

Table S4. Peak intensities of 113 proteins (light only)

Entry Name	Sample Name																
	C1	C2	C3	C4	C5	AS1	AS2	AS3	AS4	AS5	LC1	LC2	LC3	LC4	LC5	BC1	BC2
CFAH_HUMAN	1.25E+07	1.13E+07	1.08E+07	1.27E+07	1.40E+07	5.69E+06	5.21E+06	4.26E+06	2.61E+06	1.94E+06	6.57E+06	5.61E+06	6.32E+06	4.79E+06	1.63E+06	2.38E+06	3.31E+06
HBA_HUMAN	2.15E+07	2.34E+07	7.27E+06	1.30E+07	2.11E+07	4.38E+06	1.47E+06	1.86E+06	2.32E+06	1.01E+06	2.27E+07	1.06E+08	1.39E+07	2.05E+07	4.52E+06	8.94E+05	9.56E+05
CLUS_HUMAN	5.61E+07	4.94E+07	4.77E+07	3.90E+07	4.19E+07	2.79E+07	2.41E+07	1.66E+07	2.56E+07	1.44E+07	2.19E+07	2.54E+07	1.75E+07	1.55E+07	1.19E+07	1.03E+07	1.44E+07
C1QB_HUMAN	2.28E+06	1.43E+06	2.29E+06	1.85E+06	2.70E+06	2.21E+03	2.21E+03	2.21E+03	2.21E+03	2.21E+03	3.02E+06	1.07E+06	2.75E+06	2.02E+06	6.62E+05	5.05E+05	2.54E+05
ANGT_HUMAN	1.79E+07	7.94E+06	7.36E+06	8.68E+06	5.54E+06	5.52E+06	5.90E+06	4.64E+06	3.97E+06	6.93E+06	1.87E+06	2.42E+06	2.23E+06	2.30E+06	3.40E+06	9.10E+05	2.26E+06
ALBU_HUMAN	2.19E+07	1.53E+07	1.47E+07	1.23E+07	1.42E+07	5.98E+06	8.65E+06	6.61E+06	7.00E+06	1.70E+06	1.05E+07	7.63E+06	8.64E+06	5.68E+06	4.64E+06	7.82E+06	5.88E+06
THRB_HUMAN	1.20E+07	5.52E+06	6.42E+06	6.50E+06	6.44E+06	4.06E+06	3.49E+06	3.23E+06	3.11E+06	1.60E+06	7.84E+06	5.17E+06	6.20E+06	5.95E+06	3.24E+06	1.93E+06	4.37E+06
A1BG_HUMAN	4.10E+07	1.97E+07	3.61E+07	5.11E+07	3.89E+07	1.70E+07	1.35E+07	1.68E+07	1.28E+07	1.02E+07	2.74E+07	2.63E+07	1.79E+07	1.23E+07	1.14E+07	3.99E+06	5.48E+06
PROZ_HUMAN	1.99E+06	1.77E+06	1.19E+06	2.81E+06	2.14E+06	1.00E+06	8.86E+05	2.78E+05	2.21E+03	1.77E+05	4.45E+05	8.97E+05	9.46E+05	2.21E+03	3.23E+05	2.21E+03	3.06E+05
HPTR_HUMAN	6.04E+06	5.37E+06	5.31E+06	3.07E+06	2.05E+06	4.78E+06	1.30E+06	1.73E+06	1.02E+06	1.64E+06	1.67E+06	2.08E+06	6.69E+05	1.01E+06	7.41E+05	5.14E+05	1.68E+06
APOA1_HUMAN	1.29E+09	7.14E+08	8.63E+08	1.01E+09	1.03E+09	2.89E+08	2.67E+08	2.06E+08	2.18E+08	9.91E+07	1.01E+09	6.77E+08	7.56E+08	5.34E+08	2.26E+08	5.20E+08	3.36E+08
FETUA_HUMAN	5.16E+07	2.47E+07	3.27E+07	3.20E+07	3.56E+07	1.48E+07	9.09E+06	1.16E+07	4.60E+06	3.19E+06	4.39E+07	1.01E+07	3.72E+07	2.37E+07	6.69E+06	1.57E+07	1.13E+07
CO5_HUMAN	1.65E+07	1.02E+07	1.34E+07	1.48E+07	8.44E+06	1.50E+07	6.40E+06	6.68E+06	5.23E+06	6.54E+06	7.32E+06	7.13E+06	4.02E+06	2.51E+06	1.76E+06	1.01E+06	4.30E+06
A1AT_HUMAN	4.15E+08	3.27E+08	4.07E+08	2.68E+08	2.21E+08	1.76E+08	2.49E+08	2.82E+08	2.14E+08	1.81E+08	1.41E+08	9.83E+07	7.41E+07	5.55E+07	4.83E+07	3.80E+07	8.48E+07
ITIH2_HUMAN	7.87E+07	5.92E+07	4.87E+07	5.54E+07	2.95E+07	3.15E+07	1.93E+07	1.83E+07	1.23E+07	7.33E+06	1.44E+07	2.19E+07	1.31E+07	1.27E+07	9.89E+06	4.91E+06	1.08E+07
