

Uniprot Acc	proteins	121	113	ratio(113/121)	114	ratio(114/121)	115	ratio(115/121)	116	ratio(116/121)	117	ratio(117/121)	118	ratio(118/121)	119	ratio(119/121)
C0JYY2	Apolipoprot	510456.9	587089.8	1.15	722432.2	1.42	577726.5	1.13	553022.3	1.08	631905.3	1.24	553297.9	1.08	487383.6	0.95
V9HWA9	Epididymis	1086362.4	1038495.5	0.96	987768	0.91	1014543.4	0.93	1059000	0.97	1036688.9	0.95	1058344.2	0.97	1125990.4	1.04
A0A140TA	Complement	10069.7	10298.9	1.02	11246.3	1.12	10190.4	1.01	9787.3	0.97	10555.8	1.05	9545.9	0.95	7018.6	0.70
A0A140TA	Complement C4-A OS=Homo sapiens GN=C4A PE=1 SV=1															
P0C0L4	Complement	379640.7	354515.7	0.93	358550.6	0.94	353533.8	0.93	371096.1	0.98	369917.8	0.97	392042.8	1.03	352593.5	0.93
A8K5A4	cDNA FLJ7	181927.1	184669.2	1.02	167773	0.92	172794.9	0.95	189477.9	1.04	181011.4	0.99	192480.9	1.06	188787.6	1.04
P08603	Complement	127138.9	139326.8	1.10	138368.7	1.09	152055.5	1.20	147415.7	1.16	134161.2	1.06	124277.4	0.98	143670	1.13
A8K5T0	cDNA FLJ7	1595.6	2166.5	1.36	1478.4	0.93	1667	1.04	1764.5	1.11	1600.3	1.00	1443.2	0.90	1747.7	1.10
Q7Z7Q0	APOB prote	369.3	378.4		434.3	1.18	176	0.48	338.2	0.92	290.4	0.79	163	0.44	248.4	0.67
E9PFZ2	Ceruloplasm	232	305.6	1.32	1344.1	5.79	340.1	1.47	294.9	1.27	245.3	1.06	225.6	0.97	226.5	0.98
P01023	Alpha-2-mac	155893	159242.4	1.02	150022.1	0.96	148585.9	0.95	127199.8	0.82	139287.4	0.89	196290.4	1.26	150833.5	0.97
B4E1Z4	cDNA FLJ5	179991.2	168887.5	0.94	175330.4	0.97	170379.3	0.95	181366.8	1.01	186553.2	1.04	182496	1.01	179532.4	1.00
P02768	Serum album	374998.6	393627.8	1.05	415295.5	1.11	553515.2	1.48	401168.8	1.07	313800.8	0.84	273544.9	0.73	216497.6	0.58
P00747	Plasminoger	110712.8	123328	1.11	118392.6	1.07	116615.2	1.05	117254.4	1.06	115644.8	1.04	104325.5	0.94	105844.6	0.96
P02790	Hemopexin	407277.3	374618.6	0.92	386591	0.95	389342.2	0.96	396054.9	0.97	408398.8	1.00	392817.7	0.96	388013.2	0.95
P00738	Haptoglobin	185933.7	185102	1.00	162600.1	0.87	155639.9	0.84	279577.3	1.50	162525.6	0.87	258436.5	1.39	410792.3	2.21
A0A024R3I	Apolipoprot	411885.3	338583	0.82	381625.8	0.93	366565	0.89	373110.1	0.91	364223.9	0.88	375076	0.91	391751.5	0.95
P00734	Prothrombin	57305.1	69844.1	1.22	67971.2	1.19	72856.6	1.27	63897.5	1.12	65950.1	1.15	56143.8	0.98	58040.4	1.01
B4E1C2	Kininogen 1	131253.4	125717	0.96	122596.9	0.93	124775.5	0.95	130027.4	0.99	132243.9	1.01	130159.7	0.99	124968.7	0.95
P01031	Complement	93176.8	94890.7	1.02	90259.9	0.97	89862.7	0.96	87867	0.94	94067.8	1.01	91781.4	0.99	92978.4	1.00
D3DNU8	Kininogen 1	7787.9	6418.8	0.82	6744.1	0.87	6171.8	0.79	6016	0.77	6433	0.83	9059.8	1.16	8373.3	1.08
B4E1B2	cDNA FLJ5	83286.1	92039.8	1.11	81713.4	0.98	90197.9	1.08	70892.5	0.85	76534	0.92	83395	1.00	66649.1	0.80
E9KL23	Epididymis	6306.2	8613.5	1.37	5499.6	0.87	5387.8	0.85	4010.7	0.64	5925.8	0.94	6670.1	1.06	4887.9	0.78
A0A024R6I	Alpha-1-anti	165755.8	129336.8	0.78	145830.8	0.88	133687.4	0.81	163476	0.99	143911.9	0.87	158637.4	0.96	254797.8	1.54
A0A0K0K1	Epididymis	866.1	905.6	1.05	975.5	1.13	1107.8	1.28	873	1.01	818.1	0.94	1068.3	1.23	849.7	0.98
D6RF35	Vitamin D-b	203990.6	191458.5	0.94	193095.1	0.95	175613.7	0.86	171725	0.84	181511.4	0.89	200116.5	0.98	203654.9	1.00
V9HWI6	Epididymis	1938.8	1493.4	0.77	1742	0.90	1355.4	0.70	1572.2	0.81	2818.7	1.45	2236	1.15	1573.4	0.81
D3DRR6	Inter-alpha (108810.6	108481.7	1.00	98996.5	0.91	99369.2	0.91	94019.8	0.86	105151.6	0.97	106137.5	0.98	95323.2	0.88
B2RMS9	Inter-alpha (104677.5	108110.8	1.03	103576.1	0.99	102139.3	0.98	101346.1	0.97	109288.1	1.04	106762.4	1.02	106066.8	1.01
A8K2T4	cDNA FLJ7	34968	39819.6	1.14	35815.5	1.02	37899.4	1.08	38127.9	1.09	37936.9	1.08	36598.8	1.05	35816.4	1.02
B7Z539	cDNA FLJ5	53820	54625.6	1.01	50116.6	0.93	55433.1	1.03	58397.6	1.09	56789.4	1.06	51279.7	0.95	46599.2	0.87
P06727	Apolipoprot	132070.1	121784.2	0.92	122790	0.93	118549.5	0.90	111855.3	0.85	125959.3	0.95	118246.5	0.90	98547	0.75
P04003	C4b-binding	89101.1	98140.4	1.10	87061.7	0.98	94314.1	1.06	96031.3	1.08	95727.1	1.07	96432.8	1.08	93592.4	1.05
B2R6W1	cDNA, FLJ5	340	467.6	1.38	371.3	1.09	403.2	1.19	355.5	1.05	366.1	1.08	384.4	1.13	402.6	1.18
B4E1D8	cDNA FLJ5	356.1	419.3	1.18	405.5	1.14	393	1.10	441.7	1.24	454.7	1.28	390.7	1.10	344.8	0.97
P00739	Haptoglobin	2635.5	2606.9	0.99	2512.7	0.95	2255.3	0.86	2714.9	1.03	2471.1	0.94	3397.7	1.29	4403	1.67
B7Z1F8	cDNA FLJ5	5945	6148.8	1.03	5133	0.86	6345.8	1.07	5999.2	1.01	5232.9	0.88	5756.8	0.97	5222.4	0.88
P06396	Gelsolin OS	53072.9	51360.2	0.97	48208.4	0.91	50016.7	0.94	46518.3	0.88	50644.9	0.95	50760.1	0.96	44495	0.84
D9IWP9	Beta-2-glycc	95095.7	95869.3	1.01	94684.3	1.00	92089.5	0.97	92442.1	0.97	97157.4	1.02	95290.3	1.00	97045.8	1.02
A0A024R94	Serpin pepti	76446.5	81924	1.07	75341.2	0.99	73519.1	0.96	66972.7	0.88	79640.7	1.04	77286.2	1.01	59152.2	0.77
B3KS79	cDNA FLJ3	99858	93851	0.94	98190.2	0.98	91704	0.92	98831.5	0.99	103081.2	1.03	95366.8	0.96	132515.7	1.33
P13671	Complement	34641.2	36577.2	1.06	33403.3	0.96	34161.9	0.99	36176	1.04	36281.2	1.05	33438.1	0.97	34559.2	1.00
E9PGN7	Plasma prote	110273.8	100248.8	0.91	112257.1	1.02	98724.4	0.90	106678.1	0.97	114992.3	1.04	110430.4	1.00	101300.1	0.92
P02760	Protein AMI	38024	40442	1.06	36134.6	0.95	39749.4	1.05	40433.6	1.06	41020	1.08	38398	1.01	34490.9	0.91
