

Supplemental table 2: Identifications and calculation of Fold<sub>2</sub>Change of proteins shared in T50 and T90 groups obtained from LC-MS/MS of control and endotoxemic shock rat plasma samples. NA corresponds to missing values.

Accession	Peptide Unique	Coverage	Area.C90.T0.2
Q6MG90	41	80	167000
D3ZQ25	30	38	160000
A0A0G2JXI1	40	23	44900
Q8CHN8	43	53	226000
P31394	12	33	28300
B1H260	42	56	169000
P18420	15	75	27800
Q5M890	22	64	508000
Q63313	25	49	51400
P16296	11	24	137000
Q5M8C3	23	55	137000
B5DEH7	94	82	7,00E+06
M0R816	8	44	38100
Q4KM75	18	47	428000
Q5EBA7	19	26	125000
A0A0G2JYC4	25	58	632000
M0RA79	5	60	103000
Q5FWS5	33	52	1030000
Q63207	12	29	121000
Q7TQ11	28	34	2220000
Q5EBC0	113	84	6880000
F1LPW0	69	63	12900000
F1LWS4	36	62	1280000
Q5FVN3	13	68	65900
Q80ZA3	55	74	2700000
F1LTN6	22	70	2220000
P58775	4	35	404000
G3V8D4	47	90	55700000
D3ZPI8	15	86	1320000
D3ZWD6	55	78	2450000
G3V6G1	13	56	791000
D3ZFH5	17	23	84600
Q5GAM5	15	61	868000
M0R485	44	69	4780000
Q5I0M3	39	89	6840000
G3V9R2	265	93	80400000
F7FMS0	19	29	168000
A0A0G2K975	75	76	5780000
F7FHF3	12	20	303000
F1LTY5	2	28	61500
G3V9J1	36	68	4260000
Q5BK77	15	61	762000
Q5M7T5	38	54	1920000
O08619	43	46	1080000
A0A0G2K7X7	15	20	61000
P08649	11	86	NA
Q64240	20	42	NA
D4ACR1	8	83	NA
F1LXY6	11	100	NA

F1LZ11	6	55	NA
P07151	8	81	NA
M0R9U2	4	39	NA
M0R4C5	4	30	NA

	Area.C90.T0.3	Area.C90.T0.4	Area.C90.T0.5
	356000	1310000	3430000
	412000	757000	1650000
	115000	47300	701000
	1220000	553000	3560000
	230000	53300	329000
	422000	2250000	1600000
	96000	105000	352000
	3580000	4570000	4380000
	251000	179000	415000
	898000	654000	1460000
	114000	57600	725000
	10500000	9130000	32300000
NA		34000	208000
	770000	2110000	4800000
	86600	452000	926000
	322000	1680000	983000
NA	NA		253000
	1030000	371000	2140000
	116000	121000	717000
	2630000	4950000	11700000
	6250000	7420000	18600000
	19300000	2,50E+07	1,04E+08
	1340000	1340000	3390000
	27400	38400	18400
	2380000	3210000	6440000
	4670000 NA		30100000
	93300 NA		512000
	1,00E+08	1,28E+08	1,93E+08
	1100000	1740000	4260000
	1030000	2590000	7420000
	393000	1250000	3920000
	31800 NA		154000
	1020000	343000	1560000
	6860000	4420000	15400000
	7840000	9450000	18200000
	85400000	97900000	2,24E+08
	137000	91900	272000
	4330000	2030000	10400000
	156000	20500	183000
NA	NA		415000
	4110000	2690000	10100000
	457000	221000	1540000
	297000	2130000	1580000
NA		79300	297000
NA		35000 NA	
	48300	415000	3390000
	73300	241000	298000
	25800	61100	661000
	69000	363000	2750000