CBG_HUMAN	2.39E+07	1.78E+07	1.75E+07	1.77E+07	1.90E+07	9.19E+06	7.78E+06	7.32E+06	2.21E+03	3.41E+06	1.78E+07	1.23E+07	1.48E+07	1.26E+07	6.15E+06	5.89E+06	7.52E+06
APOD_HUMAN	1.38E+07	1.40E+07	1.24E+07	7.17E+06	7.79E+06	7.73E+06	3.64E+06	3.12E+06	5.48E+06	2.23E+06	3.08E+06	4.07E+06	1.54E+06	9.02E+05	9.21E+05	9.49E+05	4.34E+06
LYVE1_HUMAN	1.56E+06	1.02E+06	1.11E+06	1.10E+06	8.51E+05	3.64E+05	3.93E+05	3.67E+05	2.85E+05	1.83E+05	3.99E+05	4.29E+05	3.49E+05	2.37E+05	8.28E+04	1.97E+05	2.53E+05
CO8B_HUMAN	4.07E+06	2.91E+06	3.19E+06	3.14E+06	3.31E+06	1.69E+06	1.08E+06	1.19E+06	4.14E+05	4.21E+05	5.22E+05	1.10E+06	4.41E+05	2.37E+05	4.14E+05	3.32E+05	4.90E+05
CPN2_HUMAN	3.21E+07	2.35E+07	1.37E+07	1.59E+07	3.94E+07	4.70E+06	9.02E+06	4.46E+06	4.01E+06	1.84E+06	2.21E+03	2.76E+06	2.21E+03	1.57E+06	1.80E+06	3.95E+06	1.38E+06
CO6_HUMAN	7.19E+06	3.35E+06	4.13E+06	3.66E+06	1.09E+06	1.95E+06	9.51E+05	1.26E+06	5.08E+05	8.10E+05	3.98E+05	5.63E+05	1.30E+05	1.50E+05	4.09E+05	1.08E+05	7.70E+05
ANT3_HUMAN	5.48E+07	3.43E+07	2.44E+07	1.81E+07	2.33E+07	1.37E+07	1.71E+07	1.28E+07	1.71E+07	6.64E+06	1.39E+07	1.02E+07	1.75E+07	1.75E+07	7.52E+06	1.59E+07	1.31E+07
APOC3_HUMAN	2.59E+08	3.65E+08	1.59E+08	4.37E+08	3.33E+08	5.71E+07	4.55E+07	3.26E+07	5.42E+07	1.64E+07	4.65E+07	4.56E+07	5.15E+07	4.05E+07	5.08E+07	1.33E+07	2.16E+07
CFAB_HUMAN	2.31E+07	1.31E+07	1.36E+07	1.32E+07	1.45E+07	6.35E+06	6.21E+06	7.26E+06	2.14E+06	2.64E+06	6.53E+06	4.75E+06	2.16E+06	1.84E+06	2.54E+06	1.38E+06	2.19E+06
LDHB_HUMAN	6.12E+05	4.94E+05	6.14E+05	6.54E+05	7.74E+05	2.28E+05	4.05E+05	7.21E+05	4.44E+05	1.76E+05	4.59E+05	1.41E+05	1.64E+05	1.23E+05	6.24E+04	2.71E+05	1.13E+05
CD5L_HUMAN	6.18E+06	2.47E+06	9.03E+06	3.74E+06	4.04E+06	3.61E+06	3.66E+06	2.21E+06	1.66E+06	1.64E+06	6.78E+05	1.78E+06	1.11E+06	2.42E+06	2.58E+06	1.33E+06	1.99E+06
ITIH4_HUMAN	2.95E+06	1.88E+06	5.18E+05	2.35E+06	2.05E+06	1.81E+06	8.75E+05	9.80E+05	1.16E+06	3.86E+06	8.17E+05	5.46E+05	4.77E+05	3.26E+05	2.65E+05	1.28E+06	2.01E+06
IGHA1_HUMAN	9.80E+07	1.11E+08	1.53E+08	1.14E+08	1.13E+08	4.66E+07	5.33E+07	5.95E+07	3.71E+07	1.38E+07	1.30E+08	7.42E+07	1.68E+08	7.50E+07	7.04E+07	4.46E+07	4.95E+07
HBB_HUMAN	4.72E+06	8.95E+06	3.19E+06	6.67E+06	6.08E+06	5.38E+06	5.50E+05	6.20E+05	4.57E+05	1.18E+06	3.17E+06	2.50E+07	2.29E+06	3.12E+06	1.09E+06	4.11E+05	2.18E+06
APOA2_HUMAN	7.83E+07	5.59E+07	4.50E+07	5.17E+07	6.60E+07	2.38E+07	2.32E+07	1.99E+07	2.28E+07	7.71E+06	5.24E+07	2.21E+03	5.58E+07	4.55E+07	2.02E+07	3.78E+07	2.54E+07
