



UNIVERSITAT DE
BARCELONA

Study of extracellular matrix remodeling and the role of periostin b during zebrafish heart regeneration

Anna Garcia Puig

ADVERTIMENT. La consulta d'aquesta tesi queda condicionada a l'acceptació de les següents condicions d'ús: La difusió d'aquesta tesi per mitjà del servei TDX (www.tdx.cat) i a través del Dipòsit Digital de la UB (diposit.ub.edu) ha estat autoritzada pels titulars dels drets de propietat intel·lectual únicament per a usos privats emmarcats en activitats d'investigació i docència. No s'autoritza la seva reproducció amb finalitats de lucre ni la seva difusió i posada a disposició des d'un lloc aliè al servei TDX ni al Dipòsit Digital de la UB. No s'autoritza la presentació del seu contingut en una finestra o marc aliè a TDX o al Dipòsit Digital de la UB (framing). Aquesta reserva de drets afecta tant al resum de presentació de la tesi com als seus continguts. En la utilització o cita de parts de la tesi és obligat indicar el nom de la persona autora.

ADVERTENCIA. La consulta de esta tesis queda condicionada a la aceptación de las siguientes condiciones de uso: La difusión de esta tesis por medio del servicio TDR (www.tdx.cat) y a través del Repositorio Digital de la UB (diposit.ub.edu) ha sido autorizada por los titulares de los derechos de propiedad intelectual únicamente para usos privados enmarcados en actividades de investigación y docencia. No se autoriza su reproducción con finalidades de lucro ni su difusión y puesta a disposición desde un sitio ajeno al servicio TDR o al Repositorio Digital de la UB. No se autoriza la presentación de su contenido en una ventana o marco ajeno a TDR o al Repositorio Digital de la UB (framing). Esta reserva de derechos afecta tanto al resumen de presentación de la tesis como a sus contenidos. En la utilización o cita de partes de la tesis es obligado indicar el nombre de la persona autora.

WARNING. On having consulted this thesis you're accepting the following use conditions: Spreading this thesis by the TDX (www.tdx.cat) service and by the UB Digital Repository (diposit.ub.edu) has been authorized by the titular of the intellectual property rights only for private uses placed in investigation and teaching activities. Reproduction with lucrative aims is not authorized nor its spreading and availability from a site foreign to the TDX service or to the UB Digital Repository. Introducing its content in a window or frame foreign to the TDX service or to the UB Digital Repository is not authorized (framing). Those rights affect to the presentation summary of the thesis as well as to its contents. In the using or citation of parts of the thesis it's obliged to indicate the name of the author.

Control 1	Uncharacterized protein OS=Danio rerio GN=sich211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	94,40%	1	1	2	0,01%	6,81%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=slc2: E9QJ49_DANRE,F1QAJ7_I	39.378,90	39,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=slc2: E9QJ49_DANRE,F1QAJ7_I	39.378,90	39,20%	0	0	0	0,00%	0,00%
Control 1	Cox5aa protein OS=Danio rerio GN=cox5aa PE=1 SV=1 Q4VBU7_DANRE	15.897,10	100,00%	2	2	2	0,01%	27,70%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=vtg5 F1R255_DANRE	149.378,70	100,00%	3	3	4	0,01%	2,35%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=laml F1QE66_DANRE,F1REL9_I	197.673,00	99,90%	1	1	1	0,00%	0,70%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=laml F1QE66_DANRE,F1REL9_I	197.673,00	99,90%	1	1	1	0,00%	0,70%
Control 1	Uncharacterized protein OS=Danio rerio GN=pdha1a PE=4 SV A0A0R4IT09_DANRE,Q6P5	43.741,90	100,00%	1	1	2	0,01%	2,25%
Control 1	Uncharacterized protein OS=Danio rerio GN=pdha1a PE=4 SV A0A0R4IT09_DANRE,Q6P5	43.741,90	100,00%	1	1	2	0,01%	2,25%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttnb G1K2X0_DANRE	2.458.846,80	100,00%	3	3	4	0,01%	0,14%
Control 1	Uncharacterized protein OS=Danio rerio GN=tln1 PE=4 SV=1 A0A0R4ID28_DANRE,A0AI	270.987,00	100,00%	6	6	7	0,03%	2,75%
Control 1	Uncharacterized protein OS=Danio rerio GN=tln1 PE=4 SV=1 A0A0R4ID28_DANRE,A0AI	270.987,00	100,00%	6	6	7	0,03%	2,75%
Control 1	Aspartate aminotransferase OS=Danio rerio GN=got2a PE=1 ! Q7SYK7_DANRE	47.588,10	99,90%	1	1	2	0,01%	1,87%
Control 1	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1 A0A0R4IEB4_DANRE,A0AI	128.154,40	100,00%	2	2	2	0,01%	2,31%
Control 1	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1 A0A0R4IEB4_DANRE,A0AI	128.154,40	100,00%	2	2	2	0,01%	2,31%
Control 1	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1 A0A0R4IEB4_DANRE,A0AI	128.154,40	100,00%	2	2	2	0,01%	2,31%
Control 1	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1 A0A0R4IEB4_DANRE,A0AI	128.154,40	100,00%	2	2	2	0,01%	2,31%
Control 1	ATP synthase subunit beta OS=Danio rerio GN=atp5b PE=1 S\ A8WGC6_DANRE	55.130,90	100,00%	13	15	42	0,15%	42,20%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttnb B0S6Y0_DANRE	626.764,10	96,30%	1	1	1	0,00%	0,61%
Control 1	Hbaa1 protein OS=Danio rerio GN=hbaa1 PE=1 SV=1 HBA_DANRE,Q1JQ69_DAI	15.565,70	99,70%	1	1	1	0,00%	7,69%
Control 1	Hbaa1 protein OS=Danio rerio GN=hbaa1 PE=1 SV=1 HBA_DANRE,Q1JQ69_DAI	15.565,70	99,70%	1	1	1	0,00%	7,69%
Control 1	Hbaa1 protein OS=Danio rerio GN=hbaa1 PE=1 SV=1 HBA_DANRE,Q1JQ69_DAI	15.565,70	99,70%	1	1	1	0,00%	7,69%
Control 1	Caveolin OS=Danio rerio GN=cav1 PE=1 SV=1 Q6YLH8_DANRE,Q6YLH9_	20.652,80	89,50%	0	0	0	0,00%	0,00%
Control 1	Caveolin OS=Danio rerio GN=cav1 PE=1 SV=1 Q6YLH8_DANRE,Q6YLH9_	20.652,80	89,50%	0	0	0	0,00%	0,00%
Control 1	Tgm2b protein OS=Danio rerio GN=tgm2b PE=2 SV=1 Q1ECX3_DANRE	75.774,00	100,00%	5	6	7	0,03%	8,98%
Control 1	Uncharacterized protein OS=Danio rerio GN=krt94 PE=1 SV=1 F1R8U0_DANRE	47.787,30	100,00%	3	3	5	0,02%	8,12%
Control 1	Uncharacterized protein OS=Danio rerio GN=aifm1 PE=1 SV=: A2BGE4_DANRE,A2BGE5_	66.977,00	99,60%	1	1	1	0,00%	1,47%
Control 1	Uncharacterized protein OS=Danio rerio GN=aifm1 PE=1 SV=: A2BGE4_DANRE,A2BGE5_	66.977,00	99,60%	1	1	1	0,00%	1,47%
Control 1	Uncharacterized protein OS=Danio rerio GN=aifm1 PE=1 SV=: A2BGE4_DANRE,A2BGE5_	66.977,00	99,60%	1	1	1	0,00%	1,47%
Control 1	Protein S100 OS=Danio rerio GN=s100a10b PE=1 SV=1 Q7ZV44_DANRE	11.157,10	100,00%	1	1	2	0,01%	13,00%
Control 1	Elongation factor 1-alpha OS=Danio rerio GN=eef1a PE=2 SV= A0A0R4IE10_DANRE,E7FC	50.150,70	99,90%	1	2	2	0,01%	2,60%
Control 1	Elongation factor 1-alpha OS=Danio rerio GN=eef1a PE=2 SV= A0A0R4IE10_DANRE,E7FC	50.150,70	99,90%	1	2	2	0,01%	2,60%
Control 1	Elongation factor 1-alpha OS=Danio rerio GN=eef1a PE=2 SV= A0A0R4IE10_DANRE,E7FC	50.150,70	99,90%	1	2	2	0,01%	2,60%
Control 1	Elongation factor 1-alpha OS=Danio rerio GN=eef1a PE=2 SV= A0A0R4IE10_DANRE,E7FC	50.150,70	99,90%	1	2	2	0,01%	2,60%
Control 1	Uncharacterized protein OS=Danio rerio GN=atp5f1 PE=1 SV= B8JIS1_DANRE	28.169,80	9,80%	0	0	0	0,00%	0,00%
Control 1	Succinyl-CoA ligase subunit beta OS=Danio rerio GN=sucla2 P Q6PBI0_DANRE	50.032,30	99,70%	1	1	1	0,00%	2,15%
Control 1	Enolase 1, (Alpha) OS=Danio rerio GN=enol1a PE=1 SV=1 Q6PC12_DANRE	47.075,10	100,00%	1	1	1	0,00%	2,55%
Control 1	Uncharacterized protein OS=Danio rerio GN=wu:fi04e12 PE= F1QAU9_DANRE	251.072,30	100,00%	2	2	2	0,01%	1,04%
Control 1	Aconitate hydratase, mitochondrial (Fragment) OS=Danio rer F8W4M7_DANRE	85.591,90	100,00%	3	4	7	0,03%	5,06%
Control 1	Uncharacterized protein OS=Danio rerio GN=col4a2 PE=1 SV= A0A0R4IUK2_DANRE	165.476,80	100,00%	6	9	17	0,06%	4,85%
Control 1	Uncharacterized protein OS=Danio rerio GN=cox6c PE=1 SV= Q6ZM23_DANRE	8.480,50	99,50%	1	1	1	0,00%	17,10%
Control 1	Catenin (Cadherin-associated protein), alpha OS=Danio rerio Q9PVF8_DANRE	100.520,10	100,00%	3	3	3	0,01%	6,50%
Control 1	Malate dehydrogenase OS=Danio rerio GN=mdh2 PE=1 SV=1 Q7T334_DANRE	35.420,30	99,90%	1	1	1	0,00%	2,97%
Control 1	Uncharacterized protein OS=Danio rerio GN=ptx3a PE=1 SV=: A3KPN4_DANRE	49.273,20	100,00%	5	6	11	0,04%	12,10%
Control 1	Uncharacterized protein OS=Danio rerio GN=tln2a PE=4 SV=: A0A0R4IIA7_DANRE,A0AC	229.740,40	100,00%	2	2	2	0,01%	1,10%
Control 1	Uncharacterized protein OS=Danio rerio GN=tln2a PE=4 SV=: A0A0R4IIA7_DANRE,A0AC	229.740,40	100,00%	2	2	2	0,01%	1,10%
Control 1	Uncharacterized protein OS=Danio rerio GN=tln2a PE=4 SV=: A0A0R4IIA7_DANRE,A0AC	229.740,40	100,00%	2	2	2	0,01%	1,10%
Control 1	Uncharacterized protein OS=Danio rerio GN=hist2h3ca1 PE=: A0A0R4IYZ6_DANRE,A8KE	17.030,40	39,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=hist2h3ca1 PE=: A0A0R4IYZ6_DANRE,A8KE	17.030,40	39,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=hist2h3ca1 PE=: A0A0R4IYZ6_DANRE,A8KE	17.030,40	39,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=hist2h3ca1 PE=: A0A0R4IYZ6_DANRE,A8KE	17.030,40	39,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=hist2h3ca1 PE=: A0A0R4IYZ6_DANRE,A8KE	17.030,40	39,20%	0	0	0	0,00%	0,00%
Control 1	Radixin OS=Danio rerio GN=msna PE=1 SV=1 Q66M42_DANRE	68.035,00	100,00%	1	1	1	0,00%	2,07%
Control 1	Uncharacterized protein OS=Danio rerio GN=col12a1a PE=1 S\ X1WF98_DANRE	321.902,50	89,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=sid:key-238c7.1: Q1LX26_DANRE	21.858,50	99,70%	1	1	1	0,00%	3,83%
Control 1	Uncharacterized protein OS=Danio rerio GN=hk1 PE=4 SV=1 A0A0R4IUT7_DANRE,Q6N	102.760,50	100,00%	5	5	7	0,03%	6,09%
Control 1	Uncharacterized protein OS=Danio rerio GN=hk1 PE=4 SV=1 A0A0R4IUT7_DANRE,Q6N	102.760,50	100,00%	5	5	7	0,03%	6,09%
Control 1	Uncharacterized protein OS=Danio rerio GN=sich211-199o1. A5PMG2_DANRE	70.660,80	99,10%	1	1	1	0,00%	1,72%
Control 1	Inner membrane protein, mitochondrial (Mitofilin) OS=Danio Q6PFS4_DANRE	83.201,30	100,00%	3	3	4	0,01%	5,02%
Control 1	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IG11_DANRE,E9QI	35.784,00	21,00%	0	0	0	0,00%	0,00%
Control 1	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IG11_DANRE,E9QI	35.784,00	21,00%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=zgc:101840 PE= Q5TYR5_DANRE,Q5TYR6_	9.470,50	99,90%	1	1	1	0,00%	18,60%
Control 1	Uncharacterized protein OS=Danio rerio GN=zgc:101840 PE= Q5TYR5_DANRE,Q5TYR6_	9.470,50	99,90%	1	1	1	0,00%	18,60%
Control 1	Uncharacterized protein OS=Danio rerio GN=zgc:101840 PE= Q5TYR5_DANRE,Q5TYR6_	9.470,50	99,90%	1	1	1	0,00%	18,60%
Control 1	Uncharacterized protein OS=Danio rerio GN=fbn2b PE=1 SV= A0A0G2KQ62_DANRE	212.420,00	100,00%	13	16	28	0,10%	8,76%
Control 1	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1 B3DG37_DANRE,HBB1_D_	16.389,20	99,60%	1	1	1	0,00%	5,41%
Control 1	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1 B3DG37_DANRE,HBB1_D_	16.389,20	99,60%	1	1	1	0,00%	5,41%
Control 1	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1 B3DG37_DANRE,HBB1_D_	16.389,20	99,60%	1	1	1	0,00%	5,41%
Control 1	Uncharacterized protein OS=Danio rerio GN=myoz2b PE=4 S\ F1QBP1_DANRE,F1QKE7_	25.288,50	15,70%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=myoz2b PE=4 S\ F1QBP1_DANRE,F1QKE7_	25.288,50	15,70%	0	0	0	0,00%	0,00%
Control 1	Alpha-1,4 glucan phosphorylase OS=Danio rerio GN=pygmb F E7EXT3_DANRE	96.868,30	100,00%	2	2	2	0,01%	2,61%
Control 1	Laminin subunit gamma-1 OS=Danio rerio GN=lamc1 PE=2 SV LAMC1_DANRE	176.205,30	100,00%	9	9	11	0,04%	7,53%
Control 1	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IG11_DANRE,E9QI	35.784,00	21,00%	0	0	0	0,00%	0,00%
Control 1	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IG11_DANRE,E9QI	35.784,00	21,00%	0	0	0	0,00%	0,00%
Control 1	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IG11_DANRE,E9QI	35.784,00	21,00%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4IUT7_DANRE,E7F4	518.896,50	100,00%	1	1	1	0,00%	0,22%
Control 1	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4IUT7_DANRE,E7F4	518.896,50	100,00%	1	1	1	0,00%	0,22%
Control 1	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4IUT7_DANRE,E7F4	518.896,50	100,00%	1	1	1	0,00%	0,22%
Control 1	Uncharacterized protein OS=Danio rerio GN=ccd141 PE=4 S\ B0S755_DANRE,E7FG99_I	190.652,30	100,00%	2	2	2	0,01%	1,26%
Control 1	Uncharacterized protein OS=Danio rerio GN=ccd141 PE=4 S\ B0S755_DANRE,E7FG99_I	190.652,30	100,00%	2	2	2	0,01%	1,26%
Control 1	Nicotinamide nucleotide transhydrogenase OS=Danio rerio G Q6NYQ7_DANRE	112.785,50	35,70%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=ckma PE=1 SV=: A2BHA3_DANRE	42.814,90	100,00%	2	2	2	0,01%	6,30%
Control 1	Iso citrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=1 A0A0G2L6I6_DANRE,E7F4	50.921,70	100,00%	2	3	5	0,02%	5,78%
Control 1	Iso citrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=1 A0A0G2L6I6_DANRE,E7F4	50.921,70	100,00%	2	3	5	0,02%	5,78%
Control 1	Uncharacterized protein OS=Danio rerio GN=wfdc2 PE=4 SV= E7F3G3_DANRE	17.778,70	100,00%	2	3	7	0,03%	13,10%
Control 1	Phosphoglycerate kinase OS=Danio rerio GN=pgk1 PE=1 SV=: F1QXV8_DANRE	44.733,20	96,30%	1	1	2	0,01%	2,40%
Control 1	Uncharacterized protein OS=Danio rerio GN=vdac2 PE=1 SV= Q8AWD0_DANRE	30.284,90	100,00%	2	2	3	0,01%	7,77%
Control 1	Uncharacterized protein OS=Danio rerio GN=myom1b PE=4 S A0A0R4IGQ8_DANRE	180.977,20	100,00%	5	5	5	0,02%	3,62%
Control 1	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV= A0A0R4IP01_DANRE,B8JK	119.385,60	99,90%	1	1	1	0,00%	0,78%
Control 1	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV= A0A0R4IP01_DANRE,B8JK	119.385,60	99,90%	1	1	1	0,00%	0,78%
Control 1	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV= A0A0R4IP01_DANRE,B8JK	119.385,60	99,90%	1	1	1	0,00%	0,78%
Control 1	Uncharacterized protein OS=Danio rerio GN=tnt2a PE=1 SV= E7EYC9_DANRE,F1R0K6_I	33.942,20	39,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=tnt2a PE=1 SV= E7EYC9_DANRE,F1R0K6_I	33.942,20	39,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=tnt2a PE=1 SV= E7EYC9_DANRE,F1R0K6_I	33.942,20	39,20%	0	0	0	0,00%	0,00%

Control 1	Uncharacterized protein OS=Danio rerio GN=prdx3 PE=1 SV= A8E7G6_DANRE,A8E7G7_	25.875,70	47,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=prdx3 PE=1 SV= A8E7G6_DANRE,A8E7G7_	25.875,70	47,20%	0	0	0	0,00%	0,00%
Control 1	Fatty acid binding protein 3, muscle and heart OS=Danio rerio Q8UVG7_DANRE	14.882,40	100,00%	1	1	1	0,00%	9,77%
Control 1	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE= A0A0R4IE46_DANRE,A0A0	101.943,10	100,00%	2	2	3	0,01%	3,25%
Control 1	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE= A0A0R4IE46_DANRE,A0A0	101.943,10	100,00%	2	2	3	0,01%	3,25%
Control 1	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE= A0A0R4IE46_DANRE,A0A0	101.943,10	100,00%	2	2	3	0,01%	3,25%
Control 1	Transforming growth factor, beta-induced OS=Danio rerio G1 Q5O3K1_DANRE	74.192,20	38,00%	0	0	0	0,00%	0,00%
Control 1	Calcium-transporting ATPase OS=Danio rerio GN=atp2a2a PE A9C3Q4_DANRE,A9C3Q5_	109.519,40	100,00%	6	6	9	0,03%	8,64%
Control 1	Calcium-transporting ATPase OS=Danio rerio GN=atp2a2a PE A9C3Q4_DANRE,A9C3Q5_	109.519,40	100,00%	6	6	9	0,03%	8,64%
Control 1	Myosin light chain 2 Mlc2a OS=Danio rerio GN=myl7 PE=1 SV Q8O1M3_DANRE	19.021,00	100,00%	6	7	20	0,07%	58,10%
Control 1	Hadha protein OS=Danio rerio GN=hadhaa PE=1 SV=1 A7Y4T7_DANRE	81.877,10	100,00%	2	2	2	0,01%	2,37%
Control 1	Fructose-bisphosphate aldolase OS=Danio rerio GN=aldoaa P Q6P043_DANRE,Q803Q7_	39.740,60	100,00%	3	3	6	0,02%	10,20%
Control 1	Fructose-bisphosphate aldolase OS=Danio rerio GN=aldoaa P Q6P043_DANRE,Q803Q7_	39.740,60	100,00%	3	3	6	0,02%	10,20%
Control 1	Uncharacterized protein OS=Danio rerio GN=tns1b PE=1 SV=1 F1Q649_DANRE	185.031,00	100,00%	3	3	3	0,01%	2,05%
Control 1	Uncharacterized protein OS=Danio rerio GN=ltbp1 PE=4 SV=2 F1Q056_DANRE	164.883,20	100,00%	3	3	3	0,01%	2,82%
Control 1	Uncharacterized protein OS=Danio rerio GN=flnca PE=4 SV=1 A0A0R4IU54_DANRE,E7F1	291.642,80	100,00%	3	3	3	0,01%	1,33%
Control 1	Uncharacterized protein OS=Danio rerio GN=flnca PE=4 SV=1 A0A0R4IU54_DANRE,E7F1	291.642,80	100,00%	3	3	3	0,01%	1,33%
Control 1	Uncharacterized protein OS=Danio rerio GN=fnh5 PE=4 SV=1 A0A0R4IWY0_DANRE,B0V	32.504,70	39,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=fnh5 PE=4 SV=1 A0A0R4IWY0_DANRE,B0V	32.504,70	39,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=fnh5 PE=4 SV=1 A0A0R4IWY0_DANRE,B0V	32.504,70	39,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=fnh5 PE=4 SV=1 A0A0R4IWY0_DANRE,B0V	32.504,70	39,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV= A0A0R4IBJ3_DANRE,F1R6	72.513,60	87,60%	1	1	1	0,00%	1,19%
Control 1	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV= A0A0R4IBJ3_DANRE,F1R6	72.513,60	87,60%	1	1	1	0,00%	1,19%
Control 1	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV= A0A0R4IBJ3_DANRE,F1R6	72.513,60	87,60%	1	1	1	0,00%	1,19%
Control 1	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV= A0A0R4IBJ3_DANRE,F1R6	72.513,60	87,60%	1	1	1	0,00%	1,19%
Control 1	Transporter OS=Danio rerio GN=slc64aa PE=2 SV=1 Q1WGB5_DANRE	72.122,80	100,00%	1	1	1	0,00%	1,39%
Control 1	Sodium/potassium-transporting ATPase subunit alpha OS=Dz Q9DGL6_DANRE	113.384,80	100,00%	3	3	6	0,02%	3,31%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	3	4	0,01%	16,10%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=lmo A0A0R4ITT8_DANRE,B0U	31.021,60	24,80%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=lmo A0A0R4ITT8_DANRE,B0U	31.021,60	24,80%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=lmo A0A0R4ITT8_DANRE,B0U	31.021,60	24,80%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=cmlc1 PE=1 SV= B0R0F7_DANRE	21.865,20	100,00%	10	13	84	0,31%	51,50%
Control 1	Uncharacterized protein OS=Danio rerio GN=tnni1b PE=1 SV= E7F5J9_DANRE,F8W417_I	21.032,10	83,30%	1	1	2	0,01%	4,19%
Control 1	Uncharacterized protein OS=Danio rerio GN=tnni1b PE=1 SV= E7F5J9_DANRE,F8W417_I	21.032,10	83,30%	1	1	2	0,01%	4,19%
Control 1	Uncharacterized protein OS=Danio rerio GN=tnni1b PE=1 SV= E7F5J9_DANRE,F8W417_I	21.032,10	83,30%	1	1	2	0,01%	4,19%
Control 1	Uncharacterized protein OS=Danio rerio GN=tnni1b PE=1 SV= E7F5J9_DANRE,F8W417_I	21.032,10	83,30%	1	1	2	0,01%	4,19%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=slc: E9QDE7_DANRE	83.925,70	94,50%	1	1	1	0,00%	1,25%
Control 1	Uncharacterized protein OS=Danio rerio GN=ank3b PE=1 SV= F1QN83_DANRE	218.497,50	100,00%	4	4	5	0,02%	2,45%
Control 1	Uncharacterized protein OS=Danio rerio GN=cox5b2 PE=1 SV Q0P3U4_DANRE	14.308,50	100,00%	1	1	3	0,01%	11,10%
Control 1	Uncharacterized protein OS=Danio rerio GN=srl PE=4 SV=1 A0A0R4IRI3_DANRE,A0A0	55.028,70	100,00%	2	2	2	0,01%	4,04%
Control 1	Uncharacterized protein OS=Danio rerio GN=srl PE=4 SV=1 A0A0R4IRI3_DANRE,A0A0	55.028,70	100,00%	2	2	2	0,01%	4,04%
Control 1	Uncharacterized protein OS=Danio rerio GN=epppk1 PE=4 SV= A0A0R4IND9_DANRE,I3IS	611.724,10	98,20%	1	1	1	0,00%	0,22%
Control 1	Uncharacterized protein OS=Danio rerio GN=epppk1 PE=4 SV= A0A0R4IND9_DANRE,I3IS	611.724,10	98,20%	1	1	1	0,00%	0,22%
Control 1	Uncharacterized protein OS=Danio rerio GN=palm1b PE=4 SV B0V0Y4_DANRE	27.123,40	100,00%	3	3	7	0,03%	12,30%
Control 1	Uncharacterized protein OS=Danio rerio GN=vwf PE=4 SV=2 E7F921_DANRE	307.905,50	100,00%	1	1	2	0,01%	0,46%
Control 1	Uncharacterized protein OS=Danio rerio GN=sdprb PE=1 SV= F1Q688_DANRE,F1R665_I	51.441,60	8,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=sdprb PE=1 SV= F1Q688_DANRE,F1R665_I	51.441,60	8,20%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=vdac3 PE=1 SV= B0R197_DANRE,B0R198_I	29.812,00	100,00%	2	2	2	0,01%	9,75%
Control 1	Uncharacterized protein OS=Danio rerio GN=vdac3 PE=1 SV= B0R197_DANRE,B0R198_I	29.812,00	100,00%	2	2	2	0,01%	9,75%
Control 1	Uncharacterized protein OS=Danio rerio GN=uqcrcl1 PE=1 SV= F1QU63_DANRE	55.450,70	100,00%	2	2	3	0,01%	6,36%
Control 1	Dihydropyridyl dehydrogenase OS=Danio rerio GN=dlhd PE=1: Q6TNU6_DANRE	53.620,50	100,00%	1	1	1	0,00%	1,97%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=plek A0A0R4I85_DANRE,F1QC	135.554,10	100,00%	2	2	2	0,01%	1,95%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=plek A0A0R4I85_DANRE,F1QC	135.554,10	100,00%	2	2	2	0,01%	1,95%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=plek A0A0R4I85_DANRE,F1QC	135.554,10	100,00%	2	2	2	0,01%	1,95%
Control 1	Uncharacterized protein OS=Danio rerio GN=tfa PE=1 SV=2 B8JL43_DANRE,E7FB1_D	73.717,60	9,50%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=tfa PE=1 SV=2 B8JL43_DANRE,E7FB1_D	73.717,60	9,50%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=tfa PE=1 SV=2 B8JL43_DANRE,E7FB1_D	73.717,60	9,50%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=c4 PE=4 SV=1 F1Q4X5_DANRE	191.050,80	100,00%	2	2	2	0,01%	1,24%
Control 1	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV A0A0R4IU91_DANRE,F6NF	94.381,10	100,00%	3	3	4	0,01%	3,77%
Control 1	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV A0A0R4IU91_DANRE,F6NF	94.381,10	100,00%	3	3	4	0,01%	3,77%
Control 1	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV A0A0R4IU91_DANRE,F6NF	94.381,10	100,00%	3	3	4	0,01%	3,77%
Control 1	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV A0A0R4IU91_DANRE,F6NF	94.381,10	100,00%	3	3	4	0,01%	3,77%
Control 1	Pyruvate kinase OS=Danio rerio GN=pkma PE=1 SV=1 A0A0R4IGP6_DANRE,Q7Z	58.059,00	99,90%	1	1	1	0,00%	1,69%
Control 1	Pyruvate kinase OS=Danio rerio GN=pkma PE=1 SV=1 A0A0R4IGP6_DANRE,Q7Z	58.059,00	99,90%	1	1	1	0,00%	1,69%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=post F1QM50_DANRE	17.277,20	100,00%	2	2	3	0,01%	16,30%
Control 1	Uncharacterized protein OS=Danio rerio GN=acs1b PE=1 SV= F1QD46_DANRE	77.694,10	99,80%	1	1	1	0,00%	1,29%
Control 1	Uncharacterized protein OS=Danio rerio GN=LOC100330353 F1QWE6_DANRE	86.981,10	100,00%	2	3	3	0,01%	3,16%
Control 1	Uncharacterized protein OS=Danio rerio GN=sorbs1 PE=4 SV= A0A0R4IP30_DANRE	85.938,80	100,00%	2	3	3	0,01%	4,31%
Control 1	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=1 SV= A0A0R4IF18_DANRE,A2BC	103.798,70	100,00%	4	4	4	0,01%	6,39%
Control 1	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=1 SV= A0A0R4IF18_DANRE,A2BC	103.798,70	100,00%	4	4	4	0,01%	6,39%
Control 1	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=1 SV= A0A0R4IF18_DANRE,A2BC	103.798,70	100,00%	4	4	4	0,01%	6,39%
Control 1	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=1 SV= A0A0R4IF18_DANRE,A2BC	103.798,70	100,00%	4	4	4	0,01%	6,39%
Control 1	Uncharacterized protein OS=Danio rerio GN=tjp1b PE=1 SV=1 A0A0R4IHR7_DANRE,E7FF	205.686,40	96,30%	1	1	1	0,00%	0,65%
Control 1	Uncharacterized protein OS=Danio rerio GN=tjp1b PE=1 SV=1 A0A0R4IHR7_DANRE,E7FF	205.686,40	96,30%	1	1	1	0,00%	0,65%
Control 1	Uncharacterized protein OS=Danio rerio GN=tjp1b PE=1 SV=1 A0A0R4IHR7_DANRE,E7FF	205.686,40	96,30%	1	1	1	0,00%	0,65%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=ltbp X1WC00_DANRE	33.442,60	96,30%	1	1	2	0,01%	9,27%
Control 1	Uncharacterized protein OS=Danio rerio GN=thbs4a PE=4 SV= A0A0R4IN91_DANRE,A0A	104.325,40	100,00%	2	2	2	0,01%	3,47%
Control 1	Uncharacterized protein OS=Danio rerio GN=thbs4a PE=4 SV= A0A0R4IN91_DANRE,A0A	104.325,40	100,00%	2	2	2	0,01%	3,47%
Control 1	Uncharacterized protein OS=Danio rerio GN=thbs4a PE=4 SV= A0A0R4IN91_DANRE,A0A	104.325,40	100,00%	2	2	2	0,01%	3,47%
Control 1	ATP synthase subunit gamma OS=Danio rerio GN=atp5c1 PE= Q5RH26_DANRE	32.149,90	85,10%	1	1	1	0,00%	3,42%