	27900	43300	612000
	60300	207000	401000
NA		42400	159000
NA		88700	266000

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Area.C90.T0.7	Area.C90.T0.8	Area.C90.T90.2
21300000	23400	23700000
1,20E+07	74600	16400000
4480000	118000	4320000
14300000 NA		21400000
2620000	848000	2070000
11500000 NA		11900000
1950000 NA		1930000
54900000 NA		2,40E+07
4810000 NA		1770000
6600000	137000	4650000
2590000	255000	4270000
80600000	14600000	1,98E+08
2060000 NA		987000
29500000	3160000	10700000
3210000	245000	3090000
8650000	349000	12500000
1520000	79500	2020000
14200000	11300000	19800000
2970000	733000	2020000
48900000	13400000	35800000
5,30E+07	6610000	1,00E+08
5,04E+08	35900000	1,73E+08
19300000	1460000	1,60E+07
150000	253000	768000
1,80E+07	6710000	30500000
1,26E+08	12600000	22600000
8,00E+05	77400	3900000
7,67E+08	70300000	5,29E+08
11200000	3880000	12500000
25400000	1,20E+07	22700000
20300000	4020000	7160000
593000	920000	712000
9200000	5120000	7190000
33700000	23100000	39500000
70900000	28600000	52800000
6,94E+08	2,73E+08	6,20E+08
832000	1310000	1240000
43500000	23100000	42100000
3170000	1530000	2200000
530000	220000	369000
23200000	3120000	22600000
4140000	3820000	3630000
6320000	2790000	8710000
5,00E+06 NA		4770000
186000	259000	159000
13800000 NA		22600000
2230000	8470	10600000
278000 NA		341000
4490000	377000	1100000

NA	2050000	135000	966000
		26100	1140000
	1430000	246000	861000
	934000	206000	1290000
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	Area.C90.T90.3	Area.C90.T90.4	Area.C90.T90.5
	1080000	1410000 NA	
	468000	1210000	86900
	13700	4040000	23800
	501000	8670000	236000
	87900	2380000	175000
	411000	1160000	106000
	24700	426000 NA	
	2280000	448000	41800
	32300	1540000 NA	
	170000	4120000	779000
	618000	4550000 NA	
	6810000	65400000	15200000
	30800	139000	91300
	407000	11300000	6210000
	121000	7780000	169000
	252000	5810000	697000
NA		244000	374000
	726000	13400000	2800000
	62700	4320000	1030000
	7480000	1,50E+07	2420000
	3,30E+07	52200000	11500000
	17900000	1,68E+08	43700000
	456000	13200000	1620000
NA		7050000	1970000
	1060000	3,50E+07	1430000
	3600000 NA		1,80E+07
NA		540000	52600
	4,70E+07	1,85E+08	76700000
	1080000	14200000	2200000
	1010000	36300000	5090000
	447000	10400000	5340000
	256000	2870000	9940
	70400	22200000	6870000
	1520000	63100000	5470000
	4350000	9,60E+07	29100000
	50200000	7,76E+08	1,98E+08
	74900	4600000	542000
	1640000	56300000	12500000
	114000	7250000	1200000
	7650	726000	575000
	1040000	18400000	7340000
	26500	9060000	3630000
	831000	4080000	1020000
	169000 NA	NA	
NA		961000 NA	
	280000	831000	2,00E+05
	245000	494000 NA	
NA		988000	170000
	178000	2450000	1090000