CRAC1_HUMAN	1.03E+06	3.81E+05	4.53E+05	5.09E+05	3.18E+05	2.23E+05	9.17E+04	7.10E+04	5.29E+04	8.74E+04	9.54E+04	7.92E+04	5.96E+04	4.82E+04	2.21E+04	4.56E+04	7.04E+04
C4BPA_HUMAN	2.21E+03	1.09E+08	1.15E+08	8.57E+07	1.33E+08	2.60E+07	1.94E+07	2.44E+07	1.92E+07	7.27E+06	5.49E+07	3.54E+07	3.29E+07	2.20E+07	2.26E+07	1.45E+07	1.40E+07
PROS_HUMAN	1.25E+06	7.94E+05	5.61E+05	7.09E+05	5.19E+05	5.02E+06	2.33E+06	3.29E+06	2.11E+06	1.13E+06	2.88E+05	3.19E+05	2.21E+03	2.21E+03	2.18E+05	2.21E+03	1.26E+06
NCHL1_HUMAN	2.21E+03	2.46E+05	2.16E+05	2.16E+05	2.81E+05	1.55E+05	1.29E+05	9.09E+04	1.14E+05	6.36E+04	1.43E+05	3.07E+05	1.43E+05	1.07E+05	1.03E+05	1.30E+05	2.21E+03
PRG4_HUMAN	5.47E+05	3.13E+05	3.53E+05	8.46E+05	1.13E+06	1.12E+05	1.02E+05	3.57E+05	3.05E+05	2.48E+04	5.58E+05	4.01E+05	7.00E+05	8.42E+05	1.47E+05	2.64E+05	1.20E+05
ALS_HUMAN	7.43E+06	6.46E+06	7.33E+06	7.59E+06	6.66E+06	2.13E+06	7.88E+05	1.63E+06	8.24E+05	5.65E+05	2.10E+06	1.18E+06	1.16E+06	1.48E+06	1.52E+06	1.20E+06	1.38E+06
IGD_HUMAN	7.73E+06	4.85E+06	1.00E+07	6.38E+06	5.98E+06	1.71E+06	1.46E+06	9.53E+05	7.62E+05	5.60E+05	2.47E+06	1.33E+06	1.91E+06	1.12E+06	5.10E+05	1.03E+06	9.32E+05
LUM_HUMAN	6.80E+06	3.49E+06	3.22E+06	3.78E+06	4.27E+06	1.49E+06	1.49E+06	1.18E+06	8.61E+05	3.17E+05	2.07E+06	1.88E+06	2.19E+06	9.39E+05	6.38E+05	1.33E+06	7.14E+05
TRFE_HUMAN	4.05E+08	2.01E+08	1.67E+08	1.78E+08	2.23E+08	1.12E+08	6.30E+07	6.49E+07	2.77E+07	4.65E+07	5.64E+07	4.82E+07	4.01E+07	2.98E+07	2.56E+07	4.25E+07	4.09E+07
APOH_HUMAN	8.57E+05	1.87E+06	2.02E+06	1.72E+06	1.41E+06	1.88E+06	1.80E+06	1.50E+06	2.03E+06	1.99E+06	1.12E+06	1.04E+06	8.20E+05	6.78E+05	5.13E+05	5.14E+05	1.59E+06
CO2_HUMAN	5.80E+06	4.74E+06	4.31E+06	4.49E+06	4.88E+06	1.59E+06	1.07E+06	1.30E+06	1.08E+06	5.73E+05	2.42E+06	2.33E+06	1.26E+06	1.34E+06	1.19E+06	4.75E+06	6.55E+05
TSP1_HUMAN	2.60E+06	4.24E+06	5.75E+06	4.56E+06	3.91E+06	1.87E+06	1.56E+06	4.98E+05	1.83E+06	6.15E+05	1.57E+06	1.53E+06	2.72E+06	2.01E+06	1.22E+06	7.58E+05	5.14E+05
LYAM1_HUMAN	2.21E+03	1.89E+05	1.12E+05	1.47E+05	2.21E+03	2.21E+03	2.21E+03	2.21E+03	2.21E+03	2.21E+03	3.63E+04	3.73E+04	6.18E+04	2.21E+03	3.39E+04	6.71E+04	2.21E+03
APOB_HUMAN	2.04E+07	1.41E+07	1.30E+07	1.82E+07	1.52E+07	7.26E+06	6.58E+06	6.91E+06	6.09E+06	3.26E+06	6.64E+06	1.00E+07	3.10E+06	8.42E+06	5.51E+06	3.73E+06	8.71E+06
APOA_HUMAN	2.82E+05	1.14E+05	1.31E+05	1.72E+05	2.21E+03	5.55E+04	2.21E+03	2.24E+05	3.52E+04	2.21E+03	2.16E+05	7.65E+04	2.59E+04	2.45E+04	1.74E+05	1.13E+05	1.09E+05
CFAI_HUMAN	1.96E+07	1.42E+07	1.47E+07	2.25E+07	1.55E+07	2.47E+06	4.67E+06	5.03E+06	6.57E+06	3.57E+06	6.22E+06	2.21E+03	3.95E+06	2.62E+06	1.73E+06	7.18E+05	2.56E+06