A8K8Z4	cDNA FLJ7	91.7	91.3	1.00	102.6	1.12	84.4	0.92	91.7	1.00	76	0.83	104.5	1.14	85.7	0.93
B2R8I2	cDNA, FLJ5	53687.5	56488.4	1.05	55925.9	1.04	42986.9	0.80	48963.1	0.91	56480.5	1.05	61840.2	1.15	50180	0.93

H0YAC1	Plasma kallikrein	33082.6	34217.8	1.03	32918	1.00	34183.6	1.03	32889.7	0.99	34399	1.04	32603.1	0.99	29661.8	0.90
P04196	Histidine-rich glycoprotein	661.8	866.1	1.31	992.4	1.50	590.2	0.89	622.6	0.94	688.2	1.04	799.8	1.21	706.1	1.07
P07358	Complement factor D	25346.9	24943.9	0.98	24633.6	0.97	24581.7	0.97	24823.5	0.98	24915.6	0.98	24972.2	0.99	23582.2	0.93
P43652	Afamin OS=	60761.3	62649.1	1.03	58267.1	0.96	59622.5	0.98	58119.3	0.96	61682.8	1.02	62913.2	1.04	54287.5	0.89
P06681	Complement factor B	5096.4	5761.7	1.13	5226.7	1.03	5861.4	1.15	5486.3	1.08	5452.7	1.07	5408.6	1.06	5137.8	1.01
P04275	von Willebrand factor	10196	13089.5	1.28	11473.5	1.13	12553.1	1.23	13082	1.28	13206.5	1.30	13496.9	1.32	11111.4	1.09
A0A024R9C	Thrombospondin-1	21666.3	26614.2	1.23	21742.9	1.00	25643.9	1.18	21725	1.00	27021.2	1.25	22529.8	1.04	20044.7	0.93
P07996	Thrombospondin-1	392.8	557.4	1.42	499.6	1.27	559	1.42	467	1.19	606.6	1.54	606.1	1.54	458.5	1.17
P10909	Clusterin OS=	33598.7	50095.5	1.49	45302.4	1.35	48175.9	1.43	48440.7	1.44	47018.7	1.40	34883.3	1.04	34107.1	1.02
O75882	Attractin OS=	25522.8	27658.2	1.08	25777.2	1.01	25986.1	1.02	25151.2	0.99	26915.2	1.05	24672.1	0.97	23733.1	0.93
E7ETH0	Complement factor D	26360	28620.6	1.09	26667.1	1.01	27252.9	1.03	26287.2	1.00	27657.8	1.05	25695.2	0.97	27742.1	1.05
L8E853	von Willebrand factor															
B3KNX0	cDNA FLJ37966	24353	27673.9	1.14	25972.1	1.07	24866.1	1.02	26889	1.10	25674.2	1.05	23715.2	0.97	25775	1.06
A0A024R46	Fibronectin	8653	14350.7	1.66	12920.5	1.49	17208	1.99	13008.3	1.50	18122.1	2.09	8789.5	1.02	7557	0.87
B4DPQ0	Complement factor D	27059.9	29398.6	1.09	29623.9	1.09	30198.6	1.12	30114.2	1.11	28399.3	1.05	25563.7	0.94	28589.3	1.06
P02649	Apolipoprotein A-II	54616	60795.2	1.11	53606.3	0.98	61843.3	1.13	62405.6	1.14	66560.6	1.22	58524.1	1.07	55467.3	1.02
P07357	Complement factor D	19863.5	21076	1.06	19665.2	0.99	20194.4	1.02	19189.5	0.97	20228.2	1.02	19921.3	1.00	19774.3	1.00
A0A024R03	Complement factor D	31276.8	28186.5	0.90	29862.6	0.95	26992.6	0.86	31248.7	1.00	30833.4	0.99	31964.7	1.02	39963.3	1.28
Q6LAM1	Heavy chain of IgG1	7000	5884.2	0.84	5922.3	0.85	5458.6	0.78	5879.3	0.84	6019	0.86	6552.1	0.94	9038.4	1.29
P05546	Heparin cofactor II	40235.4	42058.8	1.05	39849.4	0.99	38495	0.96	41473.5	1.03	42452	1.06	44464.2	1.11	39529.8	0.98
Q9UNU2	Complement factor D	2086.8	2438.3	1.17	2216	1.06	2211.7	1.06	1819.1	0.87	2243	1.07	1953.8	0.94	2485	1.19
V9HWD8	Epididymis protein 1	87354.8	85465.5	0.98	85173.4	0.98	82675.7	0.95	85248.3	0.98	87032.2	1.00	83186	0.95	82018.4	0.94
A0A0S2Z4I	Protein S isoform 1	20335.7	20521.9	1.01	19479.9	0.96	20695.3	1.02	18904.4	0.93	20998.2	1.03	20174.6	0.99	20330.7	1.00
D9ZGG2	Vitronectin	45241.8	47676.1	1.05	42103.8	0.93	50280.8	1.11	50872	1.12	50370.3	1.11	44782.5	0.99	40512.9	0.90
B7Z8Q2	cDNA FLJ37966	78902.1	72510.6	0.92	71430.7	0.91	69072.4	0.88	68315.1	0.87	75454.7	0.96	69386.5	0.88	66683.6	0.85
D9YZU5	Beta-globin	26416.6	26762	1.01	25508.7	0.97	21785.1	0.82	27519.4	1.04	20787.3	0.79	30161.5	1.14	30166.4	1.14
B7Z8B6	cDNA FLJ37966	3715.2	4851.3	1.31	3182.4	0.86	3426.6	0.92	2914.7	0.78	3116.9	0.84	4486.4	1.21	3978.8	1.07
A0A1B0GU	Ig mu chain	41758.8	33261.6	0.80	33963.5	0.81	33291.6	0.80	31512.1	0.75	39637.9	0.95	55273.6	1.32	42527.6	1.02
B2R950	cDNA, FLJ37966	2893	2645.4	0.91	2669.2	0.92	2807.5	0.97	2688	0.93	3024	1.05	2831.3	0.98	3108.6	1.07
Q96PD5	N-acetylmuramidase 2B	18131.8	20449.1	1.13	16845.4	0.93	17634.1	0.97	17893.7	0.99	18152.3	1.00	17621.4	0.97	16026.7	0.88
P02763	Alpha-1-acid glycoprotein 1	21588.2	20978.4	0.97	23424.7	1.09	17359	0.80	26223.1	1.21	21535.6	1.00	24203.1	1.12	45114	2.09
B0AZL7	cDNA, FLJ37966	18319.3	22324.3	1.22	19098.3	1.04	20124.5	1.10	19414.7	1.06	21036	1.15	19426.1	1.06	16382.5	0.89
B2R815	cDNA, FLJ37966	18523.7	20326.1	1.10	19314.7	1.04	17136.1	0.93	15861.9	0.86	19134.1	1.03	18435.3	1.00	13815.6	0.75
P08697	Alpha-2-antiplasmin	16208.5	15568.9	0.96	15471.1	0.95	16057.6	0.99	15651.7	0.97	15778.6	0.97	15332.5	0.95	15477.2	0.95
P25311	Zinc-alpha-2-glycoprotein	22823.9	24197	1.06	22776.3	1.00	22025	0.96	20191.2	0.88	24868.6	1.09	21172.2	0.93	17749	0.78
Q86TT1	Full-length cDNA	628.3	731.5	1.16	612.5	0.97	656.3	1.04	623.5	0.99	686.7	1.09	818	1.30	646.6	1.03
V9HW68	Epididymis protein 1	14318	14200.7	0.99	13471.4	0.94	14091.7	0.98	14781.3	1.03	14841.2	1.04	12780.1	0.89	15179.6	1.06
Q06033	Inter-alpha-trypsin inhibitor heavy chain 1	12468.4	13567.8	1.09	12959.8	1.04	12149.5	0.97	12249.6	0.98	12309.6	0.99	12294.9	0.99	12842.5	1.03
P80108	Phosphatidylinositol 3-kinase	9864.1	11082	1.12	10784.2	1.09	10919.1	1.11	10552.3	1.07	10448.7	1.06	10629.6	1.08	9586.6	0.97
P07360	Complement factor D	10063.5	10529.2	1.05	9986.8	0.99	10975.9	1.09	10507.7	1.04	10028.2	1.00	10438.8	1.04	9234.4	0.92
P22792	Carboxypeptidase Y	16767.1	18835.2	1.12	18865.2	1.13	17056.8	1.02	16192.8	0.97	17940.2	1.07	16797.5	1.00	14238.6	0.85