	31900	369000	111000
	33500	678000	149000
	41400	224000	51300
NA		239000	0

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Area.C90.T90.7	Area.C90.T90.8	Area.L90.T0.2
7560000 NA		7320000
5920000 NA		8230000
3640000 NA		3160000
1,50E+07 NA		13900000
3200000	27300	2550000
9030000 NA		7270000
1060000 NA		721000
21900000 NA		12800000
3120000 NA		1210000
4690000 NA		1450000
2560000 NA		2150000
95100000	1940000	75800000
1300000 NA		953000
24200000	539000	7020000
4050000 NA		1680000
7890000 NA		2020000
743000 NA		335000
12200000	1580000	1,40E+07
2670000	76000	1160000
35900000	665000	43200000
75100000	626000	31100000
3,67E+08	8320000	1,27E+08
27400000 NA		11700000
585000	61900 NA	
18400000	441000	16100000
1,04E+08	1130000	26700000
728000 NA		2450000
6,88E+08	9980000	6,47E+08
8930000	395000	5290000
19800000	411000	11500000
16600000	1950000	6300000
517000 NA		533000
14100000	2250000	3360000
30100000	1770000	30400000
73500000	2410000	30100000
6,41E+08	2,80E+07	3,20E+08
1080000 NA		468000
55700000	2890000	20300000
3810000 NA		2630000
330000	19700	415000
34400000	52900	20700000
1140000	1220000	1390000
5590000	135000	3390000
1390000	91300	2640000
94000	15300	34500
6970000 NA		2480000
2680000 NA		1510000
194000 NA	NA	
3220000 NA		2140000

	1880000	NA		435000
NA		NA		102000
	854000	NA	NA	
	856000	NA		584000
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Area.L90.T0.3	Area.L90.T0.5	Area.L90.T0.6
26500000	17100000	42300000
1,60E+07	21300000	16300000
6650000	12900000	3930000
9280000	29600000	16400000
1250000	3280000	1180000
8850000	2,40E+07	12200000
634000	1090000	3310000
86800000	1,60E+08	6,50E+07
6090000	8960000	1660000
1800000	3070000	4820000
7740000	28700000	4820000
1,01E+08	1,33E+08	1,47E+08
69900	883000	1030000
9250000	12400000	17100000
2680000	10100000	5310000
7560000	22600000	6,00E+06
619000	1040000	1110000
10200000	22100000	9850000
2540000	3790000	1980000
21400000	53700000	33900000
43500000	1,67E+08	73600000
2,35E+08	3,16E+08	3,18E+08
14400000	30800000	9240000
103000	834000	202000
25100000	74900000	35800000
29400000	69800000	1,12E+08
1050000	1540000	202000
3,81E+08	1,17E+09	8,26E+08
14500000	2,50E+07	6620000
2,20E+07	41200000	8300000
9580000	1,40E+07	9590000
212000	827000	739000
1650000	10400000	3450000
27400000	91500000	60700000
2,20E+07	95600000	52900000
5,03E+08	1,14E+09	7,62E+08
1020000	4340000	653000
21700000	52900000	20100000
1070000	3350000	1570000
430000	917000	545000
20400000	57900000	23800000
4250000	7390000	3610000
25600000	7760000	4900000
15800000	9930000	14700000
125000	1180000	179000
7870000	3200000	23500000
8240000	13300000	5180000
1410000	2850000	1480000
7,00E+06	7780000	1090000

1330000	3670000	565000
170000	1640000	2390000
672000	1180000	1960000
749000	880000	2330000

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Area.L90.T0.7	Area.L90.T0.8	Area.L90.T90.2
2,00E+07	5350000	740000
14100000	2630000	1310000
8110000	903000	931000
17300000	4800000	3400000
2360000	1030000	1430000
10200000	3390000	5,00E+05
1770000 NA		202000
15500000	30300000	540000
9080000	1890000	1390000
4240000	1350000	999000
4990000	3580000	694000
1,85E+08	36200000	42400000
1740000	146000	4,00E+05
13100000	4030000	4720000
4730000	1810000	1500000
11500000	3010000	497000
24900	15000	281000
17200000	3150000	25800000
2980000	384000	1470000
65300000	14300000	10600000
6,27E+08	34100000	4,00E+07
1,70E+08	91400000	72300000
9950000	7150000	3830000
334000	207000	1300000
20100000	12200000	18900000
95900000	27900000	8660000
3080000	2040000	1,00E+07
5,33E+08	2,97E+08	2,58E+08
1,40E+07	4800000	3570000
30300000	6680000	10300000
8070000	2590000	4920000
4450000	389000	1600000
4610000	3330000	8430000
36900000	25200000	34200000
73100000	21200000	3,50E+07
9,23E+08	3,18E+08	1,36E+08
1710000	470000	1960000
65600000	16300000	16100000
4500000	739000	6410000
279000	107000	712000
13600000	15500000	16600000
4780000	1320000	869000
20400000	2570000	427000
3650000	279000	12700
960000 NA		839000
20800000	1140000	249000
5490000	2630000	241000
NA	63600	155000
635000	1360000	1560000