VWF_HUMAN	3.76E+05	2.54E+05	2.71E+06	1.55E+06	7.06E+05	8.78E+05	1.80E+06	2.46E+06	2.49E+06	1.14E+06	1.96E+06	1.54E+06	3.91E+05	4.23E+05	7.62E+05	2.19E+05	3.70E+05

AMBP_HUMAN	1.40E+07	1.52E+07	1.07E+07	1.57E+07	1.62E+06	1.33E+07	5.48E+06	7.93E+06	6.30E+06	5.73E+06	7.87E+05	2.02E+06	1.01E+06	3.83E+05	1.42E+06	2.21E+03	2.19E+06
KAIN_HUMAN	2.28E+06	1.50E+06	1.44E+06	1.61E+06	1.54E+06	1.05E+06	4.25E+05	2.80E+05	1.61E+05	2.79E+05	6.31E+05	5.49E+05	5.15E+05	3.51E+05	2.58E+05	1.91E+05	4.16E+05
IGHM_HUMAN	6.81E+07	4.00E+07	1.42E+08	6.15E+07	6.86E+07	5.09E+07	7.16E+07	2.85E+07	2.87E+07	1.75E+07	1.24E+07	2.17E+07	1.81E+07	3.52E+07	2.79E+07	2.09E+07	1.38E+07
KNGL1_HUMAN	1.03E+08	9.28E+07	8.69E+07	8.24E+07	1.15E+08	4.08E+07	2.98E+07	2.78E+07	2.33E+07	9.04E+06	3.99E+07	3.74E+07	4.08E+07	2.41E+07	3.72E+07	1.58E+07	1.31E+07
EPCR_HUMAN	5.44E+05	9.17E+05	4.13E+05	1.15E+06	3.43E+05	1.37E+05	1.07E+05	2.21E+03	4.42E+05	5.82E+04	1.14E+05	6.71E+05	2.21E+03	2.21E+03	2.21E+03	2.21E+03	2.21E+03
PGRP2_HUMAN	1.45E+06	1.16E+06	9.19E+05	7.80E+05	7.92E+05	5.72E+05	2.21E+03	2.76E+05	2.21E+03	2.21E+03	1.92E+05	2.96E+05	1.27E+05	9.58E+04	2.21E+03	2.66E+05	3.76E+05
APOC2_HUMAN	2.77E+07	3.74E+07	1.73E+07	5.82E+07	3.15E+07	6.44E+06	4.91E+06	6.26E+06	7.79E+06	1.90E+06	4.99E+06	7.17E+06	5.68E+06	5.18E+06	3.09E+06	1.80E+06	3.81E+06
CO7_HUMAN	8.12E+06	7.57E+06	7.07E+06	5.92E+06	4.63E+06	2.68E+06	1.20E+06	1.32E+06	6.80E+05	7.88E+05	1.05E+06	1.67E+06	8.08E+05	5.80E+05	7.84E+05	6.13E+05	1.21E+06
TENX_HUMAN	8.34E+05	6.66E+05	6.40E+05	6.43E+05	9.19E+05	2.42E+05	2.21E+03	2.21E+03	9.37E+04	2.21E+03	3.40E+05	3.05E+05	2.88E+05	1.72E+05	6.89E+04	2.21E+03	2.21E+03
CXCL7_HUMAN	3.66E+07	5.85E+07	3.76E+07	2.05E+07	1.78E+07	8.40E+06	7.43E+06	1.45E+06	5.71E+06	1.47E+06	1.78E+06	3.15E+06	3.32E+06	1.90E+06	3.68E+06	2.21E+06	3.75E+06
FCN2_HUMAN	1.37E+06	1.64E+06	1.06E+06	1.13E+06	6.41E+05	2.85E+05	2.21E+03	2.21E+03	2.22E+05	1.00E+05	2.21E+03	2.21E+03	2.21E+03	8.91E+04	6.59E+04	2.21E+03	1.20E+05
FA9_HUMAN	2.21E+03	3.87E+06	3.81E+06	4.49E+06	4.01E+06	2.21E+03	2.10E+06	1.81E+06	1.97E+05	7.10E+05	1.14E+06	1.65E+06	9.56E+05	6.03E+05	1.27E+06	2.78E+05	4.54E+05
CERU_HUMAN	6.52E+07	4.37E+07	4.73E+07	4.81E+07	6.34E+07	2.68E+07	3.30E+07	4.21E+07	3.31E+07	1.03E+07	9.53E+07	4.77E+07	4.13E+07	3.14E+07	2.36E+07	1.30E+07	1.28E+07
IPSP_HUMAN	1.82E+06	1.12E+06	1.67E+06	1.56E+06	1.08E+06	9.70E+05	3.12E+05	2.46E+05	4.65E+05	2.12E+05	1.27E+06	1.33E+06	1.53E+06	1.21E+06	7.02E+05	3.36E+05	3.60E+05
BTD_HUMAN	3.45E+06	3.04E+06	2.78E+06	3.30E+06	3.17E+06	1.17E+06	7.70E+05	6.03E+05	5.35E+05	2.43E+05	1.13E+06	1.09E+06	1.20E+06	1.05E+06	6.63E+05	6.09E+05	2.86E+05