Control 1	Uncharacterized protein OS=Danio rerio GN=emilin1a PE=1 S F1QC17_DANRE,Q5TZ63_	112.771,70	100,00%	6	6	10	0,04%	8,28%
Control 1	Uncharacterized protein OS=Danio rerio GN=emilin1a PE=1 S F1QC17_DANRE,Q5TZ63_	112.771,70	100,00%	6	6	10	0,04%	8,28%
Control 1	Ubiquinol-cytochrome c reductase core protein II OS=Danio r Q6IQ59_DANRE	48.122,60	56,80%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=aimf1 PE=1 SV=: A2BG64_DANRE,A2BGE5_	66.977,00	99,60%	1	1	1	0,00%	1,47%
Control 1	Uncharacterized protein OS=Danio rerio GN=fina PE=1 SV=1 E9QI62_DANRE,F1R9P5_C	272.475,80	100,00%	4	4	6	0,02%	2,59%
Control 1	Uncharacterized protein OS=Danio rerio GN=fina PE=1 SV=1 E9QI62_DANRE,F1R9P5_C	272.475,80	100,00%	4	4	6	0,02%	2,59%
Control 1	Uncharacterized protein OS=Danio rerio GN=c3a.1 PE=1 SV=:1 B8JKW4_DANRE	183.235,60	100,00%	2	2	2	0,01%	1,83%
Control 1	Uncharacterized protein OS=Danio rerio GN=ndrg4 PE=4 SV= A0A0R4ICQ1_DANRE,A0A	37.210,30	27,30%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=ndrg4 PE=4 SV= A0A0R4ICQ1_DANRE,A0A	37.210,30	27,30%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=ndrg4 PE=4 SV= A0A0R4ICQ1_DANRE,A0A	37.210,30	27,30%	0	0	0	0,00%	0,00%
Control 1	L-lactate dehydrogenase B-A chain OS=Danio rerio GN=ldhba LDHA_DANRE,LDHBA_DA	36.246,70	98,70%	1	1	2	0,01%	3,59%
Control 1	L-lactate dehydrogenase B-A chain OS=Danio rerio GN=ldhba LDHA_DANRE,LDHBA_DA	36.246,70	98,70%	1	1	2	0,01%	3,59%
Control 1	Uncharacterized protein OS=Danio rerio GN=si:dkeyp-80c12. H0WES8_DANRE	6.338,70	99,70%	1	1	1	0,00%	27,60%
Control 1	Uncharacterized protein OS=Danio rerio GN=pdhb PE=1 SV=:1 Q1LV50_DANRE,Q7T368_	39.308,70	96,30%	1	1	2	0,01%	5,42%
Control 1	Uncharacterized protein OS=Danio rerio GN=pdhb PE=1 SV=:1 Q1LV50_DANRE,Q7T368_	39.308,70	96,30%	1	1	2	0,01%	5,42%
Control 1	Uncharacterized protein OS=Danio rerio GN=slc25a12 PE=4 S A0A0R4IHT0_DANRE,A0A	72.864,20	70,60%	1	1	1	0,00%	1,76%
Control 1	Uncharacterized protein OS=Danio rerio GN=slc25a12 PE=4 S A0A0R4IHT0_DANRE,A0A	72.864,20	70,60%	1	1	1	0,00%	1,76%
Control 1	Uncharacterized protein OS=Danio rerio PE=4 SV=1 A0A0G2L836_DANRE	136.805,30	100,00%	2	2	2	0,01%	2,17%
Control 1	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, A0A0R4ITN5_DANRE,A0A	72.077,60	100,00%	2	2	2	0,01%	4,24%
Control 1	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, A0A0R4ITN5_DANRE,A0A	72.077,60	100,00%	2	2	2	0,01%	4,24%
Control 1	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, A0A0R4ITN5_DANRE,A0A	72.077,60	100,00%	2	2	2	0,01%	4,24%
Control 1	Uncharacterized protein OS=Danio rerio GN=LOC557764 PE= F1R8B2_DANRE	169.128,30	98,30%	1	2	3	0,01%	0,76%
Control 1	Uncharacterized protein OS=Danio rerio GN=atp5b1 PE=1 SV= A2ARG7_DANRE	8.316,90	100,00%	2	3	6	0,02%	33,80%
Control 1	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	100,00%	2	2	4	0,01%	3,08%
Control 1	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	100,00%	2	2	4	0,01%	3,08%
Control 1	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	100,00%	2	2	4	0,01%	3,08%
Control 1	Uncharacterized protein OS=Danio rerio GN=aldh6a1 PE=1 S Q6DHT4_DANRE	57.120,40	99,90%	1	1	1	0,00%	2,86%
Control 1	Uncharacterized protein OS=Danio rerio GN=vtmb PE=1 SV=1 F1Q5D8_DANRE	51.527,70	93,40%	1	1	1	0,00%	2,67%
Control 1	Uncharacterized protein OS=Danio rerio GN=ndufs1 PE=1 SV: F1RDA6_DANRE	79.460,90	36,70%	0	0	0	0,00%	0,00%
Control 1	Cadherin-2 OS=Danio rerio GN=cdh2 PE=2 SV=2 CADH2_DANRE,G1K2E7_I	99.051,70	100,00%	2	2	3	0,01%	3,14%
Control 1	Cadherin-2 OS=Danio rerio GN=cdh2 PE=2 SV=2 CADH2_DANRE,G1K2E7_I	99.051,70	100,00%	2	2	3	0,01%	3,14%
Control 1	Uncharacterized protein OS=Danio rerio GN=popdc2 PE=1 SV F8W5H2_DANRE,Q6JWV8	41.034,60	100,00%	2	2	8	0,03%	11,40%
Control 1	Uncharacterized protein OS=Danio rerio GN=popdc2 PE=1 SV F8W5H2_DANRE,Q6JWV8	41.034,60	100,00%	2	2	8	0,03%	11,40%
Control 1	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV A0A0R4IH03_DANRE,A0A	108.322,90	100,00%	2	2	6	0,02%	3,82%
Control 1	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV A0A0R4IH03_DANRE,A0A	108.322,90	100,00%	2	2	6	0,02%	3,82%
Control 1	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV A0A0R4IH03_DANRE,A0A	108.322,90	100,00%	2	2	6	0,02%	3,82%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=pccz: F1QP17_DANRE	79.153,60	96,30%	1	1	1	0,00%	2,08%
Control 1	Uncharacterized protein (Fragment) OS=Danio rerio GN=abli: E9QI22_DANRE,U3JB13_C	84.999,80	17,80%	0	0	0	0,00%	0,00%
Control 1	Uncharacterized protein OS=Danio rerio GN=mfap5 PE=4 SV: E9QCQ4_DANRE	14.664,20	96,30%	1	2	12	0,04%	11,90%
Control 1	Uncharacterized protein OS=Danio rerio GN=megf6a PE=4 SV E7FFP9_DANRE,E9QHA9_	154.525,00	96,30%	1	1	1	0,00%	1,23%
Control 1	Uncharacterized protein OS=Danio rerio GN=megf6a PE=4 SV E7FFP9_DANRE,E9QHA9_	154.525,00	96,30%	1	1	1	0,00%	1,23%
Control 1	Heat shock 60kD protein 1 (Chaperonin) OS=Danio rerio GN= Q803B0_DANRE	61.196,80	98,80%	1	1	1	0,00%	4,87%
Control 2	Uncharacterized protein OS=Danio rerio GN=ccd141 PE=4 S B05755_DANRE,E7FG99_I	190.652,30	100,00%	4	4	4	0,02%	2,70%
Control 2	Uncharacterized protein OS=Danio rerio GN=ccd141 PE=4 S B05755_DANRE,E7FG99_I	190.652,30	100,00%	4	4	4	0,02%	2,70%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=col4 F1Q855_DANRE,F1QG79_	128.554,50	100,00%	2	4	9	0,04%	1,52%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=col4 F1Q855_DANRE,F1QG79_	128.554,50	100,00%	2	4	9	0,04%	1,52%
Control 2	Uncharacterized protein OS=Danio rerio GN=wfdc2 PE=4 SV: E7F3G3_DANRE	17.778,70	100,00%	3	3	4	0,02%	23,80%
Control 2	Uncharacterized protein OS=Danio rerio GN=add1 PE=4 SV=:1 A0A0R4IAJ9_DANRE,A0A	81.732,80	99,90%	1	1	1	0,00%	0,97%
Control 2	Uncharacterized protein OS=Danio rerio GN=add1 PE=4 SV=:1 A0A0R4IAJ9_DANRE,A0A	81.732,80	99,90%	1	1	1	0,00%	0,97%
Control 2	Uncharacterized protein OS=Danio rerio GN=add1 PE=4 SV=:1 A0A0R4IAJ9_DANRE,A0A	81.732,80	99,90%	1	1	1	0,00%	0,97%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=col6 F1Q924_DANRE	278.174,40	100,00%	6	6	7	0,03%	3,15%
Control 2	Uncharacterized protein OS=Danio rerio GN=palm2 PE=4 SV: A0A0R4ISM7_DANRE,E7F	92.198,80	100,00%	4	5	6	0,02%	6,18%
Control 2	Uncharacterized protein OS=Danio rerio GN=palm2 PE=4 SV: A0A0R4ISM7_DANRE,E7F	92.198,80	100,00%	4	5	6	0,02%	6,18%
Control 2	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1 A8W605_DANRE,ACTB1_	41.753,80	100,00%	3	4	4	0,02%	25,90%
Control 2	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1 A8W605_DANRE,ACTB1_	41.753,80	100,00%	3	4	4	0,02%	25,90%
Control 2	Uncharacterized protein OS=Danio rerio GN=acta1a PE=3 SV: F1QUN8_DANRE,Q6DHS1	41.974,00	100,00%	11	18	39	0,15%	37,70%
Control 2	Uncharacterized protein OS=Danio rerio GN=acta1a PE=3 SV: F1QUN8_DANRE,Q6DHS1	41.974,00	100,00%	11	18	39	0,15%	37,70%
Control 2	Uncharacterized protein OS=Danio rerio GN=acta1a PE=3 SV: F1QUN8_DANRE,Q6DHS1	41.974,00	100,00%	11	18	39	0,15%	37,70%
Control 2	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1 A8W605_DANRE,ACTB1_	41.753,80	100,00%	3	4	4	0,02%	25,90%
Control 2	Uncharacterized protein OS=Danio rerio GN=acta1a PE=3 SV: F1QUN8_DANRE,Q6DHS1	41.974,00	100,00%	11	18	39	0,15%	37,70%
Control 2	Uncharacterized protein OS=Danio rerio GN=acta1a PE=3 SV: F1QUN8_DANRE,Q6DHS1	41.974,00	100,00%	11	18	39	0,15%	37,70%
Control 2	Uncharacterized protein OS=Danio rerio GN=acta1a PE=3 SV: F1QUN8_DANRE,Q6DHS1	41.974,00	100,00%	11	18	39	0,15%	37,70%
Control 2	Uncharacterized protein OS=Danio rerio GN=desma PE=1 SV: F1R8W4_DANRE	54.149,90	84,20%	1	1	1	0,00%	30,20%
Control 2	Uncharacterized protein OS=Danio rerio GN=desma PE=1 SV: F1R8W3_DANRE	55.727,70	100,00%	13	14	21	0,08%	28,70%
Control 2	Uncharacterized protein OS=Danio rerio GN=col6a1 PE=1 SV: F1Q6P3_DANRE	107.466,50	100,00%	11	11	16	0,06%	12,90%
Control 2	Uncharacterized protein OS=Danio rerio GN=fn1b PE=1 SV=:2 A2CEW3_DANRE,F1R809_	265.661,60	100,00%	27	33	56	0,22%	14,60%
Control 2	Uncharacterized protein OS=Danio rerio GN=fn1b PE=1 SV=:2 A2CEW3_DANRE,F1R809_	265.661,60	100,00%	27	33	56	0,22%	14,60%
Control 2	Uncharacterized protein OS=Danio rerio GN=myhb PE=1 SV= F1QVX3_DANRE,X1WF87_	222.608,80	99,90%	1	1	2	0,01%	6,96%
Control 2	Uncharacterized protein OS=Danio rerio GN=vmhcl PE=1 SV= F1QSE1_DANRE	223.380,90	100,00%	69	96	233	0,92%	36,90%
Control 2	Uncharacterized protein OS=Danio rerio GN=vmhc PE=1 SV=: F1QZW1_DANRE	223.237,90	99,80%	1	1	1	0,00%	15,20%
Control 2	Uncharacterized protein OS=Danio rerio GN=LOC100329813 A0A0G2L365_DANRE	224.588,80	100,00%	1	2	5	0,02%	12,40%
Control 2	Uncharacterized protein OS=Danio rerio GN=myhb PE=1 SV= F1QVX3_DANRE,X1WF87_	222.608,80	99,90%	1	2	2	0,01%	6,96%
Control 2	Uncharacterized protein OS=Danio rerio GN=vcla PE=4 SV=1 A0A0R4IYQ3_DANRE,B3D	124.046,80	22,90%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=vcla PE=4 SV=1 A0A0R4IYQ3_DANRE,B3D	124.046,80	22,90%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=vcla PE=4 SV=1 A0A0R4IYQ3_DANRE,B3D	124.046,80	22,90%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=col5a1 PE=4 SV: A0A0R4IRG2_DANRE,F6N	198.988,80	100,00%	3	4	10	0,04%	1,87%
Control 2	Uncharacterized protein OS=Danio rerio GN=col5a1 PE=4 SV: A0A0R4IRG2_DANRE,F6N	198.988,80	100,00%	3	4	10	0,04%	1,87%
Control 2	Uncharacterized protein OS=Danio rerio GN=spna2 PE=4 SV= A0A0R4IU08_DANRE	289.251,20	100,00%	23	26	37	0,15%	11,60%
Control 2	Uncharacterized protein OS=Danio rerio GN=hspg2 PE=4 SV= A0A0R4IMY4_DANRE	390.507,40	100,00%	42	48	69	0,27%	17,10%
Control 2	Uncharacterized protein OS=Danio rerio GN=fga PE=1 SV=:1 B8A5L6_DANRE	75.292,70	100,00%	18	23	53	0,21%	33,30%
Control 2	Uncharacterized protein OS=Danio rerio GN=esyt1b PE=1 SV: F1QVP9_DANRE,F6NZP4_	175.083,00	11,70%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=esyt1b PE=1 SV: F1QVP9_DANRE,F6NZP4_	175.083,00	11,70%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=ith2 PE=4 SV=:1 Q5R929_DANRE	106.292,90	100,00%	5	6	11	0,04%	6,66%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttna B05758_DANRE	843.490,10	100,00%	0	0	0	0,00%	0,53%
Control 2	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4IU71_DANRE,E7F4	518.896,50	100,00%	1	1	1	0,00%	0,49%
Control 2	Uncharacterized protein OS=Danio rerio GN=eppk1 PE=4 SV= A0A0R4IND9_DANRE,I3IS	611.724,10	41,60%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=:1 A0A0R4INC4_DANRE,A5V	543.007,50	100,00%	4	4	4	0,02%	0,90%
Control 2	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=:1 A0A0R4INC4_DANRE,A5V	543.007,50	100,00%	4	4	4	0,02%	0,90%
Control 2	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=:1 A0A0R4INC4_DANRE,A5V	543.007,50	100,00%	4	4	4	0,02%	0,90%
Control 2	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4IU71_DANRE,E7F4	518.896,50	100,00%	1	1	1	0,00%	0,49%
Control 2	Uncharacterized protein OS=Danio rerio GN=eppk1 PE=4 SV= A0A0R4IND9_DANRE,I3IS	611.724,10	41,60%	0	0	0	0,00%	0,00%

Control 2	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=:A0A0R4U71_DANRE,E7F4	518.896,50	100,00%	1	1	1	0,00%	0,49%
Control 2	Uncharacterized protein OS=Danio rerio GN=ahnak PE=1 SV=:F1QZ50_DANRE	685.184,90	100,00%	37	38	64	0,25%	7,71%
Control 2	Uncharacterized protein OS=Danio rerio GN=postna PE=1 SV=:E7F513_DANRE	97.135,00	100,00%	10	11	18	0,07%	12,00%
Control 2	Solute carrier family 25 alpha, member 5 OS=Danio rerio GN:Q8JH10_DANRE	32.764,40	100,00%	3	4	5	0,02%	8,39%
Control 2	Uncharacterized protein OS=Danio rerio GN=col1a1a PE=1 SV:F1QJC9_DANRE,F1QSC5_I	128.043,70	100,00%	15	15	30	0,12%	13,40%
Control 2	Uncharacterized protein OS=Danio rerio GN=col1a1a PE=1 SV:F1QJC9_DANRE,F1QSC5_I	128.043,70	100,00%	15	15	30	0,12%	13,40%
Control 2	Periostin OS=Danio rerio GN=postnb PE=1 SV=1 Q7U566_DANRE	85.887,00	100,00%	5	5	5	0,02%	7,54%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttbn B056Y0_DANRE	626.764,10	100,00%	1	1	1	0,00%	0,39%
Control 2	Uncharacterized protein OS=Danio rerio GN=elnb PE=4 SV=1 A0A0R4IPZ1_DANRE,F8W	171.803,90	93,10%	1	1	1	0,00%	0,79%
Control 2	Uncharacterized protein OS=Danio rerio GN=elnb PE=4 SV=1 A0A0R4IPZ1_DANRE,F8W	171.803,90	93,10%	1	1	1	0,00%	0,79%
Control 2	Uncharacterized protein OS=Danio rerio GN=elnb PE=4 SV=1 A0A0R4IPZ1_DANRE,F8W	171.803,90	93,10%	1	1	1	0,00%	0,79%
Control 2	Keratin, type II cytoskeletal 8 OS=Danio rerio GN=kr18 PE=1 S K2C8_DANRE	57.761,30	100,00%	5	8	11	0,04%	8,27%
Control 2	Uncharacterized protein OS=Danio rerio GN=vtg4 PE=1 SV=2 F1RBA0_DANRE	148.836,20	100,00%	7	7	8	0,03%	6,11%
Control 2	Uncharacterized protein OS=Danio rerio GN=vtg4 PE=1 SV=2 E9QFD8_DANRE	149.301,80	68,00%	0	0	0	0,00%	4,04%
Control 2	ATP synthase subunit alpha OS=Danio rerio GN=atp5a1 PE=1 Q08BA1_DANRE	59.745,30	100,00%	6	6	12	0,05%	14,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=tpm4a PE=1 SV:F1QK77_DANRE	32.776,90	100,00%	9	13	22	0,09%	37,70%
Control 2	Uncharacterized protein OS=Danio rerio GN=sptb PE=1 SV=1 F1QEQ5_DANRE	272.829,20	100,00%	12	17	17	0,07%	6,66%
Control 2	Uncharacterized protein OS=Danio rerio GN=sptbn1 PE=1 SV:A0A0G2L1F5_DANRE	262.630,10	100,00%	11	11	14	0,06%	5,77%
Control 2	Uncharacterized protein OS=Danio rerio GN=actn2b PE=1 SV:E9QFR8_DANRE	103.214,90	100,00%	11	13	19	0,08%	16,20%
Control 2	Cytochrome c oxidase subunit 2 OS=Danio rerio GN=COX2 PE A0A0A0V0G8_DANRE,CO	25.985,80	98,40%	0	0	0	0,00%	0,00%
Control 2	Cytochrome c oxidase subunit 2 OS=Danio rerio GN=COX2 PE A0A0A0V0G8_DANRE,CO	25.985,80	98,40%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=col1a2 PE=1 SV:A0A0R4IHC0_DANRE,E9Q	127.211,90	100,00%	17	19	71	0,28%	16,60%
Control 2	Uncharacterized protein OS=Danio rerio GN=col1a2 PE=1 SV:A0A0R4IHC0_DANRE,E9Q	127.211,90	100,00%	17	19	71	0,28%	16,60%
Control 2	Uncharacterized protein OS=Danio rerio GN=col1a2 PE=1 SV:A0A0R4IHC0_DANRE,E9Q	127.211,90	100,00%	17	19	71	0,28%	16,60%
Control 2	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 S E7E7Z9_DANRE,F1Q615_I	142.109,70	100,00%	15	15	21	0,08%	14,10%
Control 2	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 S E7E7Z9_DANRE,F1Q615_I	142.109,70	100,00%	15	15	21	0,08%	14,10%
Control 2	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 S E7E7Z9_DANRE,F1Q615_I	142.109,70	100,00%	15	15	21	0,08%	14,10%
Control 2	Uncharacterized protein OS=Danio rerio GN=col1a1b PE=1 S F1QD11_DANRE	136.914,60	100,00%	15	19	45	0,18%	9,87%
Control 2	Uncharacterized protein OS=Danio rerio GN=zgc:65851 PE=1 Q7SZP4_DANRE	64.751,30	100,00%	2	3	4	0,02%	1,60%
Control 2	Uncharacterized protein OS=Danio rerio GN=col6a2 PE=4 SV:E7FCV8_DANRE	106.916,20	100,00%	11	12	15	0,06%	13,50%
Control 2	Uncharacterized protein OS=Danio rerio GN=kr194 PE=1 SV=:F1R8U0_DANRE	47.787,30	100,00%	3	3	3	0,01%	7,19%
Control 2	Calcium-transporting ATPase OS=Danio rerio GN=atp2a2a PE A9C3Q4_DANRE,A9C3Q5	109.519,40	100,00%	5	5	7	0,03%	6,05%
Control 2	Blood vessel epicardial substance OS=Danio rerio GN=bves P POPD1_DANRE	40.341,40	100,00%	2	2	5	0,02%	8,52%
Control 2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1:A0A0N4SUD2_DANRE,B0F	17.999,60	100,00%	1	1	1	0,00%	5,53%
Control 2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1:A0A0N4SUD2_DANRE,B0F	17.999,60	100,00%	1	1	1	0,00%	5,53%
Control 2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1:A0A0N4SUD2_DANRE,B0F	17.999,60	100,00%	1	1	1	0,00%	5,53%
Control 2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1:A0A0N4SUD2_DANRE,B0F	17.999,60	100,00%	1	1	1	0,00%	5,53%
Control 2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1:A0A0N4SUD2_DANRE,B0F	17.999,60	100,00%	1	1	1	0,00%	5,53%
Control 2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1:A0A0N4SUD2_DANRE,B0F	17.999,60	100,00%	1	1	1	0,00%	5,53%
Control 2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1:A0A0N4SUD2_DANRE,B0F	17.999,60	100,00%	1	1	1	0,00%	5,53%
Control 2	Elongation factor 1-alpha OS=Danio rerio GN=eef1a11 PE=1 A0A0R4IE10_DANRE,E7FC	50.150,70	94,80%	0	0	0	0,00%	0,00%
Control 2	Elongation factor 1-alpha OS=Danio rerio GN=eef1a11 PE=1 A0A0R4IE10_DANRE,E7FC	50.150,70	94,80%	0	0	0	0,00%	0,00%
Control 2	Elongation factor 1-alpha OS=Danio rerio GN=eef1a11 PE=1 A0A0R4IE10_DANRE,E7FC	50.150,70	94,80%	0	0	0	0,00%	0,00%
Control 2	Elongation factor 1-alpha OS=Danio rerio GN=eef1a11 PE=1 A0A0R4IE10_DANRE,E7FC	50.150,70	94,80%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV:A0A0N4SUA5_DANRE,E7F	81.108,80	99,90%	1	1	1	0,00%	1,54%
Control 2	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV:A0A0N4SUA5_DANRE,E7F	81.108,80	99,90%	1	1	1	0,00%	1,54%
Control 2	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV:A0A0N4SUA5_DANRE,E7F	81.108,80	99,90%	1	1	1	0,00%	1,54%
Control 2	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV:A0A0N4SUA5_DANRE,E7F	81.108,80	99,90%	1	1	1	0,00%	1,54%
Control 2	Cox5a protein OS=Danio rerio GN=cox5a PE=1 SV=1 Q4VBU7_DANRE	15.897,10	92,30%	0	0	0	0,00%	0,00%
Control 2	Heat shock protein 1 OS=Danio rerio GN=hspb1 PE=1 SV=1 A4VAK4_DANRE	22.408,90	99,90%	1	1	1	0,00%	5,53%
Control 2	Aspartate aminotransferase OS=Danio rerio GN=got2a PE=1 Q7SYK7_DANRE	47.588,10	99,90%	1	1	2	0,01%	1,87%
Control 2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV:A0A0N4STN6_DANRE,A7H	13.590,30	97,50%	0	0	0	0,00%	0,00%
Control 2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV:A0A0N4STN6_DANRE,A7H	13.590,30	97,50%	0	0	0	0,00%	0,00%
Control 2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV:A0A0N4STN6_DANRE,A7H	13.590,30	97,50%	0	0	0	0,00%	0,00%
Control 2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV:A0A0N4STN6_DANRE,A7H	13.590,30	97,50%	0	0	0	0,00%	0,00%
Control 2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV:A0A0N4STN6_DANRE,A7H	13.590,30	97,50%	0	0	0	0,00%	0,00%
Control 2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV:A0A0N4STN6_DANRE,A7H	13.590,30	97,50%	0	0	0	0,00%	0,00%
Control 2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV:A0A0N4STN6_DANRE,A7H	13.590,30	97,50%	0	0	0	0,00%	0,00%
Control 2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV:A0A0N4STN6_DANRE,A7H	13.590,30	97,50%	0	0	0	0,00%	0,00%
Control 2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV:A0A0N4STN6_DANRE,A7H	13.590,30	97,50%	0	0	0	0,00%	0,00%
Control 2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV:A0A0N4STN6_DANRE,A7H	13.590,30	97,50%	0	0	0	0,00%	0,00%
Control 2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV:A0A0N4STN6_DANRE,A7H	13.590,30	97,50%	0	0	0	0,00%	0,00%
Control 2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV:A0A0N4STN6_DANRE,A7H	13.590,30	97,50%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=col6a3 PE=1 SV:F1Q4K1_DANRE	309.287,80	100,00%	25	26	41	0,16%	10,30%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=col6 F1QKE8_DANRE	405.906,40	100,00%	2	2	3	0,01%	6,80%
Control 2	Uncharacterized protein OS=Danio rerio GN=ttna PE=1 SV=2 F1R7N8_DANRE	3.089.806,00	100,00%	22	23	24	0,09%	1,11%
Control 2	Uncharacterized protein OS=Danio rerio GN=emilin1b PE=1 S F1Q9G3_DANRE,F8W4C7	116.801,40	100,00%	15	15	22	0,09%	15,10%
Control 2	Uncharacterized protein OS=Danio rerio GN=emilin1b PE=1 S F1Q9G3_DANRE,F8W4C7	116.801,40	100,00%	15	15	22	0,09%	15,10%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=laml F1QEB6_DANRE,F1REL9_I	197.673,00	100,00%	1	1	1	0,00%	0,82%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=laml F1QEB6_DANRE,F1REL9_I	197.673,00	100,00%	1	1	1	0,00%	0,82%
Control 2	Cadherin-2 OS=Danio rerio GN=cdh2 PE=2 SV=2 CADH2_DANRE,G1K2E7_I	99.051,70	96,60%	1	1	2	0,01%	1,46%
Control 2	Cadherin-2 OS=Danio rerio GN=cdh2 PE=2 SV=2 CADH2_DANRE,G1K2E7_I	99.051,70	96,60%	1	1	2	0,01%	1,46%
Control 2	Caveolin OS=Danio rerio GN=cav1 PE=1 SV=1 Q6YLH8_DANRE,Q6YLH9_I	20.652,80	99,00%	1	1	1	0,00%	6,76%
Control 2	Caveolin OS=Danio rerio GN=cav1 PE=1 SV=1 Q6YLH8_DANRE,Q6YLH9_I	20.652,80	99,00%	1	1	1	0,00%	6,76%
Control 2	Tgm2b protein OS=Danio rerio GN=tgm2b PE=2 SV=1 Q1ECX3_DANRE	75.774,00	100,00%	1	1	1	0,00%	2,06%
Control 2	Keratin, type I cytoskeletal 18 OS=Danio rerio GN=kr18 PE=1 K1C18_DANRE	48.631,90	96,60%	1	1	1	0,00%	2,09%
Control 2	Protein S100 OS=Danio rerio GN=S100a10b PE=1 SV=1 Q7ZV4A_DANRE	11.157,10	100,00%	1	1	2	0,01%	13,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=uqcr1 PE=1 SV:F1QUC3_DANRE	55.450,70	100,00%	1	1	1	0,00%	1,79%
Control 2	Uncharacterized protein OS=Danio rerio GN=atp5f1 PE=1 SV:B8J51_DANRE	28.169,80	19,50%	0	0	0	0,00%	0,00%
Control 2	Enolase (Fragment) OS=Danio rerio GN=enol1a PE=1 SV=1 E9QCX7_DANRE,Q6PC12_I	47.075,10	96,60%	1	1	1	0,00%	3,63%
Control 2	Enolase (Fragment) OS=Danio rerio GN=enol1a PE=1 SV=1 E9QCX7_DANRE,Q6PC12_I	47.075,10	96,60%	1	1	1	0,00%	3,63%
Control 2	Uncharacterized protein OS=Danio rerio GN=wu:fi04e12 PE=:F1QAU9_DANRE	251.072,30	100,00%	2	2	2	0,01%	1,27%
Control 2	Aconitate hydratase, mitochondrial (Fragment) OS=Danio rer F8W4M7_DANRE	85.591,90	100,00%	5	5	6	0,02%	8,99%
Control 2	Uncharacterized protein OS=Danio rerio GN=col4a2 PE=1 SV:A0A0H2UKP2_DANRE	165.476,80	100,00%	4	5	9	0,04%	2,63%
Control 2	Uncharacterized protein OS=Danio rerio GN=LOC100330353 F1QWE6_DANRE	86.981,10	100,00%	2	2	3	0,01%	3,16%
Control 2	Uncharacterized protein OS=Danio rerio GN=cox6c PE=1 SV= Q6ZM23_DANRE	8.480,50	95,20%	1	1	2	0,01%	17,10%
Control 2	Catenin (Cadherin-associated protein), alpha OS=Danio rerio Q9PVB8_DANRE	100.520,10	100,00%	2	2	2	0,01%	3,31%
Control 2	Uncharacterized protein OS=Danio rerio GN=vdac2 PE=1 SV= Q8AWD0_DANRE	30.284,90	99,90%	2	2	3	0,01%	6,36%
Control 2	Myoglobin OS=Danio rerio GN=mb PE=2 SV=3 MYG_DANRE	15.583,60	100,00%	4	5	10	0,04%	46,90%