482000	426000	154000
8420000	324000 NA	
1330000	657000	171000
341000	59800	224000

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Area.L90.T90.3	Area.L90.T90.5	Area.L90.T90.6
3910000	1040000	15700000
5540000	1770000	6870000
1160000	3320000	3670000
14400000	10900000	12600000
1930000	2220000	308000
4200000	937000	6220000
1470000	534000	2510000
4630000	3270000	23600000
3820000	2,00E+06	4370000
2540000	1780000	4290000
3240000	4250000	1640000
1,22E+08	67600000	97400000
330000	593000	627000
18200000	9500000	7700000
2550000	3660000	4080000
5040000	3170000	4560000
287000	874000	352000
27800000	19400000	8490000
1,00E+06	2080000	1390000
3,10E+07	20600000	27300000
70300000	48300000	56900000
3,52E+08	1,62E+08	1,68E+08
12200000	12300000	7100000
1240000	575000	298000
41600000	27900000	12100000
1,04E+08	82300000	47700000
1540000	811000	819000
3,20E+08	3,50E+08	4,04E+08
13800000	9410000	2750000
30100000	20600000	2470000
16800000	10700000	5900000
3010000	2550000	343000
18300000	15900000	5070000
55400000	56400000	37800000
5,50E+07	1,00E+08	32500000
8,38E+08	7,08E+08	4,76E+08
1310000	1810000	372000
66300000	46800000	1,30E+07
4370000	5880000	1120000
1200000	1020000	305000
52500000	31300000	21500000
1150000	2600000	306000
14600000	3570000	3140000
560000	26300	4110000
966000	979000	401000
4950000	405000	12100000
904000	385000	2430000
1080000	1,30E+07	1380000
4590000	2350000	761000



763000	457000	282000
597000	655000	746000
1100000	913000	1020000
1420000	935000	1140000

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	Area.L90.T90.7	Area.L90.T90.8	Log2FC.RC2.T90
	173000	607000	7,148895146
	270000	618000	6,6794801
	426000	792000	6,588172057
	2580000	3,00E+06	6,565144214
	676000	1260000	6,192684904
	292000	704000	6,137794517
	103000	179000	6,117372154
	355000	1260000	5,562062099
	1130000	775000	5,105837191
	1240000	1530000	5,084982918
	908000	1660000	4,961988272
	50800000	31600000	4,822001698
	445000	228000	4,695187182
	4650000	5630000	4,64385619
	1790000	1890000	4,627606838
	1770000	1760000	4,305859726
NA		187000	4,29363905
	11800000	9460000	4,264784188
	1600000	1510000	4,06127634
	5860000	8160000	4,011328006
	40600000	30400000	3,861447625
	7,20E+07	97400000	3,745329067
	4640000	5020000	3,64385619
	7340000	2310000	3,542755941
	8880000	11600000	3,49777793
	38100000	43800000	3,347691191
	5280000	3130000	3,271046926
	87100000	1,96E+08	3,24751849
	4610000	5730000	3,24331826
	9960000	11600000	3,211838643
	5220000	5100000	3,178209988
	3510000	1810000	3,073147673
	15100000	12200000	3,050224823
	20800000	31300000	3,04677013
	43900000	41100000	2,948469699
	4,92E+08	4,31E+08	2,947000809
	1500000	878000	2,883806982
	38100000	26300000	2,864678835
	3350000	3320000	2,860113825
NA		196000	2,584962501
	6210000	14800000	2,407397437
	890000	1160000	2,252106645
	4050000	2690000	2,181566408
NA	NA		2,142957954
	455000	122000	1,382145618
	859000	507000	NA
	231000	262000	NA
NA		672000	NA
	212000	6,00E+05	NA