DOPO_HUMAN	3.19E+05	2.94E+05	6.52E+05	3.65E+05	7.85E+05	2.21E+03	2.21E+03	2.21E+03	2.21E+03	2.21E+03	8.97E+04	1.14E+05	1.30E+05	3.75E+04	2.21E+03	5.29E+04	1.16E+05
ICOSL_HUMAN	3.75E+05	3.08E+05	2.73E+05	3.00E+05	3.26E+05	1.13E+05	1.08E+05	7.18E+04	4.91E+04	2.21E+03	1.05E+05	1.66E+05	1.04E+05	5.22E+04	2.21E+03	8.86E+04	6.83E+04
FHR1_HUMAN	2.21E+03	2.21E+03	2.21E+03	9.09E+05	1.78E+06	2.09E+05	2.21E+03	4.19E+05	2.21E+03	2.21E+03	8.26E+05	4.05E+05	1.85E+06	8.80E+05	9.63E+05	2.21E+03	1.55E+05
IL1A1_HUMAN	7.03E+05	4.87E+05	5.81E+05	5.58E+05	5.29E+05	2.21E+03	2.21E+03	2.21E+03	2.21E+03	2.21E+03	2.54E+05	1.72E+05	1.32E+05	2.38E+05	1.10E+05	2.21E+03	2.21E+03
QSOX1_HUMAN	5.98E+05	3.14E+05	3.34E+05	3.86E+05	2.51E+05	1.51E+05	1.31E+05	1.42E+05	1.88E+05	3.30E+04	1.34E+05	1.28E+05	7.87E+04	6.48E+04	7.78E+04	2.21E+03	2.21E+03
SAMP_HUMAN	1.23E+07	9.78E+06	1.05E+07	8.70E+06	2.35E+07	4.09E+06	3.58E+06	3.62E+06	5.52E+06	8.82E+05	5.84E+06	4.97E+06	5.32E+06	1.98E+06	2.71E+06	1.13E+06	1.43E+06
RET4_HUMAN	5.95E+06	7.44E+06	9.16E+06	7.93E+06	1.02E+07	9.49E+05	1.54E+06	1.15E+06	1.75E+06	1.63E+05	2.78E+06	2.14E+06	5.95E+06	2.45E+06	2.13E+06	7.13E+05	2.94E+05
MASP2_HUMAN	1.95E+06	1.38E+06	1.81E+06	1.33E+06	1.50E+06	5.30E+05	5.34E+05	6.09E+05	6.37E+05	2.66E+05	1.04E+06	4.66E+05	5.58E+05	2.22E+05	5.20E+05	2.13E+05	1.58E+05
IGF2_HUMAN	1.23E+06	8.05E+05	8.03E+05	9.16E+05	1.01E+06	2.97E+05	1.97E+05	2.79E+05	2.48E+05	9.04E+04	4.48E+05	3.42E+05	4.44E+05	3.99E+05	1.79E+05	4.25E+05	2.48E+05
IGG1_HUMAN	6.69E+06	3.66E+06	9.99E+06	4.03E+06	5.10E+06	2.81E+06	2.28E+06	2.27E+06	2.67E+06	6.03E+05	2.39E+06	2.76E+06	3.15E+06	3.86E+06	1.80E+06	1.64E+06	1.17E+06
CNTN1_HUMAN	6.02E+05	2.78E+05	3.34E+05	2.99E+05	2.21E+03	7.75E+04	1.10E+05	9.67E+04	9.63E+04	4.78E+04	1.50E+05	2.07E+05	1.09E+05	1.27E+05	5.35E+04	1.15E+05	8.00E+04
F13B_HUMAN	2.73E+06	2.05E+06	1.32E+06	1.04E+06	1.60E+06	8.61E+05	8.34E+05	1.10E+06	9.46E+05	2.49E+05	1.47E+06	9.68E+05	1.43E+06	1.15E+06	4.00E+05	1.50E+06	8.90E+05
LBP_HUMAN	5.14E+06	3.32E+06	4.18E+06	3.19E+06	3.62E+06	7.72E+05	8.47E+06	7.28E+06	3.72E+06	2.10E+06	1.55E+06	2.63E+06	9.38E+05	6.60E+05	5.74E+05	4.99E+05	7.20E+05
MASP1_HUMAN	1.75E+06	1.56E+06	1.89E+06	1.36E+06	1.62E+06	8.64E+05	3.12E+05	6.27E+05	4.84E+05	3.90E+05	1.63E+06	1.03E+06	5.51E+05	7.54E+05	5.28E+05	2.74E+05	4.87E+05
CYTC_HUMAN	2.20E+06	1.90E+06	1.53E+06	1.70E+06	1.27E+06	7.14E+05	1.34E+06	1.02E+06	1.86E+06	2.62E+05	5.05E+05	6.60E+05	3.37E+05	1.89E+05	2.76E+05	1.29E+05	2.07E+05