A0A140VK	Testis tissue	22611	25131.6	1.11	22440.4	0.99	22366.6	0.99	20490.3	0.91	23660.7	1.05	22065.7	0.98	21306.7	0.94
P51884	Lumican OS=	21478.9	21871.6	1.02	20178.2	0.94	20998.2	0.98	20168.7	0.94	21854.5	1.02	20957.9	0.98	18878.5	0.88
A0A0A0MF	Coagulation factor XIII	6480.2	8192.9	1.26	7006.1	1.08	7385.3	1.14	6954.2	1.07	8063.6	1.24	6384.5	0.99	6217.3	0.96
Q59EP2	Angiotensinogen	24089.4	24341.2	1.01	23979.5	1.00	23177.3	0.96	23539.2	0.98	25181.7	1.05	23425	0.97	22223.6	0.92
B2R582	cDNA, FLJ37966	798	921.9	1.16	782	0.98	830.3	1.04	714.9	0.90	798.1	1.00	808.3	1.01	819.5	1.03
Q4TZM4	Hemoglobin	1332.1	1405	1.05	1277.4	0.96	979.5	0.74	1322.2	0.99	935.1	0.70	1697.9	1.27	1700.3	1.28

Q6MZQ6	Putative unc															
P27169	Serum parac	26425.9	24841.5	0.94	26246	0.99	25042.9	0.95	24302	0.92	27227.6	1.03	26242.8	0.99	22453.9	0.85
D6RF20	Vitamin D-b	24.1	28.9	1.20	18.6	0.77	31.5	1.31	25.4	1.05	23.9	0.99	27.1	1.12	21.2	0.88
Q96K68	cDNA FLJ1	9509.7	9530.9	1.00	9767.8	1.03	10313.7	1.08	9694.6	1.02	9181.8	0.97	9114	0.96	12195.3	1.28
G3XAK1	Hepatocyte g	6404.4	6901.6	1.08	6318.4	0.99	6943.5	1.08	6491.8	1.01	6537.2	1.02	5956.3	0.93	5849.2	0.91
A0A024R2C	C-type lectin	7606	8337.9	1.10	7454.1	0.98	7456.1	0.98	6832.4	0.90	8007.7	1.05	7034	0.92	6269.8	0.82
B2R9F2	cDNA, FLJ5	21182.2	24324.2	1.15	24773.4	1.17	18653.9	0.88	20418.2	0.96	23084.3	1.09	23221.8	1.10	21849.1	1.03
P05543	Thyroxine-b	10522.1	10969	1.04	11223.3	1.07	10323.5	0.98	10070	0.96	11408.6	1.08	11465.7	1.09	10219.8	0.97
A0N071	Delta globin	110.6	124.5	1.13	109	0.99	104.9	0.95	128.9	1.17	97.3	0.88	127.9	1.16	133	1.20
P02753	Retinol-bind	33157.6	34673.7	1.05	31209.4	0.94	35599.5	1.07	34712.9	1.05	35446.5	1.07	33435.2	1.01	27320.1	0.82
C9JF17	Apolipoprot	24386.8	26206.3	1.07	26020.4	1.07	27458.5	1.13	23217.7	0.95	27609.1	1.13	23713.6	0.97	21141.3	0.87
A8K1K1	cDNA FLJ7	6742.3	7542.4	1.12	7262.6	1.08	7434.9	1.10	6611.6	0.98	6953.1	1.03	6538.6	0.97	5646.1	0.84
A0A1K0GX	Globin C1 C	14723.1	14157.8	0.96	14141.1	0.96	9465.6	0.64	14883	1.01	10905.9	0.74	15034.8	1.02	18249.7	1.24
D6RAR4	Hepatocyte g	76.6	88.7	1.16	66.5	0.87	89.3	1.17	80.3	1.05	82.1	1.07	88.3	1.15	60.3	0.79
Q16610	Extracellular	4476.7	4777.3	1.07	4009.6	0.90	4909.4	1.10	4730.8	1.06	4439.4	0.99	4352.6	0.97	3766.9	0.84
Q04756	Hepatocyte g	6012.3	6577.1	1.09	5836.5	0.97	6665.3	1.11	5873.9	0.98	6133.9	1.02	5968.9	0.99	5362	0.89
Q8TCD0	Uncharacter	466.9	415.3	0.89	418.3	0.90	400.5	0.86	341.9	0.73	406.3	0.87	519.1	1.11	440.6	0.94
P03951	Coagulation	4345.1	5192.9	1.20	4314.6	0.99	4788.6	1.10	4719.5	1.09	4850.2	1.12	4739	1.09	4267.9	0.98
Q03591	Complemen	1223.8	1251.5	1.02	1403.4	1.15	1466.7	1.20	1296.1	1.06	1136.9	0.93	1164.7	0.95	1201	0.98
A0A0S2Z3	Lectin galac	6950.6	7320.9	1.05	6436	0.93	6925.7	1.00	6615.1	0.95	6572.9	0.95	8034.7	1.16	8214.9	1.18
Q6PIL8	IGK@ prote	7153.1	7606.5	1.06	6323.7	0.88	5824.9	0.81	5681	0.79	5845.4	0.82	8047.9	1.13	8245.1	1.15
P02746	Complemen	8644	8778.3	1.02	8648.3	1.00	9690.7	1.12	9963.4	1.15	8865	1.03	8777.6	1.02	8280.1	0.96
A0A096LPI	Protein SAA	10462.9	10688.6	1.02	10717.2	1.02	11896	1.14	12038.4	1.15	11679.4	1.12	11065.4	1.06	11844.1	1.13
Q6MZU6	Putative unc	3853.2	4359	1.13	3634.2	0.94	4226.4	1.10	4274.9	1.11	4095.3	1.06	3280.3	0.85	3843.7	1.00
P02750	Leucine-rich	20434.1	19280.5	0.94	19204.6	0.94	19544.7	0.96	22329.2	1.09	19626.4	0.96	21509.2	1.05	27178.7	1.33
P48740	Mannan-bin	5616.9	7116.6	1.27	6236	1.11	6539.9	1.16	6207.5	1.11	6035.4	1.07	6160.2	1.10	5467.1	0.97
P02775	Platelet basi	15640.4	17782.9	1.14	15010.3	0.96	17397.2	1.11	17472.2	1.12	19286.4	1.23	15447.8	0.99	15413.3	0.99
A6XNE2	Complemen	3098.3	3661.6	1.18	3079	0.99	3909.2	1.26	4144.5	1.34	3542.2	1.14	3244.9	1.05	2871.5	0.93
P02747	Complemen	8282.3	8248.2	1.00	8431.2	1.02	8152.3	0.98	8062.2	0.97	7885.1	0.95	7807.7	0.94	8072.6	0.97
P15169	Carboxypep	6584.9	6941.6	1.05	6477.7	0.98	7257	1.10	7312.7	1.11	7140.8	1.08	6848.2	1.04	6112.5	0.93
P00742	Coagulation	8124.9	8976.6	1.10	8245.7	1.01	8527.9	1.05	8393.9	1.03	8808	1.08	7701.3	0.95	7101.8	0.87
P19652	Alpha-1-acic	19263.5	19165	0.99	20571.4	1.07	17406.3	0.90	19854.6	1.03	18726.5	0.97	18554.9	0.96	27901.1	1.45
Q14520	Hyaluronan-	11039.1	11316.7	1.03	10532.6	0.95	11084.5	1.00	11173.5	1.01	11337.9	1.03	10612	0.96	10238.1	0.93
F2RM37	Coagulation	6388.3	6699.2	1.05	6153.5	0.96	6191.1	0.97	6563.2	1.03	6383.7	1.00	6621.1	1.04	6072.5	0.95
Q96IY4	Carboxypep	11789.6	12269.4	1.04	11550.9	0.98	13003.2	1.10	12811.5	1.09	12370	1.05	12796.3	1.09	10423.8	0.88
A2NUT2	Lambda-cha	14755.2	15187.4	1.03	14275.1	0.97	15722.5	1.07	15169.7	1.03	16572.9	1.12	14111.3	0.96	15719	1.07
O75636	Ficolin-3 OS	15367.2	17638.8	1.15	18165.6	1.18	16665	1.08	14385.8	0.94	16004	1.04	16769.9	1.09	14579.7	0.95
Q9NPP6	Immunoglot	439.2	454.8	1.04	457.1	1.04	412.1	0.94	401.7	0.91	470.5	1.07	460.3	1.05	526.6	1.20
Q567P1	IGL@ protei	1174.7	1092.4	0.93	1442.3	1.23	1467.9	1.25	1287.8	1.10	1469.6	1.25	959	0.82	1225	1.04
A0A024R6	Serpin pepti	3529	4175.3	1.18	3766.7	1.07	4461.6	1.26	4485.9	1.27	4064	1.15	4083.4	1.16	2912.8	0.83
H6VRG1	Keratin 1 OS	2159.3	3451.6	1.60	3228.1	1.49	2797.8	1.30	3184.2	1.47	3565.9	1.65	2341.6	1.08	2726	1.26
A0A0R7FJ	Coagulation	2366.7	2998.3	1.27	2976.7	1.26	2986.4	1.26	2883.8	1.22	2602.2	1.10	2378.2	1.00	2213.3	0.94