40700	129000	NA
1630000	128000	NA
249000	1020000	NA
1280000	199000	NA

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	Log2FC.RC3.T90	Log2FC.RC4.T90	Log2FC.RC5.T90
	1,601082166	0,106128351 NA	
	0,183864192	0,676641842	-4,246966037
	-3,069386063	6,416371299	-4,880380966
	-1,283998639	3,970680608	-3,915018476
	-1,387698791	5,48068223	-0,910732662
	-0,038104605	-0,955800196	-3,915935735
	-1,958523364	2,020464103 NA	
	-0,650925763	-3,350623528	-6,711284117
	-2,958081294	3,104898858 NA	
	-2,401180699	2,655281797	-0,906273136
	2,438573014	6,303653923 NA	
	-0,624662625	2,84060387	-1,087462841
NA		2,031478231	-1,187896763
	-0,919829651	2,421007869	0,371558863
	0,482568117	4,105375477	-2,453988947
	-0,353636955	1,790076931	-0,49603276
NA	NA		0,563900885
	-0,504602884	5,174670004	0,387816031
	-0,887587457	5,15795236	0,522599313
	1,507975471	1,59946207	-2,273429577
	2,40053793	2,814558715	-0,69366876
	-0,10864126	2,748461233	-1,250878344
	-1,555127271	3,300233024	-1,06529146
NA		7,520373136	6,742346053
	-1,166897309	3,44670972	-2,171045541
	-0,375425643 NA		-0,74176658
NA	NA		-3,283009106
	-1,089267338	0,531381461	-1,331302365
	-0,026472211	3,028731719	-0,953349907
	-0,028289044	3,808945545	-0,543753531
	0,185745519	3,056583528	0,445986088
	3,00904514 NA		-3,953540689
	-3,856849913	6,01620729	2,13876407
	-2,174137253	3,83552173	-1,493317613
	-0,849838253	3,344648171	0,677080703
	-0,766548706	2,986675887	-0,177998302
	-0,871138269	5,645425189	0,9946862
	-1,40067121	4,79358329	0,265344567
	-0,452512205	8,466213275	2,713118852
NA	NA		0,47045062
	-1,982554866	2,774027688	-0,460503325
	-4,1081299	5,357392776	1,237039197
	1,484385546	0,937715722	-0,631355406
NA	NA	NA	
NA		4,779109604 NA	
	2,535331733	1,001737141	-4,083213368
	1,740896646	1,035477895 NA	
NA		4,015266757	-1,959115525
	1,367208974	2,754740296	-1,335103484

	0,193291302	3,091181886	-2,462971976
	-0,847996907	1,711654506	-1,428289906
NA		2,401362562	-1,631996035
NA		1,430004609	#NOM?

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Log2FC.RC7.T90	Log2FC.RC8.T90	Log2FC.RL2.T90
-1,494395291 NA		-3,306246473
-1,019365325 NA		-2,651325619
-0,299560282 NA		-1,763071486
0,068947354 NA		-2,031478231
0,288505093	-4,957091409	-0,8344821
-0,348835968 NA		-3,861955364
-0,879409859 NA		-1,835643966
-1,325875279 NA		-4,567040593
-0,624490865 NA		0,200077835
-0,492878102 NA		-0,537496317
-0,016808288 NA		-1,631329092
0,238665502	-2,911839812	-0,838133584
-0,664132714 NA		-1,252476214
-0,285707907	-2,55156738	-0,572684171
0,335348611 NA		-0,163498732
-0,132674833 NA		-2,023037536
-1,032637208 NA		-0,253590965
-0,219009782	-2,838326309	0,881944238
-0,153623189	-3,269741875	0,34169135
-0,445850621	-4,33273485	-2,026967048
0,502820548	-3,400415709	0,36308542
-0,457643671	-2,109328411	-0,812760945
0,505575046 NA		-1,611092233
1,963474124	-2,03112607 NA	
0,03170886	-3,927462206	0,231325546
-0,276840205	-3,479029056	-1,624400812
-0,13606155 NA		2,029146346
-0,156818013	-2,816412969	-1,326394647
-0,326766652	-3,296132094	-0,567343648
-0,359328067	-4,867752202	-0,158989524
-0,290296486	-1,043721377	-0,356693513
-0,197867824 NA		1,585864467
0,615989396	-1,186218809	1,327071398
-0,162985104	-3,706071586	0,169925001
0,051958622	-3,568910095	0,217591435
-0,114611306	-3,285402219	-1,234465254
0,376375879 NA		2,066273219
0,356661927	-2,998751454	-0,334419039
0,265308157 NA		1,285261557
-0,683526335	-3,481235989	0,778765905
0,56828376	-5,882134497	-0,318447526
-1,860596943	-1,64669149	-0,677656801
-0,177076275	-4,36923381	-2,988977298
-1,846843212 NA		-7,699565622
-0,984569959	-4,08134854	4,604002544
-0,985437706 NA		-3,316122473
0,265189291 NA		-2,647443498
-0,519028231 NA	NA	
-0,479654757 NA		-0,456064768