CHLE_HUMAN	6.90E+06	4.14E+06	2.43E+06	4.31E+06	5.23E+06	9.81E+05	7.79E+05	3.62E+05	4.80E+05	3.50E+05	1.23E+06	1.27E+06	1.27E+06	9.81E+05	5.41E+05	4.84E+05	4.73E+05
PZP_HUMAN	2.75E+06	3.84E+05	3.83E+06	8.11E+05	3.62E+05	3.19E+05	2.21E+03	4.91E+05	5.32E+05	1.62E+05	1.47E+06	1.48E+06	7.00E+05	1.29E+06	9.71E+05	8.09E+05	8.75E+05
CAD13_HUMAN	3.73E+05	2.14E+05	3.70E+05	2.46E+05	2.40E+05	1.84E+05	1.43E+05	1.23E+05	9.50E+04	9.10E+04	2.21E+03	2.09E+05	2.20E+05	2.21E+03	2.21E+03	1.62E+05	1.58E+05
SAA2_HUMAN	2.21E+03	2.21E+03	2.21E+03	2.21E+03	2.21E+03	2.21E+03	2.50E+07	2.65E+07	1.24E+07	4.45E+06	3.11E+05	2.21E+03	2.21E+03	2.21E+03	8.12E+04	2.21E+03	3.24E+05
ITIH3_HUMAN	8.61E+06	6.52E+06	8.90E+06	6.19E+06	5.77E+06	1.99E+06	4.20E+06	4.51E+06	7.01E+06	3.13E+06	8.48E+06	5.75E+06	2.05E+06	2.10E+06	2.53E+06	2.67E+06	2.17E+06
AFAM_HUMAN	1.38E+06	9.02E+05	6.24E+05	8.81E+05	8.49E+05	4.31E+05	5.61E+05	4.09E+05	5.07E+05	3.21E+05	3.01E+05	3.99E+05	3.16E+05	3.67E+05	1.83E+05	3.58E+05	4.98E+05
LCAT_HUMAN	8.26E+05	8.28E+05	8.45E+05	1.03E+06	5.31E+05	8.08E+05	3.68E+05	3.10E+05	2.08E+05	2.70E+05	2.21E+05	2.28E+05	1.72E+05	2.30E+05	1.81E+05	8.27E+04	5.36E+05
TETN_HUMAN	8.71E+06	4.61E+06	5.62E+06	4.93E+06	5.92E+06	1.61E+06	1.24E+06	8.32E+05	8.86E+05	6.16E+05	1.49E+06	2.37E+06	1.25E+06	1.25E+06	5.84E+05	7.53E+05	1.35E+06
FINC_HUMAN	3.35E+06	6.73E+06	2.25E+07	3.24E+07	2.54E+07	1.43E+07	9.29E+06	1.49E+07	9.35E+06	6.75E+06	1.55E+07	1.42E+07	1.62E+07	7.39E+06	7.80E+06	1.48E+07	
FHR5_HUMAN	5.70E+05	5.52E+05	4.00E+05	4.30E+05	4.69E+05	2.13E+05	2.21E+03	4.12E+05	2.51E+05	2.21E+03	2.99E+05	2.93E+05	2.12E+05	1.79E+05	2.21E+03	2.21E+03	2.21E+03
TTHY_HUMAN	4.70E+07	2.82E+07	2.51E+07	2.16E+07	1.92E+07	1.88E+07	8.55E+06	7.33E+06	6.91E+06	1.43E+07	5.66E+06	1.15E+07	5.50E+06	6.87E+06	9.04E+06	4.68E+06	2.93E+07
IGKC_HUMAN	2.27E+08	2.21E+03	2.21E+03	2.21E+03	5.21E+08	2.21E+03	1.23E+08	1.42E+08	1.57E+08	4.10E+07	2.30E+08	2.02E+08	2.69E+08	2.03E+08	1.22E+08	9.23E+07	2.21E+03
ZA2G_HUMAN	1.47E+06	1.02E+06	4.38E+06	1.37E+06	1.53E+06	5.70E+05	7.28E+05	7.63E+05	1.32E+06	3.53E+05	1.24E+06	7.93E+05	1.26E+06	7.26E+05	4.61E+05	4.33E+05	2.56E+05
IGA2_HUMAN	4.90E+07	6.13E+07	6.61E+07	3.13E+07	5.18E+07	7.29E+06	1.30E+07	1.64E+07	9.95E+06	4.27E+06	3.29E+07	2.16E+07	1.67E+07	2.35E+07	3.21E+07	1.41E+07	1.12E+07
TFR1_HUMAN	2.21E+03	2.21E+03	3.38E+05	3.78E+05	3.29E+05	2.21E+03	2.21E+03	2.21E+03	2.21E+03	2.21E+03	2.65E+05	2.64E+05	2.21E+03	1.01E+05	5.30E+04	5.12E+05	2.21E+03
HRG_HUMAN	3.85E+07	3.00E+07	2.08E+07	2.22E+07	1.34E+07	1.36E+07	6.93E+06	1.03E+07	5.08E+06	9.62E+06	1.45E+07	1.18E+07	1.70E+07	1.07E+07	3.89E+06	1.04E+07	1.47E+07