P20851	C4b-binding	8738.8	9027.5	1.03	8504	0.97	9754.4	1.12	9465.3	1.08	9543.2	1.09	9385.3	1.07	9386.6	1.07
P63261	Actin, cytop	3679	5034.2	1.37	4421.8	1.20	4569	1.24	4498.6	1.22	4666.4	1.27	4301.8	1.17	3602.9	0.98
S6AWE6	IgG L chain	443.9	446.9	1.01	440.7	0.99	498.9	1.12	495.6	1.12	479.3	1.08	490	1.10	532	1.20
A0A024R0	Apolipoprot	20202.7	21308.9	1.05	26544.7	1.31	20263.1	1.00	17356.9	0.86	22290.7	1.10	21967.1	1.09	21581	1.07
O00391	Sulfhydryl o	2526.4	3092.7	1.22	2603.2	1.03	2911.4	1.15	2915	1.15	2811.2	1.11	2625.9	1.04	2246.1	0.89

P02743	Serum amyl	12930.8	13581.8	1.05	12697.6	0.98	13609.2	1.05	12915.9	1.00	14808.5	1.15	11096.7	0.86	9575.8	0.74
P02776	Platelet fact	7708.1	7328	0.95	14120.9	1.83	14224.7	1.85	9230.1	1.20	9519.2	1.23	9673.7	1.26	7847.5	1.02
P09172	Dopamine b	2072.2	2206.3	1.06	2263.1	1.09	2213	1.07	2303	1.11	1895.8	0.91	2180.4	1.05	1778.4	0.86
P43251	Biotinidase	3353.8	4160.3	1.24	3812.2	1.14	3998.6	1.19	3618	1.08	3739.1	1.11	3523.6	1.05	2741.5	0.82
P17936	Insulin-like	5061.2	5551.8	1.10	4713.9	0.93	4955.8	0.98	5015.6	0.99	5662.5	1.12	5154.3	1.02	4313.3	0.85
A6XGL1	Transthyreti	6699.7	7120.2	1.06	5982.7	0.89	6096.6	0.91	5738.3	0.86	6744	1.01	7894.5	1.18	6887.9	1.03
V9GYM3	Apolipoprot	58257.6	44115.6	0.76	54074.9	0.93	48466.6	0.83	40529.6	0.70	47555	0.82	51115.9	0.88	44102	0.76
B4DPQ3	cDNA FLJ5	3019.2	3156	1.05	2875.5	0.95	3008.6	1.00	3181.3	1.05	3251	1.08	3124.7	1.03	2837.3	0.94
Q1HP67	Lipoprotein,	4008	3572.9	0.89	3229.5	0.81	5065.3	1.26	3792.2	0.95	3698.1	0.92	3973.9	0.99	3843.8	0.96
B2R888	Monocyte di	1767.4	2126.2	1.20	1838	1.04	2245	1.27	2263	1.28	2124.1	1.20	1927.7	1.09	1746.3	0.99
Q6GMV7	Uncharacter	874.1	719.1	0.82	817	0.93	835.9	0.96	834.4	0.95	988	1.13	992.9	1.14	973.6	1.11
P13645	Keratin, typ	4187.6	6299.9	1.50	5141.4	1.23	5083.2	1.21	5122.4	1.22	5879.9	1.40	4691.1	1.12	4037.6	0.96
A0A024R1C	Apolipoprot	5734.4	6244	1.09	5517.6	0.96	7231.8	1.26	5956.6	1.04	5785.2	1.01	6265	1.09	6103	1.06
P36980	Complemen	655.8	697.3	1.06	723	1.10	663.2	1.01	683.4	1.04	651.3	0.99	658.5	1.00	640.3	0.98
O95445	Apolipoprot	8414.8	8296.5	0.99	9042	1.07	8840.9	1.05	8126.8	0.97	8617.6	1.02	8163.2	0.97	7562.5	0.90
Q6MZX7	Putative unc	1228.3	1342.6	1.09	1183.7	0.96	1099.8	0.90	1043.5	0.85	1109.7	0.90	938.7	0.76	1199.3	0.98
P22352	Glutathione	9294	10412	1.12	9504.2	1.02	9810.5	1.06	9756.1	1.05	9806.6	1.06	9337.8	1.00	8045.1	0.87
P04278	Sex hormon	2653.1	3130.4	1.18	2779.4	1.05	2783.5	1.05	2998.7	1.13	2665.9	1.00	2874.3	1.08	2493.8	0.94
K7ER74	Protein APC	13980.3	13740.2	0.98	12868.2	0.92	16333.2	1.17	13350.9	0.95	13390.5	0.96	13986.8	1.00	16334.8	1.17
Q8NF17	FLJ00385 p	242.4	320.1	1.32	247.7	1.02	331.5	1.37	279.1	1.15	251.7	1.04	370.4	1.53	292.4	1.21
S6BGE0	IgG H chain															
A0A0S2Z4I	Complemen	3322.3	4095.2	1.23	3644.8	1.10	3830.2	1.15	3920.4	1.18	4101.1	1.23	4187.3	1.26	3549.1	1.07
D3DNN4	Carboxylic e	5130.5	5064.1	0.99	5240.3	1.02	5522.4	1.08	4966.8	0.97	5218	1.02	4797.5	0.94	4317.1	0.84
A0A161I20	Lactoferrin	879.4	1147.8	1.31	1147.8	1.31	1211.5	1.38	1145.7	1.30	1138.4	1.29	1107.8	1.26	981.9	1.12
P00915	Carbonic an	2002.4	2162.4	1.08	2609.2	1.30	1957.8	0.98	2055.7	1.03	1812.4	0.91	2027.6	1.01	2103.9	1.05
Q9UGM5	Fetuin-B OS	4107.9	4351.3	1.06	3769.4	0.92	4123.9	1.00	4226.1	1.03	4171.8	1.02	4422.2	1.08	3868.4	0.94
A0A0F7G8J	Plasminoger															
Q8TCZ8	Apolipoprot	380.9	476.4	1.25	343.4	0.90	456.2	1.20	890	2.34	571.4	1.50	559.4	1.47	698.9	1.83
P11226	Mannose-bin	1549.9	2010.3	1.30	1675.5	1.08	1836.5	1.18	1840.5	1.19	1926	1.24	1449.9	0.94	1492.7	0.96
P14780	Matrix meta	1462.7	1945.1	1.33	1718.1	1.17	1892.8	1.29	1825.1	1.25	1878.1	1.28	1733.5	1.19	1533.9	1.05
Q9NZP8	Complemen	2150.2	2606.4	1.21	2149.6	1.00	2431	1.13	2787.5	1.30	2422.7	1.13	2219.6	1.03	2302	1.07
A8K3K1	cDNA FLJ7	62.2	82.1	1.32	62.4	1.00	92.7	1.49	91.6	1.47	73.8	1.19	70.5	1.13	64.1	1.03
P35527	Keratin, typ	3567.9	4828.9	1.35	4184.2	1.17	4681.3	1.31	4297.7	1.20	4160.6	1.17	3806.3	1.07	3832.4	1.07
A8KAJ3	cDNA FLJ7	1343.2	1837.5	1.37	1648.2	1.23	1889.4	1.41	1779.4	1.32	1797	1.34	1394.9	1.04	1262.5	0.94
Q9UHG3	Prenylcyste	1396.4	1623	1.16	1448	1.04	1676.7	1.20	1553.9	1.11	1683.2	1.21	1543.2	1.11	1253.5	0.90
P10720	Platelet fact															
S6B2A1	IgG L chain	15.4	16.1	1.05	15.7	1.02	18.2	1.18	14.9	0.97	14	0.91	20.7	1.34	13.1	0.85
B0YIW2	Apolipoprot	15931.1	15946.8	1.00	14299.1	0.90	16740.7	1.05	12479.1	0.78	14347.2	0.90	18360.1	1.15	14244.5	0.89
Q13790	Apolipoprot	2324.8	2383.7	1.03	2210.3	0.95	2269.3	0.98	2033.8	0.87	2229.3	0.96	2090	0.90	1990.1	0.86
P35908	Keratin, typ	701.5	1259.7	1.80	955	1.36	943.1	1.34	1154.5	1.65	1302.1	1.86	818.6	1.17	752.4	1.07
A0A024RD	Lymphocyte	1247.4	1464.3	1.17	1267.8	1.02	1430.2	1.15	1427	1.14	1398	1.12	1284.8	1.03	1092.9	0.88
P40197	Platelet gly	882	992.6	1.13	1028.8	1.17	1166.6	1.32	1082.7	1.23	1030.5	1.17	1056.7	1.20	824.5	0.93
P16070	CD44 antige	3109.9	3521.4	1.13	3226.7	1.04	3652	1.17	3836.3	1.23	3838	1.23	3254.7	1.05	2930.4	0.94
A0A024R93	Proteoglycar	3044.2	4220.7	1.39	3217.9	1.06	3556.5	1.17	3585.4	1.18	3689.6	1.21	3751.3	1.23	3316.8	1.09
A2KBC2	Anti-(ED-B)	1576.5	1521.9	0.97	1577	1.00	1624.3	1.03	1472.1	0.93	1518.2	0.96	1839.6	1.17	1784.3	1.13