	-0,124891248 NA		-1,49808505
NA	NA	NA	
	-0,743707172 NA	NA	
	-0,125811753 NA		-1,382469637

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Log2FC.RL3.T90	Log2FC.RL5.T90	Log2FC.RL6T90
-2,760751847	-4,039340892	-1,429893104
-1,530114024	-3,589032165	-1,24648996
-2,519229535	-1,958115919	-0,098749249
0,633872101	-1,441269041	-0,380272081
0,626672753	-0,563136138	-1,937784604
-1,075288127	-4,678841548	-0,971894662
1,21326141	-1,029416488	-0,399143853
-4,228610944	-5,612637459	-1,461652859
-0,67286959	-2,163498732	1,396450038
0,49683159	-0,786361414	-0,168055499
-1,256339753	-2,755515991	-1,555337332
0,272525855	-0,976331094	-0,593822478
2,239101664	-0,574381333	-0,716106989
0,97641318	-0,384340702	-1,151065974
-0,071735754	-1,464439739	-0,380142709
-0,584962501	-2,833768027	-0,395928676
-1,108888673	-0,250878344	-1,656912343
1,446515731	-0,187989717	-0,214359171
-1,344828497	-0,86561432	-0,510415547
0,534657419	-1,382277751	-0,312384322
0,692509288	-1,789753009	-0,371277114
0,582914672	-0,963930745	-0,920565533
-0,239187664	-1,324272035	-0,380073827
3,589623878	-0,536485428	0,560957038
0,728896164	-1,424700597	-1,56495254
1,822695468	0,237665394	-1,231437561
0,552541023	-0,925156531	2,019508159
-0,251719093	-1,741081703	-1,031786489
-0,071384633	-1,409661467	-1,267399598
0,452259963	-1	-1,748600295
0,810363672	-0,387816031	-0,700815861
3,827627317	1,624538013	-1,107365788
3,471305719	0,612443237	0,555389385
1,015710083	-0,698076581	-0,683310282
1,321928095	0,064917477	-0,702828004
0,736391844	-0,687212559	-0,678829424
0,36099766	-1,261705345	-0,81178037
1,611313828	-0,176759193	-0,628683878
2,030022483	0,811655059	-0,487265827
1,480625841	0,153575513	-0,837446987
1,363748271	-0,887400691	-0,146624914
-1,88582898	-1,507062741	-3,560395279
-0,810175441	-1,120132578	-0,64201719
-4,818353921	-8,560587108	-1,838605856
2,950095094	-0,269406095	1,163642649
-0,668935111	-2,982078092	-0,957653709
-3,18824966	-5,11042399	-1,091995784
-0,38466385	2,189477799	-0,100928909
-0,608860768	-1,727109398	-0,518359776



-0,801671284	-3,005513993	-1,002555705
1,812196185	-1,324129003	-1,679763083
0,710970386	-0,370100094	-0,942284502
0,922853306	0,087462841	-1,03129613