Control 2	Fgg protein OS=Danio rerio GN=fgg PE=1 SV=1	Q7ZVG7_DANRE	48.833,70	100,00%	13	17	50	0,20%	36,70%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttnb G1K2X0_DANRE		2.458.846,80	100,00%	9	9	9	0,04%	0,70%
Control 2	Alpha-1,4 glucan phosphorylase OS=Danio rerio GN=pygmb F E7EXT3_DANRE		96.868,30	100,00%	5	5	5	0,02%	5,58%
Control 2	Uncharacterized protein OS=Danio rerio GN=ttni1b PE=1 SV=1	E7F5J9_DANRE,F8W417_I	21.032,10	99,80%	1	1	1	0,00%	4,19%
Control 2	Uncharacterized protein OS=Danio rerio GN=ttni1b PE=1 SV=1	E7F5J9_DANRE,F8W417_I	21.032,10	99,80%	1	1	1	0,00%	4,19%
Control 2	Uncharacterized protein OS=Danio rerio GN=ttni1b PE=1 SV=1	E7F5J9_DANRE,F8W417_I	21.032,10	99,80%	1	1	1	0,00%	4,19%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=mid1 F1R998_DANRE,F1RAG3_		130.098,40	99,30%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=mid1 F1R998_DANRE,F1RAG3_		130.098,40	99,30%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=col6 F6NRG0_DANRE		88.827,90	100,00%	2	2	4	0,02%	11,50%
Control 2	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1	B3DG37_DANRE,HBB1_D	16.389,20	99,80%	1	1	1	0,00%	5,41%
Control 2	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1	B3DG37_DANRE,HBB1_D	16.389,20	99,80%	1	1	1	0,00%	5,41%
Control 2	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1	B3DG37_DANRE,HBB1_D	16.389,20	99,80%	1	1	1	0,00%	5,41%
Control 2	Uncharacterized protein OS=Danio rerio GN=utrn PE=1 SV=1	A0A0H2UKN4_DANRE	299.775,50	100,00%	1	1	1	0,00%	0,38%
Control 2	Uncharacterized protein OS=Danio rerio GN=plekha7a PE=4 S A0A0R4I185_DANRE,F1Q7		135.554,10	23,10%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=plekha7a PE=4 S A0A0R4I185_DANRE,F1Q7		135.554,10	23,10%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=plekha7a PE=4 S A0A0R4I185_DANRE,F1Q7		135.554,10	23,10%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=plekha7a PE=4 S A0A0R4I185_DANRE,F1Q7		135.554,10	23,10%	0	0	0	0,00%	0,00%
Control 2	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IG11_DANRE,G3P_		35.784,00	99,20%	1	1	1	0,00%	2,38%
Control 2	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IG11_DANRE,G3P_		35.784,00	99,20%	1	1	1	0,00%	2,38%
Control 2	Uncharacterized protein OS=Danio rerio GN=ptx3a PE=1 SV=1	A3KPN4_DANRE	49.273,20	100,00%	1	1	1	0,00%	2,46%
Control 2	Fibrinogen, B beta polypeptide OS=Danio rerio GN=fgb PE=1	Q6NEY1_DANRE	55.407,60	100,00%	10	11	28	0,11%	31,80%
Control 2	Fructose-bisphosphate aldolase OS=Danio rerio GN=aldoaa P Q803Q7_DANRE		39.740,60	100,00%	4	4	5	0,02%	16,50%
Control 2	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=1 SV=1	A0A0R4IF18_DANRE,A2B	103.798,70	100,00%	4	4	4	0,02%	5,07%
Control 2	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=1 SV=1	A0A0R4IF18_DANRE,A2B	103.798,70	100,00%	4	4	4	0,02%	5,07%
Control 2	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=1 SV=1	A0A0R4IF18_DANRE,A2B	103.798,70	100,00%	4	4	4	0,02%	5,07%
Control 2	Tropomyosin alpha-1 chain OS=Danio rerio GN=tpma PE=2 S F6NVA3_DANRE,F6P435_		32.724,00	100,00%	2	3	5	0,02%	18,30%
Control 2	Tropomyosin alpha-1 chain OS=Danio rerio GN=tpma PE=2 S F6NVA3_DANRE,F6P435_		32.724,00	100,00%	2	3	5	0,02%	18,30%
Control 2	Tropomyosin alpha-1 chain OS=Danio rerio GN=tpma PE=2 S F6NVA3_DANRE,F6P435_		32.724,00	100,00%	2	3	5	0,02%	18,30%
Control 2	Laminin subunit gamma-1 OS=Danio rerio GN=lamc1 PE=2 SV=LAMC1_DANRE		176.206,30	100,00%	7	7	10	0,04%	6,09%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:cf E9QDE7_DANRE		83.925,70	100,00%	2	2	2	0,01%	2,37%
Control 2	Iso citrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=1	A0A0G2L616_DANRE,E7F4	50.921,70	100,00%	1	1	1	0,00%	3,78%
Control 2	Iso citrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=1	A0A0G2L616_DANRE,E7F4	50.921,70	100,00%	1	1	1	0,00%	3,78%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:cf F1QT86_DANRE		147.365,60	100,00%	3	3	7	0,03%	2,76%
Control 2	Uncharacterized protein OS=Danio rerio GN=sorbs1 PE=4 SV=1	A0A0R4IP30_DANRE	85.938,80	100,00%	6	6	7	0,03%	9,79%
Control 2	Uncharacterized protein OS=Danio rerio GN=tln2a PE=4 SV=1	A0A0R4IIA7_DANRE,A0AC	271.774,00	100,00%	2	2	2	0,01%	1,10%
Control 2	Uncharacterized protein OS=Danio rerio GN=tln2a PE=4 SV=1	A0A0R4IIA7_DANRE,A0AC	271.774,00	100,00%	2	2	2	0,01%	1,10%
Control 2	Uncharacterized protein OS=Danio rerio GN=LOC557764 PE=1	F1RB82_DANRE	169.128,30	100,00%	3	3	3	0,01%	2,96%
Control 2	Radixin OS=Danio rerio GN=msna PE=1 SV=1	Q66I42_DANRE	68.035,00	100,00%	2	2	2	0,01%	2,76%
Control 2	Uncharacterized protein OS=Danio rerio GN=flna PE=1 SV=1	E9QI62_DANRE,F1R9P5_I	272.475,80	100,00%	5	5	5	0,02%	2,78%
Control 2	Uncharacterized protein OS=Danio rerio GN=flna PE=1 SV=1	E9QI62_DANRE,F1R9P5_I	272.475,80	100,00%	5	5	5	0,02%	2,78%
Control 2	Fatty acid binding protein 3, muscle and heart OS=Danio rerio Q8UVG7_DANRE		14.882,40	100,00%	1	1	1	0,00%	9,77%
Control 2	Sodium/potassium-transporting ATPase subunit alpha OS=Dz Q9DGL6_DANRE		113.384,80	100,00%	6	6	7	0,03%	8,37%
Control 2	Uncharacterized protein OS=Danio rerio GN=tln1 PE=4 SV=1	A0A0R4ID28_DANRE,A0AI	270.987,00	100,00%	6	6	6	0,02%	2,91%
Control 2	Uncharacterized protein OS=Danio rerio GN=tln1 PE=4 SV=1	A0A0R4ID28_DANRE,A0AI	270.987,00	100,00%	6	6	6	0,02%	2,91%
Control 2	Phosphoglycerate kinase OS=Danio rerio GN=pgk1 PE=1 SV=1	F1QXV8_DANRE	44.733,20	99,80%	1	1	2	0,01%	2,40%
Control 2	Inner membrane protein, mitochondrial (Mitofilin) OS=Danio Q6PF54_DANRE		83.201,30	99,90%	1	1	1	0,00%	2,11%
Control 2	Uncharacterized protein OS=Danio rerio GN=ckma PE=1 SV=1	A2BHA3_DANRE	42.814,90	100,00%	2	2	2	0,01%	7,35%
Control 2	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE=	A0A0R4IE46_DANRE,A0AI	101.943,10	96,50%	1	1	1	0,00%	1,84%
Control 2	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE=	A0A0R4IE46_DANRE,A0AI	101.943,10	96,50%	1	1	1	0,00%	1,84%
Control 2	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE=	A0A0R4IE46_DANRE,A0AI	101.943,10	96,50%	1	1	1	0,00%	1,84%
Control 2	Uncharacterized protein OS=Danio rerio GN=acs1b PE=1 SV=1	F1QDA6_DANRE	77.694,10	100,00%	1	1	1	0,00%	1,15%
Control 2	Uncharacterized protein OS=Danio rerio GN=hk1 PE=4 SV=1	A0A0R4IUT7_DANRE,Q6N	102.760,50	100,00%	4	4	5	0,02%	5,33%
Control 2	Uncharacterized protein OS=Danio rerio GN=hk1 PE=4 SV=1	A0A0R4IUT7_DANRE,Q6N	102.760,50	100,00%	4	4	5	0,02%	5,33%
Control 2	Uncharacterized protein OS=Danio rerio GN=zgc:101560 PE=	A0A0R4IBE8_DANRE,A0AI	21.007,60	100,00%	3	3	5	0,02%	17,50%
Control 2	Uncharacterized protein OS=Danio rerio GN=zgc:101560 PE=	A0A0R4IBE8_DANRE,A0AI	21.007,60	100,00%	3	3	5	0,02%	17,50%
Control 2	Myosin light chain 2 Mlc2a OS=Danio rerio GN=myl7 PE=1 SV=1	Q801M3_DANRE	19.021,00	100,00%	4	5	14	0,06%	30,80%
Control 2	Uncharacterized protein OS=Danio rerio GN=emilin1a PE=1 SV=1	F1QC17_DANRE,Q5T263_	112.771,70	100,00%	8	8	10	0,04%	9,57%
Control 2	Uncharacterized protein OS=Danio rerio GN=emilin1a PE=1 SV=1	F1QC17_DANRE,Q5T263_	112.771,70	100,00%	8	8	10	0,04%	9,57%
Control 2	Uncharacterized protein OS=Danio rerio GN=fn1a PE=1 SV=1	B0S602_DANRE,F1RBP8_I	271.523,90	96,80%	0	0	0	0,00%	0,69%
Control 2	Uncharacterized protein OS=Danio rerio GN=fn1a PE=1 SV=1	B0S602_DANRE,F1RBP8_I	271.523,90	96,80%	0	0	0	0,00%	0,69%
Control 2	Calcium-transporting ATPase OS=Danio rerio GN=atp2a2a PE=4 SV=1	A9C3Q4_DANRE,A9C3Q5_	109.519,40	100,00%	5	5	7	0,03%	6,05%
Control 2	Uncharacterized protein OS=Danio rerio GN=flnca PE=1 SV=1	E7F1L7_DANRE	291.642,80	100,00%	1	2	2	0,01%	1,40%
Control 2	Uncharacterized protein OS=Danio rerio GN=pdlim5b PE=4 S A0A0R4IS44_DANRE,A0AI		66.521,00	96,60%	1	1	1	0,00%	5,61%
Control 2	Uncharacterized protein OS=Danio rerio GN=pdlim5b PE=4 S A0A0R4IS44_DANRE,A0AI		66.521,00	96,60%	1	1	1	0,00%	5,61%
Control 2	Uncharacterized protein OS=Danio rerio GN=pdlim5b PE=4 S A0A0R4IS44_DANRE,A0AI		66.521,00	96,60%	1	1	1	0,00%	5,61%
Control 2	Uncharacterized protein OS=Danio rerio GN=pdlim5b PE=4 S A0A0R4IS44_DANRE,A0AI		66.521,00	96,60%	1	1	1	0,00%	5,61%
Control 2	ATP synthase subunit beta OS=Danio rerio GN=atp5b PE=1 SV=1	A8WGC6_DANRE	55.130,60	100,00%	6	8	19	0,08%	15,90%
Control 2	Uncharacterized protein OS=Danio rerio GN=fhl2b PE=4 SV=1	B0V234_DANRE,Q6AZB7_	32.504,70	100,00%	1	1	1	0,00%	6,09%
Control 2	Uncharacterized protein OS=Danio rerio GN=fhl2b PE=4 SV=1	B0V234_DANRE,Q6AZB7_	32.504,70	100,00%	1	1	1	0,00%	6,09%
Control 2	Uncharacterized protein OS=Danio rerio GN=fbn2b PE=1 SV=1	A0A0G2KQ62_DANRE	212.420,00	100,00%	16	17	28	0,11%	12,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=dnah12 PE=4 SV=1	E7FGS8_DANRE,E9QBH9_	438.810,60	98,70%	1	1	1	0,00%	0,20%
Control 2	Uncharacterized protein OS=Danio rerio GN=dnah12 PE=4 SV=1	E7FGS8_DANRE,E9QBH9_	438.810,60	98,70%	1	1	1	0,00%	0,20%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=eno F1QBW7_DANRE		51.320,60	94,10%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=cmlc1 PE=1 SV=1	B0RF07_DANRE	21.865,20	100,00%	6	9	21	0,08%	41,80%
Control 2	Uncharacterized protein OS=Danio rerio GN=col12a1a PE=1 S X1WF98_DANRE		321.902,50	100,00%	1	1	1	0,00%	0,37%
Control 2	Uncharacterized protein OS=Danio rerio GN=sdprb PE=1 SV=1	F1R665_DANRE	51.441,60	43,00%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=lama4 PE=1 SV=1	F1QYU0_DANRE	207.483,80	11,80%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=lmo7a PE=4 SV=1	A0A0R4ITT8_DANRE,B0U	207.438,50	100,00%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=lmo7a PE=4 SV=1	A0A0R4ITT8_DANRE,B0U	207.438,50	100,00%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=cox5b2 PE=1 SV=1	Q0P3U4_DANRE	14.308,50	99,90%	1	1	2	0,01%	11,10%
Control 2	Uncharacterized protein OS=Danio rerio GN=popdc2 PE=1 SV=1	F8W5H2_DANRE,Q6JWV8	41.034,60	96,60%	1	1	1	0,00%	4,09%
Control 2	Uncharacterized protein OS=Danio rerio GN=popdc2 PE=1 SV=1	F8W5H2_DANRE,Q6JWV8	41.034,60	96,60%	1	1	1	0,00%	4,09%
Control 2	Uncharacterized protein OS=Danio rerio GN=palm1b PE=4 SV=1	B0V0Y4_DANRE	27.123,40	100,00%	3	3	6	0,02%	14,80%
Control 2	Histone H4 OS=Danio rerio GN=si:dkey-108k21.23 PE=1 SV=1	A3KPR4_DANRE,X1WGM	11.478,70	99,80%	1	1	2	0,01%	9,71%
Control 2	Histone H4 OS=Danio rerio GN=si:dkey-108k21.23 PE=1 SV=1	A3KPR4_DANRE,X1WGM	11.478,70	99,80%	1	1	2	0,01%	9,71%
Control 2	Uncharacterized protein OS=Danio rerio GN=slc25a12 PE=4 S A0A0R4IHT0_DANRE,A0AI		72.864,20	100,00%	2	2	2	0,01%	2,93%
Control 2	Uncharacterized protein OS=Danio rerio GN=slc25a12 PE=4 S A0A0R4IHT0_DANRE,A0AI		72.864,20	100,00%	2	2	2	0,01%	2,93%
Control 2	Malate dehydrogenase OS=Danio rerio GN=mdh2 PE=1 SV=1	Q7T334_DANRE	35.420,30	100,00%	1	1	1	0,00%	3,86%
Control 2	Uncharacterized protein OS=Danio rerio GN=myom1b PE=4 S A0A0R4IGQ8_DANRE		180.977,20	100,00%	4	4	4	0,02%	2,83%
Control 2	Nicotinamide nucleotide transhydrogenase OS=Danio rerio G Q6NYQ7_DANRE		112.785,50	100,00%	1	1	1	0,00%	1,11%
Control 2	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1	A0A0R4IEB4_DANRE,A0AI	128.154,40	100,00%	2	2	3	0,01%	2,31%
Control 2	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1	A0A0R4IEB4_DANRE,A0AI	128.154,40	100,00%	2	2	3	0,01%	2,31%
Control 2	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1	A0A0R4IEB4_DANRE,A0AI	128.154,40	100,00%	2	2	3	0,01%	2,31%
Control 2	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1	F1Q775_DANRE	130.911,60	100,00%	2	2	3	0,01%	2,31%

Control 2	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1 A0A0R4IEB4_DANRE,A0AI	128.154,40	100,00%	2	2	3	0,01%	2,31%
Control 2	Dihydropolyl dehydrogenase OS=Danio rerio GN=dlhd PE=1: Q6TNUU_DANRE	53.620,50	96,90%	1	1	1	0,00%	1,97%
Control 2	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV A0A0R4IU91_DANRE,F6NH	94.381,10	100,00%	4	4	6	0,02%	6,20%
Control 2	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV A0A0R4IU91_DANRE,F6NH	94.381,10	100,00%	4	4	6	0,02%	6,20%
Control 2	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV A0A0R4IU91_DANRE,F6NH	94.381,10	100,00%	4	4	6	0,02%	6,20%
Control 2	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV A0A0R4IU91_DANRE,F6NH	94.381,10	100,00%	4	4	6	0,02%	6,20%
Control 2	Uncharacterized protein OS=Danio rerio GN=pkma PE=4 SV:- A0A0R4IGP6_DANRE,QZ7	58.059,00	99,90%	1	1	1	0,00%	2,07%
Control 2	Uncharacterized protein OS=Danio rerio GN=pkma PE=4 SV:- A0A0R4IGP6_DANRE,QZ7	58.059,00	99,90%	1	1	1	0,00%	2,07%
Control 2	Uncharacterized protein OS=Danio rerio GN=pdha1a PE=4 SV A0A0R4IT09_DANRE,Q6P	43.741,90	99,80%	1	1	1	0,00%	2,25%
Control 2	Uncharacterized protein OS=Danio rerio GN=pdha1a PE=4 SV A0A0R4IT09_DANRE,Q6P	43.741,90	99,80%	1	1	1	0,00%	2,25%
Control 2	Uncharacterized protein OS=Danio rerio GN=thbs4a PE=4 SV: A0A0R4I9N1_DANRE,A0A	104.325,40	100,00%	2	2	2	0,01%	3,58%
Control 2	Uncharacterized protein OS=Danio rerio GN=thbs4a PE=4 SV: A0A0R4I9N1_DANRE,A0A	104.325,40	100,00%	2	2	2	0,01%	3,58%
Control 2	Uncharacterized protein OS=Danio rerio GN=thbs4a PE=4 SV: A0A0R4I9N1_DANRE,A0A	104.325,40	100,00%	2	2	2	0,01%	3,58%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=post F1QM50_DANRE	17.277,20	100,00%	2	2	3	0,01%	19,60%
Control 2	Uncharacterized protein OS=Danio rerio GN=tjp1b PE=1 SV=1 A0A0R4IHR7_DANRE,E7F	205.686,40	90,40%	1	1	1	0,00%	0,59%
Control 2	Uncharacterized protein OS=Danio rerio GN=tjp1b PE=1 SV=1 A0A0R4IHR7_DANRE,E7F	205.686,40	90,40%	1	1	1	0,00%	0,59%
Control 2	Uncharacterized protein OS=Danio rerio GN=tjp1b PE=1 SV=1 A0A0R4IHR7_DANRE,E7F	205.686,40	90,40%	1	1	1	0,00%	0,59%
Control 2	Uncharacterized protein OS=Danio rerio GN=tjp1b PE=1 SV=1 A0A0R4IHR7_DANRE,E7F	205.686,40	90,40%	1	1	1	0,00%	0,59%
Control 2	Uncharacterized protein OS=Danio rerio GN=tjp1b PE=1 SV=1 A0A0R4IHR7_DANRE,E7F	205.686,40	90,40%	1	1	1	0,00%	0,59%
Control 2	Uncharacterized protein OS=Danio rerio GN=lamb2 PE=4 SV= B7ZDA6_DANRE	196.598,90	61,70%	0	0	0	0,00%	0,00%
Control 2	Citrate synthase OS=Danio rerio GN=csc PE=1 SV=1 B2GTW6_DANRE,CISY_DA	51.752,20	37,20%	0	0	0	0,00%	0,00%
Control 2	Citrate synthase OS=Danio rerio GN=csc PE=1 SV=1 B2GTW6_DANRE,CISY_DA	51.752,20	37,20%	0	0	0	0,00%	0,00%
Control 2	Succinyl-CoA ligase subunit beta OS=Danio rerio GN=sucla2 P Q6PBI0_DANRE	50.032,30	93,70%	1	1	1	0,00%	2,15%
Control 2	Uncharacterized protein OS=Danio rerio GN=anxa11b PE=4 S F8W2N0_DANRE,Q804G3	51.462,90	100,00%	1	1	1	0,00%	1,86%
Control 2	Uncharacterized protein OS=Danio rerio GN=anxa11b PE=4 S F8W2N0_DANRE,Q804G3	51.462,90	100,00%	1	1	1	0,00%	1,86%
Control 2	Uncharacterized protein OS=Danio rerio GN=tns1b PE=1 SV:- F1Q6A9_DANRE	185.031,00	100,00%	1	1	1	0,00%	0,59%
Control 2	Glutamate dehydrogenase 1 OS=Danio rerio GN=glud1b PE=: Q6P3L9_DANRE	59.978,90	96,80%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	20.097,20	65,50%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	20.097,20	65,50%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	20.097,20	65,50%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	20.097,20	65,50%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=vwf PE=4 SV=2 E7F921_DANRE	307.905,50	100,00%	1	1	1	0,00%	0,46%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=mm F1R6V5_DANRE	122.540,40	100,00%	3	3	3	0,01%	2,67%
Control 2	Hbaa1 protein OS=Danio rerio GN=hbaa1 PE=1 SV=1 HBA_DANRE,Q1JQ69_DAI	15.565,70	75,80%	1	1	1	0,00%	7,69%
Control 2	Hbaa1 protein OS=Danio rerio GN=hbaa1 PE=1 SV=1 HBA_DANRE,Q1JQ69_DAI	15.565,70	75,80%	1	1	1	0,00%	7,69%
Control 2	Hbaa1 protein OS=Danio rerio GN=hbaa1 PE=1 SV=1 HBA_DANRE,Q1JQ69_DAI	15.565,70	75,80%	1	1	1	0,00%	7,69%
Control 2	Uncharacterized protein OS=Danio rerio GN=hadhab PE=1 SV F1QYX8_DANRE	82.848,80	87,20%	0	0	0	0,00%	0,00%
Control 2	Fhla protein OS=Danio rerio GN=fhl1a PE=1 SV=1 Q6GQL7_DANRE	34.215,70	100,00%	1	1	2	0,01%	5,05%
Control 2	Uncharacterized protein OS=Danio rerio GN=c3a.1 PE=1 SV=1 B8JKW4_DANRE	183.235,60	100,00%	3	3	3	0,01%	2,50%
Control 2	Uncharacterized protein OS=Danio rerio GN=acadvl PE=1 SV= F1Q8I4_DANRE	71.085,10	99,90%	1	1	1	0,00%	1,67%
Control 2	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV A0A0R4IH03_DANRE,A0A	108.322,90	100,00%	3	3	5	0,02%	4,95%
Control 2	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV A0A0R4IH03_DANRE,A0A	108.322,90	100,00%	3	3	5	0,02%	4,95%
Control 2	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV A0A0R4IH03_DANRE,A0A	108.322,90	100,00%	3	3	5	0,02%	4,95%
Control 2	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV A0A0R4IH03_DANRE,A0A	108.322,90	100,00%	3	3	5	0,02%	4,95%
Control 2	L-lactate dehydrogenase B-A chain OS=Danio rerio GN=ldhba LDHBA_DANRE	36.246,70	100,00%	3	4	5	0,02%	11,40%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=aldh F8W3W6_DANRE,F8W4L7	57.120,40	100,00%	1	1	1	0,00%	5,16%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=aldh F8W3W6_DANRE,F8W4L7	57.120,40	100,00%	1	1	1	0,00%	5,16%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=aldh F8W3W6_DANRE,F8W4L7	57.120,40	100,00%	1	1	1	0,00%	5,16%
Control 2	Uncharacterized protein OS=Danio rerio GN=ank3b PE=1 SV= F1QN3B_DANRE	218.497,50	100,00%	4	4	4	0,02%	2,45%
Control 2	Uncharacterized protein OS=Danio rerio GN=mfap2 PE=4 SV: F1QSF1_DANRE	17.997,30	96,60%	1	2	5	0,02%	6,33%
Control 2	Heat shock cognate 71 kDa protein OS=Danio rerio GN=hsc70 Q6PGK4_DANRE	70.948,00	100,00%	2	2	3	0,01%	3,71%
Control 2	Uncharacterized protein OS=Danio rerio GN=apoc1 PE=1 SV= E9QD11_DANRE,E9QFX0_I	9.785,40	99,90%	1	1	1	0,00%	16,50%
Control 2	Uncharacterized protein OS=Danio rerio GN=apoc1 PE=1 SV= E9QD11_DANRE,E9QFX0_I	9.785,40	99,90%	1	1	1	0,00%	16,50%
Control 2	Uncharacterized protein OS=Danio rerio GN=c4 PE=4 SV=1 F1Q4X5_DANRE	191.050,80	99,70%	1	1	2	0,01%	0,65%
Control 2	Uncharacterized protein OS=Danio rerio GN=ndrg4 PE=4 SV= A0A0R4ICQ1_DANRE,A0A	37.210,30	100,00%	2	2	3	0,01%	10,90%
Control 2	Uncharacterized protein OS=Danio rerio GN=ndrg4 PE=4 SV= A0A0R4ICQ1_DANRE,A0A	37.210,30	100,00%	2	2	3	0,01%	10,90%
Control 2	Uncharacterized protein OS=Danio rerio GN=ndrg4 PE=4 SV= A0A0R4ICQ1_DANRE,A0A	37.210,30	100,00%	2	2	3	0,01%	10,90%
Control 2	Uncharacterized protein OS=Danio rerio GN=ndrg4 PE=4 SV= A0A0R4ICQ1_DANRE,A0A	37.210,30	100,00%	2	2	3	0,01%	10,90%
Control 2	Uncharacterized protein OS=Danio rerio PE=4 SV=1 A0A0G2L836_DANRE	136.805,30	26,10%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=dlst PE=1 SV=1 F1QF06_DANRE	48.683,10	100,00%	2	2	2	0,01%	5,68%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:d A0A0R4IER4_DANRE,A0AI	145.574,80	60,10%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:d A0A0R4IER4_DANRE,A0AI	145.574,80	60,10%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:d A0A0R4IER4_DANRE,A0AI	145.574,80	60,10%	0	0	0	0,00%	0,00%
Control 2	Annexin OS=Danio rerio GN=anxa2a PE=1 SV=1 Q6P603_DANRE	38.140,50	100,00%	2	2	2	0,01%	6,82%
Control 2	Uncharacterized protein OS=Danio rerio PE=1 SV=1 A0A0G2KHA1_DANRE	68.436,80	96,60%	1	1	1	0,00%	1,91%
Control 2	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, A0A0R4ITN5_DANRE,A0A	72.077,60	98,90%	0	0	0	0,00%	0,00%
Control 2	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, A0A0R4ITN5_DANRE,A0A	72.077,60	98,90%	0	0	0	0,00%	0,00%
Control 2	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, A0A0R4ITN5_DANRE,A0A	72.077,60	98,90%	0	0	0	0,00%	0,00%
Control 2	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, A0A0R4ITN5_DANRE,A0A	72.077,60	98,90%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=srl PE=4 SV=1 A0A0R4IRI3_DANRE,A0AC	55.028,70	81,20%	1	1	1	0,00%	2,77%
Control 2	Uncharacterized protein OS=Danio rerio GN=srl PE=4 SV=1 A0A0R4IRI3_DANRE,A0AC	55.028,70	81,20%	1	1	1	0,00%	2,77%
Control 2	Uncharacterized protein OS=Danio rerio GN=srl PE=4 SV=1 A0A0R4IRI3_DANRE,A0AC	55.028,70	81,20%	1	1	1	0,00%	2,77%
Control 2	Uncharacterized protein OS=Danio rerio GN=vdac3 PE=1 SV= B0R197_DANRE,B0R198_I	29.812,00	100,00%	2	2	3	0,01%	9,75%
Control 2	Uncharacterized protein OS=Danio rerio GN=vdac3 PE=1 SV= B0R197_DANRE,B0R198_I	29.812,00	100,00%	2	2	3	0,01%	9,75%
Control 2	Uncharacterized protein OS=Danio rerio GN=atp5b PE=1 SV= A2ARG7_DANRE	8.316,90	100,00%	3	3	3	0,01%	35,20%
Control 2	Transporter OS=Danio rerio GN=slc6a4a PE=2 SV=1 Q1WGB5_DANRE	72.122,80	96,60%	1	1	1	0,00%	2,17%
Control 2	Uncharacterized protein OS=Danio rerio GN=cox4i1 PE=1 SV: F1R6I1_DANRE	20.331,60	95,20%	1	1	1	0,00%	6,94%
Control 2	Uncharacterized protein OS=Danio rerio GN=vtbn PE=1 SV=1 F1Q5D8_DANRE	51.527,70	88,30%	1	1	1	0,00%	2,67%
Control 2	Uncharacterized protein OS=Danio rerio GN=uqrc2a PE=4 S A0A0R4IG82_DANRE	49.074,30	73,90%	1	1	1	0,00%	3,05%
Control 2	Uncharacterized protein OS=Danio rerio GN=ndufs1 PE=1 SV: F1RD46_DANRE	79.460,90	17,50%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6PI	29.220,20	99,90%	1	1	1	0,00%	7,95%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6PI	29.220,20	99,90%	1	1	1	0,00%	7,95%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6PI	29.220,20	99,90%	1	1	1	0,00%	7,95%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6PI	29.220,20	99,90%	1	1	1	0,00%	7,95%
Control 2	Uncharacterized protein OS=Danio rerio GN=ddx3b PE=1 SV: F1RCB2_DANRE,X1WHB2	24.761,00	18,50%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=ltbp X1WCC0_DANRE	33.442,60	100,00%	2	2	4	0,02%	9,27%
Control 2	Uncharacterized protein (Fragment) OS=Danio rerio GN=pccc F1QPL7_DANRE	79.153,60	100,00%	2	2	2	0,01%	4,02%
Control 2	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV= A0A0R4IP01_DANRE,B8JK	119.385,60	7,20%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV= A0A0R4IP01_DANRE,B8JK	119.385,60	7,20%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV= A0A0R4IP01_DANRE,B8JK	119.385,60	7,20%	0	0	0	0,00%	0,00%
Control 2	Uncharacterized protein OS=Danio rerio GN=mfap5 PE=4 SV: E9QC24_DANRE	14.664,20	96,60%	1	1	2	0,01%	11,90%
Control 2	Uncharacterized protein OS=Danio rerio GN=megf6a PE=4 SV E7FFP9_DANRE,E9QHA9_	154.525,00	100,00%	2	2	2	0,01%	2,53%
Control 2	Uncharacterized protein OS=Danio rerio GN=megf6a PE=4 SV E7FFP9_DANRE,E9QHA9_	154.525,00	100,00%	2	2	2	0,01%	2,53%