A5YAK2	Apolipoprot	3983.5	4272.5	1.07	3741.9	0.94	4575.6	1.15	4034	1.01	4428.9	1.11	4371	1.10	4010.6	1.01
G3V2W1	Protein Z-de	1874.1	2292.1	1.22	2045.7	1.09	2003.8	1.07	1918.6	1.02	2069.7	1.10	2209.2	1.18	2075.1	1.11

A0A140VK	Testicular se	2477.8	2812	1.13	2487.7	1.00	2736.8	1.10	2375.6	0.96	2635.4	1.06	2525.6	1.02	2230.7	0.90
O43866	CD5 antigen	1243.5	1335.5	1.07	1384.5	1.11	1606.8	1.29	1391.1	1.12	1358.8	1.09	1626.1	1.31	1351.8	1.09
P01591	Immunoglob	3550.8	3113	0.88	3322.8	0.94	3143.9	0.89	3267.3	0.92	3614.4	1.02	3594.7	1.01	3742.6	1.05
A8K3I0	cDNA FLJ7	549.7	652.6	1.19	540.7	0.98	621.8	1.13	590.1	1.07	595.1	1.08	527.4	0.96	506.8	0.92
Q9H804	cDNA FLJ1	156.4	188.7	1.21	150.8	0.96	254.6	1.63	222.8	1.42	105.8	0.68	138.6	0.89	212.9	1.36
B0YJC6	Vitamin K-d	933.3	1103.8	1.18	1108.9	1.19	1160.9	1.24	1103.9	1.18	913.5	0.98	890.5	0.95	832.8	0.89
Q6UXB8	Peptidase in	818.1	905.3	1.11	840.5	1.03	932.9	1.14	822	1.00	835.3	1.02	645.3	0.79	631.8	0.77
B3KWB5	cDNA FLJ4															
P08709	Coagulation	328	407.9	1.24	288.2	0.88	410	1.25	429.3	1.31	361	1.10	314.7	0.96	263.8	0.80
V9HWB4	Epididymis	597.3	779.4	1.30	658	1.10	736.6	1.23	713.6	1.19	670.1	1.12	707.2	1.18	577.6	0.97
P02741	C-reactive p	1082.9	830.7	0.77	1080.4	1.00	1074.1	0.99	1604.3	1.48	1017.9	0.94	930.1	0.86	4798.1	4.43
Q9UL72	Myosin-reac	364.5	481.2	1.32	387	1.06	447.7	1.23	412.4	1.13	424.7	1.17	408.7	1.12	401.4	1.10
P49908	Selenoprotei	3441.9	3795	1.10	3274.3	0.95	3369.5	0.98	3224.2	0.94	3638.3	1.06	3596.3	1.04	2561.1	0.74
V9HW12	Epididymis	1538.3	1592.7	1.04	1603.5	1.04	1412.8	0.92	1548.9	1.01	1461.7	0.95	1574.2	1.02	1641.4	1.07
P07359	Platelet glyco	2084.5	2648.4	1.27	2443.9	1.17	2770	1.33	2679.9	1.29	2718.9	1.30	2386.6	1.14	2117.4	1.02
D3DQH8	Secreted pro	2098	2082.9	0.99	1926.3	0.92	2272.4	1.08	2198	1.05	1998.8	0.95	1964.4	0.94	1970.9	0.94
P08779	Keratin, type	1452.8	1891.4	1.30	1570.9	1.08	1774.5	1.22	1739.5	1.20	1762.6	1.21	1580.2	1.09	1378	0.95
P61626	Lysozyme C	1103.7	1304.7	1.18	1215.2	1.10	1329.5	1.20	1310.4	1.19	1321.3	1.20	1230.1	1.11	1218.8	1.10
A0A024R8Z	L-selectin O	1228.5	1334.2	1.09	1161.3	0.95	1309.8	1.07	1228.2	1.00	1343.6	1.09	1204.4	0.98	1086.6	0.88
S6BAQ4	IgG H chain															
Q59E93	Membrane a	406	515.6	1.27	400.2	0.99	502.8	1.24	474.2	1.17	465	1.15	389	0.96	363.5	0.90
A0A193CH	10E8 heavy	157.8	158.8	1.01	163.2	1.03	155.1	0.98	171.8	1.09	181.3	1.15	163	1.03	182.6	1.16
Q15166	Serum parac	355.2	388.4	1.09	362.1	1.02	443.9	1.25	424	1.19	404.1	1.14	387.6	1.09	306.9	0.86
B2RA39	cDNA, FLJ9	781.4	981	1.26	877	1.12	968.8	1.24	973.7	1.25	916.8	1.17	847.9	1.09	950.7	1.22
P0DJ19	Serum amyl	294.1	332.3	1.13	257.1	0.87	325.2	1.11	329	1.12	287.9	0.98	283.8	0.96	467.6	1.59
P0DJ18	Serum amyl	389.3	384.7	0.99	312.6	0.80	424.2	1.09	400.2	1.03	334.5	0.86	316.6	0.81	683.4	1.76
A0A068LKI	Ig heavy cha															
Q15582	Transformin	1222.9	1217.3	1.00	1181.9	0.97	1282.9	1.05	1272.7	1.04	1157.6	0.95	1193.1	0.98	1078.2	0.88
A0A024R8C	Prostaglandi	727.8	868.5	1.19	780.4	1.07	887.8	1.22	877.6	1.21	863.8	1.19	666	0.92	610.6	0.84
O00187	Mannan-bin	1989	2314.3	1.16	2064	1.04	2289	1.15	2051	1.03	2010.4	1.01	1745.3	0.88	1793.4	0.90
Q65ZC9	Single-chain															
P18065	Insulin-like	301.5	411.4	1.36	349.1	1.16	408.5	1.35	403.6	1.34	332.6	1.10	324.6	1.08	270.7	0.90
A0A140TA	Tenascin-X	362.7	402.7	1.11	380.1	1.05	427.2	1.18	385.6	1.06	348.8	0.96	371.3	1.02	319.7	0.88
P18428	Lipopolysac	996.7	1117.8	1.12	1038.8	1.04	1148.7	1.15	1059.3	1.06	991.4	0.99	1082.1	1.09	1446.9	1.45
A0A1B1CY	Vitamin D b	10424.2	10328.3	0.99	9334.3	0.90	10642.8	1.02	10623.2	1.02	7147.6	0.69	8648.4	0.83	3808.9	0.37
A2NYQ9	Anti-folate b	53.9	45.6	0.85	42	0.78	53.7	1.00	41.3	0.77	43.9	0.81	53.3	0.99	44.4	0.82
Q9Y5Y7	Lymphatic v	1364.9	1481	1.09	1276.5	0.94	1382.1	1.01	1324.6	0.97	1456	1.07	1268.8	0.93	964.9	0.71
P02671	Fibrinogen a	771.5	989.7	1.28	850.9	1.10	726.5	0.94	790.5	1.02	773.6	1.00	947.4	1.23	939.9	1.22
P02533	Keratin, type	55.3	86.1	1.56	62.7	1.13	80	1.45	78.1	1.41	73.8	1.33	79	1.43	51	0.92
Q6EMK4	Vasorin OS-	363.3	507	1.40	411	1.13	474.9	1.31	446.2	1.23	485.5	1.34	398	1.10	338	0.93
A0A0S2Z42	HCG203981	406.5	557.3	1.37	425.9	1.05	473.6	1.17	446.2	1.10	492.1	1.21	391.6	0.96	358.9	0.88
B7Z4R8	cDNA FLJ5															
B3KUE5	Phospholipic	901.5	1026.7	1.14	1001.6	1.11	1132.5	1.26	1000.6	1.11	1019.8	1.13	992.1	1.10	662.1	0.73
A2NB46	Cold aggluti	397.6	438.5	1.10	393.2	0.99	457.3	1.15	431.9	1.09	398.7	1.00	419.1	1.05	400.9	1.01
B2R773	cDNA, FLJ5	546.2	835.1	1.53	1129.4	2.07	751.6	1.38	746	1.37	632.2	1.16	663.9	1.22	539.3	0.99
A8K6K4	cDNA FLJ7	1308	1248.6	0.95	1077.3	0.82	1346	1.03	1536.6	1.17	1482.1	1.13	1344.5	1.03	1075.5	0.82
A0A087WX	Cadherin-1	167.8	177.2	1.06	154.2	0.92	206.2	1.23	194.4	1.16	167	1.00	155.5	0.93	132.6	0.79

A8K6A6	cDNA FLJ7	640.8	759.3	1.18	704	1.10	749.6	1.17	701.3	1.09	759.5	1.19	878.3	1.37	610.8	0.95
B3KQF4	cDNA FLJ9	166.7	222.9	1.34	170.1	1.02	180	1.08	186.7	1.12	152.8	0.92	165.9	1.00	169.3	1.02
A8K430	Fructose-bis	111.1	159.5	1.44	123.1	1.11	169.1	1.52	169.2	1.52	125.7	1.13	152.2	1.37	103.7	0.93
A0A0K0K1	Cystatin OS	801.8	842.1	1.05	825.4	1.03	805.6	1.00	821	1.02	888.1	1.11	813	1.01	778.2	0.97
A8K335	cDNA FLJ7	626.5	727.5	1.16	654.2	1.04	707.8	1.13	763.3	1.22	715.1	1.14	602	0.96	560	0.89