APOE_HUMAN	2.92E+07	1.29E+07	6.80E+06	1.44E+07	8.64E+06	3.43E+06	5.75E+06	3.71E+06	7.83E+06	2.54E+06	3.12E+06	3.69E+06	1.68E+06	2.63E+06	1.98E+06	1.69E+06	1.62E+06
CO4A_HUMAN	1.83E+07	5.97E+06	7.79E+06	7.67E+06	1.00E+07	2.01E+06	1.92E+06	3.29E+06	2.73E+06	1.16E+06	1.06E+07	1.90E+07	6.91E+06	9.97E+06	2.47E+06	2.31E+06	1.92E+06
MA1A1_HUMAN	1.03E+06	7.87E+05	8.68E+05	9.20E+05	9.03E+05	4.45E+05	1.06E+06	1.22E+06	8.81E+05	2.81E+05	7.04E+05	6.77E+05	7.06E+05	5.32E+05	3.87E+05	7.24E+05	5.95E+05
CSF1R_HUMAN	2.21E+03	1.61E+05	9.24E+04	1.33E+05	2.21E+03	1.74E+05	1.69E+05	2.15E+05	2.25E+05	3.74E+04	4.36E+05	2.22E+05	1.60E+05	1.69E+05	2.21E+03	7.73E+05	1.88E+05
IGLL1_HUMAN	1.55E+06	8.86E+05	8.37E+05	8.88E+05	1.09E+06	3.81E+05	5.39E+05	5.14E+05	5.25E+05	1.66E+05	1.23E+06	9.70E+05	3.18E+06	1.05E+06	5.88E+05	4.97E+05	9.09E+05

[illegible]

	BC3	BC4	BC5
CFAH_HUMAN	3.44E+06	2.98E+06	2.35E+06
HBA_HUMAN	5.59E+05	2.55E+06	4.18E+05
CLUS_HUMAN	1.24E+07	8.94E+06	9.49E+06
C1QB_HUMAN	5.16E+05	3.97E+05	2.12E+05
ANGT_HUMAN	1.22E+06	1.10E+06	1.30E+06
ALBU_HUMAN	9.44E+06	7.38E+06	7.01E+06
THRB_HUMAN	4.12E+06	3.35E+06	2.48E+06
A1BG_HUMAN	7.35E+06	5.15E+06	2.45E+06
PROZ_HUMAN	2.21E+03	2.21E+03	2.21E+03
HPTR_HUMAN	3.08E+06	1.81E+06	1.80E+06
APOA1_HUMAN	3.57E+08	3.47E+08	2.86E+08
FETUA_HUMAN	2.07E+07	1.53E+07	1.13E+07
CO5_HUMAN	2.95E+06	2.08E+06	1.55E+06
A1AT_HUMAN	7.37E+07	7.01E+07	5.44E+07
ITIH2_HUMAN	1.07E+07	7.69E+06	7.93E+06
CBG_HUMAN	9.19E+06	6.10E+06	5.20E+06
APOD_HUMAN	1.50E+06	1.46E+06	1.32E+06
LYVE1_HUMAN	2.97E+05	1.92E+05	1.80E+05
CO8B_HUMAN	6.14E+05	4.36E+05	3.24E+05
CPN2_HUMAN	2.21E+03	2.21E+03	2.01E+06
CO6_HUMAN	5.67E+05	4.54E+05	3.49E+05
ANT3_HUMAN	1.63E+07	1.33E+07	9.38E+06
APOC3_HUMAN	2.01E+07	1.57E+07	1.41E+07
CFAB_HUMAN	3.72E+06	2.48E+06	1.39E+06
LDHB_HUMAN	1.93E+05	1.32E+05	1.29E+05
CD5L_HUMAN	2.96E+06	1.95E+06	1.66E+06
ITIH4_HUMAN	1.00E+06	9.24E+05	1.12E+06
IGHA1_HUMAN	4.80E+07	4.72E+07	5.41E+07
HBB_HUMAN	4.50E+05	2.34E+06	4.78E+05
APOA2_HUMAN	2.94E+07	3.10E+07	2.23E+07
CRAC1_HUMAN	5.47E+04	3.88E+04	4.05E+04
C4BPA_HUMAN	1.86E+07	1.58E+07	9.05E+06
PROS_HUMAN	6.07E+05	4.98E+05	3.89E+05
NCHL1_HUMAN	1.51E+05	8.55E+04	6.73E+04
PRG4_HUMAN	2.84E+05	4.41E+05	2.00E+05
ALS_HUMAN	2.03E+06	1.46E+06	1.11E+06
IGD_HUMAN	1.89E+06	1.00E+06	1.29E+06
LUM_HUMAN	1.14E+06	6.03E+05	5.04E+05
TRFE_HUMAN	6.14E+07	3.79E+07	2.84E+07
APOH_HUMAN	8.44E+05	7.91E+05	6.79E+05
CO2_HUMAN	6.13E+05	5.49E+05	3.84E+05
TSP1_HUMAN	1.02E+06	6.07E+05	7.65E+05
LYAM1_HUMAN	2.21E+03	7.80E+04	4.84E+04
APOB_HUMAN	9.13E+06	7.69E+06	7.56E+06
APOA_HUMAN	5.97E+04	7.84E+04	5.68E+04
CFAI_HUMAN	2.46E+06	1.08E+06	1.70E+06