Control 2	Uncharacterized protein OS=Danio rerio GN=ddx3b PE=1 SV= F1RCB2_DANRE,X1WHB2_	24.761,00	18,50%	0	0	0	0,00%	0,00%	
Control 2	Heat shock 60kD protein 1 (Chaperonin) OS=Danio rerio GN= Q803B0_DANRE	61.196,80	96,60%	1	1	1	0,00%	4,35%	
Control 2	A0A0R4ITJ5-DECOY	A0A0R4ITJ5-DECOY,F1QKI	18	90,80%	1	1	0,00%	NaN%	
Control 2	A0A0R4ITJ5-DECOY	A0A0R4ITJ5-DECOY,F1QKI	18	90,80%	1	1	0,00%	NaN%	
7dpa_1	PDZ and LIM domain 3b OS=Danio rerio GN=pdlim3b PE=2 SV= Q0PWB8_DANRE	34.712,10	100,00%	2	2	2	0,01%	7,94%	
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ahnk PE=1 SV= F1QZ50_DANRE	685.184,90	100,00%	5	5	5	0,02%	1,25%	
7dpa_1	Fgg protein OS=Danio rerio GN=fgg PE=1 SV=1	Q7ZVZ7_DANRE	48.833,70	100,00%	31	48	405	1,49%	70,50%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ttna PE=1 SV=2	F1R7N8_DANRE	3.089.806,00	100,00%	9	9	9	0,03%	0,52%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttna B05758_DANRE	843.490,10	100,00%	2	2	2	0,01%	0,70%	
7dpa_1	Radixin OS=Danio rerio GN=msna PE=1 SV=1	Q66I42_DANRE	68.035,00	94,10%	1	1	2	0,01%	1,55%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=col6a1 PE=1 SV=1	F1Q6P3_DANRE	107.466,50	100,00%	14	16	31	0,11%	16,00%
7dpa_1	Fibrinogen, B beta polypeptide OS=Danio rerio GN=fgb PE=1	Q6NYE1_DANRE	54.407,60	100,00%	13	55	356	1,31%	68,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=fga PE=1 SV=1	B8A5L6_DANRE,E9QCD1_	49.680,00	100,00%	30	43	190	0,70%	41,80%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=fga PE=1 SV=1	B8A5L6_DANRE,E9QCD1_	49.680,00	100,00%	30	43	190	0,70%	41,80%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=1	A0A0R4I71_DANRE,E7F4'	518.896,50	100,00%	1	1	1	0,00%	0,22%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=1	A0A0R4INC4_DANRE,ASV	543.007,50	95,20%	1	1	1	0,00%	0,22%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=1	A0A0R4INC4_DANRE,ASV	543.007,50	95,20%	1	1	1	0,00%	0,22%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=1	A0A0R4INC4_DANRE,ASV	543.007,50	95,20%	1	1	1	0,00%	0,22%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=1	A0A0R4I71_DANRE,E7F4'	518.896,50	100,00%	1	1	1	0,00%	0,22%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=1	A0A0R4I71_DANRE,E7F4'	518.896,50	100,00%	1	1	1	0,00%	0,22%
7dpa_1	Annexin OS=Danio rerio GN=anxa11b PE=2 SV=1	F8W2N0_DANRE,Q804G3	51.462,90	100,00%	2	2	2	0,01%	6,60%
7dpa_1	Annexin OS=Danio rerio GN=anxa2a PE=1 SV=1	Q6P603_DANRE	38.140,50	100,00%	3	3	9	0,03%	11,30%
7dpa_1	Annexin OS=Danio rerio GN=anxa11b PE=2 SV=1	F8W2N0_DANRE,Q804G3	51.462,90	100,00%	2	2	2	0,01%	6,60%
7dpa_1	ATP synthase subunit alpha OS=Danio rerio GN=atp5a1 PE=1	Q08BA1_DANRE	59.745,30	100,00%	5	6	15	0,06%	13,60%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=col1a1a PE=1 SV=1	F1QJ9C_DANRE,F1QSC5_I	128.043,70	100,00%	13	16	73	0,27%	8,85%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=col1a1a PE=1 SV=1	F1QJ9C_DANRE,F1QSC5_I	128.043,70	100,00%	13	16	73	0,27%	8,85%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=fn1b PE=1 SV=2	A2CEW3_DANRE,F1R809_	265.661,60	100,00%	74	97	476	1,75%	46,90%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=fn1b PE=1 SV=2	A2CEW3_DANRE,F1R809_	265.661,60	100,00%	74	97	476	1,75%	46,90%
7dpa_1	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1	A8W605_DANRE,ACTB1_	41.753,80	100,00%	3	4	10	0,04%	35,70%
7dpa_1	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1	A8W605_DANRE,ACTB1_	41.753,80	100,00%	3	4	10	0,04%	35,70%
7dpa_1	Actin, alpha, cardiac muscle 1a OS=Danio rerio GN=si:ch73-1f1	F1QUN8_DANRE,Q6IQL9_	41.976,10	100,00%	16	20	48	0,18%	54,90%
7dpa_1	Actin, alpha, cardiac muscle 1a OS=Danio rerio GN=si:ch73-1f1	F1QUN8_DANRE,Q6IQL9_	41.976,10	100,00%	16	20	48	0,18%	54,90%
7dpa_1	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1	A8W605_DANRE,ACTB1_	41.753,80	100,00%	3	4	10	0,04%	35,70%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:86725 PE=4	A0A0R4IBT5_DANRE,F1RC	41.976,90	99,50%	1	1	1	0,00%	32,70%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:86725 PE=4	A0A0R4IBT5_DANRE,F1RC	41.976,90	99,50%	1	1	1	0,00%	32,70%
7dpa_1	Actin, alpha, cardiac muscle 1a OS=Danio rerio GN=si:ch73-1f1	F1QUN8_DANRE,Q6IQL9_	41.976,10	100,00%	16	20	48	0,18%	54,90%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=emilin1a PE=1 SV=1	F1QC17_DANRE,Q5T263_	112.771,70	99,90%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=emilin1a PE=1 SV=1	F1QC17_DANRE,Q5T263_	112.771,70	99,90%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=palm1b PE=4 SV=1	B0V0Y4_DANRE	27.123,40	100,00%	2	2	3	0,01%	9,84%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=col6a3 PE=1 SV=1	F1Q4K1_DANRE	309.287,80	100,00%	26	29	42	0,15%	13,30%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=c3a.1 PE=1 SV=1	B8JKW4_DANRE	183.235,60	100,00%	14	18	26	0,10%	11,90%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=spna2 PE=1 SV=1	F1R443_DANRE,F1R446_I	284.932,80	90,50%	1	1	1	0,00%	11,50%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=spna2 PE=4 SV=1	A0A0R4IU08_DANRE	289.251,20	100,00%	22	26	33	0,12%	11,70%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=spna2 PE=1 SV=1	F1R443_DANRE,F1R446_I	284.932,80	90,50%	1	1	1	0,00%	11,50%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=c4 PE=4 SV=1	F1Q4X5_DANRE	191.050,80	99,40%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=tnnb G1K2X0_DANRE	2.458.846,80	100,00%	2	2	2	0,01%	0,16%	
7dpa_1	Uncharacterized protein OS=Danio rerio GN=hspg2 PE=1 SV=1	A0A0R4I2A6_DANRE,F1R5	373.247,70	100,00%	21	25	37	0,14%	9,26%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=hspg2 PE=1 SV=1	A0A0R4I2A6_DANRE,F1R5	373.247,70	100,00%	21	25	37	0,14%	9,26%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=hspg2 PE=1 SV=1	A0A0R4I2A6_DANRE,F1R5	373.247,70	100,00%	21	25	37	0,14%	9,26%
7dpa_1	Histone H3 OS=Danio rerio GN=h3f3b.1 PE=1 SV=1	A8KBJ5_DANRE,B2GSF5_I	17.030,40	99,90%	1	1	1	0,00%	6,62%
7dpa_1	Histone H3 OS=Danio rerio GN=h3f3b.1 PE=1 SV=1	A8KBJ5_DANRE,B2GSF5_I	17.030,40	99,90%	1	1	1	0,00%	6,62%
7dpa_1	Histone H3 OS=Danio rerio GN=h3f3b.1 PE=1 SV=1	A8KBJ5_DANRE,B2GSF5_I	17.030,40	99,90%	1	1	1	0,00%	6,62%
7dpa_1	Histone H3 OS=Danio rerio GN=h3f3b.1 PE=1 SV=1	A8KBJ5_DANRE,B2GSF5_I	17.030,40	99,90%	1	1	1	0,00%	6,62%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=palm2 PE=4 SV=1	A0A0R4ISM7_DANRE,E7F	92.198,80	100,00%	2	3	3	0,01%	4,24%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=palm2 PE=4 SV=1	A0A0R4ISM7_DANRE,E7F	92.198,80	100,00%	2	3	3	0,01%	4,24%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=elnb PE=4 SV=1	A0A0R4IP21_DANRE,F8W	171.803,90	100,00%	1	1	1	0,00%	0,79%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=elnb PE=4 SV=1	A0A0R4IP21_DANRE,F8W	171.803,90	100,00%	1	1	1	0,00%	0,79%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=elnb PE=4 SV=1	A0A0R4IP21_DANRE,F8W	171.803,90	100,00%	1	1	1	0,00%	0,79%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=col6 F1Q924_DANRE	278.174,40	100,00%	10	10	15	0,06%	5,58%	
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ank3b PE=1 SV=1	F1QN83_DANRE	218.497,50	100,00%	4	4	4	0,01%	2,30%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=vmhcl PE=1 SV=1	F1QSE1_DANRE	223.380,90	100,00%	82	98	309	1,13%	46,10%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=sptb PE=1 SV=1	F1QOE5_DANRE	272.829,20	100,00%	9	9	9	0,03%	6,66%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=sptbn1 PE=1 SV=1	A0A0G2L1F5_DANRE	262.630,10	100,00%	10	10	12	0,04%	6,07%
7dpa_1	Periostin OS=Danio rerio GN=postb PE=1 SV=1	Q7SU66_DANRE	85.887,00	100,00%	14	17	39	0,14%	30,40%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=tpm4a PE=1 SV=1	F1QK67_DANRE	32.726,90	100,00%	8	9	13	0,05%	35,60%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=itih2 PE=4 SV=1	Q5RH29_DANRE	106.292,90	100,00%	9	9	18	0,07%	13,80%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=LOC100329813	A0A0G2L365_DANRE	224.588,80	100,00%	3	3	5	0,02%	17,30%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=aifm1 PE=1 SV=1	A2BGE5_DANRE,F1RBU0_	66.977,00	58,80%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=aifm1 PE=1 SV=1	A2BGE5_DANRE,F1RBU0_	66.977,00	58,80%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=aifm1 PE=1 SV=1	A2BGE5_DANRE,F1RBU0_	66.977,00	58,80%	0	0	0	0,00%	0,00%
7dpa_1	Alpha-1,4 glucan phosphorylase OS=Danio rerio GN=pygb PE=1 SV=1	E7EXT3_DANRE,E7F2Z5_C	96.935,30	45,80%	0	0	0	0,00%	0,00%
7dpa_1	Alpha-1,4 glucan phosphorylase OS=Danio rerio GN=pygb PE=1 SV=1	E7EXT3_DANRE,E7F2Z5_C	96.935,30	45,80%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=col4a2 PE=1 SV=1	A0A0H2UKP2_DANRE	165.476,80	100,00%	2	2	3	0,01%	1,56%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=col1a2 PE=1 SV=1	A0A0R4IHCO_DANRE,E9Q	127.211,90	100,00%	15	16	55	0,20%	14,10%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=col1a2 PE=1 SV=1	A0A0R4IHCO_DANRE,E9Q	127.211,90	100,00%	15	16	55	0,20%	14,10%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=col1a2 PE=1 SV=1	A0A0R4IHCO_DANRE,E9Q	127.211,90	100,00%	15	16	55	0,20%	14,10%
7dpa_1	Elongation factor 1-alpha OS=Danio rerio GN=eef1a PE=2 SV=1	EF1A_DANRE	50.048,20	100,00%	1	1	1	0,00%	5,19%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=slc2: E9QJ49_DANRE,F1QAJ7_I	39.378,90	99,90%	1	1	1	0,00%	4,30%	
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=slc2: E9QJ49_DANRE,F1QAJ7_I	39.378,90	99,90%	1	1	1	0,00%	4,30%	
7dpa_1	Uncharacterized protein OS=Danio rerio GN=vtg1 PE=1 SV=1	Q1LWN2_DANRE	149.238,80	100,00%	5	6	8	0,03%	4,63%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=laml F1QE86_DANRE,F1RE9L_I	197.673,00	96,20%	1	1	1	0,00%	0,53%	
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=laml F1QE86_DANRE,F1RE9L_I	197.673,00	96,20%	1	1	1	0,00%	0,53%	
7dpa_1	Uncharacterized protein OS=Danio rerio GN=desma PE=1 SV=1	F1R8W3_DANRE,F1R8W4	54.149,90	100,00%	13	13	23	0,08%	32,60%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=desma PE=1 SV=1	F1R8W3_DANRE,F1R8W4	54.149,90	100,00%	13	13	23	0,08%	32,60%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=hmcn2 PE=1 SV=1	E7F3V2_DANRE	454.987,50	100,00%	2	2	2	0,01%	0,57%
7dpa_1	ATP synthase subunit beta OS=Danio rerio GN=atp5b PE=1 SV=1	A8W6G6_DANRE	55.130,90	100,00%	10	12	18	0,07%	35,20%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=fn1a PE=1 SV=1	B0S602_DANRE	271.953,50	100,00%	9	9	12	0,04%	7,86%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=emilin1b PE=1 SV=1	F1Q9G3_DANRE	117.470,20	100,00%	7	8	8	0,03%	8,12%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=col1a1b PE=1 SV=1	F1QD11_DANRE	136.914,60	100,00%	11	13	31	0,11%	8,35%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=lmo7a PE=4 SV=1	A0A0R4ITT8_DANRE,B0U1	207.438,50	100,00%	2	2	2	0,01%	2,26%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=cttna1 PE=4 SV=1	A0A0R4IF47_DANRE,Q9P1	100.520,10	100,00%	2	2	2	0,01%	2,77%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=cttna1 PE=4 SV=1	A0A0R4IF47_DANRE,Q9P1	100.520,10	100,00%	2	2	2	0,01%	2,77%
7dpa_1	Hbaa1 protein OS=Danio rerio GN=hbaa1 PE=1 SV=1	HBA_DANRE,Q1JQ69_DAI	15.565,70	100,00%	1	1	2	0,01%	12,60%
7dpa_1	Hbaa1 protein OS=Danio rerio GN=hbaa1 PE=1 SV=1	HBA_DANRE,Q1JQ69_DAI	15.565,70	100,00%	1	1	2	0	

7dpa_1	Hbaa1 protein OS=Danio rerio GN=hbaa1 PE=1 SV=1	HBA_DANRE,Q1JQ69_DAI	15.565,70	100,00%	1	1	2	0,01%	12,60%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=actn2b PE=1 SV=1	E9QFR8_DANRE	103.214,90	100,00%	12	12	16	0,06%	17,40%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=srl PE=4 SV=1	A0A0R4IRI3_DANRE,A0AC	55.028,70	10,40%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=srl PE=4 SV=1	A0A0R4IRI3_DANRE,A0AC	55.028,70	10,40%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=srl PE=4 SV=1	A0A0R4IRI3_DANRE,A0AC	55.028,70	10,40%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 SV=1	E7EZ79_DANRE,F1Q615_I	142.109,70	100,00%	10	10	13	0,05%	12,50%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 SV=1	E7EZ79_DANRE,F1Q615_I	142.109,70	100,00%	10	10	13	0,05%	12,50%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 SV=1	E7EZ79_DANRE,F1Q615_I	142.109,70	100,00%	10	10	13	0,05%	12,50%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ank3a PE=4 SV=1	F1QNBI_DANRE	455.578,80	99,50%	1	1	1	0,00%	0,80%
7dpa_1	Malate dehydrogenase OS=Danio rerio GN=mdh2 PE=1 SV=1	Q7T334_DANRE	35.420,30	100,00%	6	7	11	0,04%	29,10%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=eno F1QBW7_DANRE		51.320,60	100,00%	4	4	4	0,01%	15,00%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=col4 F1Q855_DANRE,F1QG79_		128.554,50	100,00%	3	3	12	0,04%	3,47%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=col4 F1Q855_DANRE,F1QG79_		128.554,50	100,00%	3	3	12	0,04%	3,47%
7dpa_1	Keratin, type II cytoskeletal 8 OS=Danio rerio GN=krts PE=1 SV=1	S K2C8_DANRE	57.761,30	100,00%	2	2	2	0,01%	6,92%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=krts PE=1 SV=1	F1QV31_DANRE,F1R5A5_	58.787,80	97,20%	1	1	1	0,00%	3,71%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=krts PE=4 SV=1	F1QK60_DANRE	53.998,30	100,00%	2	2	2	0,01%	6,64%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=krts PE=1 SV=1	F1QV31_DANRE,F1R5A5_	58.787,80	97,20%	1	1	1	0,00%	3,71%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=mm F1R6V5_DANRE		122.540,40	99,90%	1	1	1	0,00%	1,56%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=vdac2 PE=1 SV=1	Q8AWD0_DANRE	30.284,90	100,00%	6	6	7	0,03%	26,90%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=postna PE=1 SV=1	E7F5I3_DANRE	97.135,00	100,00%	12	12	19	0,07%	22,30%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=col12a1a PE=1 SV=1	X1WF98_DANRE	321.902,50	100,00%	6	6	9	0,03%	2,92%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=post F1QM50_DANRE		17.277,20	100,00%	1	1	1	0,00%	12,40%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=fnb2b PE=1 SV=1	A0A0G2K062_DANRE	212.420,00	100,00%	6	6	6	0,02%	3,99%
7dpa_1	Solute carrier family 25 alpha, member 5 OS=Danio rerio GN=Q8JHI0_DANRE		32.764,40	100,00%	4	5	7	0,03%	12,40%
7dpa_1	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1	B3DG37_DANRE,HBB1_D_	16.389,00	97,20%	1	1	2	0,01%	48,00%
7dpa_1	Hemoglobin subunit beta-2 OS=Danio rerio GN=ba2 PE=1 SV=1	HBB2_DANRE	16.389,20	100,00%	5	6	10	0,04%	48,00%
7dpa_1	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1	B3DG37_DANRE,HBB1_D_	16.389,00	97,20%	1	1	2	0,01%	48,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=fnca PE=1 SV=1	A0A0R4IU54_DANRE,E7FJ	291.642,80	100,00%	2	2	2	0,01%	1,43%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=fnca PE=1 SV=1	A0A0R4IU54_DANRE,E7FJ	291.642,80	100,00%	2	2	2	0,01%	1,43%
7dpa_1	Alpha-1,4 glucan phosphorylase OS=Danio rerio GN=pygb PE=1 SV=1	E7EXT3_DANRE,E7F2Z5_C	96.935,30	45,80%	0	0	0	0,00%	0,00%
7dpa_1	Alpha-1,4 glucan phosphorylase OS=Danio rerio GN=pygb PE=1 SV=1	E7EXT3_DANRE,E7F2Z5_C	96.935,30	45,80%	0	0	0	0,00%	0,00%
7dpa_1	Myoglobin OS=Danio rerio GN=mb PE=1 SV=1	H0WE38_DANRE,MYG_D_	15.583,60	100,00%	4	4	9	0,03%	65,10%
7dpa_1	Myoglobin OS=Danio rerio GN=mb PE=1 SV=1	H0WE38_DANRE,MYG_D_	15.583,60	100,00%	4	4	9	0,03%	65,10%
7dpa_1	Fructose-bisphosphate aldolase OS=Danio rerio GN=aldoaa P Q803Q7_DANRE		39.740,60	100,00%	3	3	3	0,01%	11,80%
7dpa_1	Isocitrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=1	A0A0G2L616_DANRE,E7F4	50.921,70	100,00%	2	2	2	0,01%	6,67%
7dpa_1	Isocitrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=1	A0A0G2L616_DANRE,E7F4	50.921,70	100,00%	2	2	2	0,01%	6,67%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=col6a2 PE=4 SV=1	E7FCV8_DANRE	106.916,20	100,00%	12	15	30	0,11%	16,10%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:cf F1QT86_DANRE		147.365,60	100,00%	4	4	5	0,02%	3,09%
7dpa_1	Cytochrome c oxidase subunit 2 OS=Danio rerio GN=COX2 PE A0A0A0VG08_DANRE,CO_		25.985,80	97,20%	0	0	0	0,00%	0,00%
7dpa_1	Cytochrome c oxidase subunit 2 OS=Danio rerio GN=COX2 PE A0A0A0VG08_DANRE,CO_		25.985,80	97,20%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=4 SV=1	A0A0R4IF18_DANRE,A2B_	110.503,30	100,00%	3	3	3	0,01%	4,45%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=4 SV=1	A0A0R4IF18_DANRE,A2B_	110.503,30	100,00%	3	3	3	0,01%	4,45%
7dpa_1	Malate dehydrogenase 1a, NAD (Soluble) OS=Danio rerio GN Q7T3D9_DANRE,Q801E6_		36.258,40	100,00%	2	2	2	0,01%	8,20%
7dpa_1	Malate dehydrogenase 1a, NAD (Soluble) OS=Danio rerio GN Q7T3D9_DANRE,Q801E6_		36.258,40	100,00%	2	2	2	0,01%	8,20%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ctnnb2 PE=1 SV=1	E7FC99_DANRE,F1QY76_I	81.108,80	100,00%	2	2	2	0,01%	6,33%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ctnnb2 PE=1 SV=1	E7FC99_DANRE,F1QY76_I	81.108,80	100,00%	2	2	2	0,01%	6,33%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=uqrcr2a PE=4 SV=1	A0A0R4IAK9_DANRE,A0A_	49.189,50	100,00%	2	2	2	0,01%	7,19%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=uqrcr2a PE=4 SV=1	A0A0R4IAK9_DANRE,A0A_	49.189,50	100,00%	2	2	2	0,01%	7,19%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=uqrcr2a PE=4 SV=1	A0A0R4IAK9_DANRE,A0A_	49.189,50	100,00%	2	2	2	0,01%	7,19%
7dpa_1	Phosphoglycerate kinase OS=Danio rerio GN=pgk1 PE=1 SV=1	F1QXV8_DANRE	44.733,20	100,00%	2	2	2	0,01%	7,91%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=hk1 PE=4 SV=1	A0A0R4IUT7_DANRE,Q6N_	102.760,50	100,00%	2	2	2	0,01%	2,72%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=hk1 PE=4 SV=1	A0A0R4IUT7_DANRE,Q6N_	102.760,50	100,00%	2	2	2	0,01%	2,72%
7dpa_1	E9QD93-DECOY	E9QD93-DECOY,Q5RH26_	18	27,30%	0	0	0	0,00%	NaN%
7dpa_1	Anion exchange protein OS=Danio rerio GN=slc4a1a PE=1 SV=1	Q7ZZI7_DANRE	101.463,30	100,00%	2	2	4	0,01%	3,65%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=cox4i1 PE=1 SV=1	F1R6I1_DANRE	20.331,60	99,90%	1	1	1	0,00%	6,36%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE=	A0A0R4IE46_DANRE,A0A_	101.943,10	100,00%	3	4	5	0,02%	3,03%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE=	A0A0R4IE46_DANRE,A0A_	101.943,10	100,00%	3	4	5	0,02%	3,03%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE=	A0A0R4IE46_DANRE,A0A_	101.943,10	100,00%	3	4	5	0,02%	3,03%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=flna PE=1 SV=1	E9QI62_DANRE,F1R9P5_C	272.475,80	100,00%	3	3	3	0,01%	1,84%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=flna PE=1 SV=1	E9QI62_DANRE,F1R9P5_C	272.475,80	100,00%	3	3	3	0,01%	1,84%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=paplnb PE=4 SV=1	F1QR48_DANRE	60.121,30	100,00%	2	2	2	0,01%	3,83%
7dpa_1	Aspartate aminotransferase OS=Danio rerio GN=got2a PE=1 SV=1	Q7YK71_DANRE	47.588,10	99,90%	1	1	1	0,00%	3,04%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=slc25a12 PE=4 SV=1	A0A0R4IHT0_DANRE,A0A_	72.864,20	100,00%	1	1	1	0,00%	1,91%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=slc25a12 PE=4 SV=1	A0A0R4IHT0_DANRE,A0A_	72.864,20	100,00%	1	1	1	0,00%	1,91%
7dpa_1	Tropomyosin alpha-1 chain OS=Danio rerio GN=tpma PE=2 SV=1	E7FBZ3_DANRE,TPM1_D_	32.724,00	100,00%	1	1	1	0,00%	13,00%
7dpa_1	Tropomyosin alpha-1 chain OS=Danio rerio GN=tpma PE=2 SV=1	E7FBZ3_DANRE,TPM1_D_	32.724,00	100,00%	1	1	1	0,00%	13,00%
7dpa_1	Aconitate hydratase, mitochondrial (Fragment) OS=Danio rer F8W4M7_DANRE		85.591,90	100,00%	4	4	4	0,01%	7,22%
7dpa_1	Sodium/potassium-transporting ATPase subunit alpha OS=D_	Q9DGL6_DANRE	113.384,80	100,00%	4	4	4	0,01%	5,16%
7dpa_1	Uncharacterized protein OS=Danio rer GN=f13a1b PE=1 SV=1	F1R1R7_DANRE	83.603,80	100,00%	2	2	2	0,01%	2,43%
7dpa_1	Keratin, type I cytoskeletal 18 OS=Danio rerio GN=krts18 PE=1 SV=1	K1C18_DANRE	48.631,90	100,00%	2	2	2	0,01%	4,87%
7dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=1	A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	2	3	0,01%	21,00%
7dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=1	A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	2	3	0,01%	21,00%
7dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=1	A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	2	3	0,01%	21,00%
7dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=1	A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	2	3	0,01%	21,00%
7dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=1	A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	2	3	0,01%	21,00%
7dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=1	A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	2	3	0,01%	21,00%
7dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=1	A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	2	3	0,01%	21,00%
7dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=1	A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	2	3	0,01%	21,00%
7dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=1	A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	2	3	0,01%	21,00%
7dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=1	A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	2	3	0,01%	21,00%
7dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=1	A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	2	2	3	0,01%	21,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=cmlc1 PE=1 SV=1	B0RF07_DANRE	21.865,20	100,00%	7	9	12	0,04%	41,30%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV=1	A0A0R4IP01_DANRE,B8JK	119.385,60	48,90%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV=1	A0A0R4IP01_DANRE,B8JK	119.385,60	48,90%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV=1	A0A0R4IP01_DANRE,B8JK	119.385,60	48,90%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV=1	A0A0R4IBJ3_DANRE,F1R6	72.513,60	100,00%	3	3	3	0,01%	5,21%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV=1	A0A0R4IBJ3_DANRE,F1R6	72.513,60	100,00%	3	3	3	0,01%	5,21%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV=1	A0A0R4IBJ3_DANRE,F1R6	72.513,60	100,00%	3	3	3	0,01%	5,21%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV=1	A0A0R4IBJ3_DANRE,F1R6	72.513,60	100,00%	3	3	3	0,01%	5,21%

7dpa_1	Uncharacterized protein OS=Danio rerio GN=cox5b2 PE=1 SV= QQP3U4_DANRE	14.308,50	62,20%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=nid1a PE=1 SV=:; F1RAG3_DANRE	130.098,40	100,00%	5	6	7	0,03%	5,96%
7dpa_1	Histone H4 OS=Danio rerio GN=si:dkcy-108K21.23 PE=1 SV=1 A3KPR4_DANRE,X1WGM:	11.478,70	100,00%	2	2	2	0,01%	17,50%
7dpa_1	Histone H4 OS=Danio rerio GN=si:dkcy-108K21.23 PE=1 SV=1 A3KPR4_DANRE,X1WGM:	11.478,70	100,00%	2	2	2	0,01%	17,50%
7dpa_1	Superoxide dismutase OS=Danio rerio GN=sod2 PE=1 SV=1 Q6P980_DANRE	25.008,40	99,40%	1	1	1	0,00%	6,70%
7dpa_1	Heat shock protein 1 OS=Danio rerio GN=hspb1 PE=1 SV=1 A4VAK4_DANRE	22.408,90	100,00%	3	3	4	0,01%	18,60%
7dpa_1	Glutamate dehydrogenase 1 OS=Danio rerio GN=glud1b PE=: Q6P319_DANRE	59.978,90	100,00%	1	1	1	0,00%	1,48%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=megf6a PE=4 SV= E7FFP9_DANRE,E9QHA9_	154.525,00	100,00%	2	2	2	0,01%	1,92%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=megf6a PE=4 SV= E7FFP9_DANRE,E9QHA9_	154.525,00	100,00%	2	2	2	0,01%	1,92%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=vda: B0R198_DANRE	29.812,00	99,90%	1	1	2	0,01%	6,81%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=tnnt2a PE=1 SV: E7EY9C_DANRE,F1R0K6_I	33.942,20	100,00%	2	2	7	0,03%	11,40%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=tnnt2a PE=1 SV: E7EY9C_DANRE,F1R0K6_I	33.942,20	100,00%	2	2	7	0,03%	11,40%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=tnnt2a PE=1 SV: E7EY9C_DANRE,F1R0K6_I	33.942,20	100,00%	2	2	7	0,03%	11,40%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=tnl1 PE=4 SV=1 A0A0R4ID28_DANRE,A0A	271.258,30	100,00%	5	5	5	0,02%	3,38%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=tnl1 PE=4 SV=1 A0A0R4ID28_DANRE,A0A	271.258,30	100,00%	5	5	5	0,02%	3,38%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=tnl1 PE=4 SV=1 A0A0R4ID28_DANRE,A0A	271.258,30	100,00%	5	5	5	0,02%	3,38%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=tnl2a PE=4 SV=1 A0A0R4IIA7_DANRE,A0AC	271.774,00	99,90%	1	1	1	0,00%	1,53%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=tnl2a PE=4 SV=1 A0A0R4IIA7_DANRE,A0AC	271.774,00	99,90%	1	1	1	0,00%	1,53%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=LOC100333064 A0A0G2L2Q2_DANRE	7.222,40	100,00%	1	1	1	0,00%	24,20%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=col5 F6NPA4_DANRE	198.988,80	100,00%	3	3	5	0,02%	1,44%
7dpa_1	Laminin subunit gamma-1 OS=Danio rerio GN=lamc1 PE=2 SV LAMC1_DANRE	176.206,30	100,00%	4	4	5	0,02%	3,08%
7dpa_1	Citrate synthase OS=Danio rerio GN=csc PE=1 SV=1 B2GTW6_DANRE,CISY_DA	51.752,20	100,00%	2	2	2	0,01%	6,41%
7dpa_1	Citrate synthase OS=Danio rerio GN=csc PE=1 SV=1 B2GTW6_DANRE,CISY_DA	51.752,20	100,00%	2	2	2	0,01%	6,41%
7dpa_1	Alpha-1,4 glucan phosphorylase OS=Danio rerio GN=pygb PE: E7EXT3_DANRE,E7F2Z5_C	96.935,30	45,80%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=sorbs1 PE=4 SV: A0A0R4IP30_DANRE	85.938,80	99,00%	1	1	1	0,00%	1,96%
7dpa_1	Protein S100 OS=Danio rerio GN=S100A10b PE=1 SV=1 Q17VAA_DANRE	11.157,10	100,00%	2	2	2	0,01%	24,00%
7dpa_1	Novel beta globin OS=Danio rerio GN=OTTDARPO000000200: Q6ZM12_DANRE	16.294,80	100,00%	2	2	2	0,01%	15,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=emilin2a PE=4 S F1R1C5_DANRE	122.219,70	100,00%	1	1	1	0,00%	1,18%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ltbp1 PE=4 SV=2 F1QQ56_DANRE,X1WBQ9	131.384,10	99,60%	1	1	1	0,00%	1,31%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ltbp1 PE=4 SV=2 F1QQ56_DANRE,X1WBQ9	131.384,10	99,60%	1	1	1	0,00%	1,31%
7dpa_1	Fhlh protein OS=Danio rerio GN=fhl1a PE=1 SV=1 Q6GQL7_DANRE	34.215,70	46,40%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=uqrc1 PE=1 SV: F1QUQE_DANRE	55.450,70	93,90%	1	1	1	0,00%	1,99%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=tuba1b PE=1 SV B8A516_DANRE	49.969,40	100,00%	2	2	2	0,01%	6,46%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=myom1b PE=4 S A0A0R4IGQ8_DANRE,A0A	161.642,80	100,00%	1	1	1	0,00%	1,04%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=myom1b PE=4 S A0A0R4IGQ8_DANRE,A0A	161.642,80	100,00%	1	1	1	0,00%	1,04%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=col6 F6NRG0_DANRE	88.827,90	87,90%	1	1	1	0,00%	10,60%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=vwf PE=4 SV=2 E7F9Z1_DANRE,F1QHS6_I	225.634,30	97,20%	1	1	2	0,01%	0,46%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=vwf PE=4 SV=2 E7F9Z1_DANRE,F1QHS6_I	225.634,30	97,20%	1	1	2	0,01%	0,46%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1 A0A0R4IEB4_DANRE,A0AI	128.154,40	100,00%	3	3	4	0,01%	3,99%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1 A0A0R4IEB4_DANRE,A0AI	128.154,40	100,00%	3	3	4	0,01%	3,99%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1 A0A0R4IEB4_DANRE,A0AI	128.154,40	100,00%	3	3	4	0,01%	3,99%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1 A0A0R4IEB4_DANRE,A0AI	128.154,40	100,00%	3	3	4	0,01%	3,99%
7dpa_1	Cox5aa protein OS=Danio rerio GN=cox5aa PE=1 SV=1 Q4VBUR7_DANRE	15.897,10	100,00%	2	2	2	0,01%	17,70%
7dpa_1	Ubiquinol-cytochrome c reductase core protein II OS=Danio r Q6IQ59_DANRE	48.122,60	99,80%	1	1	1	0,00%	2,20%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=lama4 PE=1 SV: F1QYUO_DANRE	207.483,80	100,00%	2	2	2	0,01%	0,53%
7dpa_1	Enolase 1, (Alpha) OS=Danio rerio GN=eno1a PE=1 SV=1 Q6PC12_DANRE	47.075,10	100,00%	2	2	3	0,01%	6,48%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=utrn PE=1 SV=1 A0A0H2UKN4_DANRE	299.775,50	100,00%	2	2	2	0,01%	0,91%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:101560 PE=: A0A0R4IBE8_DANRE,A0AI	21.007,60	18,70%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:101560 PE=: A0A0R4IBE8_DANRE,A0AI	21.007,60	18,70%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=lamb2 PE=4 SV= B7ZDA6_DANRE	196.598,90	100,00%	1	1	1	0,00%	0,95%
7dpa_1	Tgm2b protein OS=Danio rerio GN=tgm2b PE=2 SV=1 Q1ECX3_DANRE	75.774,00	100,00%	1	2	2	0,01%	1,62%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=pdha1a PE=4 SV A0A0R4IT09_DANRE,Q6P:	43.741,90	100,00%	2	2	2	0,01%	5,75%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=pdha1a PE=4 SV A0A0R4IT09_DANRE,Q6P:	43.741,90	100,00%	2	2	2	0,01%	5,75%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV A0A0R4I919_DANRE,F6NF	94.381,10	97,20%	1	1	1	0,00%	1,58%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV A0A0R4I919_DANRE,F6NF	94.381,10	97,20%	1	1	1	0,00%	1,58%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV A0A0R4I919_DANRE,F6NF	94.381,10	97,20%	1	1	1	0,00%	1,58%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=acadv1 PE=1 SV: F1Q8I4_DANRE	71.085,10	99,40%	1	1	1	0,00%	2,28%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=acsl1b PE=1 SV: F1QDA6_DANRE	77.694,10	100,00%	1	1	1	0,00%	1,87%
7dpa_1	Dihydropyridyl dehydrogenase OS=Danio rerio GN=dldh PE=: Q6TNUU_DANRE	53.620,50	99,60%	1	1	1	0,00%	2,56%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=col12a1b PE=1 S E7FG81_DANRE,I3IT32_D_	242.762,40	100,00%	2	2	2	0,01%	0,88%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=col12a1b PE=1 S E7FG81_DANRE,I3IT32_D_	242.762,40	100,00%	2	2	2	0,01%	0,88%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ccd141 PE=: B0S755_DANRE,E7FG99_I	190.652,30	100,00%	3	3	4	0,01%	2,28%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ccd141 PE=: B0S755_DANRE,E7FG99_I	190.652,30	100,00%	3	3	4	0,01%	2,28%
7dpa_1	L-lactate dehydrogenase B-A chain OS=Danio rerio GN=ldhba LDHBA_DANRE	36.246,70	100,00%	2	3	5	0,02%	8,08%
7dpa_1	Mut protein OS=Danio rerio GN=mut PE=1 SV=1 A5D653_DANRE	83.519,60	99,90%	1	1	1	0,00%	1,45%
7dpa_1	Heat shock cognate 71 kDa protein OS=Danio rerio GN=hsc7: Q6PGX4_DANRE	70.948,00	99,40%	1	1	1	0,00%	1,85%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=mfap2 PE=4 SV: F1QSF1_DANRE	17.997,30	97,20%	1	1	1	0,00%	6,33%
7dpa_1	Ckmt2 protein OS=Danio rerio GN=ckmt2b PE=2 SV=1 A8E5J9_DANRE,F6NMS7_	46.281,40	100,00%	3	4	4	0,01%	11,10%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ckmt2a PE=2 SV Q7ZU7U_DANRE	20.487,80	97,20%	1	1	2	0,01%	24,20%
7dpa_1	Ckmt2 protein OS=Danio rerio GN=ckmt2b PE=2 SV=1 A8E5J9_DANRE,F6NMS7_	46.281,40	100,00%	3	4	4	0,01%	11,10%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=sorbs2a PE=1 SV F1QI09_DANRE	148.129,10	100,00%	1	1	1	0,00%	1,50%
7dpa_1	Nicotinamide nucleotide transhydrogenase OS=Danio rerio G Q6NYP7_DANRE	112.785,50	84,90%	1	1	1	0,00%	1,11%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=thbs4a PE=4 SV: A0A0R4I9N1_DANRE,A0A	104.325,40	77,00%	1	1	1	0,00%	1,16%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=thbs4a PE=4 SV: A0A0R4I9N1_DANRE,A0A	104.325,40	77,00%	1	1	1	0,00%	1,16%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=thbs4a PE=4 SV: A0A0R4I9N1_DANRE,A0A	104.325,40	77,00%	1	1	1	0,00%	1,16%
7dpa_1	Hydroxyacyl-Coenzyme A dehydrogenase OS=Danio rerio GN Q6DI22_DANRE	33.336,90	100,00%	2	2	2	0,01%	9,39%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=hadhb PE=1 SV F1QYX8_DANRE	82.848,80	100,00%	2	2	3	0,01%	3,28%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ndrg4 PE=4 SV= A0A0R4IDM0_DANRE,A6)	37.210,30	100,00%	2	2	2	0,01%	10,90%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ndrg4 PE=4 SV= A0A0R4IDM0_DANRE,A6)	37.210,30	100,00%	2	2	2	0,01%	10,90%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=esyt1b PE=1 SV: F1QVPP9_DANRE,F6NZP4_	175.083,00	26,40%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=esyt1b PE=1 SV: F1QVPP9_DANRE,F6NZP4_	175.083,00	26,40%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ptx3a PE=1 SV=: A3KPN4_DANRE	49.273,20	100,00%	3	5	6	0,02%	8,28%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=atp5b PE=1 SV= A2ZAG7_DANRE	8.316,90	100,00%	2	3	3	0,01%	36,60%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=cox6c PE=1 SV= Q6ZM23_DANRE	8.480,50	10,00%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=hnrr F8W446_DANRE,Q6NYU8	37.111,30	100,00%	1	1	1	0,00%	4,82%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=hnrr F8W446_DANRE,Q6NYU8	37.111,30	100,00%	1	1	1	0,00%	4,82%
7dpa_1	Serotransferrin OS=Danio rerio GN=tfa PE=1 SV=1 B8JL43_DANRE,F1R858_D	73.717,60	100,00%	2	2	3	0,01%	3,86%
7dpa_1	Serotransferrin OS=Danio rerio GN=tfa PE=1 SV=1 B8JL43_DANRE,F1R858_D	73.717,60	100,00%	2	2	3	0,01%	3,86%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=popdc2 PE=1 SV F8W5H2_DANRE,Q6JWV8	41.034,60	99,90%	1	1	1	0,00%	3,54%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=popdc2 PE=1 SV F8W5H2_DANRE,Q6JWV8	41.034,60	99,90%	1	1	1	0,00%	3,54%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	20.097,20	41,30%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	20.097,20	41,30%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	20.097,20	41,30%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	20.097,20	41,30%	0	0	0	0,00%	0,00%