Q9NQ79	Cartilage aci	792.2	908.4	1.15	874.4	1.10	1227.9	1.55	829.9	1.05	874.3	1.10	746.8	0.94	679.4	0.86
Q15485	Ficolin-2 OS	1232.5	1636.6	1.33	1319.3	1.07	1375.2	1.12	1231.1	1.00	1469.1	1.19	1458.5	1.18	1153.2	0.94
B4E0X1	Beta-2-micro	1181.8	1324.5	1.12	1277	1.08	1268.4	1.07	1283.3	1.09	1219.7	1.03	1453.9	1.23	1226.4	1.04
P05362	Intercellular	202.9	238.6	1.18	210.4	1.04	221.3	1.09	230.1	1.13	208.3	1.03	214.7	1.06	218.3	1.08
Q76LX8	A disintegrin	311.2	543.8	1.75	327.7	1.05	425.6	1.37	395.9	1.27	310.8	1.00	349.8	1.12	260.2	0.84
A0A075B7E	Protein IGH															
P21333	Filamin-A C	123.6	169.3	1.37	148	1.20	168.1	1.36	169.7	1.37	144.1	1.17	141.4	1.14	110.8	0.90
Q92496	Complemen	1422.9	1447.6	1.02	1316	0.92	1489.3	1.05	1467.8	1.03	1465.4	1.03	1237.4	0.87	1561.3	1.10
A0A024RD	Phospholipa	431.5	565	1.31	425.7	0.99	544.9	1.26	555.5	1.29	541	1.25	451.2	1.05	424.7	0.98
Q86VB7	Scavenger re	167.3	208.9	1.25	181	1.08	222.9	1.33	208.4	1.25	189.1	1.13	157.1	0.94	166.1	0.99
P07737	Profilin-1 O	247.6	387.8	1.57	353.3	1.43	350.2	1.41	331.5	1.34	356.3	1.44	337.3	1.36	253.1	1.02
P37802	Transgelin-2	445.5	690.4	1.55	625.6	1.40	751.4	1.69	676.2	1.52	673.6	1.51	605.5	1.36	526.5	1.18
Q13103	Secreted phc	286.8	285.6	1.00	310.5	1.08	260.6	0.91	242.5	0.85	241.1	0.84	256.7	0.90	215.2	0.75
B1AHL2	Fibulin-1 OS	181.8	239.3	1.32	197.7	1.09	215.8	1.19	222.6	1.22	261.5	1.44	166	0.91	171.8	0.94
Q59EA3	Cadherin 5,	314	558.2	1.78	497.9	1.59	596.1	1.90	503.1	1.60	496.3	1.58	329.5	1.05	293.5	0.93
Q0IIN1	Keratin 77 C															
Q16853	Membrane p	99.9	129.3	1.29	94.8	0.95	137.7	1.38	148	1.48	126.3	1.26	94.6	0.95	76.7	0.77
P02745	Complemen	2193.5	2073.6	0.95	2290.7	1.04	2089.9	0.95	2080.8	0.95	2050.6	0.93	2014.3	0.92	2287.5	1.04
A0N5G5	Rheumatoid	94.5	136.1	1.44	97.1	1.03	132.6	1.40	124.5	1.32	110.3	1.17	101.2	1.07	108.4	1.15
Q9HCC1	Single chain	19	31.1	1.64	15.9	0.84	22.6	1.19	19.8	1.04	18.6	0.98	20.8	1.09	18.8	0.99
Q9Y490	Talin-1 OS=															
P33908	Mannosyl-ol	296.9	388.2	1.31	349.4	1.18	419.1	1.41	421.2	1.42	385.4	1.30	302.6	1.02	327	1.10
A8K6C9	cDNA FLJ7	934.1	1228.8	1.32	985.8	1.06	1209.8	1.30	986.1	1.06	1181.6	1.26	934.7	1.00	818.7	0.88
A0A087WT	Neural cell a	888.9	1126.6	1.27	994.6	1.12	1094.7	1.23	1013.7	1.14	997.6	1.12	1054.1	1.19	759.1	0.85
V9H1C1	Gelsolin exo															
Q9HDC9	Adipocyte p	668.2	696.8	1.04	716.8	1.07	782.3	1.17	793.5	1.19	763.3	1.14	642.7	0.96	620.9	0.93
A0A068LLC	Ig heavy cha	59.3	55.8	0.94	60.9	1.03	56.5	0.95	48	0.81	77.1	1.30	57.8	0.97	80.2	1.35
O00533	Neural cell a	308.4	401.8	1.30	325.1	1.05	460.4	1.49	348.5	1.13	365.9	1.19	330.3	1.07	276.8	0.90
P35443	Thrombospc	35.7	47.3	1.32	37.8	1.06	54.5	1.53	40.4	1.13	45.5	1.27	37.1	1.04	36.5	1.02
P23470	Receptor-tyr	147.9	180.8	1.22	144	0.97	201.5	1.36	200.5	1.36	186.1	1.26	134.1	0.91	104.6	0.71
E9PR17	CD59 glyco	192.7	191.7	0.99	175.9	0.91	212	1.10	214.1	1.11	197.9	1.03	177	0.92	156.6	0.81
B2R9V7	Superoxide c	103	123.8	1.20	106.3	1.03	144.8	1.41	116.1	1.13	109.3	1.06	109.1	1.06	73.1	0.71
A0A125QY	GCT-A9 lig	100.7	97.4	0.97	98.7	0.98	124.6	1.24	111.1	1.10	94.1	0.93	115.4	1.15	270.5	2.69
V9HW31	ATP synthas	83.6	149.2	1.78	140.3	1.68	154.3	1.85	152	1.82	138.6	1.66	89.4	1.07	122.9	1.47
M9MML0	Fc of IgG lo	525.7	649.1	1.23	555.7	1.06	596.8	1.14	601.8	1.14	549.8	1.05	534.9	1.02	510.7	0.97
Q13201	Multimerin-	484.6	578.8	1.19	527.2	1.09	603.9	1.25	553.1	1.14	542.8	1.12	563.8	1.16	501.9	1.04
P05164	Myeloperoxi	150.6	216.2	1.44	219.4	1.46	213.8	1.42	211.2	1.40	238.4	1.58	216	1.43	165.9	1.10
B2R7D2	cDNA, FLJC	180	201.9	1.12	179.8	1.00	203.8	1.13	180.4	1.00	175.6	0.98	167.6	0.93	145.5	0.81
G9K388	YWHAE/FA	139.1	382.4	2.75	307.1	2.21	367.6	2.64	384.3	2.76	322.6	2.32	149.6	1.08	340.3	2.45
A0A125QY	IBM-B2 hea	338.5	351.7	1.04	286.3	0.85	377.4	1.11	331.4	0.98	326.2	0.96	328.2	0.97	296.8	0.88
A0A125QY	GCT-A7 hea															
Q8NBP7	Proprotein c	138.5	190.5	1.38	139.5	1.01	186.7	1.35	178	1.29	184.5	1.33	130.9	0.95	92.9	0.67

A4D2D2	Procollagen	134.5	157.4	1.17	139.2	1.03	137.5	1.02	118.9	0.88	136	1.01	122.5	0.91	96.7	0.72
P07195	L-lactate de	187.3	219	1.17	207.3	1.11	205.5	1.10	176	0.94	197.2	1.05	192.4	1.03	191.6	1.02
E7EQ64	Trypsin-1 O	329.5	485.9	1.47	434	1.32	340.8	1.03	333.8	1.01	442.1	1.34	322.1	0.98	229.1	0.70
Q9UL85	Myosin-reac	1213.3	1238.1	1.02	1267.8	1.04	1233.3	1.02	1110.3	0.92	1252.6	1.03	1142.2	0.94	1319.2	1.09
P19320	Vascular cel	119.6	165.9	1.39	124.6	1.04	135.8	1.14	129.9	1.09	127.7	1.07	132.7	1.11	109.2	0.91
A0A140VJF	Testicular tis	106.7	133.4	1.25	108.5	1.02	133.7	1.25	124.3	1.16	118.2	1.11	114.2	1.07	84.4	0.79
P05019	Insulin-like g															
Q15430	Coagulation															
A2J1N5	Rheumatoid	112.9	128.6	1.14	126.1	1.12	218.9	1.94	147	1.30	127.4	1.13	123.3	1.09	107.3	0.95
Q07954	Prolow-dens	22.3	23.3	1.04	22.5	1.01	29.5	1.32	30	1.35	27.7	1.24	25.7	1.15	31.1	1.39
P24592	Insulin-like g	120.4	139.1	1.16	119	0.99	147.8	1.23	160	1.33	139.5	1.16	141	1.17	102.9	0.85
B2RBF5	cDNA, FLJ5	95.9	110	1.15	98.6	1.03	119.6	1.25	98.5	1.03	117	1.22	103.3	1.08	121.3	1.26
P12111	Collagen alp	83.9	112.8	1.34	84.9	1.01	109	1.30	100.6	1.20	97.6	1.16	87.7	1.05	70.1	0.84
B7Z9B1	cDNA FLJ5	183.5	213	1.16	182	0.99	223.4	1.22	245.4	1.34	200.6	1.09	175	0.95	158.9	0.87