VWF_HUMAN	1.61E+05	3.81E+05	1.68E+05

AMBP_HUMAN	7.56E+05	1.00E+06	1.05E+06
KAIN_HUMAN	6.91E+05	3.30E+05	1.89E+05
IGHM_HUMAN	2.71E+07	2.04E+07	1.39E+07
KNIG1_HUMAN	1.63E+07	1.34E+07	1.13E+07
EPCR_HUMAN	2.21E+03	2.21E+03	2.21E+03
PGRP2_HUMAN	4.94E+05	2.21E+03	2.21E+03
APOC2_HUMAN	3.14E+06	1.92E+06	1.79E+06
CO7_HUMAN	5.32E+05	6.35E+05	7.71E+05
TENX_HUMAN	1.46E+05	1.37E+05	2.21E+03
CXCL7_HUMAN	3.52E+06	1.96E+06	2.38E+06
FCN2_HUMAN	1.34E+05	1.33E+05	1.21E+05
FA9_HUMAN	5.93E+05	4.78E+05	3.26E+05
CERU_HUMAN	1.33E+07	1.39E+07	7.77E+06
IPSP_HUMAN	2.84E+05	2.58E+05	3.26E+05
BTD_HUMAN	6.09E+05	4.16E+05	2.86E+05
DOPO_HUMAN	4.59E+04	3.61E+04	5.14E+04
ICOSL_HUMAN	6.86E+04	5.05E+04	5.42E+04
FHR1_HUMAN	2.21E+03	2.21E+03	2.21E+03
IL1AP_HUMAN	2.21E+03	2.21E+03	2.21E+03
QSOX1_HUMAN	5.56E+04	5.04E+04	4.16E+04
SAMP_HUMAN	2.28E+06	2.40E+06	8.80E+05
RET4_HUMAN	7.47E+05	3.50E+05	2.38E+05
MASP2_HUMAN	2.21E+03	2.18E+05	2.21E+03
IGF2_HUMAN	3.56E+05	2.40E+05	3.01E+05
IGG1_HUMAN	1.39E+06	1.35E+06	9.65E+05
CNTN1_HUMAN	1.18E+05	9.09E+04	7.16E+04
F13B_HUMAN	1.24E+06	1.03E+06	1.08E+06
LBP_HUMAN	8.37E+05	1.41E+06	8.99E+05
MASP1_HUMAN	4.96E+05	2.80E+05	2.43E+05
CYTC_HUMAN	2.55E+05	2.13E+05	1.73E+05
CHLE_HUMAN	6.39E+05	5.49E+05	4.32E+05
PZP_HUMAN	1.24E+06	9.88E+05	1.33E+06
CAD13_HUMAN	1.56E+05	1.46E+05	1.55E+05
SAA2_HUMAN	2.21E+03	6.13E+05	1.19E+05
ITIH3_HUMAN	2.36E+06	1.57E+06	1.11E+06
AFAM_HUMAN	3.61E+05	3.41E+05	4.29E+05
LCAT_HUMAN	3.05E+05	2.66E+05	1.85E+05
TETN_HUMAN	1.68E+06	8.92E+05	9.61E+05
FINC_HUMAN	1.22E+07	1.15E+07	4.73E+06
FHR5_HUMAN	2.21E+03	2.21E+03	2.21E+03
TTHY_HUMAN	1.88E+07	1.23E+07	1.85E+07
IGKC_HUMAN	6.39E+07	3.66E+07	2.21E+03
ZA2G_HUMAN	4.05E+05	3.13E+05	2.91E+05
IGA2_HUMAN	1.32E+07	2.38E+07	1.71E+07
TFR1_HUMAN	2.32E+05	1.63E+05	1.22E+05
HRG_HUMAN	1.49E+07	8.33E+06	1.02E+07
APOE_HUMAN	2.03E+06	1.86E+06	2.57E+06
CO4A_HUMAN	2.60E+06	2.35E+06	1.31E+06
MA1A1_HUMAN	8.57E+05	9.10E+05	6.36E+05
CSF1R_HUMAN	2.21E+03	3.40E+05	3.09E+05
IGLL1_HUMAN	8.24E+05	8.17E+05	6.77E+05

EGLN_HUMAN	2.21E+03	2.21E+03	2.21E+03
PTGDS_HUMAN	1.52E+05	2.21E+03	2.21E+03
C1S_HUMAN	2.21E+03	2.21E+03	2.21E+03
FIBB_HUMAN	2.21E+03	2.21E+03	2.21E+03
VTDB_HUMAN	2.21E+03	2.21E+03	2.21E+03
IBP7_HUMAN	2.21E+03	2.21E+03	2.21E+03
APOA4_HUMAN	2.21E+03	2.21E+03	2.21E+03
PRDX2_HUMAN	2.21E+03	2.21E+03	2.21E+03
PCSK9_HUMAN	2.21E+03	2.21E+03	2.21E+03
PGK1_HUMAN	2.21E+03	2.21E+03	2.21E+03
CAH2_HUMAN	2.21E+03	2.21E+03	2.21E+03
BGH3_HUMAN	2.21E+03	2.21E+03	2.21E+03
CADH1_HUMAN	2.21E+03	2.21E+03	2.21E+03
BLVRB_HUMAN	2.21E+03	2.21E+03	2.21E+03
PEPD_HUMAN	2.21E+03	2.21E+03	2.21E+03