7dpa_1	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,AOAC	20.097,20	41,30%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ndufs1 PE=1 SV: F1RDA6_DANRE	79.460,90	78,90%	1	1	1	0,00%	2,60%
7dpa_1	Ceruloplasmin OS=Danio rerio GN=cp PE=1 SV=1 Q6P3G1_DANRE	124.388,00	100,00%	2	2	2	0,01%	2,58%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV A0A0R4IH07_DANRE,A0A	342.958,30	22,90%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV A0A0R4IH07_DANRE,A0A	342.958,30	22,90%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV A0A0R4IH07_DANRE,A0A	342.958,30	22,90%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV A0A0R4IH07_DANRE,A0A	342.958,30	22,90%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV A0A0R4IH07_DANRE,A0A	342.958,30	22,90%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV A0A0R4IH07_DANRE,A0A	342.958,30	22,90%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=itih3b PE=4 SV= A0A0R4INP2_DANRE,A0A	100.008,80	99,80%	1	1	1	0,00%	2,13%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=itih3b PE=4 SV= A0A0R4INP2_DANRE,A0A	100.008,80	99,80%	1	1	1	0,00%	2,13%
7dpa_1	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IG11_DANRE,G3P_	35.784,00	100,00%	2	2	3	0,01%	9,52%
7dpa_1	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IG11_DANRE,G3P_	35.784,00	100,00%	2	2	3	0,01%	9,52%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=Imo7a PE=4 SV= A0A0R4IT8_DANRE,BOU	207.438,50	100,00%	2	2	2	0,01%	2,26%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=myoz2b PE=1 SV E9QE52_DANRE,F1QB1_	25.288,50	89,70%	1	1	2	0,01%	14,60%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=myoz2b PE=1 SV E9QE52_DANRE,F1QB1_	25.288,50	89,70%	1	1	2	0,01%	14,60%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=myoz2b PE=1 SV E9QE52_DANRE,F1QB1_	25.288,50	89,70%	1	1	2	0,01%	14,60%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=pccb PE=1 SV=1 A0A0R4IGR9_DANRE,BOV	60.927,60	100,00%	2	2	2	0,01%	6,49%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=pccb PE=1 SV=1 A0A0R4IGR9_DANRE,BOV	60.927,60	100,00%	2	2	2	0,01%	6,49%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=pccb PE=1 SV=1 A0A0R4IGR9_DANRE,BOV	60.927,60	100,00%	2	2	2	0,01%	6,49%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:cf E9QC7_DANRE	83.925,70	91,40%	1	1	1	0,00%	1,87%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=PE=4 SV=1 A0A0G2L836_DANRE	136.805,30	97,20%	1	1	1	0,00%	1,21%
7dpa_1	Transforming growth factor, beta-induced OS=Danio rerio G Q503K1_DANRE	74.192,20	100,00%	1	1	2	0,01%	2,36%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=si:dkryp-80C12. H0WES8_DANRE	6.338,70	96,00%	1	1	1	0,00%	27,60%
7dpa_1	Myosin light chain 2 Mic2a OS=Danio rerio GN=myl7 PE=1 SV Q801M3_DANRE	19.021,00	100,00%	5	6	14	0,05%	57,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=vtmb PE=1 SV=1 F1Q5D8_DANRE	51.527,70	50,70%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV A0A0R4IH03_DANRE,F1R	108.322,90	100,00%	3	3	3	0,01%	5,78%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV A0A0R4IH03_DANRE,F1R	108.322,90	100,00%	3	3	3	0,01%	5,78%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV A0A0R4IH03_DANRE,F1R	108.322,90	100,00%	3	3	3	0,01%	5,78%
7dpa_1	Acetyl-CoA acetyltransferase, mitochondrial OS=Danio rerio A0A0R4I176_DANRE,THIL	44.342,90	10,40%	0	0	0	0,00%	0,00%
7dpa_1	Acetyl-CoA acetyltransferase, mitochondrial OS=Danio rerio A0A0R4I176_DANRE,THIL	44.342,90	10,40%	0	0	0	0,00%	0,00%
7dpa_1	Blood vessel epicardial substance OS=Danio rerio GN=bves P POPD1_DANRE	40.341,40	100,00%	2	2	2	0,01%	9,38%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=c9 PE=1 SV=1 B8A565_DANRE	72.713,70	84,20%	1	1	1	0,00%	2,08%
7dpa_1	Pyruvate dehydrogenase (Lipoamide) beta OS=Danio rerio GI Q7T368_DANRE	39.308,70	100,00%	1	1	1	0,00%	4,46%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=ablim1b PE=4 SV U3J1B3_DANRE	84.999,80	99,70%	1	1	1	0,00%	2,11%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=mfap5 PE=4 SV= E9QC4_DANRE	14.664,20	97,20%	1	1	14	0,05%	11,90%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=dlst PE=1 SV=1 F1QFQ6_DANRE	48.683,10	97,20%	1	1	2	0,01%	4,59%
7dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:df F1Q6R2_DANRE	193.577,50	97,20%	1	1	1	0,00%	1,16%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=PE=3 SV=1 A0A0G2KML7_DANRE,BO	41.786,40	97,20%	1	1	2	0,01%	8,44%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=PE=3 SV=1 A0A0G2KML7_DANRE,BO	41.786,40	97,20%	1	1	2	0,01%	8,44%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=pdlim5b PE=1 SV F1QGC3_DANRE	21.781,00	78,90%	1	1	1	0,00%	10,70%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=PE=1 SV=1 A0A0G2KHA1_DANRE	68.436,80	91,20%	1	1	1	0,00%	3,34%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=tns1b PE=1 SV=: F1Q649_DANRE,F1QJY1_I	77.272,20	24,20%	0	0	0	0,00%	0,00%
7dpa_1	Uncharacterized protein OS=Danio rerio GN=tns1b PE=1 SV=: F1Q649_DANRE,F1QJY1_I	77.272,20	24,20%	0	0	0	0,00%	0,00%
7dpa_1	E9QD93-DECOY	E9QD93-DECOY,Q5RH26_	18	27,30%	0	0	0,00%	NaNa%
7dpa_1	E9QD93-DECOY	E9QD93-DECOY,Q5RH26_	18	27,30%	0	0	0,00%	NaNa%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=vtg1 PE=1 SV=1 Q1LWN2_DANRE	149.238,80	100,00%	2	2	3	0,01%	1,91%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=fga PE=1 SV=1 B8A5L6_DANRE,E9QCD1_	49.680,00	100,00%	24	33	70	0,24%	38,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=fga PE=1 SV=1 B8A5L6_DANRE,E9QCD1_	49.680,00	100,00%	24	33	70	0,24%	38,00%
7dpa_2	Fgg protein OS=Danio rerio GN=fgg PE=1 SV=1 Q7ZV77_DANRE	48.833,70	100,00%	27	46	280	0,96%	65,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=ctnna1 PE=4 SV: A0A0R4IF47_DANRE,Q9P	100.520,10	100,00%	3	3	3	0,01%	7,38%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=ctnna1 PE=4 SV: A0A0R4IF47_DANRE,Q9P	100.520,10	100,00%	3	3	3	0,01%	7,38%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=vcla PE=4 SV=1 A0A0R4IYQ3_DANRE,B3D	124.046,80	94,80%	1	1	1	0,00%	1,31%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=vcla PE=4 SV=1 A0A0R4IYQ3_DANRE,B3D	124.046,80	94,80%	1	1	1	0,00%	1,31%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=vcla PE=4 SV=1 A0A0R4IYQ3_DANRE,B3D	124.046,80	94,80%	1	1	1	0,00%	1,31%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=col6a3 PE=1 SV: F1Q4X1_DANRE	309.287,80	100,00%	7	10	17	0,06%	4,10%
7dpa_2	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1 A8W605_DANRE,ACTB1_	41.753,80	100,00%	3	3	3	0,01%	22,40%
7dpa_2	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1 A8W605_DANRE,ACTB1_	41.753,80	100,00%	3	3	3	0,01%	22,40%
7dpa_2	Actin, alpha, cardiac muscle 1a OS=Danio rerio GN=si:ch73-11 Q6IQR3_DANRE	41.976,10	98,00%	1	2	2	0,01%	43,50%
7dpa_2	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1 A8W605_DANRE,ACTB1_	41.753,80	100,00%	3	3	3	0,01%	22,40%
7dpa_2	Actin OS=Danio rerio GN=act1b PE=1 SV=1 Q6XN18_DANRE	41.976,10	100,00%	11	13	29	0,10%	43,50%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=sorbs1 PE=4 SV: A0A0R4IP30_DANRE	85.938,80	95,30%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=sptbn1 PE=1 SV: A0A0G2L1F5_DANRE	262.630,10	100,00%	4	4	5	0,02%	2,68%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=flna PE=1 SV=1 E9QI62_DANRE,F1R9P5_I	272.475,80	100,00%	1	1	1	0,00%	0,55%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=flna PE=1 SV=1 E9QI62_DANRE,F1R9P5_I	272.475,80	100,00%	1	1	1	0,00%	0,55%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=fn1b PE=1 SV=2 A2CEW3_DANRE,F1R809_	265.661,60	100,00%	42	55	116	0,40%	29,10%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=fn1b PE=1 SV=2 A2CEW3_DANRE,F1R809_	265.661,60	100,00%	42	55	116	0,40%	29,10%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tnl1 PE=4 SV=1 A0A0R4ID28_DANRE,A0A	271.258,30	100,00%	3	3	3	0,01%	2,20%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tnl1 PE=4 SV=1 A0A0R4ID28_DANRE,A0A	271.258,30	100,00%	3	3	3	0,01%	2,20%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tnl1 PE=4 SV=1 A0A0R4ID28_DANRE,A0A	271.258,30	100,00%	3	3	3	0,01%	2,20%
7dpa_2	Fibrinogen, B beta polypeptide OS=Danio rerio GN=fgb PE=1 Q6NEY1_DANRE	54.407,60	100,00%	17	31	179	0,62%	39,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 SV E7EZ79_DANRE,F1Q615_I	142.109,70	100,00%	2	2	2	0,01%	1,97%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 SV E7EZ79_DANRE,F1Q615_I	142.109,70	100,00%	2	2	2	0,01%	1,97%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tnna PE=1 SV=2 F1R7N8_DANRE	3.089.800,00	100,00%	3	3	2	0,01%	0,16%
7dpa_2	ATP synthase subunit beta OS=Danio rerio GN=atp5b PE=1 SV A8W6C6_DANRE	55.130,90	100,00%	7	8	17	0,06%	23,40%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=postna PE=1 SV: E7F513_DANRE	97.135,00	100,00%	8	10	13	0,04%	15,30%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a1a PE=1 SV F1QJC9_DANRE,F1QSC5_I	128.043,70	100,00%	4	5	5	0,02%	3,80%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a1a PE=1 SV F1QJ11_DANRE	136.914,60	100,00%	7	7	13	0,04%	6,69%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a1a PE=1 SV F1QJ11_DANRE,F1QSC5_I	128.043,70	100,00%	4	5	5	0,02%	3,80%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=Imo A0A0R4IT8_DANRE,BOU	207.438,50	74,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=Imo A0A0R4IT8_DANRE,BOU	207.438,50	74,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=vmhcl PE=1 SV= F1QSE1_DANRE	223.380,90	100,00%	55	65	116	0,40%	33,50%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=fn1a PE=1 SV=1 B0S602_DANRE	271.953,50	100,00%	6	6	8	0,03%	5,40%
7dpa_2	Hemoglobin subunit beta-2 OS=Danio rerio GN=ba2 PE=1 SV: HBB2_DANRE	16.389,20	100,00%	2	2	2	0,01%	14,90%
7dpa_2	Anion exchange protein OS=Danio rerio GN=slc4a1a PE=1 SV: Q7ZZJ7_DANRE	101.463,30	99,90%	1	1	1	0,00%	1,55%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=desma PE=1 SV: F1R8W3_DANRE	55.727,70	100,00%	6	7	9	0,03%	12,50%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=tnnb B0S6Y0_DANRE	626.764,10	94,30%	1	1	2	0,01%	0,36%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=1 A0A0R4INC4_DANRE,ASV	543.007,50	99,80%	1	1	1	0,00%	0,35%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=1 A0A0R4INC4_DANRE,ASV	543.007,50	99,80%	1	1	1	0,00%	0,35%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=1 A0A0R4INC4_DANRE,ASV	543.007,50	99,80%	1	1	1	0,00%	0,35%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=mfap5 PE=4 SV: E9QC4_DANRE	14.664,20	99,90%	1	2	2	0,01%	11,90%
7dpa_2	Solute carrier family 25 alpha, member 5 OS=Danio rerio GN: Q8JH10_DANRE	32.764,40	100,00%	3	3	3	0,01%	10,40%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=tnna B0S758_DANRE	843.490,10	99,40%	1	1	1	0,00%	0,30%

7dpa_2	Heat shock cognate 71 kDa protein OS=Danio rerio GN=hspa1 A0A0R4IMF8_DANRE,A0A	71.182,10	100,00%	2	2	2	0,01%	4,44%
7dpa_2	Heat shock cognate 71 kDa protein OS=Danio rerio GN=hspa1 A0A0R4IMF8_DANRE,A0A	71.182,10	100,00%	2	2	2	0,01%	4,44%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=zc:112265 PE= A0A0R4IE46_DANRE,A0A	101.943,10	100,00%	4	4	5	0,02%	5,84%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=zc:112265 PE= A0A0R4IE46_DANRE,A0A	101.943,10	100,00%	4	4	5	0,02%	5,84%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=zc:112265 PE= A0A0R4IE46_DANRE,A0A	101.943,10	100,00%	4	4	5	0,02%	5,84%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV= A0A0R4IBJ3_DANRE,F1R6	72.513,60	99,80%	1	1	1	0,00%	1,19%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV= A0A0R4IBJ3_DANRE,F1R6	72.513,60	99,80%	1	1	1	0,00%	1,19%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV= A0A0R4IBJ3_DANRE,F1R6	72.513,60	99,80%	1	1	1	0,00%	1,19%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV= A0A0R4IBJ3_DANRE,F1R6	72.513,60	99,80%	1	1	1	0,00%	1,19%
7dpa_2	Cadherin-2 OS=Danio rerio GN=cdh2 PE=2 SV=2 CADH2_DANRE,G1K2E7_I	99.051,70	98,90%	1	1	1	0,00%	1,01%
7dpa_2	Cadherin-2 OS=Danio rerio GN=cdh2 PE=2 SV=2 CADH2_DANRE,G1K2E7_I	99.051,70	98,90%	1	1	1	0,00%	1,01%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV= A0A0R4IH03_DANRE,A0A	108.322,90	100,00%	2	2	2	0,01%	3,61%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV= A0A0R4IH03_DANRE,A0A	108.322,90	100,00%	2	2	2	0,01%	3,61%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV= A0A0R4IH03_DANRE,A0A	108.322,90	100,00%	2	2	2	0,01%	3,61%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=cmlc1 PE=1 SV= B0R0F7_DANRE	21.865,20	100,00%	10	12	20	0,07%	60,70%
7dpa_2	Inner membrane protein, mitochondrial (Mitofilin) OS=Danio Q6PF54_DANRE	83.201,30	55,20%	0	0	0	0,00%	0,00%
7dpa_2	HEAT repeat-containing protein 5A OS=Danio rerio GN=heatr E7F1I7_DANRE,E9QH09_I	216.706,90	99,80%	1	1	1	0,00%	0,55%
7dpa_2	HEAT repeat-containing protein 5A OS=Danio rerio GN=heatr E7F1I7_DANRE,E9QH09_I	216.706,90	99,80%	1	1	1	0,00%	0,55%
7dpa_2	HEAT repeat-containing protein 5A OS=Danio rerio GN=heatr E7F1I7_DANRE,E9QH09_I	216.706,90	99,80%	1	1	1	0,00%	0,55%
7dpa_2	HEAT repeat-containing protein 5A OS=Danio rerio GN=heatr E7F1I7_DANRE,E9QH09_I	216.706,90	99,80%	1	1	1	0,00%	0,55%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=glud1a PE=1 SV= A0A0R4IU7_DANRE,B8A	59.978,90	14,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=glud1a PE=1 SV= A0A0R4IU7_DANRE,B8A	59.978,90	14,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=glud1a PE=1 SV= A0A0R4IU7_DANRE,B8A	59.978,90	14,60%	0	0	0	0,00%	0,00%
7dpa_2	Laminin subunit gamma-1 OS=Danio rerio GN=lamc1 PE=2 SV= LAMC1_DANRE	176.206,30	99,20%	1	1	1	0,00%	1,32%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=vdac3 PE=1 SV= B0R197_DANRE,B0R198_I	29.812,00	39,30%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=vdac3 PE=1 SV= B0R197_DANRE,B0R198_I	29.812,00	39,30%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV= A0A0R4IU91_DANRE,F6NF	94.381,10	97,70%	1	1	1	0,00%	1,70%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV= A0A0R4IU91_DANRE,F6NF	94.381,10	97,70%	1	1	1	0,00%	1,70%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV= A0A0R4IU91_DANRE,F6NF	94.381,10	97,70%	1	1	1	0,00%	1,70%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV= A0A0R4IU91_DANRE,F6NF	94.381,10	97,70%	1	1	1	0,00%	1,70%
7dpa_2	Elongation factor 1-alpha OS=Danio rerio GN=eef1a PE=2 SV= EF1A_DANRE	50.048,20	100,00%	2	2	2	0,01%	8,44%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tln2a PE=4 SV= A0A0R4IIA7_DANRE	271.364,80	97,80%	1	1	1	0,00%	1,18%
7dpa_2	Dihydropyridyl dehydrogenase OS=Danio rerio GN=dlhd PE=1 SV= Q6TNU6_DANRE	53.620,50	70,20%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=myoz2b PE=1 SV= F1QKE7_DANRE	25.288,50	17,90%	0	0	0	0,00%	0,00%
7dpa_2	Alpha-1,4 glucan phosphorylase OS=Danio rerio GN=pygmb F E7EXT3_DANRE	96.868,30	45,90%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=sorbs2a PE=1 SV= F1QI09_DANRE	148.129,10	99,90%	1	1	1	0,00%	1,50%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=uqrcr1 PE=1 SV= F1QUE3_DANRE	55.450,70	99,10%	1	1	1	0,00%	1,99%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=hmcn2 PE=1 SV= E7F3V2_DANRE	454.987,50	100,00%	2	2	2	0,01%	6,24%
7dpa_2	Calcium-transporting ATPase OS=Danio rerio GN=atp2a2a PE= A9C3Q4_DANRE,A9C3Q5	109.519,40	99,90%	1	1	1	0,00%	2,21%
7dpa_2	Calcium-transporting ATPase OS=Danio rerio GN=atp2a2a PE= A9C3Q4_DANRE,A9C3Q5	109.519,40	99,90%	1	1	1	0,00%	2,21%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=paplnb PE=4 SV= F1QRA8_DANRE	60.121,30	100,00%	1	1	1	0,00%	2,19%
7dpa_2	Aconitate hydratase, mitochondrial (Fragment) OS=Danio rer F8W4M7_DANRE	85.591,90	100,00%	1	1	1	0,00%	2,15%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=ptx3a PE=1 SV= A3KPN4_DANRE	49.273,20	100,00%	1	1	1	0,00%	3,36%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=myom1b PE=4 SV= A0A0R4IGQ8_DANRE,A0A	161.642,80	93,50%	1	1	1	0,00%	0,80%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=myom1b PE=4 SV= A0A0R4IGQ8_DANRE,A0A	161.642,80	93,50%	1	1	1	0,00%	0,80%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=col6a1 PE=1 SV= F1Q6P3_DANRE	107.466,50	100,00%	6	6	7	0,02%	8,04%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=eno F1QBW7_DANRE	51.320,60	100,00%	1	1	1	0,00%	5,57%
7dpa_2	Histone H4 OS=Danio rerio GN=si:dkey-108k21.23 PE=1 SV=1 A3KPR4_DANRE,X1WGM1	11.478,70	100,00%	1	1	2	0,01%	11,70%
7dpa_2	Histone H4 OS=Danio rerio GN=si:dkey-108k21.23 PE=1 SV=1 A3KPR4_DANRE,X1WGM1	11.478,70	100,00%	1	1	2	0,01%	11,70%
7dpa_2	Malate dehydrogenase OS=Danio rerio GN=mdh2 PE=1 SV=1 Q7T334_DANRE	35.420,30	100,00%	2	2	2	0,01%	8,90%
7dpa_2	Succinyl-CoA ligase subunit beta OS=Danio rerio GN=sucla2 P Q6PB10_DANRE	50.032,30	79,90%	1	1	1	0,00%	5,58%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=c9 PE=1 SV=1 B8A565_DANRE	72.713,70	99,80%	1	1	2	0,01%	3,12%
7dpa_2	Sodium/potassium-transporting ATPase subunit alpha OS=D A0A0R4IRT1_DANRE,Q9D	113.384,80	90,70%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV= A0A0R4IP01_DANRE,B8JK	119.385,60	99,90%	1	1	1	0,00%	1,17%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV= A0A0R4IP01_DANRE,B8JK	119.385,60	99,90%	1	1	1	0,00%	1,17%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV= A0A0R4IP01_DANRE,B8JK	119.385,60	99,90%	1	1	1	0,00%	1,17%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=col12a1b PE=1 SV= E7FG81_DANRE,I3IT32_D	242.762,40	22,30%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=col12a1b PE=1 SV= E7FG81_DANRE,I3IT32_D	242.762,40	22,30%	0	0	0	0,00%	0,00%
7dpa_2	ATP synthase subunit gamma OS=Danio rerio GN=atp5c1 PE= Q5RH26_DANRE	32.149,90	99,40%	1	1	1	0,00%	10,30%
7dpa_2	Myosin light chain 2 Mlc2a OS=Danio rerio GN=myl7 PE=1 SV= Q801M3_DANRE	19.021,00	100,00%	5	5	7	0,02%	41,30%
7dpa_2	Tgm2b protein OS=Danio rerio GN=tgm2b PE=2 SV=1 Q1ECX3_DANRE	75.774,00	100,00%	1	1	1	0,00%	1,62%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=mfap2 PE=4 SV= F1QSF1_DANRE	17.997,30	92,90%	1	1	1	0,00%	6,33%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=col4a2 PE=1 SV= A0A0H2UKP2_DANRE,F1C	103.249,70	100,00%	3	3	3	0,01%	3,89%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=col4a2 PE=1 SV= A0A0H2UKP2_DANRE,F1C	103.249,70	100,00%	3	3	3	0,01%	3,89%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=nid1a PE=1 SV= F1RA63_DANRE	130.098,40	100,00%	2	2	3	0,01%	2,47%
7dpa_2	Ubiquinol-cytochrome c reductase core protein II OS=Danio r Q6IQ59_DANRE	48.122,60	47,30%	0	0	0	0,00%	0,00%
7dpa_2	Myoglobin OS=Danio rerio GN=mb PE=1 SV=1 H0WE38_DANRE,MYG_D	15.583,60	100,00%	1	1	5	0,02%	13,20%
7dpa_2	Myoglobin OS=Danio rerio GN=mb PE=1 SV=1 H0WE38_DANRE,MYG_D	15.583,60	100,00%	1	1	5	0,02%	13,20%
7dpa_2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	98,00%	1	1	1	0,00%	12,10%
7dpa_2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	98,00%	1	1	1	0,00%	12,10%
7dpa_2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	98,00%	1	1	1	0,00%	12,10%
7dpa_2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	98,00%	1	1	1	0,00%	12,10%
7dpa_2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	98,00%	1	1	1	0,00%	12,10%
7dpa_2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	98,00%	1	1	1	0,00%	12,10%
7dpa_2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	98,00%	1	1	1	0,00%	12,10%
7dpa_2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	98,00%	1	1	1	0,00%	12,10%
7dpa_2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	98,00%	1	1	1	0,00%	12,10%
7dpa_2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	98,00%	1	1	1	0,00%	12,10%
7dpa_2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	98,00%	1	1	1	0,00%	12,10%
7dpa_2	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	98,00%	1	1	1	0,00%	12,10%
7dpa_2	Pyruvate dehydrogenase (Lipoamide) beta OS=Danio rerio G1Q7T368_DANRE	39.308,70	99,90%	1	1	1	0,00%	4,46%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tpm4a PE=1 SV= F1QKQ7_DANRE	32.726,90	100,00%	5	5	6	0,02%	25,40%
7dpa_2	Annexin OS=Danio rerio GN=anxa2a PE=1 SV=1 Q6P6Q3_DANRE	38.140,50	100,00%	2	2	2	0,01%	7,42%
7dpa_2	Protein S100 OS=Danio rerio GN=S100A10b PE=1 SV=1 Q7ZV44_DANRE	11.157,10	98,00%	1	1	1	0,00%	13,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=fbn2b PE=1 SV= A0A0G2KQ62_DANRE	212.420,00	100,00%	2	2	3	0,01%	1,95%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=krtr94 PE=1 SV= F1R8U0_DANRE,Q1RLR3	50.932,10	21,00%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=krtr94 PE=1 SV= F1R8U0_DANRE,Q1RLR3	50.932,10	21,00%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=popdc2 PE=1 SV= F8W5H2_DANRE,Q6JWV8	41.034,60	44,20%	0	0	0	0,00%	0,00%