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	Log2FC.RL7.T90	Log2FC.RL8.T90
	-6,853084152	-3,13977047
	-5,706591945	-2,089384056
	-4,250776579	-0,189225557
	-2,745329067	-0,678071905
	-1,803691708	0,290779396
	-5,126456973	-2,267637939
	-4,103033118 NA	
	-5,448305381	-4,587822155
	-3,006369525	-1,286118019
	-1,773724144	0,180572246
	-2,458275613	-1,108776346
	-1,864624869	-0,196065139
	-1,967210065	0,643065455
	-1,49426419	0,482355084
	-1,401880596	0,062396537
	-2,699812596	-0,774188058
NA		3,640003864
	-0,543621705	1,586488355
	-0,897240426	1,975370333
	-3,478110422	-0,80937409
	-3,948913811	-0,165700416
	-1,239465935	0,091727607
	-1,10057172	-0,510255878
	4,457860055	3,480190179
	-1,17856392	-0,072756342
	-1,331739818	0,650665748
	0,777607579	0,617593505
	-2,613390909	-0,599609277
	-1,602588171	0,255500733
	-1,605100146	0,796204798
	-0,628518867	0,977545149
	-0,342334306	2,218147637
	1,711709894	1,873287065
	-0,827037288	0,312738923
	-0,735650466	0,955074129
	-0,907672332	0,438661104
	-0,189033824	0,901560183
	-0,783904817	0,690190835
	-0,425763906	2,167536972
NA		0,873242858
	-1,130941478	-0,06667104
	-2,425133377	-0,186413124
	-2,332575339	0,065837813
NA	NA	
	-1,077167861 NA	
	-4,597781587	-1,168976172
	-4,570841392	-3,327424083
NA		3,401362562
	-1,582692327	-1,180572246

-3,565932447	-1,723482365
-2,368948269	-1,339850003
-2,417208598	0,634603877
1,908300166	1,734551041