B2R4R0	Histone H4	266.4	875.9	3.29	676.1	2.54	749.1	2.81	753.8	2.83	704.1	2.64	320.6	1.20	564.9	2.12
A8K6C1	cDNA FLJ7	679.2	730.8	1.08	691	1.02	887	1.31	885.6	1.30	756.5	1.11	766.5	1.13	666.3	0.98
A0N5G1	Rheumatoid	236.4	293.3	1.24	244.9	1.04	279.1	1.18	241.5	1.02	264.9	1.12	263.1	1.11	246.1	1.04
X6RBG4	Uromodulin	174.1	213.9	1.23	141	0.81	221.9	1.27	221.6	1.27	205.7	1.18	147.3	0.85	149.8	0.86
P49746	Thrombospc	626.2	765.2	1.22	612.7	0.98	777.6	1.24	651.6	1.04	854.1	1.36	638.2	1.02	558.4	0.89
A0A0X9UV	MS-D4 heav	644.4	710.5	1.10	592.1	0.92	723.2	1.12	664	1.03	740.1	1.15	668.3	1.04	688.6	1.07
P00918	Carbonic an	122.8	163.1	1.33	160	1.30	141.6	1.15	152	1.24	162.7	1.32	170	1.38	149.8	1.22
P08253	72 kDa type															
J3KPA1	Cysteine-ric	1673.4	2029.3	1.21	1726.2	1.03	1542.2	0.92	1556.9	0.93	1752.8	1.05	1671	1.00	1279.2	0.76
P22692	Insulin-like g															
Q4ZG40	Macrophage	171.9	200.9	1.17	143.7	0.84	187.4	1.09	181.2	1.05	159.4	0.93	157.2	0.91	131.7	0.77
E2RVJ0	Anion excha	23.2	45.7	1.97	41.6	1.79	51.8	2.23	36.3	1.56	32.8	1.41	41.9	1.81	22.8	0.98
A6NC48	ADP-ribosyl	236.2	275.1	1.16	235.9	1.00	310.2	1.31	251.1	1.06	259.8	1.10	236.7	1.00	225.3	0.95
Q6MZL2	Putative unc	569.9	701.4	1.23	643.4	1.13	639.9	1.12	593	1.04	599.9	1.05	603.8	1.06	543.8	0.95
P04040	Catalase OS	182.4	325.4	1.78	234.3	1.28	271	1.49	208.5	1.14	178	0.98	199.2	1.09	194.1	1.06
D3YTG3	Target of Ne	153.2	157.3	1.03	163.3	1.07	221.3	1.44	176.3	1.15	176	1.15	133.7	0.87	123.8	0.81
Q9UL89	Myosin-reac	74.9	83.3	1.11	66	0.88	98.2	1.31	70.9	0.95	86	1.15	68.2	0.91	64.1	0.86
X6R8F3	Neutrophil g	264.9	369.7	1.40	353.7	1.34	377.1	1.42	359.3	1.36	354	1.34	316.2	1.19	289.9	1.09
A0A024RD	SPARC-like	67.4	108.7	1.61	78	1.16	88.3	1.31	113.9	1.69	86.9	1.29	81.4	1.21	62.4	0.93
P35916	Vascular enc	44.5	45.6	1.02	35.5	0.80	48.1	1.08	50.6	1.14	34.5	0.78	40.7	0.91	31.2	0.70
B7ZMD7	Alpha-amyl	35.3	56.2	1.59	35.9	1.02	53.3	1.51	50.1	1.42	44	1.25	49	1.39	42.1	1.19
J3KPS3	Fructose-bis	133.9	197.8	1.48	164.3	1.23	206.9	1.55	205.3	1.53	173.7	1.30	144.9	1.08	133.8	1.00
Q14126	Desmoglein-															
P23284	Peptidyl-pro															
Q8WZ75	Roundabout	152.6	221.9	1.45	170.2	1.12	207.2	1.36	220.8	1.45	185.4	1.21	179.8	1.18	158.2	1.04
A0A087X1I	Calcium-dep	298	326.4	1.10	282.7	0.95	391.4	1.31	337.5	1.13	297.2	1.00	291.3	0.98	284.7	0.96
A0A0B4J1Y	Immunoglob															
Q86YW5	Trem-like tr	115.4	143.9	1.25	116.3	1.01	143.2	1.24	132.4	1.15	126.9	1.10	121.4	1.05	99.9	0.87
A0A024R9J	Collectin sul															
Q6UY14	ADAMTS-l	42.9	62.8	1.46	54.2	1.26	65.4	1.52	56.2	1.31	53.3	1.24	47	1.10	39.6	0.92
Q6UX71	Plexin doma	179.8	226.6	1.26	197.4	1.10	227.5	1.27	181.4	1.01	199.7	1.11	211.7	1.18	150.5	0.84
P03950	Angiogenin	363.2	420	1.16	390.8	1.08	489.1	1.35	518.6	1.43	472.9	1.30	389.7	1.07	343.2	0.94
Q05639	Elongation f	40.3	127.6	3.17	111.4	2.76	84.2	2.09	85.3	2.12	136.9	3.40	56.2	1.39	104.1	2.58

Q59ED3	Intercellular	68.1	87.6	1.29	66	0.97	99.7	1.46	102.8	1.51	70	1.03	66	0.97	56.5	0.83
Q8NGE6	Olfactory re	1201.7	1274.4	1.06	1087.1	0.90	1695.4	1.41	1970.5	1.64	1579.6	1.31	898.4	0.75	1144.2	0.95
P07333	Macrophage	32.6	52.1	1.60	37.1	1.14	42.2	1.29	44.6	1.37	33.9	1.04	37.4	1.15	31.8	0.98
Q9H8L6	Multimerin-	160.8	227.9	1.42	155.2	0.97	220	1.37	188.5	1.17	186.5	1.16	178.5	1.11	142.2	0.88
P04066	Tissue alpha	124.2	152.2	1.23	118.8	0.96	179.1	1.44	166.6	1.34	134.4	1.08	125	1.01	104	0.84
A8K7T4	cDNA FLJ7															
A0A140VJC	Testicular tis	134.5	139.9	1.04	137	1.02	167.7	1.25	161.9	1.20	160.1	1.19	141.1	1.05	137.8	1.02
A0A1C9J6F	B cell recept	61.6	87.8	1.43	61.3	1.00	96.2	1.56	87.2	1.42	86.2	1.40	79	1.28	58.1	0.94
B4DZM1	cDNA FLJ5	1025.2	1136	1.11	998.7	0.97	907.2	0.88	924.7	0.90	1086.3	1.06	1158.3	1.13	1028.7	1.00
A0A024R6V	Dipeptidase															
Q4LE64	NUMA1 var	802.5	1029.2	1.28	821	1.02	1000.5	1.25	1075	1.34	1117.1	1.39	684	0.85	571.4	0.71
P11279	Lysosome-ac	238.8	315.2	1.32	278.2	1.16	311.7	1.31	266.4	1.12	250.5	1.05	229.6	0.96	221.9	0.93
Q99784	Noelin OS=	236.4	230.4	0.97	224.4	0.95	214.6	0.91	214.5	0.91	211.6	0.90	203.3	0.86	219.5	0.93
Q9UIU0	Dihydropyri	64.3	86.9	1.35	72.3	1.12	95.2	1.48	65.3	1.02	68.7	1.07	75.9	1.18	56.5	0.88
P07998	Ribonucleas	34.2	36.7	1.07	32	0.94	38.4	1.12	41.7	1.22	42.1	1.23	36.3	1.06	29.5	0.86
K4DIA0	ICOS ligand	522.3	570.7	1.09	574.2	1.10	552.6	1.06	517.5	0.99	534.7	1.02	498.4	0.95	476.9	0.91
Q9ULV0	Unconventic	53.7	63.5	1.18	46.6	0.87	53.9	1.00	82.8	1.54	56	1.04	64.5	1.20	84.6	1.58
O14786	Neuropilin-1															
B7Z3G0	cDNA FLJ5	295.9	219.8	0.74	169.4	0.57	259.1	0.88	320.3	1.08	272.7	0.92	313.1	1.06	191.9	0.65
Q99969	Retinoic aci	40.7	55.7	1.37	50.7	1.25	68.1	1.67	56.8	1.40	50.2	1.23	45.7	1.12	42.4	1.04
Q9HCI6	E3 SUMO-f	247.6	303.5	1.23	614.6	2.48	350.2	1.41	420.3	1.70	379.6	1.53	224.3	0.91	259.3	1.05
A0A087WV	Receptor-tyr	58	75	1.29	66.9	1.15	70.1	1.21	55	0.95	58.1	1.00	70	1.21	55	0.95
P67936	Tropomyosin	328.1	474	1.44	453.2	1.38	532.2	1.62	517.2	1.58	458.3	1.40	433.9	1.32	343.7	1.05
A0A024RC	Peroxisome	446.8	477.6	1.07	483.8	1.08	540.7	1.21	509.1	1.14	603.2	1.35	356.8	0.80	456.9	1.02
P98160	Basement m	91.9	180.2	1.96	126.8	1.38	126.6	1.38	132.4	1.44	138.4	1.51	159.1	1.73	95.7	1.04
A0A0S4XQ	MHC Class															