7dpa_2	Uncharacterized protein OS=Danio rerio GN=popdc2 PE=1 SV F8W5H2_DANRE,Q6JWV8	41.034,60	44,20%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=elnb PE=4 SV=1 A0A0R4IPZ1_DANRE,F8W	171.803,90	98,00%	1	1	1	0,00%	0,79%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=elnb PE=4 SV=1 A0A0R4IPZ1_DANRE,F8W	171.803,90	98,00%	1	1	1	0,00%	0,79%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=elnb PE=4 SV=1 A0A0R4IPZ1_DANRE,F8W	171.803,90	98,00%	1	1	1	0,00%	0,79%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=1 SV= A0A0R4IF18_DANRE,A2Bc	103.798,70	99,40%	1	1	1	0,00%	1,22%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=1 SV= A0A0R4IF18_DANRE,A2Bc	103.798,70	99,40%	1	1	1	0,00%	1,22%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=1 SV= A0A0R4IF18_DANRE,A2Bc	103.798,70	99,40%	1	1	1	0,00%	1,22%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=col4a1 PE=1 SV: F1Q079_DANRE	128.554,50	98,00%	1	2	3	0,01%	1,06%
7dpa_2	Heat shock cognate 71 kDa protein OS=Danio rerio GN=hspa1 A0A0R4IMF8_DANRE,A0A	71.182,10	100,00%	2	2	2	0,01%	4,44%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=atp5f1 PE=1 SV: B8J5I1_DANRE	28.169,80	99,90%	1	1	1	0,00%	6,35%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=emi F1Q105_DANRE,F1R1C5_L	122.219,70	53,40%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=emi F1Q105_DANRE,F1R1C5_L	122.219,70	53,40%	0	0	0	0,00%	0,00%
7dpa_2	Sodium/potassium-transporting ATPase subunit alpha OS=D: A0A0R4IRT1_DANRE,Q9D	113.384,80	99,70%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=hadhab PE=1 SV F1QYX8_DANRE	82.848,80	24,20%	0	0	0	0,00%	0,00%
7dpa_2	Transforming growth factor, beta-induced OS=Danio rerio GN= Q503K1_DANRE	74.192,20	98,00%	1	1	1	0,00%	2,36%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=si:dkeyp-80c12. H0WES8_DANRE	6.338,70	98,00%	1	2	2	0,01%	27,60%
7dpa_2	Aspartate aminotransferase OS=Danio rerio GN=get2a PE=1 ! Q7SYK7_DANRE	47.588,10	31,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tuba1b PE=1 SV A0A0R4IG84_DANRE,B8A	50.149,80	99,70%	1	1	1	0,00%	4,45%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tuba1b PE=1 SV A0A0R4IG84_DANRE,B8A	50.149,80	99,70%	1	1	1	0,00%	4,45%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tuba1b PE=1 SV A0A0R4IG84_DANRE,B8A	50.149,80	99,70%	1	1	1	0,00%	4,45%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tuba1b PE=1 SV A0A0R4IG84_DANRE,B8A	50.149,80	99,70%	1	1	1	0,00%	4,45%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tuba1b PE=1 SV A0A0R4IG84_DANRE,B8A	50.149,80	99,70%	1	1	1	0,00%	4,45%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tuba1b PE=1 SV A0A0R4IG84_DANRE,B8A	50.149,80	99,70%	1	1	1	0,00%	4,45%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tuba1b PE=1 SV A0A0R4IG84_DANRE,B8A	50.149,80	99,70%	1	1	1	0,00%	4,45%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=tuba1b PE=1 SV A0A0R4IG84_DANRE,B8A	50.149,80	99,70%	1	1	1	0,00%	4,45%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV A0A0R4IH07_DANRE,B6IC	347.704,80	41,20%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV A0A0R4IH07_DANRE,B6IC	347.704,80	41,20%	0	0	0	0,00%	0,00%
7dpa_2	Phosphoglycerate kinase OS=Danio rerio GN=pgk1 PE=1 SV=: F1QXV8_DANRE	44.733,20	98,40%	1	1	1	0,00%	3,84%
7dpa_2	Hydroxyacyl-Coenzyme A dehydrogenase OS=Danio rerio GN Q6D122_DANRE	33.336,90	34,40%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=lmo A0A0R4ITT8_DANRE,B0U	207.438,50	74,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=slc2 E9QJ49_DANRE,F1QAJ7_L	39.378,90	97,40%	1	1	1	0,00%	5,73%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=slc2 E9QJ49_DANRE,F1QAJ7_L	39.378,90	97,40%	1	1	1	0,00%	5,73%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=sdprb PE=1 SV= F1R665_DANRE	51.441,60	98,00%	1	1	1	0,00%	3,59%
7dpa_2	Blood vessel epicardial substance OS=Danio rerio GN=bves P POPD1_DANRE	40.341,40	98,00%	1	2	2	0,01%	5,11%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=serbp1a PE=1 SV F1Q5Q3_DANRE	43.221,10	100,00%	1	1	1	0,00%	8,93%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=acaa2 PE=1 SV= B055C5_DANRE	41.786,40	99,80%	1	1	1	0,00%	5,04%
7dpa_2	Caveolin OS=Danio rerio GN=cav1 PE=1 SV=1 Q6YLH8_DANRE,Q6YLH9_	20.652,80	94,00%	1	1	1	0,00%	14,20%
7dpa_2	Caveolin OS=Danio rerio GN=cav1 PE=1 SV=1 Q6YLH8_DANRE,Q6YLH9_	20.652,80	94,00%	1	1	1	0,00%	14,20%
7dpa_2	Acetyl-CoA acetyltransferase, mitochondrial OS=Danio rerio A0A0R4I76_DANRE,THIL_	44.342,90	25,40%	0	0	0	0,00%	0,00%
7dpa_2	Heat shock 60kD protein 1 (Chaperonin) OS=Danio rerio GN= Q083B0_DANRE	61.196,20	0,760%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	86.013,20	16,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	86.013,20	16,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	86.013,20	16,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	86.013,20	16,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	86.013,20	16,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	86.013,20	16,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	86.013,20	16,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	86.013,20	16,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	86.013,20	16,60%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4ILE0_DANRE,A0AC	86.013,20	16,60%	0	0	0	0,00%	0,00%
7dpa_2	Acetyl-CoA acetyltransferase, mitochondrial OS=Danio rerio A0A0R4I76_DANRE,THIL_	44.342,90	25,40%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6P	29.220,20	30,50%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6P	29.220,20	30,50%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6P	29.220,20	30,50%	0	0	0	0,00%	0,00%
7dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6P	29.220,20	30,50%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=slc25a12 PE=4 S A0A0R4IHT0_DANRE,A0A	72.864,20	99,90%	1	1	1	0,00%	1,61%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=slc25a12 PE=4 S A0A0R4IHT0_DANRE,A0A	72.864,20	99,90%	1	1	1	0,00%	1,61%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=col6 F1Q924_DANRE	278.174,40	100,00%	14	14	23	0,07%	8,73%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4IJ71_DANRE,E7F4	518.896,50	17,30%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=: A0A0R4INC4_DANRE,A5V	543.007,50	99,80%	1	1	1	0,00%	0,18%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=: A0A0R4INC4_DANRE,A5V	543.007,50	99,80%	1	1	1	0,00%	0,18%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4IJ71_DANRE,E7F4	518.896,50	17,30%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4IJ71_DANRE,E7F4	518.896,50	17,30%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttnb G1K2X0_DANRE	2.458.846,80	100,00%	11	11	14	0,04%	0,85%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ndufs1 PE=1 SV: F1RDA6_DANRE	79.460,90	15,00%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=add1 PE=4 SV=: A0A0R4IAJ9_DANRE,A0AC	81.732,80	76,60%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=add1 PE=4 SV=: A0A0R4IAJ9_DANRE,A0AC	81.732,80	76,60%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=add1 PE=4 SV=: A0A0R4IAJ9_DANRE,A0AC	81.732,80	76,60%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=col12a1a PE=1 S X1WF98_DANRE	321.902,50	96,90%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:cf F1QTR6_DANRE	147.365,40	100,00%	3	3	3	0,01%	1,18%
14dpa_1	Malate dehydrogenase 1a, NAD (Soluble) OS=Danio rerio GN Q7T3D9_DANRE,Q801E6_	36.258,40	96,30%	0	0	0	0,00%	0,00%
14dpa_1	Malate dehydrogenase 1a, NAD (Soluble) OS=Danio rerio GN Q7T3D9_DANRE,Q801E6_	36.258,40	96,30%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ttna PE=1 SV=2 F1R7N8_DANRE	3.089.806,00	100,00%	13	13	14	0,04%	0,77%
14dpa_1	Inner membrane protein, mitochondrial (Mitofilin) OS=Danio Q6PFS4_DANRE	83.201,30	99,80%	1	1	1	0,00%	1,32%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6P	29.220,20	34,20%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6P	29.220,20	34,20%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6P	29.220,20	34,20%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=acaa2 PE=1 SV= B055C5_DANRE	41.786,40	98,10%	1	1	1	0,00%	5,04%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=fn1b PE=1 SV=2 A2CEW3_DANRE,F1R809_	265.661,60	100,00%	41	56	205	0,66%	28,50%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=fn1b PE=1 SV=2 A2CEW3_DANRE,F1R809_	265.661,60	100,00%	41	56	205	0,66%	28,50%
14dpa_1	Fructose-bisphosphate aldolase OS=Danio rerio GN=aldoaa P Q803Q7_DANRE	39.740,60	100,00%	2	3	3	0,01%	9,34%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ahnk PE=1 SV= F1QZ50_DANRE	685.184,90	100,00%	2	2	2	0,01%	0,61%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=col6a2 PE=4 SV: E7FCV8_DANRE	106.916,20	100,00%	5	5	18	0,06%	10,40%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttna B05758_DANRE	843.490,10	100,00%	2	2	2	0,01%	0,34%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=utrn PE=1 SV=: A0A0H2UKN4_DANRE	299.775,50	99,20%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=myom1b PE=4 S A0A0R4IGQ8_DANRE	180.977,20	100,00%	1	1	1	0,00%	0,68%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=fga PE=1 SV=: B8A5L6_DANRE,E9QCD1_	49.680,00	100,00%	12	15	37	0,12%	19,20%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=fga PE=1 SV=: B8A5L6_DANRE,E9QCD1_	49.680,00	100,00%	12	15	37	0,12%	19,20%
14dpa_1	Fgg protein OS=Danio rerio GN=fgg PE=1 SV=: Q7ZV7G_DANRE	48.833,70	100,00%	17	23	106	0,34%	59,20%
14dpa_1	Fibrinogen, B beta polypeptide OS=Danio rerio GN=fgb PE=1 Q6NYE1_DANRE	54.407,60	100,00%	15	24	104	0,33%	49,50%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 S E7EZ79_DANRE	142.417,20	100,00%	9	9	14	0,04%	10,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=crip2 PE=1 SV=: F1QZB0_DANRE,Q5BKV7_	9.470,50	11,00%	0	0	0	0,00%	0,00%

14dpa_1	Uncharacterized protein OS=Danio rerio GN=col1a1a PE=1 SV=1 F1QJ9C_DANRE,F1QSC5_I	128.043,70	100,00%	6	7	103	0,33%	4,35%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=actn2b PE=1 SV=1 E9QFR8_DANRE	103.214,90	100,00%	7	8	15	0,05%	12,50%
14dpa_1	Dihydropyridyl dehydrogenase OS=Danio rerio GN=dldh PE=1 Q6TNU6_DANRE	53.620,50	99,90%	1	1	2	0,01%	2,56%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=vwf PE=4 SV=2 E7F921_DANRE,F1QHS6_I	225.634,30	86,70%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=vwf PE=4 SV=2 E7F921_DANRE,F1QHS6_I	225.634,30	86,70%	0	0	0	0,00%	0,00%
14dpa_1	Lamin B2 OS=Danio rerio GN=lmbn2 PE=1 SV=1 B3DFN3_DANRE	65.934,60	99,90%	1	1	1	0,00%	3,09%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=palm2 PE=4 SV= A0A0R4ISM7_DANRE,E7F1	92.198,80	100,00%	3	3	4	0,01%	4,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=palm2 PE=4 SV= A0A0R4ISM7_DANRE,E7F1	92.198,80	100,00%	3	3	4	0,01%	4,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ccdc141 PE=4 SV= B0S755_DANRE,E7FG99_I	190.652,30	98,70%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ccdc141 PE=4 SV= B0S755_DANRE,E7FG99_I	190.652,30	98,70%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ank3b PE=1 SV= F1QN83_DANRE	218.497,50	100,00%	1	1	1	0,00%	0,65%
14dpa_1	Aspartate aminotransferase OS=Danio rerio GN=got2a PE=1 F1QCDA_DANRE,Q75YK7_	47.588,10	20,60%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=tpm4a PE=1 SV= F1QK7G_DANRE	32.726,90	100,00%	4	4	5	0,02%	19,00%
14dpa_1	Heat shock cognate 71 kDa protein OS=Danio rerio GN=hspa1 HSP7C_DANRE	70.975,60	99,70%	1	1	1	0,00%	1,54%
14dpa_1	Cytochrome c oxidase subunit 2 OS=Danio rerio GN=COX2 PE A0A0A0VG08_DANRE,CO	25.985,80	100,00%	1	1	4	0,01%	6,96%
14dpa_1	Cytochrome c oxidase subunit 2 OS=Danio rerio GN=COX2 PE A0A0A0VG08_DANRE,CO	25.985,80	100,00%	1	1	4	0,01%	6,96%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ablim1b PE=4 SV= A0A0R4ITJ6_DANRE,B8A6	84.999,80	94,60%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ablim1b PE=4 SV= A0A0R4ITJ6_DANRE,B8A6	84.999,80	94,60%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ablim1b PE=4 SV= A0A0R4ITJ6_DANRE,B8A6	84.999,80	94,60%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ablim1b PE=4 SV= A0A0R4ITJ6_DANRE,B8A6	84.999,80	94,60%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=tnl1 PE=4 SV=1 A0A0R4IDZ8_DANRE,A0A1	271.258,30	100,00%	4	4	9	0,03%	2,75%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=tnl1 PE=4 SV=1 A0A0R4IDZ8_DANRE,A0A1	271.258,30	100,00%	4	4	9	0,03%	2,75%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=tnl1 PE=4 SV=1 A0A0R4IDZ8_DANRE,A0A1	271.258,30	100,00%	4	4	9	0,03%	2,75%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=tnl2a PE=4 SV=1 A0A0R4IIA7_DANRE,A0AC	271.774,00	99,70%	1	1	1	0,00%	1,77%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=tnl2a PE=4 SV=1 A0A0R4IIA7_DANRE,A0AC	271.774,00	99,70%	1	1	1	0,00%	1,77%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=uqcrc2a PE=4 SV= A0A0R4IAK9_DANRE,A0A1	49.189,50	0,70%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=flna PE=1 SV=1 E9QI62_DANRE,F1R9P5_I	272.475,80	100,00%	3	3	3	0,01%	1,76%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=flna PE=1 SV=1 E9QI62_DANRE,F1R9P5_I	272.475,80	100,00%	3	3	3	0,01%	1,76%
14dpa_1	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IGI1_DANRE,G3P_	35.784,00	100,00%	3	3	5	0,02%	15,50%
14dpa_1	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IGI1_DANRE,G3P_	35.784,00	100,00%	3	3	5	0,02%	15,50%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=flnca PE=4 SV=1 A0A0R4IU54_DANRE,E7F1	291.642,80	100,00%	1	1	1	0,00%	0,30%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=flnca PE=4 SV=1 A0A0R4IU54_DANRE,E7F1	291.642,80	100,00%	1	1	1	0,00%	0,30%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=eef1a1a PE=4 SV= A0A0R4IE10_DANRE,A0A1	50.339,50	32,40%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=eef1a1a PE=4 SV= A0A0R4IE10_DANRE,A0A1	50.339,50	32,40%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=eef1a1a PE=4 SV= A0A0R4IE10_DANRE,A0A1	50.339,50	32,40%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=eef1a1a PE=4 SV= A0A0R4IE10_DANRE,A0A1	50.339,50	32,40%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cox6c PE=1 SV= Q6ZM23_DANRE	8.480,50	52,50%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=pdha1a PE=4 SV= A0A0R4IT09_DANRE,Q6P1	43.741,90	97,80%	1	1	2	0,01%	2,25%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=pdha1a PE=4 SV= A0A0R4IT09_DANRE,Q6P1	43.741,90	97,80%	1	1	2	0,01%	2,25%
14dpa_1	Serotransferrin OS=Danio rerio GN=fta PE=1 SV=1 B8JL43_DANRE,F1R858_D	73.717,60	100,00%	2	2	2	0,01%	4,46%
14dpa_1	Serotransferrin OS=Danio rerio GN=fta PE=1 SV=1 B8JL43_DANRE,F1R858_D	73.717,60	100,00%	2	2	2	0,01%	4,46%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=slc2 E9QJ49_DANRE,F1QAJ7_I	39.378,90	94,10%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=slc2 E9QJ49_DANRE,F1QAJ7_I	39.378,90	94,10%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=lamb1b PE=1 SV= F1REL9_DANRE	197.673,00	97,70%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=sorbs2a PE=1 SV= F1QI09_DANRE	148.129,10	100,00%	1	1	3	0,01%	1,35%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=fn1a PE=1 SV=1 B0S602_DANRE	271.953,50	100,00%	1	1	1	0,00%	1,69%
14dpa_1	Tubulin alpha 8-like 3a protein OS=Danio rerio GN=tuba8l3 P Q6DC49_DANRE	50.066,90	99,80%	1	1	2	0,01%	4,44%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:56235 PE=1 Q7VZ39_DANRE	30.045,40	100,00%	1	1	3	0,01%	8,19%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=lama4 PE=1 SV= F1QYU0_DANRE	207.483,80	52,70%	0	0	0	0,00%	0,00%
14dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	1	1	1	0,00%	12,10%
14dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	1	1	1	0,00%	12,10%
14dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	1	1	1	0,00%	12,10%
14dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	1	1	1	0,00%	12,10%
14dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	1	1	1	0,00%	12,10%
14dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	1	1	1	0,00%	12,10%
14dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	1	1	1	0,00%	12,10%
14dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	1	1	1	0,00%	12,10%
14dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	1	1	1	0,00%	12,10%
14dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	1	1	1	0,00%	12,10%
14dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	1	1	1	0,00%	12,10%
14dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV= A0A0N4STN6_DANRE,A7H	13.590,30	100,00%	1	1	1	0,00%	12,10%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=hk1 PE=4 SV=1 A0A0R4IUT7_DANRE,Q6N1	102.760,50	100,00%	2	2	2	0,01%	2,39%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=hk1 PE=4 SV=1 A0A0R4IUT7_DANRE,Q6N1	102.760,50	100,00%	2	2	2	0,01%	2,39%
14dpa_1	Pyruvate kinase OS=Danio rerio GN=pkma PE=1 SV=1 Q7VZ72_DANRE	58.059,00	99,80%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=acs11b PE=1 SV= F1QDA6_DANRE	77.694,10	98,90%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=emilin1a PE=1 SV= F1QC17_DANRE,Q5T263_	112.771,70	100,00%	2	2	3	0,01%	1,97%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=emilin1a PE=1 SV= F1QC17_DANRE,Q5T263_	112.771,70	100,00%	2	2	3	0,01%	1,97%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=lmo7a PE=4 SV= A0A0R4ITT8_DANRE	156.646,40	93,90%	1	1	1	0,00%	1,24%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=myoz2b PE=4 SV= F1QBP1_DANRE,F1QKE7_	25.288,50	97,80%	1	1	1	0,00%	5,06%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=myoz2b PE=4 SV= F1QBP1_DANRE,F1QKE7_	25.288,50	97,80%	1	1	1	0,00%	5,06%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=popdc2 PE=1 SV= F8W5H2_DANRE,Q6JVV8	41.034,60	99,20%	1	1	2	0,01%	4,09%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=popdc2 PE=1 SV= F8W5H2_DANRE,Q6JVV8	41.034,60	99,20%	1	1	2	0,01%	4,09%
14dpa_1	Keratin, type I cytoskeletal 18 OS=Danio rerio GN=krt18 PE=1 K1C18_DANRE	48.631,90	100,00%	1	1	1	0,00%	2,78%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=vdac B0R198_DANRE	29.812,00	99,30%	1	1	2	0,01%	6,81%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=eno: F1QBW7_DANRE	51.320,60	100,00%	1	1	1	0,00%	4,28%
14dpa_1	Aconitate hydratase, mitochondrial (Fragment) OS=Danio rer F8W4M7_DANRE	85.591,90	100,00%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE= A0A0R4IE46_DANRE,A0A1	101.943,10	99,10%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE= A0A0R4IE46_DANRE,A0A1	101.943,10	99,10%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE= A0A0R4IE46_DANRE,A0A1	101.943,10	99,10%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=col4a2 PE=1 SV= A0A0H2UKP2_DANRE	165.476,80	100,00%	2	2	4	0,01%	1,38%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=col4 F1Q855_DANRE,F1QG79_	128.554,50	99,80%	1	2	8	0,03%	1,96%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=col4 F1Q855_DANRE,F1QG79_	128.554,50	99,80%	1	2	8	0,03%	1,96%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV= A0A0R4IH07_DANRE,B6IC	347.704,80	100,00%	1	1	1	0,00%	0,35%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV= A0A0R4IH07_DANRE,B6IC	347.704,80	100,00%	1	1	1	0,00%	0,35%
14dpa_1	Catenin (Cadherin-associated protein), alpha OS=Danio rerio A0A0R4IF47_DANRE,Q9P1	100.520,10	100,00%	2	2	2	0,01%	4,08%
14dpa_1	Catenin (Cadherin-associated protein), alpha OS=Danio rerio A0A0R4IF47_DANRE,Q9P1	100.520,10	100,00%	2	2	2	0,01%	4,08%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=vdac2 PE=1 SV= Q8AWD0_DANRE	30.284,90	97,80%	1	1	11	0,04%	6,71%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cmlc1 PE=1 SV= B0R0F7_DANRE	21.865,20	100,00%	6	7	19	0,06%	48,00%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=post F1QM50_DANRE	17.277,20	52,70%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=aifm1 PE=1 SV= A2BG5E_DANRE,F1RBU0_	66.977,00	100,00%	2	2	2	0,01%	2,54%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=aifm1 PE=1 SV= A2BG5E_DANRE,F1RBU0_	66.977,00	100,00%	2	2	2	0,01%	2,54%

14dpa_1	Uncharacterized protein OS=Danio rerio GN=afim1 PE=1 SV=-: A2BGE5_DANRE,F1RBU0_	66.977,00	100,00%	2	2	2	0,01%	2,54%
14dpa_1	Aspartate aminotransferase OS=Danio rerio GN=got2a PE=1 !: F1QCDD_DANRE,Q7SYK7_	47.588,10	20,60%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=uqrc2a PE=4 S\ A0A0R4IAK9_DANRE,A0AI	49.189,50	97,70%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=uqrc2a PE=4 S\ A0A0R4IAK9_DANRE,A0AI	49.189,50	97,70%	0	0	0	0,00%	0,00%
14dpa_1	Cox5a protein OS=Danio rerio GN=cox5a PE=1 SV=1 Q4VBU7_DANRE	15.897,10	91,10%	0	0	0	0,00%	0,00%
14dpa_1	Phosphoglycerate kinase OS=Danio rerio GN=pgk1 PE=1 SV=-: F1QXV8_DANRE	44.733,20	22,30%	0	0	0	0,00%	0,00%
14dpa_1	Keratin, type II cytoskeletal 8 OS=Danio rerio GN=kr8 PE=1 S K2C8_DANRE	57.761,30	100,00%	6	7	10	0,03%	12,10%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=vtnb PE=1 SV=1 F1Q5D8_DANRE	51.527,70	96,00%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=eef1a1a PE=4 S\ A0A0R4IE10_DANRE,A0AI	50.339,50	32,40%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=eef1a1a PE=4 S\ A0A0R4IE10_DANRE,A0AI	50.339,50	32,40%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=eef1a1a PE=4 S\ A0A0R4IE10_DANRE,A0AI	50.339,50	32,40%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=eef1a1a PE=4 S\ A0A0R4IE10_DANRE,A0AI	50.339,50	32,40%	0	0	0	0,00%	0,00%
14dpa_1	Malate dehydrogenase OS=Danio rerio GN=mdh2 PE=1 SV=1 Q7T334_DANRE	35.420,30	100,00%	3	3	10	0,03%	12,50%
14dpa_1	Caveolin OS=Danio rerio GN=cav1 PE=1 SV=1 Q6YLH8_DANRE,Q6YLH9_	20.652,80	68,40%	0	0	0	0,00%	0,00%
14dpa_1	Caveolin OS=Danio rerio GN=cav1 PE=1 SV=1 Q6YLH8_DANRE,Q6YLH9_	20.652,80	68,40%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cox5b2 PE=1 SV Q0P3U4_DANRE	14.308,50	95,30%	0	0	0	0,00%	0,00%
14dpa_1	Hadha protein OS=Danio rerio GN=hadhaa PE=1 SV=1 A7YT47_DANRE	81.877,10	100,00%	1	1	1	0,00%	1,84%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=vdac1 PE=1 SV= Q6NWC1_DANRE	30.627,50	100,00%	2	2	4	0,01%	14,80%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=nid1a PE=1 SV=-: F1RAG3_DANRE	130.098,40	99,70%	1	1	1	0,00%	1,45%
14dpa_1	Ckmt2 protein OS=Danio rerio GN=ckmt2b PE=2 SV=1 A8E5J9_DANRE,F6NMS7_	46.281,40	86,00%	1	1	1	0,00%	9,20%
14dpa_1	Ckmt2 protein OS=Danio rerio GN=ckmt2b PE=2 SV=1 A8E5J9_DANRE,F6NMS7_	46.281,40	86,00%	1	1	1	0,00%	9,20%
14dpa_1	Radixin OS=Danio rerio GN=msna PE=1 SV=1 Q6GI42_DANRE	68.035,00	100,00%	3	3	3	0,01%	5,35%
14dpa_1	Blood vessel epicardial substance OS=Danio rerio GN=bves P POPD1_DANRE	40.341,40	91,50%	1	1	1	0,00%	5,11%
14dpa_1	ATP synthase subunit beta OS=Danio rerio GN=atp5b PE=1 S\ A8WGC6_DANRE	55.130,90	100,00%	4	4	14	0,04%	14,30%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=c9 PE=1 SV=1 B8A565_DANRE	72.713,70	97,80%	1	1	1	0,00%	3,12%
14dpa_1	Heat shock protein 1 OS=Danio rerio GN=hspb1 PE=1 SV=1 A4VAK4_DANRE	22.408,90	99,50%	1	2	2	0,01%	4,52%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:65851 PE=1 Q7S2P4_DANRE	64.751,30	95,60%	1	1	1	0,00%	1,60%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1 A0A0R4IEB4_DANRE,A0AI	128.154,40	98,40%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1 A0A0R4IEB4_DANRE,A0AI	128.154,40	98,40%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1 A0A0R4IEB4_DANRE,A0AI	128.154,40	98,40%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cilp2 PE=4 SV=1 A0A0R4IEB4_DANRE,A0AI	128.154,40	98,40%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=anxa11b PE=4 S\ F8W2N0_DANRE,Q804G3	51.462,90	89,90%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=anxa11b PE=4 S\ F8W2N0_DANRE,Q804G3	51.462,90	89,90%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	99,90%	1	2	6	0,02%	2,31%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	99,90%	1	2	6	0,02%	2,31%
14dpa_1	Histone H4 OS=Danio rerio GN=si:dkey-108K21.23 PE=1 SV=1 A3KPR4_DANRE,X1WGM:	11.478,70	99,80%	1	1	1	0,00%	9,71%
14dpa_1	Histone H4 OS=Danio rerio GN=si:dkey-108K21.23 PE=1 SV=1 A3KPR4_DANRE,X1WGM:	11.478,70	99,80%	1	1	1	0,00%	9,71%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=mfpap5 PE=4 SV: E9QC4_DANRE	14.664,20	98,90%	1	1	2	0,01%	11,90%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=aldh6a1 PE=1 S\ Q6DHT4_DANRE	57.120,40	77,80%	0	0	0	0,00%	0,00%
14dpa_1	Enolase 1, (Alpha) OS=Danio rerio GN=eno1a PE=1 SV=1 Q6PC12_DANRE	47.075,10	100,00%	1	1	1	0,00%	2,31%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=jupa PE=1 SV=1 F1QTT3_DANRE	80.020,20	100,00%	2	2	2	0,01%	4,12%
14dpa_1	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1 B3DG37_DANRE,HBB1_D_	16.389,00	90,80%	1	1	1	0,00%	10,80%
14dpa_1	Hemoglobin subunit beta-2 OS=Danio rerio GN=ba2 PE=1 SV: HBB2_DANRE	16.389,20	73,90%	0	0	0	0,00%	0,00%
14dpa_1	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1 B3DG37_DANRE,HBB1_D_	16.389,00	90,80%	1	1	1	0,00%	10,80%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4IPE6_DANRE,E9QI	29.642,20	100,00%	2	2	2	0,01%	4,71%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4IPE6_DANRE,E9QI	29.642,20	100,00%	2	2	2	0,01%	4,71%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4IPE6_DANRE,E9QI	29.642,20	100,00%	2	2	2	0,01%	4,71%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4IPE6_DANRE,E9QI	29.642,20	100,00%	2	2	2	0,01%	4,71%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4IPE6_DANRE,E9QI	29.642,20	100,00%	2	2	2	0,01%	4,71%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4IPE6_DANRE,E9QI	29.642,20	100,00%	2	2	2	0,01%	4,71%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cast PE=1 SV=1 A0A0R4IPE6_DANRE,E9QI	29.642,20	100,00%	2	2	2	0,01%	4,71%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=atp5b PE=1 SV: A2AR67_DANRE	8.316,90	93,50%	1	1	1	0,00%	16,90%
14dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=pcc: F1QP17_DANRE	79.153,60	93,40%	0	0	0	0,00%	0,00%
14dpa_1	Isocitrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=1 A0A0G2L6I6_DANRE,E7F4	50.921,70	99,20%	1	1	1	0,00%	2,89%
14dpa_1	Isocitrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=1 A0A0G2L6I6_DANRE,E7F4	50.921,70	99,20%	1	1	1	0,00%	2,89%
14dpa_1	Hbaa1 protein OS=Danio rerio GN=hbaa1 PE=1 SV=1 HBA_DANRE,Q1JQ69_DAI	15.565,70	58,60%	0	0	0	0,00%	0,00%
14dpa_1	Hbaa1 protein OS=Danio rerio GN=hbaa1 PE=1 SV=1 HBA_DANRE,Q1JQ69_DAI	15.565,70	58,60%	0	0	0	0,00%	0,00%
14dpa_1	Hbaa1 protein OS=Danio rerio GN=hbaa1 PE=1 SV=1 HBA_DANRE,Q1JQ69_DAI	15.565,70	58,60%	0	0	0	0,00%	0,00%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=cox4i1 PE=1 SV F1R6I1_DANRE	20.331,60	97,80%	1	1	2	0,01%	6,94%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	99,90%	1	2	6	0,02%	2,31%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	99,90%	1	2	6	0,02%	2,31%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ptx3a PE=1 SV: A3KPN4_DANRE	49.273,20	99,00%	1	1	1	0,00%	2,91%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=prdx3 PE=1 SV: A8E7G6_DANRE,A8E7G7_	25.875,70	97,50%	1	1	1	0,00%	4,98%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=prdx3 PE=1 SV: A8E7G6_DANRE,A8E7G7_	25.875,70	97,50%	1	1	1	0,00%	4,98%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ndrg4 PE=4 SV: A0A0R4ICQ1_DANRE,A6X	37.210,30	98,90%	1	1	1	0,00%	6,53%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=ndrg4 PE=4 SV: A0A0R4ICQ1_DANRE,A6X	37.210,30	98,90%	1	1	1	0,00%	6,53%
14dpa_1	Ceruloplasmin OS=Danio rerio GN=cp PE=1 SV=1 Q6P3G1_DANRE	124.388,00	88,50%	1	1	1	0,00%	1,20%
14dpa_1	Myoglobin OS=Danio rerio GN=mb PE=1 SV=1 H0WE38_DANRE,MYG_D_	15.583,60	100,00%	2	2	25	0,08%	31,10%
14dpa_1	Myoglobin OS=Danio rerio GN=mb PE=1 SV=1 H0WE38_DANRE,MYG_D_	15.583,60	100,00%	2	2	25	0,08%	31,10%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=si:dkeyp-80C12. H0WE38_DANRE	6.338,70	97,80%	1	1	2	0,01%	27,60%
14dpa_1	Hydroxyacyl-Coenzyme A dehydrogenase OS=Danio rerio GN=Q6DI22_DANRE	33.336,90	100,00%	1	1	1	0,00%	5,50%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=tnni1b PE=1 SV: E7F5J9_DANRE,F8W417_I	21.032,10	8,50%	0	0	0	0,00%	0,00%
14dpa_1	Pyruvate dehydrogenase (Lipoamide) beta OS=Danio rerio GI Q7T368_DANRE	39.308,70	99,80%	1	1	1	0,00%	4,46%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	97,80%	1	2	2	0,01%	6,81%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	97,80%	1	2	2	0,01%	6,81%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	97,80%	1	2	2	0,01%	6,81%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	97,80%	1	2	2	0,01%	6,81%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	97,80%	1	2	2	0,01%	6,81%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	97,80%	1	2	2	0,01%	6,81%
14dpa_1	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	97,80%	1	2	2	0,01%	6,81%
14dpa_1	Acetyl-CoA acetyltransferase, mitochondrial OS=Danio rerio (A0A0R4I76_DANRE,THIL_	44.342,90	97,50%	1	1	1	0,00%	6,43%
14dpa_1	Acetyl-CoA acetyltransferase, mitochondrial OS=Danio rerio (A0A0R4I76_DANRE,THIL_	44.342,90	97,50%	1	1	1	0,00%	6,43%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=ccd141 PE=4 S\ B0S755_DANRE,E7FG99_I	190.652,30	96,70%	1	1	1	0,00%	0,78%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=ccd141 PE=4 S\ B0S755_DANRE,E7FG99_I	190.652,30	96,70%	1	1	1	0,00%	0,78%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=col6a3 PE=1 SV: F1Q4K1_DANRE	309.287,80	100,00%	18	20	42	0,14%	9,46%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=fga PE=1 SV=1 B8A5L6_DANRE,E9QCD1_	49.680,00	100,00%	13	16	31	0,11%	24,10%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=fga PE=1 SV=1 B8A5L6_DANRE,E9QCD1_	49.680,00	100,00%	13	16	31	0,11%	24,10%
14dpa_2	Actin, cytoplasmic 2 OS=Danio rerio GN=actbb PE=2 SV=2 A8WG05_DANRE,ACTB1_	41.753,80	100,00%	2	2	4	0,01%	17,90%
14dpa_2	Actin, cytoplasmic 2 OS=Danio rerio GN=actbb PE=2 SV=2 A8WG05_DANRE,ACTB1_	41.753,80	100,00%	2	2	4	0,01%	17,90%
14dpa_2	Actin, alpha, cardiac muscle 1a OS=Danio rerio GN=si:ch73-1: F1QUH8_DANRE,F8W3L1_	41.976,10	100,00%	7	12	26	0,09%	34,20%
14dpa_2	Actin, alpha, cardiac muscle 1a OS=Danio rerio GN=si:ch73-1: F1QUH8_DANRE,F8W3L1_	41.976,10	100,00%	7	12	26	0,09%	34,20%
14dpa_2	Actin, cytoplasmic 2 OS=Danio rerio GN=actbb PE=2 SV=2 A8WG05_DANRE,ACTB1_	41.753,80	100,00%	2	2	4	0,01%	17,90%