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Description
Complement C4B (Chido blood group) OS=Rattus norvegicus (Rat) OX=10116 GN=C4b PE=1 SV=1
Fibulin-1 OS=Rattus norvegicus (Rat) OX=10116 GN=Fbln1 PE=1 SV=1
F5/8 type C domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=F5 PE=1 SV=1
Mannan-binding lectin serine protease 1 OS=Rattus norvegicus (Rat) OX=10116 GN=Masp1 PE=1 SV=2
Vitamin K-dependent protein C OS=Rattus norvegicus (Rat) OX=10116 GN=Proc PE=2 SV=1
Coagulation factor XIII B chain OS=Rattus norvegicus (Rat) OX=10116 GN=F13b PE=1 SV=1
Proteasome subunit alpha type-1 OS=Rattus norvegicus (Rat) OX=10116 GN=Psma1 PE=1 SV=2
Apolipoprotein N OS=Rattus norvegicus (Rat) OX=10116 GN=Apon PE=1 SV=1
Lipopolysaccharide-binding protein OS=Rattus norvegicus (Rat) OX=10116 GN=Lbp PE=2 SV=1
Coagulation factor IX OS=Rattus norvegicus (Rat) OX=10116 GN=F9 PE=2 SV=2
SERPIN domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=Serpina4 PE=1 SV=1
C1r protein OS=Rattus norvegicus (Rat) OX=10116 GN=C1r PE=1 SV=1
Ig-like domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=ENSRNOG00000055193 PE=1 SV=1
CD5 antigen-like OS=Rattus norvegicus (Rat) OX=10116 GN=Cd5l PE=1 SV=1
HGF activator OS=Rattus norvegicus (Rat) OX=10116 GN=Hgfac PE=1 SV=1
Complement factor H-related protein B OS=Rattus norvegicus (Rat) OX=10116 GN=LOC100361907 PE=1 SV=1
Ig-like domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=ENSRNOG00000048425 PE=1 SV=1
Lysyl oxidase-like 1 OS=Rattus norvegicus (Rat) OX=10116 GN=Loxl1 PE=2 SV=1
Coagulation factor X OS=Rattus norvegicus (Rat) OX=10116 GN=F10 PE=2 SV=1
SMB domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=Vtn PE=1 SV=1
Inter alpha-trypsin inhibitor heavy chain 4 OS=Rattus norvegicus (Rat) OX=10116 GN=Itih4 PE=1 SV=1
Immunoglobulin heavy constant mu OS=Rattus norvegicus (Rat) OX=10116 GN=Ighm PE=1 SV=1
Uncharacterized protein OS=Rattus norvegicus (Rat) OX=10116 GN=Cfhr2 PE=1 SV=3
SCY domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=Ccl9 PE=1 SV=1
SERPIN domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=Serpinf1 PE=1 SV=1
Uncharacterized protein OS=Rattus norvegicus (Rat) OX=10116 GN=ENSRNOG00000049829 PE=1 SV=2
Tropomyosin beta chain OS=Rattus norvegicus (Rat) OX=10116 GN=Tpm2 PE=1 SV=1
Apolipoprotein C-II (Predicted) OS=Rattus norvegicus (Rat) OX=10116 GN=Apoc2 PE=1 SV=1
Lipocln_cytosolic_FA-bd_dom domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=C8g PE=1 SV=1
MACPF domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=C8a PE=1 SV=1
Immunoglobulin joining chain OS=Rattus norvegicus (Rat) OX=10116 GN=Jchain PE=1 SV=1
Inter-alpha-trypsin inhibitor heavy chain 2 OS=Rattus norvegicus (Rat) OX=10116 GN=Itih2 PE=1 SV=3
RNAse_Pc domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=Ang2 PE=1 SV=1
PGRP domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=Pglyrp2 PE=1 SV=2
Complement component factor h-like 1 OS=Rattus norvegicus (Rat) OX=10116 GN=Cfhr1 PE=1 SV=1
Complement factor H OS=Rattus norvegicus (Rat) OX=10116 GN=Cfh PE=1 SV=1
Macrophage-stimulating 1 OS=Rattus norvegicus (Rat) OX=10116 GN=Mst1 PE=3 SV=1
Similar to complement factor H-related protein OS=Rattus norvegicus (Rat) OX=10116 GN=RGD1564614 PE=1 SV=1
SERPIN domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=Serpinf2 PE=1 SV=1
Ig-like domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=ENSRNOG00000048167 PE=1 SV=1
Uncharacterized protein OS=Rattus norvegicus (Rat) OX=10116 GN=ENSRNOG00000042228 PE=1 SV=3
Retinoic acid receptor responder (Tazarotene induced) 2 OS=Rattus norvegicus (Rat) OX=10116 GN=Rarres2 PE=1 SV=1
SERPIN domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=Serpinc1 PE=1 SV=1
Coagulation factor XIII A chain OS=Rattus norvegicus (Rat) OX=10116 GN=F13a1 PE=2 SV=3
Complement C7 OS=Rattus norvegicus (Rat) OX=10116 GN=C7 PE=1 SV=1
Complement C4 OS=Rattus norvegicus (Rat) OX=10116 GN=C4 PE=1 SV=3
Protein AMBP OS=Rattus norvegicus (Rat) OX=10116 GN=Ambp PE=1 SV=1
Ig-like domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=ENSRNOG00000042996 PE=1 SV=1
Ig-like domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=ENSRNOG00000058824 PE=1 SV=1

Ig-like domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=ENSRNOG00000042319 PE=1 SV=  
Beta-2-microglobulin OS=Rattus norvegicus (Rat) OX=10116 GN=B2m PE=1 SV=1  
Ig-like domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=ENSRNOG00000050708 PE=1 SV=  
Ig-like domain-containing protein OS=Rattus norvegicus (Rat) OX=10116 GN=ENSRNOG00000047915 PE=4 SV=

=2

=1

. SV=1

SV=1

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E=1 SV=1

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