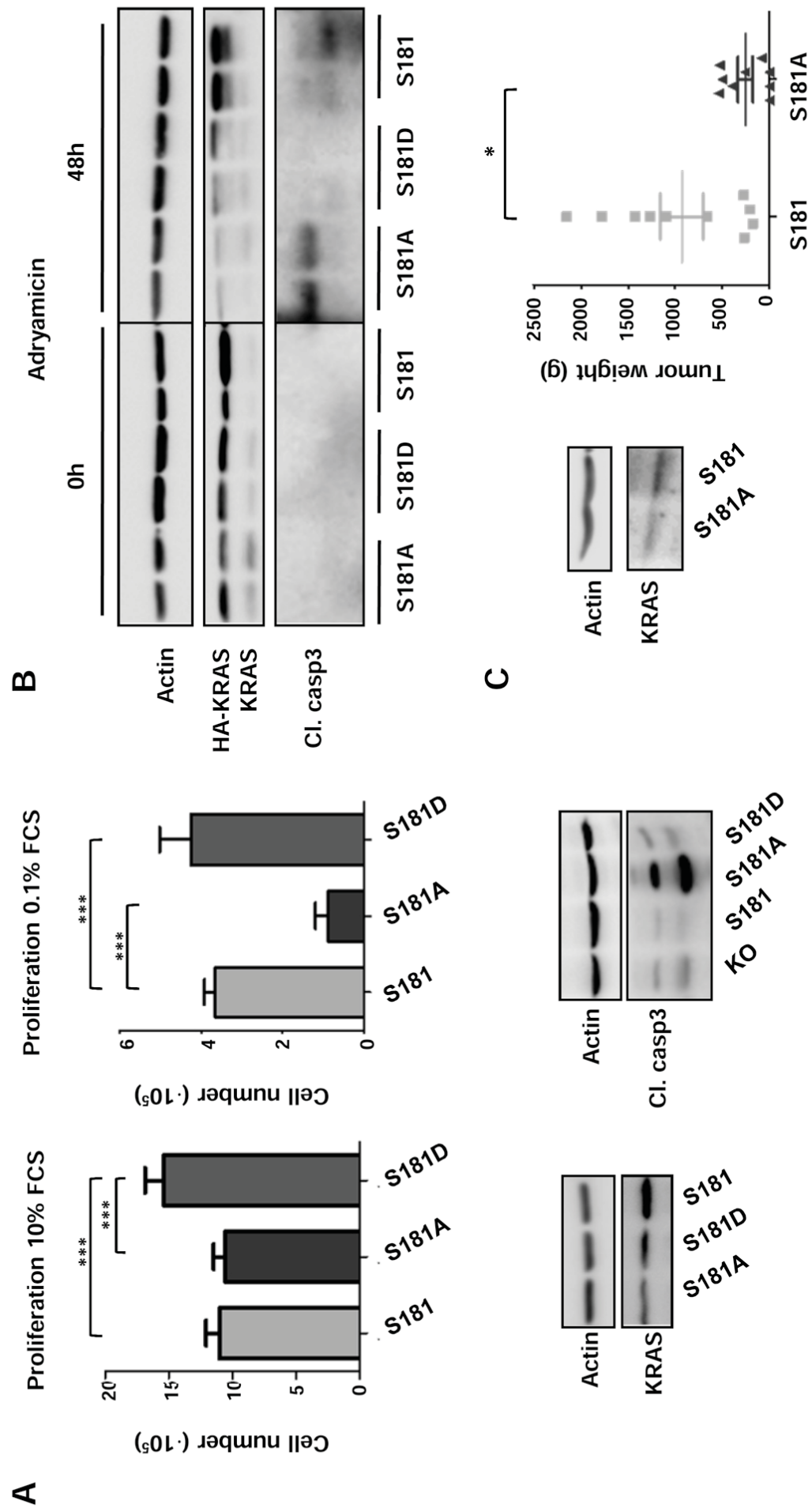
603   control.

Fig 1

Fig 2

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