14dpa_2	Actin, alpha, cardiac muscle 1a OS=Danio rerio GN=si:ch73-1 F1QUN8_DANRE,F8W3L1,	41.976,10	100,00%	7	12	26	0,09%	34,20%
14dpa_2	Actin, alpha, cardiac muscle 1a OS=Danio rerio GN=si:ch73-1 F1QUN8_DANRE,F8W3L1,	41.976,10	100,00%	7	12	26	0,09%	34,20%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=zgc:86725 PE=4 A0A0R4IBT5_DANRE,F1RC	41.976,90	98,40%	1	1	1	0,00%	20,60%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=zgc:86725 PE=4 A0A0R4IBT5_DANRE,F1RC	41.976,90	98,40%	1	1	1	0,00%	20,60%
14dpa_2	Actin, alpha, cardiac muscle 1a OS=Danio rerio GN=si:ch73-1 F1QUN8_DANRE,F8W3L1,	41.976,10	100,00%	7	12	26	0,09%	34,20%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=ttna PE=1 SV=2 F1R7N8_DANRE	3.089.806,00	100,00%	13	13	13	0,04%	0,63%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=1 A0A0R4INC4_DANRE,ASV	543.007,50	96,90%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=1 A0A0R4INC4_DANRE,ASV	543.007,50	96,90%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=1 A0A0R4INC4_DANRE,ASV	543.007,50	96,90%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=vmhcl PE=1 SV= F1QSE1_DANRE	223.380,90	100,00%	63	83	169	0,57%	36,60%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttbn G1K2X0_DANRE	2.458.846,80	100,00%	10	10	10	0,03%	0,73%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=fn1b PE=1 SV=2 A2CEW3_DANRE,F1R809_	265.661,60	100,00%	46	58	116	0,39%	31,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=fn1b PE=1 SV=2 A2CEW3_DANRE,F1R809_	265.661,60	100,00%	46	58	116	0,39%	31,00%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=col5 F6NPA4_DANRE	198.988,80	100,00%	4	4	5	0,02%	3,63%
14dpa_2	Fibrinogen, B beta polypeptide OS=Danio rerio GN=fgb PE=1 Q6NVE1_DANRE	54.407,60	100,00%	14	23	66	0,22%	37,70%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=sptb PE=1 SV=1 F1QCE5_DANRE	272.829,20	100,00%	6	6	6	0,02%	3,69%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=tnnt2a PE=1 SV: E7EY9C9_DANRE,F1R0K6_I	33.942,20	100,00%	3	3	3	0,01%	11,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=tnnt2a PE=1 SV: E7EY9C9_DANRE,F1R0K6_I	33.942,20	100,00%	3	3	3	0,01%	11,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=tnnt2a PE=1 SV: E7EY9C9_DANRE,F1R0K6_I	33.942,20	100,00%	3	3	3	0,01%	11,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=tnnt2a PE=1 SV: E7EY9C9_DANRE,F1R0K6_I	33.942,20	100,00%	3	3	3	0,01%	11,00%
14dpa_2	Fgg protein OS=Danio rerio GN=fgg PE=1 SV=1 Q7ZV77_DANRE	48.833,70	100,00%	15	22	83	0,28%	41,50%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 SV: E7EZ79_DANRE,F1Q615_I	142.109,70	100,00%	9	9	10	0,03%	9,40%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 SV: E7EZ79_DANRE,F1Q615_I	142.109,70	100,00%	9	9	10	0,03%	9,40%
14dpa_2	Blood vessel epicardial substance OS=Danio rerio GN=bves P POPD1_DANRE	40.341,40	97,00%	1	1	2	0,01%	5,11%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=sorbs1 PE=4 SV: A0A0R4IP30_DANRE	85.938,80	98,50%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=postna PE=1 SV: E7F5I3_DANRE	97.135,00	100,00%	6	6	6	0,02%	12,40%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=emilin1b PE=1 S F1Q9G3_DANRE,F8W4C7_	116.801,40	100,00%	6	6	9	0,03%	6,23%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=emilin1b PE=1 S F1Q9G3_DANRE,F8W4C7_	116.801,40	100,00%	6	6	9	0,03%	6,23%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a2 PE=1 SV: A0A0R4IHCO_DANRE,ESQ	127.211,90	100,00%	11	15	37	0,13%	11,50%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a2 PE=1 SV: A0A0R4IHCO_DANRE,ESQ	127.211,90	100,00%	11	15	37	0,13%	11,50%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a2 PE=1 SV: A0A0R4IHCO_DANRE,ESQ	127.211,90	100,00%	11	15	37	0,13%	11,50%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=kr94 PE=1 SV: F1R8U0_DANRE,Q1RLR3_	50.932,10	100,00%	1	1	1	0,00%	2,09%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=kr94 PE=1 SV: F1R8U0_DANRE,Q1RLR3_	50.932,10	100,00%	1	1	1	0,00%	2,09%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=ank3b PE=1 SV= F1QN3B_DANRE	218.497,50	59,90%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a1a PE=1 SV: F1QJ9C9_DANRE,F1Q5CS_I	128.043,70	100,00%	7	8	11	0,04%	7,26%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a1a PE=1 SV: F1QJ9C9_DANRE,F1Q5CS_I	128.043,70	100,00%	7	8	11	0,04%	7,26%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=cox4i1 PE=1 SV: F1R6I1_DANRE	20.331,60	100,00%	1	1	1	0,00%	5,78%
14dpa_2	ATP synthase subunit alpha OS=Danio rerio GN=atp5a1 PE=1 Q08B41_DANRE	59.745,30	100,00%	6	6	12	0,04%	13,20%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=col6a2 PE=4 SV: E7FCV8_DANRE	106.916,20	100,00%	9	9	12	0,04%	10,50%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=mfp5 PE=4 SV: E9QC4_DANRE	14.664,20	98,00%	1	1	1	0,00%	11,90%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=sorbs2a PE=1 SV: F1QI09_DANRE	148.129,10	99,70%	1	1	1	0,00%	1,50%
14dpa_2	Solute carrier family 25 alpha, member 5 OS=Danio rerio GN: Q8JH10_DANRE	32.764,40	100,00%	1	2	4	0,01%	3,02%
14dpa_2	Enolase 1, (Alpha) OS=Danio rerio GN=eno1a PE=1 SV=1 Q6PC12_DANRE	47.075,10	98,10%	1	1	1	0,00%	4,40%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=eno F1QBW7_DANRE	51.320,60	100,00%	2	3	3	0,01%	10,10%
14dpa_2	Aconitate hydratase, mitochondrial (Fragment) OS=Danio rer F8W4M7_DANRE	85.591,90	100,00%	2	2	2	0,01%	3,80%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	90,80%	1	1	1	0,00%	3,83%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	90,80%	1	1	1	0,00%	3,83%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	90,80%	1	1	1	0,00%	3,83%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	90,80%	1	1	1	0,00%	3,83%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	90,80%	1	1	1	0,00%	3,83%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	90,80%	1	1	1	0,00%	3,83%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=si:ch211-202a1: A0A0N4SUD2_DANRE,B0F	17.999,60	90,80%	1	1	1	0,00%	3,83%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=col6 F1Q924_DANRE	278.174,40	100,00%	11	13	19	0,06%	6,88%
14dpa_2	Cox5aa protein OS=Danio rerio GN=cox5aa PE=1 SV=1 Q4VBU7_DANRE	15.897,10	94,40%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=dspa PE=1 SV=2 F1RBA4_DANRE,F8W4Q1_	235.150,30	16,50%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=dspa PE=1 SV=2 F1RBA4_DANRE,F8W4Q1_	235.150,30	16,50%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=tln2a PE=4 SV=1 A0A0R4IIA7_DANRE	271.364,80	100,00%	2	2	2	0,01%	1,53%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=lama4 PE=1 SV= F1QYU0_DANRE	207.483,80	8,20%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=fn1a PE=1 SV=1 B0S602_DANRE,F1RBP8_I	271.523,90	100,00%	2	2	2	0,01%	1,78%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=fn1a PE=1 SV=1 B0S602_DANRE,F1RBP8_I	271.523,90	100,00%	2	2	2	0,01%	1,78%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=ahnak PE=1 SV= F1QZ50_DANRE	685.184,90	100,00%	10	10	11	0,04%	2,07%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=vtg1 PE=1 SV=1 Q1LWN2_DANRE	149.238,80	100,00%	1	1	1	0,00%	1,32%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a1b PE=1 SV: F1QD11_DANRE	136.914,60	100,00%	9	10	17	0,06%	8,07%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=sptbn1 PE=1 SV: A0A0G2L1F5_DANRE	262.630,10	100,00%	5	5	8	0,03%	3,17%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=vwf PE=4 SV=2 E7F921_DANRE,F1QHS6_I	225.634,30	99,90%	1	1	1	0,00%	0,46%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=vwf PE=4 SV=2 E7F921_DANRE,F1QHS6_I	225.634,30	99,90%	1	1	1	0,00%	0,46%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:cf F1Q786_DANRE	147.365,60	15,70%	0	0	0	0,00%	0,00%
14dpa_2	Keratin, type II cytoskeletal 8 OS=Danio rerio GN=kr8 PE=1 S K2C8_DANRE	57.761,30	100,00%	3	3	3	0,01%	5,96%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=spna2 PE=4 SV= A0A0R4IU08_DANRE	289.251,20	100,00%	14	14	15	0,05%	9,34%
14dpa_2	Keratin, type I cytoskeletal 18 OS=Danio rerio GN=kr18 PE=1 K1C18_DANRE	48.631,90	97,00%	1	1	1	0,00%	3,71%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE=: A0A0R4IE46_DANRE,A0AI	101.943,10	100,00%	2	2	3	0,01%	2,16%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE=: A0A0R4IE46_DANRE,A0AI	101.943,10	100,00%	2	2	3	0,01%	2,16%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE=: A0A0R4IE46_DANRE,A0AI	101.943,10	100,00%	2	2	3	0,01%	2,16%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttna B0S758_DANRE	843.490,10	100,00%	0	0	0	0,00%	0,49%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=hspg2 PE=4 SV= A0A0R4I2A6_DANRE,F1R5	369.970,80	100,00%	19	21	30	0,10%	8,78%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=hspg2 PE=4 SV= A0A0R4I2A6_DANRE,F1R5	369.970,80	100,00%	19	21	30	0,10%	8,78%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4IU71_DANRE,E7F4	518.896,50	100,00%	3	3	3	0,01%	0,98%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4IU71_DANRE,E7F4	518.896,50	100,00%	3	3	3	0,01%	0,98%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4IU71_DANRE,E7F4	518.896,50	100,00%	3	3	3	0,01%	0,98%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=palm1b PE=4 SV B0V0Y4_DANRE	27.123,40	78,90%	1	1	1	0,00%	6,56%
14dpa_2	Myoglobin OS=Danio rerio GN=mb PE=2 SV=3 MYG_DANRE	15.583,60	100,00%	1	1	2	0,01%	9,52%
14dpa_2	Radixin OS=Danio rerio GN=msna PE=1 SV=1 Q6GI42_DANRE	68.035,00	98,90%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=dlst PE=1 SV=1 F1QFQ6_DANRE,F1QY42_	25.587,50	44,50%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=dlst PE=1 SV=1 F1QFQ6_DANRE,F1QY42_	25.587,50	44,50%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=post F1QM50_DANRE	17.277,20	100,00%	2	2	2	0,01%	20,90%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=pkma PE=4 SV=: A0A0R4IGP6_DANRE,Q7Z	58.059,00	99,80%	1	1	1	0,00%	1,50%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=pkma PE=4 SV=: A0A0R4IGP6_DANRE,Q7Z	58.059,00	99,80%	1	1	1	0,00%	1,50%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	98,50%	1	1	1	0,00%	1,28%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	98,50%	1	1	1	0,00%	1,28%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	98,50%	1	1	1	0,00%	1,28%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	98,50%	1	1	1	0,00%	1,28%
14dpa_2	Pyruvate dehydrogenase (Lipoamide) beta OS=Danio rerio GI Q7T368_DANRE	39.308,70	100,00%	1	1	1	0,00%	3,06%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttbn B0S6Y0_DANRE	626.764,10	99,50%	1	1	1	0,00%	0,21%

14dpa_2	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1	B3DG37_DANRE,HBB1_D	16.389,20	100,00%	1	1	2	0,01%	8,78%
14dpa_2	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1	B3DG37_DANRE,HBB1_D	16.389,20	100,00%	1	1	2	0,01%	8,78%
14dpa_2	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1	B3DG37_DANRE,HBB1_D	16.389,20	100,00%	1	1	2	0,01%	8,78%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=hk1 PE=4 SV=1	A0A0R4IUT7_DANRE,Q6N	102.760,50	98,70%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=hk1 PE=4 SV=1	A0A0R4IUT7_DANRE,Q6N	102.760,50	98,70%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=c3a.1 PE=1 SV=1	B8JKW4_DANRE	183.235,60	100,00%	4	4	7	0,02%	3,53%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=slc2 E9QJ49_DANRE,F1QAJ7_I	E9QJ49_DANRE,F1QAJ7_I	39.378,90	29,50%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=slc2 E9QJ49_DANRE,F1QAJ7_I	E9QJ49_DANRE,F1QAJ7_I	39.378,90	29,50%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=col6 F6NRG0_DANRE	88.827,90	99,70%	1	1	2	0,01%	7,04%	
14dpa_2	Uncharacterized protein OS=Danio rerio GN=ptx3a PE=1 SV=1	A3KPN4_DANRE	49.273,20	60,90%	0	0	0	0,00%	0,00%
14dpa_2	Calcium-transporting ATPase OS=Danio rerio GN=atp2a2a PE A9C3Q4_DANRE,A9C3Q5_	A9C3Q4_DANRE,A9C3Q5_	109.519,40	100,00%	2	2	2	0,01%	3,26%
14dpa_2	Calcium-transporting ATPase OS=Danio rerio GN=atp2a2a PE A9C3Q4_DANRE,A9C3Q5_	A9C3Q4_DANRE,A9C3Q5_	109.519,40	100,00%	2	2	2	0,01%	3,26%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=ckma PE=1 SV=1	A2BHA3_DANRE	42.814,90	99,00%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:d1 X1WC76_DANRE	X1WC76_DANRE	17.030,40	97,00%	1	1	1	0,00%	7,28%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=vdac2 PE=1 SV=1	Q8AWD0_DANRE	30.284,90	100,00%	2	2	2	0,01%	11,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=col6a1 PE=1 SV=1	F1Q6P3_DANRE	107.466,50	100,00%	7	7	9	0,03%	10,40%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV A0A0R4IH07_DANRE,B6IC	A0A0R4IH07_DANRE,B6IC	347.704,80	93,50%	1	1	1	0,00%	0,55%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV A0A0R4IH07_DANRE,B6IC	A0A0R4IH07_DANRE,B6IC	347.704,80	93,50%	1	1	1	0,00%	0,55%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=uqrc1 PE=1 SV=1	F1QUE3_DANRE	55.450,70	100,00%	2	2	2	0,01%	3,78%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=flna PE=1 SV=1	E9QI62_DANRE,F1R9P5_I	272.475,80	100,00%	1	1	1	0,00%	0,51%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=flna PE=1 SV=1	E9QI62_DANRE,F1R9P5_I	272.475,80	100,00%	1	1	1	0,00%	0,51%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=zgc:101560 PE= A0A0R4IBE8_DANRE,A0AI	A0A0R4IBE8_DANRE,A0AI	21.007,60	95,00%	1	1	1	0,00%	4,37%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=zgc:101560 PE= A0A0R4IBE8_DANRE,A0AI	A0A0R4IBE8_DANRE,A0AI	21.007,60	95,00%	1	1	1	0,00%	4,37%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=myom1b PE=4 S A0A0R4IGQ8_DANRE	A0A0R4IGQ8_DANRE	180.977,20	100,00%	6	6	6	0,02%	4,36%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=eppk1 PE=4 SV= A0A0R4IND9_DANRE,I3IS	A0A0R4IND9_DANRE,I3IS	611.724,10	97,00%	1	1	1	0,00%	0,18%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=eppk1 PE=4 SV= A0A0R4IND9_DANRE,I3IS	A0A0R4IND9_DANRE,I3IS	611.724,10	97,00%	1	1	1	0,00%	0,18%
14dpa_2	Heat shock cognate 71 kDa protein OS=Danio rerio GN=hsc7l A0A0R4IMF8_DANRE,A0A	A0A0R4IMF8_DANRE,A0A	70.948,00	74,80%	0	0	0	0,00%	0,00%
14dpa_2	Heat shock cognate 71 kDa protein OS=Danio rerio GN=hsc7l A0A0R4IMF8_DANRE,A0A	A0A0R4IMF8_DANRE,A0A	70.948,00	74,80%	0	0	0	0,00%	0,00%
14dpa_2	Heat shock cognate 71 kDa protein OS=Danio rerio GN=hsc7l A0A0R4IMF8_DANRE,A0A	A0A0R4IMF8_DANRE,A0A	70.948,00	74,80%	0	0	0	0,00%	0,00%
14dpa_2	Heat shock cognate 71 kDa protein OS=Danio rerio GN=hsc7l A0A0R4IMF8_DANRE,A0A	A0A0R4IMF8_DANRE,A0A	70.948,00	74,80%	0	0	0	0,00%	0,00%
14dpa_2	Heat shock cognate 71 kDa protein OS=Danio rerio GN=hsc7l A0A0R4IMF8_DANRE,A0A	A0A0R4IMF8_DANRE,A0A	70.948,00	74,80%	0	0	0	0,00%	0,00%
14dpa_2	Heat shock cognate 71 kDa protein OS=Danio rerio GN=hsc7l A0A0R4IMF8_DANRE,A0A	A0A0R4IMF8_DANRE,A0A	70.948,00	74,80%	0	0	0	0,00%	0,00%
14dpa_2	Fructose-bisphosphate aldolase OS=Danio rerio GN=aldoaa P Q803Q7_DANRE	Q803Q7_DANRE	39.740,60	100,00%	3	3	3	0,01%	12,40%
14dpa_2	Heat shock protein 1 OS=Danio rerio GN=hspb1 PE=1 SV=1	A4VAK4_DANRE	22.408,90	99,80%	1	1	1	0,00%	8,04%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=cox5b2 PE=2 SV A8E500_DANRE,Q0P3U4_	A8E500_DANRE,Q0P3U4_	14.308,50	99,60%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=cox5b2 PE=2 SV A8E500_DANRE,Q0P3U4_	A8E500_DANRE,Q0P3U4_	14.308,50	99,60%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=desma PE=1 SV=1	F1R8W3_DANRE	55.727,70	100,00%	7	9	17	0,06%	16,40%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=acs11b PE=1 SV=1	F1QDA6_DANRE	77.694,10	100,00%	2	2	2	0,01%	4,73%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:c1 E9QDE7_DANRE	E9QDE7_DANRE	83.925,70	15,50%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=fbn2b PE=1 SV=1	A0A0R4IKQ62_DANRE	212.420,00	100,00%	2	2	2	0,01%	1,89%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=emilin1a PE=1 S F1QC17_DANRE,Q5T263_	F1QC17_DANRE,Q5T263_	112.771,70	100,00%	4	4	4	0,01%	4,44%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=emilin1a PE=1 S F1QC17_DANRE,Q5T263_	F1QC17_DANRE,Q5T263_	112.771,70	100,00%	4	4	4	0,01%	4,44%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=vwa1 PE=1 SV=1	E7F710_DANRE	54.924,60	99,60%	1	1	1	0,00%	1,59%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=glud1a PE=1 SV=1	A0A0R4IUG7_DANRE,B8A	59.978,90	96,60%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=glud1a PE=1 SV=1	A0A0R4IUG7_DANRE,B8A	59.978,90	96,60%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=glud1a PE=1 SV=1	A0A0R4IUG7_DANRE,B8A	59.978,90	96,60%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=utrn PE=1 SV=1	A0A0H2UKN4_DANRE	299.775,50	100,00%	2	2	2	0,01%	0,99%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=cmlc1 PE=1 SV=1	B0RF07_DANRE	21.865,20	100,00%	9	11	19	0,06%	56,60%
14dpa_2	Hadha protein OS=Danio rerio GN=hadha PE=1 SV=1	A7YT47_DANRE	81.877,10	93,30%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=hmcn2 PE=1 SV=1	E7F3V2_DANRE	454.987,50	79,00%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=add1 PE=4 SV=1	A0A0R4IAJ9_DANRE,A0AC	81.732,80	95,30%	1	1	2	0,01%	3,14%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV=1	A0A0R4IBJ3_DANRE,F1R6	72.513,60	100,00%	2	2	2	0,01%	2,83%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=add1 PE=4 SV=1	A0A0R4IAJ9_DANRE,A0AC	81.732,80	95,30%	1	1	2	0,01%	3,14%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=add1 PE=4 SV=1	A0A0R4IAJ9_DANRE,A0AC	81.732,80	95,30%	1	1	2	0,01%	3,14%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV=1	A0A0R4IBJ3_DANRE,F1R6	72.513,60	100,00%	2	2	2	0,01%	2,83%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV=1	A0A0R4IBJ3_DANRE,F1R6	72.513,60	100,00%	2	2	2	0,01%	2,83%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV=1	A0A0R4IBJ3_DANRE,F1R6	72.513,60	100,00%	2	2	2	0,01%	2,83%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=actn2b PE=1 SV=1	E9QFR8_DANRE	103.214,90	100,00%	5	5	6	0,02%	7,38%
14dpa_2	Isocitrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=1	A0A0G2L6I6_DANRE,E7F4	50.921,70	97,00%	1	1	1	0,00%	3,78%
14dpa_2	Isocitrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=1	A0A0G2L6I6_DANRE,E7F4	50.921,70	97,00%	1	1	1	0,00%	3,78%
14dpa_2	Cadherin-2 OS=Danio rerio GN=cdh2 PE=2 SV=2	CADH2_DANRE,G1K2E7_I	99.051,70	99,70%	0	0	0	0,00%	0,00%
14dpa_2	Cadherin-2 OS=Danio rerio GN=cdh2 PE=2 SV=2	CADH2_DANRE,G1K2E7_I	99.051,70	99,70%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=tln1 PE=4 SV=1	A0A0R4IDZ8_DANRE,A0AI	271.258,30	100,00%	2	2	2	0,01%	1,69%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=tln1 PE=4 SV=1	A0A0R4IDZ8_DANRE,A0AI	271.258,30	100,00%	2	2	2	0,01%	1,69%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=tln1 PE=4 SV=1	A0A0R4IDZ8_DANRE,A0AI	271.258,30	100,00%	2	2	2	0,01%	1,69%
14dpa_2	Histone H4 OS=Danio rerio GN=si:dkey-108k21.23 PE=1 SV=1	A3KPR4_DANRE,X1WGM1	11.478,70	89,80%	1	1	1	0,00%	7,77%
14dpa_2	Histone H4 OS=Danio rerio GN=si:dkey-108k21.23 PE=1 SV=1	A3KPR4_DANRE,X1WGM1	11.478,70	89,80%	1	1	1	0,00%	7,77%
14dpa_2	Malate dehydrogenase OS=Danio rerio GN=mdh2 PE=1 SV=1	Q7T334_DANRE	35.420,30	100,00%	3	3	8	0,03%	13,10%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=pdha1a PE=4 SV A0A0R4IT09_DANRE,E7F9	A0A0R4IT09_DANRE,E7F9	43.741,90	14,80%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=pdha1a PE=4 SV A0A0R4IT09_DANRE,E7F9	A0A0R4IT09_DANRE,E7F9	43.741,90	14,80%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=pdha1a PE=4 SV A0A0R4IT09_DANRE,E7F9	A0A0R4IT09_DANRE,E7F9	43.741,90	14,80%	0	0	0	0,00%	0,00%
14dpa_2	Nicotinamide nucleotide transhydrogenase OS=Danio rerio G Q6NYQ7_DANRE	Q6NYQ7_DANRE	112.785,50	99,80%	1	1	1	0,00%	1,02%
14dpa_2	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IG11_DANRE,G3P_	A0A0R4IG11_DANRE,G3P_	35.784,00	97,60%	0	0	0	0,00%	0,00%
14dpa_2	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IG11_DANRE,G3P_	A0A0R4IG11_DANRE,G3P_	35.784,00	97,60%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=mm F1R6V5_DANRE	F1R6V5_DANRE	122.540,40	9,20%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=cilp PE=4 SV=1	F1Q775_DANRE	130.911,60	99,80%	1	1	1	0,00%	1,37%
14dpa_2	Serotransferrin OS=Danio rerio GN=tfa PE=1 SV=1	B8JL43_DANRE,F1R858_D	73.717,60	99,40%	1	1	1	0,00%	1,49%
14dpa_2	Serotransferrin OS=Danio rerio GN=tfa PE=1 SV=1	B8JL43_DANRE,F1R858_D	73.717,60	99,40%	1	1	1	0,00%	1,49%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=pdha1a PE=4 SV A0A0R4IT09_DANRE,E7F9	A0A0R4IT09_DANRE,E7F9	43.741,90	14,80%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=pdha1a PE=4 SV A0A0R4IT09_DANRE,E7F9	A0A0R4IT09_DANRE,E7F9	43.741,90	14,80%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=flnca PE=4 SV=1	A0A0R4IUS4_DANRE,E7F1	291.642,80	100,00%	3	3	3	0,01%	2,18%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=flnca PE=4 SV=1	A0A0R4IUS4_DANRE,E7F1	291.642,80	100,00%	3	3	3	0,01%	2,18%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=eef1a111 PE=4 S A0A0R4IE10_DANRE,EF1A	A0A0R4IE10_DANRE,EF1A	50.048,20	100,00%	2	2	2	0,01%	5,19%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=eef1a111 PE=4 S A0A0R4IE10_DANRE,EF1A	A0A0R4IE10_DANRE,EF1A	50.048,20	100,00%	2	2	2	0,01%	5,19%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=palm2 PE=4 SV=1	A0A0R4ISM7_DANRE,E7F1	92.198,80	99,00%	1	1	1	0,00%	1,21%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=palm2 PE=4 SV=1	A0A0R4ISM7_DANRE,E7F1	92.198,80	99,00%	1	1	1	0,00%	1,21%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=si:ch211-114n2 E7F244_DANRE,Q6NWWJ5_	E7F244_DANRE,Q6NWWJ5_	50.034,50	99,50%	1	1	1	0,00%	4,43%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=si:ch211-114n2 E7F244_DANRE,Q6NWWJ5_	E7F244_DANRE,Q6NWWJ5_	50.034,50	99,50%	1	1	1	0,00%	4,43%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=elnb PE=4 SV=1	A0A0R4IPZ1_DANRE,F8W	171.803,90	97,00%	1	2	3	0,01%	0,79%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=elnb PE=4 SV=1	A0A0R4IPZ1_DANRE,F8W	171.803,90	97,00%	1	2	3	0,01%	0,79%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=elnb PE=4 SV=1	A0A0R4IPZ1_DANRE,F8W	171.803,90	97,00%	1	2	3	0,01%	0,79%
14dpa_2	ATP synthase subunit beta OS=Danio rerio GN=atp5b PE=1 S A8WGC6_DANRE	A8WGC6_DANRE	55.130,90	100,00%	9	11	21	0,07%	31,10%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=LOC10030353 F1QW6E_DANRE	F1QW6E_DANRE	86.981,10	18,40%	0	0	0	0,00%	0,00%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=plekha7a PE=4 S A0A0R4I85_DANRE,F1QC	A0A0R4I85_DANRE,F1QC	135.554,10	98,00%	1	1	1	0,00%	1,37%
14dpa_2	Uncharacterized protein OS=Danio rerio GN=plekha7a PE=4 S A0A0R4I85_DANRE,F1QC	A0A0R4I85_DANRE,F1QC	135.554,10	98,00%	1	1	1	0,00%	1,37%

30dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=	A0A0N4STN6_DANRE,A7H	13.590,30	99,60%	1	2	3	0,01%	8,87%
30dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=	A0A0N4STN6_DANRE,A7H	13.590,30	99,60%	1	2	3	0,01%	8,87%
30dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=	A0A0N4STN6_DANRE,A7H	13.590,30	99,60%	1	2	3	0,01%	8,87%
30dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=	A0A0N4STN6_DANRE,A7H	13.590,30	99,60%	1	2	3	0,01%	8,87%
30dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=	A0A0N4STN6_DANRE,A7H	13.590,30	99,60%	1	2	3	0,01%	8,87%
30dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=	A0A0N4STN6_DANRE,A7H	13.590,30	99,60%	1	2	3	0,01%	8,87%
30dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=	A0A0N4STN6_DANRE,A7H	13.590,30	99,60%	1	2	3	0,01%	8,87%
30dpa_1	Histone H2B OS=Danio rerio GN=si:dkey-261m9.19 PE=1 SV=	A0A0N4STN6_DANRE,A7H	13.590,30	99,60%	1	2	3	0,01%	8,87%
30dpa_1	Aconitase hydratase, mitochondrial (Fragment) OS=Danio rer	F8W4M7_DANRE	85.591,90	100,00%	2	2	3	0,01%	3,04%
30dpa_1	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1	A8WG05_DANRE,ACTB1_	41.753,80	100,00%	2	3	8	0,03%	24,00%
30dpa_1	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1	A8WG05_DANRE,ACTB1_	41.753,80	100,00%	2	3	8	0,03%	24,00%
30dpa_1	Actin, alpha, cardiac muscle 1a OS=Danio rerio GN=si:ch73-1	!Q6IQR3_DANRE	41.976,10	100,00%	10	16	40	0,14%	33,40%
30dpa_1	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1	A8WG05_DANRE,ACTB1_	41.753,80	100,00%	2	3	8	0,03%	24,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=cilp PE=4 SV=1	F1Q775_DANRE	130.911,60	100,00%	2	2	2	0,01%	3,25%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=fga PE=1 SV=1	B8A5L6_DANRE	75.292,70	100,00%	15	24	48	0,16%	26,80%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=col1a1b PE=1 SV	!F1QD1L_DANRE	136.914,60	100,00%	10	11	26	0,09%	8,21%
30dpa_1	Keratin, type I cytoskeletal 18 OS=Danio rerio GN=krt18 PE=1	K1C18_DANRE	48.631,90	72,50%	1	1	1	0,00%	2,32%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=vwa1 PE=1 SV=1	E7F10_DANRE	54.924,60	23,30%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=col4a2 PE=1 SV	A0A0H2UKP2_DANRE	165.476,80	100,00%	4	4	5	0,02%	3,89%
30dpa_1	Radixin OS=Danio rerio GN=msna PE=1 SV=1	Q66I42_DANRE	68.035,00	100,00%	3	3	3	0,01%	5,53%
30dpa_1	Catenin (Cadherin-associated protein), alpha OS=Danio rerio	A0A0R4IF47_DANRE,Q9P	100.520,10	100,00%	1	1	1	0,00%	1,76%
30dpa_1	Catenin (Cadherin-associated protein), alpha OS=Danio rerio	A0A0R4IF47_DANRE,Q9P	100.520,10	100,00%	1	1	1	0,00%	1,76%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=emilin1b PE=1 S	F1Q9G3_DANRE,F8W4C7_	116.801,40	100,00%	7	8	9	0,03%	7,65%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=emilin1b PE=1 S	F1Q9G3_DANRE,F8W4C7_	116.801,40	100,00%	7	8	9	0,03%	7,65%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=hspg2 PE=4 SV=	A0A0R4IZA6_DANRE,F1R5	369.970,80	100,00%	18	19	28	0,10%	7,77%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=hspg2 PE=4 SV=	A0A0R4IZA6_DANRE,F1R5	369.970,80	100,00%	18	19	28	0,10%	7,77%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=c3a.1 PE=1 SV=	!B8JKW4_DANRE	183.235,60	100,00%	2	2	2	0,01%	2,37%
30dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=slc2	E9QJ49_DANRE,F1QAJ7_I	39.378,90	99,70%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=slc2	E9QJ49_DANRE,F1QAJ7_I	39.378,90	99,70%	0	0	0	0,00%	0,00%
30dpa_1	Myoglobin OS=Danio rerio GN=mb PE=2 SV=3	MYG_DANRE	15.583,60	100,00%	3	3	5	0,02%	30,60%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=pkma PE=4 SV=	A0A0R4IGP6_DANRE,Q7Z	58.059,00	100,00%	2	2	2	0,01%	6,20%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=pkma PE=4 SV=	A0A0R4IGP6_DANRE,Q7Z	58.059,00	100,00%	2	2	2	0,01%	6,20%
30dpa_1	Annexin OS=Danio rerio GN=anxa2a PE=1 SV=1	Q6P603_DANRE	38.140,50	100,00%	1	2	4	0,01%	4,75%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=col6a2 PE=4 SV	E7FCV8_DANRE	106.916,20	100,00%	11	12	20	0,07%	12,90%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=fbn2b PE=1 SV=	A0A0G2KQ62_DANRE	212.420,00	100,00%	2	2	2	0,01%	1,74%
30dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttna	B05758_DANRE	843.490,10	100,00%	2	2	2	0,01%	0,50%
30dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=col6	F6NR90_DANRE	88.827,90	92,10%	0	0	0	0,00%	11,00%
30dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=tnb	B056Y0_DANRE	626.764,10	100,00%	1	1	1	0,00%	0,48%
30dpa_1	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1	B3DG37_DANRE,HBB1_D	16.389,00	100,00%	4	4	6	0,02%	34,50%
30dpa_1	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1	B3DG37_DANRE,HBB1_D	16.389,00	100,00%	4	4	6	0,02%	34,50%
30dpa_1	Aspartate aminotransferase OS=Danio rerio GN=got2a PE=1	!Q7SYK7_DANRE	47.588,10	100,00%	1	1	1	0,00%	1,87%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=tjp1b PE=1 SV=	!X1WHN9_DANRE	205.686,40	98,80%	1	1	1	0,00%	1,19%
30dpa_1	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio	A0A0R4IG1I_DANRE,G3P_	35.784,00	99,60%	1	1	1	0,00%	4,46%
30dpa_1	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio	A0A0R4IG1I_DANRE,G3P_	35.784,00	99,60%	1	1	1	0,00%	4,46%
30dpa_1	Elongation factor 1-alpha OS=Danio rerio GN=eef1a PE=2 SV	EF1A_DANRE	50.048,20	98,80%	1	1	1	0,00%	2,60%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=ptx3a PE=1 SV=	A3KPN4_DANRE	49.273,20	100,00%	0	0	0	0,00%	0,00%
30dpa_1	Fructose-bisphosphate aldolase OS=Danio rerio GN=aldoaa	P Q803Q7_DANRE	39.740,60	100,00%	2	2	3	0,01%	6,04%
30dpa_1	Calcium-transporting ATPase OS=Danio rerio GN=atp2a2a PE	A9C3Q4_DANRE,A9C3Q5_	109.519,40	100,00%	3	4	7	0,02%	4,41%
30dpa_1	Calcium-transporting ATPase OS=Danio rerio GN=atp2a2a PE	A9C3Q4_DANRE,A9C3Q5_	109.519,40	100,00%	3	4	7	0,02%	4,41%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=ckma PE=1 SV=	!A2BHA3_DANRE	42.814,90	33,70%	0	0	0	0,00%	0,00%
30dpa_1	Iso citrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=	A0A0G2L6I6_DANRE,E7F4	50.921,70	100,00%	1	2	2	0,01%	2,89%
30dpa_1	Iso citrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=	A0A0G2L6I6_DANRE,E7F4	50.921,70	100,00%	1	2	2	0,01%	2,89%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=LOC557764 PE=	F1R8B2_DANRE	169.128,30	52,60%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=vwf PE=4 SV=2	E7F9Z1_DANRE	307.995,50	100,00%	1	1	2	0,01%	0,46%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=wfdc2 PE=4 SV=	E7F3G3_DANRE	17.778,70	100,00%	1	1	2	0,01%	11,90%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=vdac2 PE=1 SV=	Q8AWD0_DANRE	30.284,90	100,00%	1	1	1	0,00%	6,71%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:56235 PE=1	Q7VZ39_DANRE	30.045,40	66,10%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV=	A0A0R4IBJ3_DANRE,F1R6	72.513,60	100,00%	1	1	2	0,01%	1,64%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV=	A0A0R4IBJ3_DANRE,F1R6	72.513,60	100,00%	1	1	2	0,01%	1,64%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=add3a PE=1 SV=	A0A0R4IBJ3_DANRE,F1R6	72.513,60	100,00%	1	1	2	0,01%	1,64%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=flnca PE=4 SV=	!A0A0R4IU54_DANRE,E7F1	291.642,80	100,00%	3	3	3	0,01%	2,33%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=flnca PE=4 SV=	!A0A0R4IU54_DANRE,E7F1	291.642,80	100,00%	3	3	3	0,01%	2,33%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=acs11b PE=1 SV=	F1QD46_DANRE	77.694,10	56,10%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:cf	E9QDE7_DANRE	83.925,70	20,30%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:101560 PE=	A0A0R4IBE8_DANRE,A0A	21.007,60	99,90%	1	1	1	0,00%	4,37%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:101560 PE=	A0A0R4IBE8_DANRE,A0A	21.007,60	99,90%	1	1	1	0,00%	4,37%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=ltpb1 PE=4 SV=	!F1QQ56_DANRE,X1WBQ9	131.384,10	100,00%	2	2	2	0,01%	2,10%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=ltpb1 PE=4 SV=	!F1QQ56_DANRE,X1WBQ9	131.384,10	100,00%	2	2	2	0,01%	2,10%
30dpa_1	ATP synthase subunit beta OS=Danio rerio GN=atp5b PE=1 S	A8WGG6_DANRE	55.130,90	100,00%	5	5	12	0,04%	19,90%
30dpa_1	Sodium/potassium-transporting ATPase subunit alpha OS=D	A0A0R4IRT1_DANRE,Q9D	113.384,80	100,00%	2	3	3	0,01%	3,99%
30dpa_1	Sodium/potassium-transporting ATPase subunit alpha OS=D	A0A0R4IRT1_DANRE,Q9D	113.384,80	100,00%	2	3	3	0,01%	3,99%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV	A0A0R4IH07_DANRE,B6IC	347.704,80	54,10%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=frem2b PE=4 SV	A0A0R4IH07_DANRE,B6IC	347.704,80	54,10%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=palm2 PE=4 SV=	A0A0R4ISM7_DANRE,E7F	92.198,80	100,00%	2	3	3	0,01%	2,67%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=palm2 PE=4 SV=	A0A0R4ISM7_DANRE,E7F	92.198,80	100,00%	2	3	3	0,01%	2,67%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=LOC10032905	A0A0G2KJ47_DANRE	41.855,30	99,20%	0	0	0	0,00%	0,00%
30dpa_1	Nicotinamide nucleotide transhydrogenase OS=Danio rerio	G Q6NYQ7_DANRE	112.785,50	97,10%	1	1	1	0,00%	0,83%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=pdhb PE=1 SV=	!Q1LV50_DANRE,Q7T368_	39.308,70	91,50%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=pdhb PE=1 SV=	!Q1LV50_DANRE,Q7T368_	39.308,70	91,50%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV=	A0A0R4IP01_DANRE,B8JK	119.385,60	13,90%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV=	A0A0R4IP01_DANRE,B8JK	119.385,60	13,90%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=ogdha PE=1 SV=	A0A0R4IP01_DANRE,B8JK	119.385,60	13,90%	0	0	0	0,00%	0,00%
30dpa_1	Laminin subunit gamma-1 OS=Danio rerio GN=lamc1 PE=2 SV	LAMC1_DANRE	176.206,30	100,00%	4	5	8	0,03%	5,02%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=cmlc1 PE=1 SV=	B0R0F7_DANRE	21.865,20	100,00%	6	6	10	0,03%	37,20%
30dpa_1	Hadha protein OS=Danio rerio GN=hadhaa PE=1 SV=1	A7Y747_DANRE,F1QYX8_I	82.848,80	16,10%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=	!A0A0R4INC4_DANRE,A5V	543.007,50	100,00%	2	2	2	0,01%	1,16%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=	!A0A0R4INC4_DANRE,A5V	543.007,50	100,00%	2	2	2	0,01%	1,16%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=	!A0A0R4INC4_DANRE,A5V	543.007,50	100,00%	2	2	2	0,01%	1,16%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=cox5b2 PE=1 SV	Q0P3U4_DANRE	14.308,50	99,90%	1	1	1	0,00%	10,30%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE=	A0A0R4IE46_DANRE,A0A	101.943,10	100,00%	2	2	2	0,01%	2,16%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE=	A0A0R4IE46_DANRE,A0A	101.943,10	100,00%	2	2	2	0,01%	2,16%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE=	A0A0R4IE46_DANRE,A0A	101.943,10	100,00%	2	2	2	0,01%	2,16%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=palm1b PE=4 SV	B0V0Y4_DANRE	27.123,40	97,70%	0	0	0	0,00%	0,00%
30dpa_1	Heat shock cognate 71 kDa protein OS=Danio rerio GN=hsc7	Q6PGX4_DANRE	70.948,00	99,70%	1	1	1	0,00%	1,24%