Q8IY21	Probable AT	795	976.1	1.23	858.9	1.08	1005.7	1.27	930.4	1.17	1047.1	1.32	730.4	0.92	754.9	0.95
A0A0U1RR	Histone H2A	218.4	257.8	1.18	229.1	1.05	272.6	1.25	266.6	1.22	263.5	1.21	232.6	1.07	186	0.85
G3XAI2	Laminin sub	99.6	118.5	1.19	108.6	1.09	150.7	1.51	133.7	1.34	113.4	1.14	107.6	1.08	80.6	0.81
C9JC84	Fibrinogen g															
Q9H7P6	Multivesicul															
B7Z351	Secreted phc	40.3	47.7	1.18	36.1	0.90	40.2	1.00	46.4	1.15	41.8	1.04	46.3	1.15	34.7	0.86
Q9BWP8	Collectin-11	18.3	36	1.97	28.1	1.54	40.2	2.20	40.5	2.21	40.3	2.20	21.1	1.15	16.9	0.92
A0A024QZ	Serine/threo	359.7	297.4	0.83	286.2	0.80	380.1	1.06	538.4	1.50	430	1.20	246.8	0.69	223.6	0.62
A0A024QZ	Proteoglycar	132.4	161.4	1.22	127.2	0.96	176.4	1.33	128.5	0.97	136.2	1.03	156.7	1.18	108.9	0.82
A8K9X5	cDNA FLJ7	1282.7	1183.1	0.92	1057.9	0.82	1665	1.30	1951.7	1.52	1903.3	1.48	2038.2	1.59	880.6	0.69
P11717	Cation-indep															
A0A024R2V	Dystroglycar															
D6W5L6	Pulmonary s	27.1	35.8	1.32	23.6	0.87	32.4	1.20	34.2	1.26	28.6	1.06	30.3	1.12	38.6	1.42
P20023	Complement															
A0A060VC	MHC class I	1005.6	990.7	0.99	1154.2	1.15	874.8	0.87	858.5	0.85	929.5	0.92	1310.8	1.30	899.2	0.89
A8K2T7	Receptor pro															
A0A024R1C	Contactin 1,	41.9	50	1.19	47.3	1.13	56.2	1.34	40.4	0.96	46.9	1.12	43.8	1.05	29.5	0.70
Q9BTY2	Plasma alph	237.3	282.1	1.19	242.6	1.02	252.4	1.06	280.7	1.18	273.7	1.15	242.1	1.02	220.1	0.93
Q8IXL6	Extracellular	121.2	88.7	0.73	82.9	0.68	112.4	0.93	78.9	0.65	103.9	0.86	92.9	0.77	48.6	0.40
A0A075B6F	Dynein heav	130.9	147.4	1.13	89.2	0.68	150.5	1.15	147.8	1.13	144.5	1.10	106.1	0.81	77	0.59
A0A087WX	IgGfc-bindin	260.4	310.1	1.19	245.6	0.94	327.3	1.26	268	1.03	250	0.96	265.7	1.02	235.5	0.90

Q99727	Metalloprote	135	93.4	0.69	109.7	0.81	130.5	0.97	142.1	1.05	111.3	0.82	116.6	0.86	118.5	0.88
Q0ZCH6	Immunglobu	71.2	85.4	1.20	69.3	0.97	100	1.40	97.9	1.38	77.4	1.09	76.1	1.07	67.1	0.94
Q9UNN8	Endothelial	148.8	146.2	0.98	153.2	1.03	181.9	1.22	174.8	1.17	162.8	1.09	146.9	0.99	150.4	1.01
A0A0C4DG	Poliovirus re	336.2	336	1.00	313.3	0.93	359.9	1.07	385.4	1.15	333.7	0.99	411	1.22	253.4	0.75
J3KNB4	Cathelicidin	316.2	351.3	1.11	340.1	1.08	421	1.33	395.6	1.25	385.8	1.22	382.4	1.21	289.6	0.92
Q96RZ2	N-acetylgluc	45	70.6	1.57	65.6	1.46	59.8	1.33	47.2	1.05	49.8	1.11	40	0.89	42.2	0.94
Q4LE33	TNC variant															
A8K486	Peptidyl-pro	9.5	24.4	2.57	19	2.00	13.1	1.38	15.1	1.59	13.1	1.38	13.8	1.45	8.2	0.86
B2R6V9	cDNA, FLJ5	52.7	69.4	1.32	56.9	1.08	69.2	1.31	56.4	1.07	69.9	1.33	60.9	1.16	37.2	0.71
Q16706	Alpha-mann															
H3BQD0	Cerebellin-1	13	13.6	1.05	18.9	1.45	16.2	1.25	12	0.92	15.3	1.18	18.1	1.39	12.7	0.98
Q86U17	Serpin A11	54	74	1.37	58.2	1.08	76.4	1.41	76.2	1.41	78.9	1.46	89.4	1.66	54	1.00
A0A140VK	Leukotriene	31.4	35.1	1.12	29	0.92	31.9	1.02	27.7	0.88	33.2	1.06	24	0.76	28.4	0.90
A0A140GX	Platelet mem	40.9	62.2	1.52	51.3	1.25	81.4	1.99	44.4	1.09	39.6	0.97	41.9	1.02	35.7	0.87
Q96S96	Phosphatidy	32.3	34.8	1.08	29	0.90	43.1	1.33	35	1.08	33.3	1.03	46.8	1.45	26.7	0.83
O00151	PDZ and LI	78.2	103.2	1.32	93.2	1.19	99.7	1.27	104.3	1.33	97.3	1.24	94.3	1.21	70.4	0.90
A0A068B0	Cis-AB glyc	74.1	92.9	1.25	74.9	1.01	93.7	1.26	95.4	1.29	75.6	1.02	124.2	1.68	72.3	0.98
Q6YHK3	CD109 antig	163.1	197.5	1.21	155.1	0.95	190.1	1.17	219.3	1.34	176.1	1.08	195.4	1.20	149.5	0.92
B3KSS6	Phosphodies	385.6	316.3	0.82	291.1	0.75	329.9	0.86	343.6	0.89	295.2	0.77	417.6	1.08	321.5	0.83
B4DQY7	cDNA FLJ5	15.6	22.3	1.43	20.1	1.29	20.8	1.33	17.9	1.15	17.2	1.10	20	1.28	22.6	1.45
A0A024RD	Periostin, os															
P06733	Alpha-enola	49.7	71.8	1.44	57.6	1.16	92.1	1.85	66.6	1.34	70	1.41	62.9	1.27	52.2	1.05
Q86SQ4	Adhesion G-															
Q8TER0	Sushi, nidog															
A1L4H1	Soluble scav															
Q13740	CD166 antig	39.8	60	1.51	43.9	1.10	54.1	1.36	50.8	1.28	48.3	1.21	47	1.18	37.2	0.93
O94818	Nucleolar pr	60.9	66.8	1.10	54.2	0.89	83	1.36	70.8	1.16	68.8	1.13	56.3	0.92	44.9	0.74
P06703	Protein S100															
D3DPK5	SH3 domain	63.5	99	1.56	106.2	1.67	123.2	1.94	78.5	1.24	81.8	1.29	83.4	1.31	85	1.34
E9PK25	Cofilin-1 OS															
A0A125U0	GCT-A2 hea	18.9	24.4	1.29	19.9	1.05	30.3	1.60	26.6	1.41	20.8	1.10	26.9	1.42	15.2	0.80
D6W5K2	Thymosin, b	88	215	2.44	183.4	2.08	144.4	1.64	169.2	1.92	204.7	2.33	181.3	2.06	76.5	0.87
Q6PIY7	Poly(A) RN	67.2	60.9	0.91	63.3	0.94	61.6	0.92	56.5	0.84	52.3	0.78	61.6	0.92	43	0.64
Q96AN5	Transmembr	90	156.9	1.74	268.8	2.99	443.2	4.92	179.8	2.00	136.5	1.52	83.9	0.93	93.9	1.04
Q6P4F1	Alpha-(1,3)-															
Q9UL82	Myosin-reac															
Q7Z7M0	Multiple epi															
Q2KHR2	DNA-bindin															
Q75KY0	Putative unc															
V9HVY1	Epididymis	51.6	85.5	1.66	127.8	2.48	209.8	4.07	85.8	1.66	64.8	1.26	51.9	1.01	68	1.32
Q59EB6	Complemen	49.4	66.1	1.34	46.1	0.93	55.1	1.12	58.8	1.19	50.3	1.02	63.9	1.29	59.5	1.20
Q13131	5'-AMP-acti															
Q96RR4	Calcium/cal															
A0A0R4J2F	Spermatogei	22.3	78.2	3.51	53.8	2.41	56.5	2.53	70.6	3.17	68.9	3.09	36.4	1.63	52.8	2.37
P25391	Laminin sub	67.6	83.6	1.24	71.2	1.05	80.7	1.19	80.8	1.20	84.7	1.25	74	1.09	49.1	0.73
A0A0G2JN	Transcriptio															
E5RG27	Casein kinas	115.8	125	1.08	109	0.94	168.1	1.45	173.4	1.50	152.4	1.32	80	0.69	88.9	0.77