30dpa_1	Hydroxyacyl-Coenzyme A dehydrogenase OS=Danio rerio GN Q6D122_DANRE	33.336,90	23,90%	0	0	0	0,00%	0,00%
30dpa_1	Ceruloplasmin OS=Danio rerio GN=cp PE=1 SV=1 Q6P3G1_DANRE	124.388,00	57,90%	0	0	0	0,00%	0,00%
30dpa_1	Acetyl-CoA acetyltransferase, mitochondrial OS=Danio rerio GN=A0A0R4I176_DANRE,THIL_	44.342,90	100,00%	1	1	1	0,00%	3,33%
30dpa_1	Acetyl-CoA acetyltransferase, mitochondrial OS=Danio rerio GN=A0A0R4I176_DANRE,THIL_	44.342,90	100,00%	1	1	1	0,00%	3,33%
30dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6P1	29.220,20	23,20%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6P1	29.220,20	23,20%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6P1	29.220,20	23,20%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein (Fragment) OS=Danio rerio GN=parv A0A0R4IY82_DANRE,Q6P1	29.220,20	23,20%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=aca2 PE=1 SV= B055C5_DANRE	41.786,40	100,00%	2	2	2	0,01%	8,82%
30dpa_1	Tubulin alpha 8-like 3a protein OS=Danio rerio GN=tuba8l3 P A0A0R4INE6_DANRE,E7E1	50.158,70	15,30%	0	0	0	0,00%	0,00%
30dpa_1	Phosphoglycerate kinase OS=Danio rerio GN=pgk1 PE=1 SV=1 F1QXV8_DANRE	44.733,20	98,10%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=sdprb PE=1 SV=1 F1R665_DANRE	51.441,60	99,50%	1	1	1	0,00%	4,22%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=c9 PE=1 SV=1 B8A565_DANRE	72.713,70	19,40%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=si:dkey-238c7.1 Q1LX26_DANRE	21.858,50	31,50%	0	0	0	0,00%	0,00%
30dpa_1	Uncharacterized protein OS=Danio rerio GN=zgc:101840 PE= Q5TYR6_DANRE	21.949,50	10,20%	0	0	0	0,00%	0,00%
30dpa_1	E7F2R3-DECOY A0A0R4I171-DECOY,E7F2F	0	18,30%	0	0	0	0,00%	0,00%
30dpa_1	E7F2R3-DECOY A0A0R4I171-DECOY,E7F2F	0	18,30%	0	0	0	0,00%	0,00%
30dpa_1	E7F2R3-DECOY A0A0R4I171-DECOY,E7F2F	0	18,30%	0	0	0	0,00%	0,00%
30dpa_1	E7F2R3-DECOY A0A0R4I171-DECOY,E7F2F	0	18,30%	0	0	0	0,00%	0,00%
30dpa_1	A0A0R4I1J5-DECOY A0A0R4I1J5-DECOY	18	98,90%	1	1	1	0,00%	NaNa%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ttna PE=1 SV=2 F1R7N8_DANRE	3.089.806,00	100,00%	12	12	15	0,05%	0,72%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttna B05758_DANRE	843.490,10	99,80%	1	1	1	0,00%	0,26%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttnb B056V0_DANRE	626.764,10	98,70%	1	1	1	0,00%	0,52%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=sorbs1 PE=4 SV= A0A0R4IP30_DANRE	85.938,80	100,00%	3	3	4	0,01%	6,53%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4I171_DANRE,E7F4'	518.896,50	100,00%	3	3	3	0,01%	1,26%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4I171_DANRE,E7F4'	518.896,50	100,00%	3	3	3	0,01%	1,26%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=plecb PE=1 SV=: A0A0R4I171_DANRE,E7F4'	518.896,50	100,00%	3	3	3	0,01%	1,26%
30dpa_2	Radixin OS=Danio rerio GN=msna PE=1 SV=1 Q66I42_DANRE	68.035,00	100,00%	2	2	2	0,01%	2,25%
30dpa_2	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1 A8W605_DANRE,ACTB1_	41.753,80	99,90%	1	2	8	0,03%	18,40%
30dpa_2	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1 A8W605_DANRE,ACTB1_	41.753,80	99,90%	1	2	8	0,03%	18,40%
30dpa_2	Actin, alpha, cardiac muscle 1a OS=Danio rerio GN=si:ch73-1f Q6IQR3_DANRE	41.976,10	100,00%	9	13	32	0,11%	27,90%
30dpa_2	Bactin2 protein OS=Danio rerio GN=actb2 PE=1 SV=1 A8W605_DANRE,ACTB1_	41.753,80	99,90%	1	2	8	0,03%	18,40%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=fn1b PE=1 SV=2 A2CEW3_DANRE,F1R809_	265.661,60	100,00%	43	63	141	0,50%	25,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=fn1b PE=1 SV=2 A2CEW3_DANRE,F1R809_	265.661,60	100,00%	43	63	141	0,50%	25,00%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=ttnb G1K2X0_DANRE	2.458.846,80	100,00%	9	9	13	0,05%	0,86%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=col6a3 PE=1 SV= F1Q4X1_DANRE	309.287,80	100,00%	18	20	46	0,16%	10,60%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=col5 F6NPAA_DANRE	198.988,80	100,00%	2	2	2	0,01%	1,44%
30dpa_2	Periostin OS=Danio rerio GN=postnb PE=1 SV=1 Q75U66_DANRE	85.887,00	100,00%	8	11	18	0,06%	19,80%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=hist2h3ca1 PE=: A0A0R4IY26_DANRE,A8KE	17.030,40	97,70%	1	1	1	0,00%	6,62%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=hist2h3ca1 PE=: A0A0R4IY26_DANRE,A8KE	17.030,40	97,70%	1	1	1	0,00%	6,62%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=hist2h3ca1 PE=: A0A0R4IY26_DANRE,A8KE	17.030,40	97,70%	1	1	1	0,00%	6,62%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=hist2h3ca1 PE=: A0A0R4IY26_DANRE,A8KE	17.030,40	97,70%	1	1	1	0,00%	6,62%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=hist2h3ca1 PE=: A0A0R4IY26_DANRE,A8KE	17.030,40	97,70%	1	1	1	0,00%	6,62%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=hist2h3ca1 PE=: A0A0R4IY26_DANRE,A8KE	17.030,40	97,70%	1	1	1	0,00%	6,62%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=cox4i1 PE=1 SV: F1R6I1_DANRE	20.331,60	64,50%	0	0	0	0,00%	0,00%
30dpa_2	Catenin (Cadherin-associated protein), alpha OS=Danio rerio A0A0R4IF47_DANRE,Q9P1	100.520,10	100,00%	1	1	1	0,00%	1,76%
30dpa_2	Catenin (Cadherin-associated protein), alpha OS=Danio rerio A0A0R4IF47_DANRE,Q9P1	100.520,10	100,00%	1	1	1	0,00%	1,76%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV A0A0R4IH03_DANRE,F1R4'	108.322,90	100,00%	3	3	3	0,01%	3,20%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV A0A0R4IH03_DANRE,F1R4'	108.322,90	100,00%	3	3	3	0,01%	3,20%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=slc8a1a PE=1 SV A0A0R4IH03_DANRE,F1R4'	108.322,90	100,00%	3	3	3	0,01%	3,20%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:c F1Q7R6_DANRE	147.365,60	100,00%	4	4	4	0,01%	2,89%
30dpa_2	ATP synthase subunit beta OS=Danio rerio GN=atp5b PE=1 SV: A8W6C6_DANRE	55.130,90	100,00%	4	5	8	0,03%	16,40%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ahnak PE=1 SV= F1QZ50_DANRE	685.184,90	100,00%	7	7	9	0,03%	1,42%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a1a PE=1 SV: F1QJQ9_DANRE,F1QSC5_I	128.043,70	100,00%	2	2	3	0,01%	3,39%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a1a PE=1 SV: F1QJQ9_DANRE,F1QSC5_I	128.043,70	100,00%	2	2	3	0,01%	3,39%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=spna2 PE=1 SV= F1R443_DANRE,F1R446_I	284.932,80	100,00%	22	29	46	0,16%	13,60%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=spna2 PE=1 SV= F1R443_DANRE,F1R446_I	284.932,80	100,00%	22	29	46	0,16%	13,60%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ank3a PE=4 SV= F1QN81_DANRE,F8W371_	194.040,10	100,00%	2	2	2	0,01%	1,58%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ank3a PE=4 SV= F1QN81_DANRE,F8W371_	194.040,10	100,00%	2	2	2	0,01%	1,58%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=kr194 PE=1 SV=: F1R8U0_DANRE	47.787,30	99,90%	1	1	1	0,00%	2,32%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=postna PE=1 SV: E7F513_DANRE	97.135,00	100,00%	3	3	5	0,02%	4,88%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=1 SV= A0A0R4IF18_DANRE,A2B1	103.798,70	100,00%	2	2	3	0,01%	2,94%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=1 SV= A0A0R4IF18_DANRE,A2B1	103.798,70	100,00%	2	2	3	0,01%	2,94%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=arvcfb PE=1 SV= A0A0R4IF18_DANRE,A2B1	103.798,70	100,00%	2	2	3	0,01%	2,94%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=vmhcl PE=1 SV= F1QSE1_DANRE	223.380,90	100,00%	54	74	176	0,62%	34,70%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=col6a1 PE=1 SV: F1QP63_DANRE	107.466,50	100,00%	2	2	3	0,01%	2,58%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=sptb PE=1 SV=1 F1QQE5_DANRE	272.829,20	100,00%	17	22	28	0,10%	11,30%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=sptbn1 PE=1 SV: A0A0G2L1F5_DANRE	262.630,10	100,00%	13	16	23	0,08%	8,98%
30dpa_2	ATP synthase subunit alpha OS=Danio rerio GN=atp5a1 PE=1 Q08B81_DANRE	59.745,30	100,00%	1	1	3	0,01%	2,90%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=LOC100332905 A0A0G2KJ47_DANRE,A0A	41.826,10	99,10%	1	1	1	0,00%	2,32%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=LOC100332905 A0A0G2KJ47_DANRE,A0A	41.826,10	99,10%	1	1	1	0,00%	2,32%
30dpa_2	Fgg protein OS=Danio rerio GN=fgg PE=1 SV=1 Q7ZV67_DANRE	48.833,70	100,00%	10	12	63	0,22%	33,40%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	99,10%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	99,10%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ctnnb1 PE=1 SV: A0A0N4SUA5_DANRE,E7F	81.108,80	99,10%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=desma PE=1 SV: F1R8W3_DANRE	55.727,70	100,00%	9	11	19	0,07%	19,30%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=mfap5 PE=4 SV= E9QC43_DANRE	14.664,20	98,40%	1	1	1	0,00%	11,90%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=myhb PE=1 SV=: F1QVQ3_DANRE,X1WF87_	222.608,80	93,80%	1	1	1	0,00%	5,11%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=myhb PE=1 SV=: F1QVQ3_DANRE,X1WF87_	222.608,80	93,80%	1	1	1	0,00%	5,11%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a1b PE=1 SV: F1QDL1_DANRE	136.914,60	100,00%	8	10	19	0,07%	7,25%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=mfap2 PE=4 SV= F1QSF1_DANRE	17.997,30	100,00%	1	1	2	0,01%	6,33%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ablml1b PE=4 S U3J1B3_DANRE	84.999,80	99,60%	1	1	1	0,00%	1,85%
30dpa_2	Blood vessel epicardial substance OS=Danio rerio GN=bves P POPD1_DANRE	40.341,40	100,00%	2	5	5	0,02%	9,38%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=: A0A0R4INC4_DANRE,A5W	543.007,50	100,00%	2	2	2	0,01%	0,98%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=: A0A0R4INC4_DANRE,A5W	543.007,50	100,00%	2	2	2	0,01%	0,98%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=pleca PE=1 SV=: A0A0R4INC4_DANRE,A5W	543.007,50	100,00%	2	2	2	0,01%	0,98%
30dpa_2	Fibrinogen, B beta polypeptide OS=Danio rerio GN=fgb PE=1 Q6NYE1_DANRE	54.407,60	100,00%	9	16	52	0,18%	31,30%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=si:ch211-113a1 R4GE02_DANRE	27.150,10	100,00%	1	1	1	0,00%	4,44%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=fn1a PE=1 SV=1 B05602_DANRE	271.953,50	100,00%	5	5	5	0,02%	5,60%
30dpa_2	Aconitate hydratase, mitochondrial (Fragment) OS=Danio rer F8W4M7_DANRE	85.591,90	100,00%	3	3	3	0,01%	5,06%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rer GN=col4 F1Q855_DANRE,F1Q679_	128.554,50	100,00%	4	6	15	0,05%	3,54%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rer GN=col4 F1Q855_DANRE,F1Q679_	128.554,50	100,00%	4	6	15	0,05%	3,54%
30dpa_2	Solute carrier family 25 alpha, member 5 OS=Danio rerio GN= Q8JH10_DANRE	32.764,40	100,00%	3	3	3	0,01%	9,06%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=fga PE=1 SV=1 B8A5L6_DANRE,E9QCD1_	49.680,00	100,00%	13	19	35	0,12%	23,00%

30dpa_2	Uncharacterized protein OS=Danio rerio GN=fga PE=1 SV=1 B8A5L6_DANRE,E9QCD1_	49.680,00	100,00%	13	19	35	0,12%	23,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=vtg1 PE=1 SV=1 Q1LWN2_DANRE	149.238,80	100,00%	1	1	1	0,00%	1,32%
30dpa_2	Fructose-bisphosphate aldolase OS=Danio rerio GN=aldoa P Q803Q7_DANRE	39.740,60	100,00%	2	2	2	0,01%	4,67%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=col6 F6NRG0_DANRE	88.827,90	72,30%	0	0	0	0,00%	4,73%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=tltn2a PE=4 SV=1 A0A0R4IIA7_DANRE	271.364,80	100,00%	2	2	2	0,01%	2,08%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=hspg2 PE=4 SV= A0A0R4I2A6_DANRE,F1R5	369.970,80	100,00%	12	12	19	0,07%	5,46%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=hspg2 PE=4 SV= A0A0R4I2A6_DANRE,F1R5	369.970,80	100,00%	12	12	19	0,07%	5,46%
30dpa_2	Keratin, type I cytoskeletal 18 OS=Danio rerio GN=kr18 PE=1 K1C18_DANRE	48.631,90	98,70%	1	1	1	0,00%	2,32%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ccdc141 PE=4 S1 B0S755_DANRE,E7FG99_I	190.652,30	99,90%	1	1	1	0,00%	0,66%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ccdc141 PE=4 S1 B0S755_DANRE,E7FG99_I	190.652,30	99,90%	1	1	1	0,00%	0,66%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=col4a2 PE=1 SV= A0A0H2UKP2_DANRE	165.476,80	100,00%	3	3	3	0,01%	2,34%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 S1 E7EZ79_DANRE,F1Q615_I	142.109,70	100,00%	5	6	6	0,02%	7,13%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 S1 E7EZ79_DANRE,F1Q615_I	142.109,70	100,00%	5	6	6	0,02%	7,13%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=mybpc3 PE=1 S1 E7EZ79_DANRE,F1Q615_I	142.109,70	100,00%	5	6	6	0,02%	7,13%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a2 PE=1 SV= A0A0R4IHCO_DANRE,E9Q	127.211,90	100,00%	12	14	37	0,13%	12,70%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a2 PE=1 SV= A0A0R4IHCO_DANRE,E9Q	127.211,90	100,00%	12	14	37	0,13%	12,70%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=col1a2 PE=1 SV= A0A0R4IHCO_DANRE,E9Q	127.211,90	100,00%	12	14	37	0,13%	12,70%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=dspa PE=1 SV=2 F1RBA4_DANRE	276.745,50	99,80%	1	1	1	0,00%	0,46%
30dpa_2	Annexin OS=Danio rerio GN=anxa2a PE=1 SV=1 Q6P603_DANRE	38.140,50	100,00%	2	2	5	0,02%	7,12%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=plekha7a PE=4 S A0A0R4II85_DANRE,F1Q7	135.554,10	99,80%	1	1	1	0,00%	0,88%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=plekha7a PE=4 S A0A0R4II85_DANRE,F1Q7	135.554,10	99,80%	1	1	1	0,00%	0,88%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=plekha7a PE=4 S A0A0R4II85_DANRE,F1Q7	135.554,10	99,80%	1	1	1	0,00%	0,88%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=plekha7a PE=4 S A0A0R4II85_DANRE,F1Q7	135.554,10	99,80%	1	1	1	0,00%	0,88%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ddx3a PE=1 SV= A0A0H2UJY3_DANRE,B0S	75.834,30	66,00%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ddx3a PE=1 SV= A0A0H2UJY3_DANRE,B0S	75.834,30	66,00%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ddx3a PE=1 SV= A0A0H2UJY3_DANRE,B0S	75.834,30	66,00%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ddx3a PE=1 SV= A0A0H2UJY3_DANRE,B0S	75.834,30	66,00%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=itih2 PE=4 SV=1 Q5RH29_DANRE	106.292,90	100,00%	4	4	8	0,03%	8,14%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=col6a2 PE=4 SV= E7FCV8_DANRE	106.916,20	100,00%	10	10	12	0,04%	14,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=wu:fi04e12 PE= F1QAU9_DANRE	251.072,30	99,90%	1	1	1	0,00%	0,73%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ptx3a PE=1 SV= A3KPN4_DANRE	49.273,20	100,00%	1	1	1	0,00%	2,91%
30dpa_2	Laminin subunit gamma-1 OS=Danio rerio GN=lamc1 PE=2 SV LAMC1_DANRE	176.206,30	100,00%	2	3	5	0,02%	2,20%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=tjp1b PE=1 SV=1 X1WHN9_DANRE	205.686,40	99,60%	1	1	1	0,00%	1,19%
30dpa_2	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IG1I_DANRE,G3P_	35.784,00	96,90%	0	0	0	0,00%	0,00%
30dpa_2	Glyceraldehyde-3-phosphate dehydrogenase OS=Danio rerio A0A0R4IG1I_DANRE,G3P_	35.784,00	96,90%	0	0	0	0,00%	0,00%
30dpa_2	Calcium-transporting ATPase OS=Danio rerio GN=atp2a2a PE A9C3Q4_DANRE,A9C3Q5_	109.519,40	100,00%	2	3	7	0,02%	3,36%
30dpa_2	Calcium-transporting ATPase OS=Danio rerio GN=atp2a2a PE A9C3Q4_DANRE,A9C3Q5_	109.519,40	100,00%	2	3	7	0,02%	3,36%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=si:ct E9QD77_DANRE	83.925,70	99,60%	1	1	1	0,00%	1,12%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=flnca PE=4 SV=1 A0A0R4IU54_DANRE,E7F1	291.642,80	100,00%	2	2	2	0,01%	1,22%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=flnca PE=4 SV=1 A0A0R4IU54_DANRE,E7F1	291.642,80	100,00%	2	2	2	0,01%	1,22%
30dpa_2	Iso citrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=1 A0A0G2L6I6_DANRE,E7F4	50.921,70	100,00%	2	3	5	0,02%	6,67%
30dpa_2	Iso citrate dehydrogenase [NADP] OS=Danio rerio PE=1 SV=1 A0A0G2L6I6_DANRE,E7F4	50.921,70	100,00%	2	3	5	0,02%	6,67%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=emilin1b PE=1 S F1Q9G3_DANRE,F8W4C7_	116.801,40	100,00%	5	6	7	0,02%	5,48%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=emilin1b PE=1 S F1Q9G3_DANRE,F8W4C7_	116.801,40	100,00%	5	6	7	0,02%	5,48%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=wfdc2 PE=4 SV= E7F3G3_DANRE	17.778,70	97,70%	1	1	2	0,01%	11,90%
30dpa_2	Histone H4 OS=Danio rerio GN=si:dkey-108k21.23 PE=1 SV=1 A3KPR4_DANRE,X1WGM1_	11.478,70	98,60%	0	0	0	0,00%	0,00%
30dpa_2	Histone H4 OS=Danio rerio GN=si:dkey-108k21.23 PE=1 SV=1 A3KPR4_DANRE,X1WGM1_	11.478,70	98,60%	0	0	0	0,00%	0,00%
30dpa_2	Phosphoglycerate kinase (Fragment) OS=Danio rerio GN=pgk E9QF19_DANRE,F1QXV8_	44.733,20	30,50%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=aca2 PE=1 SV= B0S5C5_DANRE	41.786,40	100,00%	2	2	2	0,01%	8,31%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=vdac2 PE=1 SV= Q8AWD0_DANRE	30.284,90	12,70%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=zgc:101560 PE= A0A0R4IBE8_DANRE,A0AI	21.007,60	99,50%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=zgc:101560 PE= A0A0R4IBE8_DANRE,A0AI	21.007,60	99,50%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ltbp1 PE=4 SV=2 F1QQ56_DANRE,X1WBQ9	131.384,10	100,00%	1	2	2	0,01%	0,79%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=ltbp1 PE=4 SV=2 F1QQ56_DANRE,X1WBQ9	131.384,10	100,00%	1	2	2	0,01%	0,79%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=emilin1a PE=1 S F1QC17_DANRE,Q5T263_	112.771,70	100,00%	5	5	5	0,02%	6,61%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=emilin1a PE=1 S F1QC17_DANRE,Q5T263_	112.771,70	100,00%	5	5	5	0,02%	6,61%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=lmo7a PE=4 SV= A0A0R4ITT8_DANRE	156.646,40	100,00%	2	3	4	0,01%	2,48%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=col6 F1Q924_DANRE	278.174,40	100,00%	7	7	12	0,04%	4,60%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=acs11b PE=1 SV= F1QDA6_DANRE	77.694,10	12,50%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=post F1QM50_DANRE	17.277,20	80,10%	1	1	1	0,00%	12,40%
30dpa_2	Nicotinamide nucleotide transhydrogenase OS=Danio rerio G Q6NYQ7_DANRE	112.785,50	99,30%	1	1	1	0,00%	0,83%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=slc25a12 PE=4 S A0A0R4IHT0_DANRE,A0AI	72.864,20	74,10%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=slc25a12 PE=4 S A0A0R4IHT0_DANRE,A0AI	72.864,20	74,10%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=cmlc1 PE=1 SV= B0R0F7_DANRE	21.865,20	100,00%	4	4	9	0,03%	26,50%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=vwf PE=4 SV=2 E7F921_DANRE,F1QHS6_I	225.634,30	100,00%	2	2	4	0,01%	1,39%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=vwf PE=4 SV=2 E7F921_DANRE,F1QHS6_I	225.634,30	100,00%	2	2	4	0,01%	1,39%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=cox5b2 PE=1 SV QQP3U4_DANRE	14.308,50	97,60%	1	1	1	0,00%	10,30%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=tpm4a PE=1 SV= F1QKQ7_DANRE	32.726,90	100,00%	5	7	8	0,03%	25,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=fbn2b PE=1 SV= A0A0G2K062_DANRE	212.420,00	100,00%	2	2	2	0,01%	1,23%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=lamb2 PE=4 SV= B7ZDA6_DANRE	196.598,90	99,80%	1	1	1	0,00%	2,02%
30dpa_2	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1 B3DG37_DANRE,HBB1_D_	16.389,00	100,00%	3	3	4	0,01%	28,40%
30dpa_2	Ba1 protein OS=Danio rerio GN=ba1 PE=1 SV=1 B3DG37_DANRE,HBB1_D_	16.389,00	100,00%	3	3	4	0,01%	28,40%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=tnnt E7EY99_DANRE,F1R0K6_I	33.942,20	97,70%	1	1	1	0,00%	6,79%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=tnnt E7EY99_DANRE,F1R0K6_I	33.942,20	97,70%	1	1	1	0,00%	6,79%
30dpa_2	Uncharacterized protein (Fragment) OS=Danio rerio GN=tnnt E7EY99_DANRE,F1R0K6_I	33.942,20	97,70%	1	1	1	0,00%	6,79%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV F6NHQ1_DANRE,F6NHV5_	94.381,10	99,70%	1	1	1	0,00%	2,07%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV F6NHQ1_DANRE,F6NHV5_	94.381,10	99,70%	1	1	1	0,00%	2,07%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=slmapa PE=4 SV F6NHQ1_DANRE,F6NHV5_	94.381,10	99,70%	1	1	1	0,00%	2,07%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=utrn PE=1 SV=1 A0A0H2UKN4_DANRE	299.775,50	100,00%	2	2	2	0,01%	1,18%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=col12a1a PE=1 S X1WF98_DANRE	321.902,50	100,00%	1	1	1	0,00%	0,71%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=lama4 PE=1 SV= F1QYU0_DANRE	207.483,80	97,80%	1	1	1	0,00%	0,64%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=palm2 PE=4 SV= A0A0R4ISM7_DANRE,E7F	92.198,80	100,00%	2	4	4	0,01%	2,67%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=palm2 PE=4 SV= A0A0R4ISM7_DANRE,E7F	92.198,80	100,00%	2	4	4	0,01%	2,67%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=anxa11b PE=4 S F8W2N0_DANRE,Q804G3	51.462,90	99,20%	1	1	1	0,00%	3,30%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=anxa11b PE=4 S F8W2N0_DANRE,Q804G3	51.462,90	99,20%	1	1	1	0,00%	3,30%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=actn2b PE=1 SV= E9QFR8_DANRE	103.214,90	100,00%	7	7	10	0,04%	10,50%
30dpa_2	Malate dehydrogenase OS=Danio rerio GN=mdh2 PE=1 SV=1 Q7T334_DANRE	35.420,30	99,10%	0	0	0	0,00%	0,00%
30dpa_2	Keratin, type II cytoskeletal 8 OS=Danio rerio GN=kr8 PE=1 S K2C8_DANRE	57.761,30	100,00%	2	2	3	0,01%	4,23%
30dpa_2	Myosin light chain 2 Mlc2a OS=Danio rerio GN=myl7 PE=1 SV Q801M3_DANRE	19.021,00	100,00%	3	3	6	0,02%	23,80%
30dpa_2	Inner membrane protein, mitochondrial (Mitofilin) OS=Danio rerio Q6PFS4_DANRE	83.201,30	55,40%	0	0	0	0,00%	0,00%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=sdprb PE=1 SV= F1R665_DANRE	51.441,60	100,00%	0	0	0	0,00%	0,00%
30dpa_2	Myoglobin OS=Danio rerio GN=mb PE=1 SV=1 HOWE38_DANRE,MYG_D_	15.583,60	100,00%	3	3	6	0,02%	42,50%
30dpa_2	Myoglobin OS=Danio rerio GN=mb PE=1 SV=1 HOWE38_DANRE,MYG_D_	15.583,60	100,00%	3	3	6	0,02%	42,50%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE= A0A0R4IE46_DANRE,A0AI	101.943,10	99,70%	1	1	1	0,00%	1,08%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE= A0A0R4IE46_DANRE,A0AI	101.943,10	99,70%	1	1	1	0,00%	1,08%
30dpa_2	Uncharacterized protein OS=Danio rerio GN=zgc:112265 PE= A0A0R4IE46_DANRE,A0AI	101.943,10	99,70%	1	1	1	0,00%	1,08%

