

Supplementary Table 3. 22DS group&gt;Control group

Accession	Description	MW [kDa]	Area					
			#01 Control1	#02 Control2	#03 Control3	#04 22DS1_1	#05 22DS2_1	#06 22DS3_1
Q16798	NADP-dependent malic enzyme, mitochondrial OS=Homo sapiens GN=ME3 PE=2 SV=2 - [MAON_HUMAN]	67.0	0.000E0	4.162E6	0.000E0	3.751E7	3.507E7	5.581E7
P19174	1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase gamma-1 OS=Homo sapiens GN=PLCG1 PE=1 SV=1 - [PLCG1_HUMAN]	148.4	1.091E7	3.897E7	2.546E7	1.581E9	1.781E9	2.336E9
O76076	WNT1-inducible-signaling pathway protein 2 OS=Homo sapiens GN=WISP2 PE=1 SV=1 - [WISP2_HUMAN]	26.8	0.000E0	0.000E0	0.000E0	6.133E7	8.950E7	8.469E7
Q6ZR08	Dynein heavy chain 12, axonemal OS=Homo sapiens GN=DNAH12 PE=2 SV=2 - [DYH12_HUMAN]	356.7	0.000E0	0.000E0	0.000E0	3.241E7	3.492E7	4.802E7
P02686	Myelin basic protein OS=Homo sapiens GN=MBP PE=1 SV=3 - [MBP_HUMAN]	33.1	0.000E0	0.000E0	0.000E0	1.129E7	1.806E7	1.458E7
Q9UJC5	SH3 domain-binding glutamic acid-rich-like protein 2 OS=Homo sapiens GN=SH3BGL2 PE=1 SV=2 - [SH3L2_HUMAN]	12.3	0.000E0	0.000E0	0.000E0	1.032E7	1.893E7	1.841E7
Q9BQI0	Allograft inflammatory factor 1-like OS=Homo sapiens GN=AIF1L PE=1 SV=1 - [AIF1L_HUMAN]	17.1	0.000E0	0.000E0	0.000E0	3.770E6	8.329E6	7.844E6
P08572	Collagen alpha-2(IV) chain OS=Homo sapiens GN=COL4A2 PE=1 SV=4 - [CO4A2_HUMAN]	167.4	0.000E0	0.000E0	0.000E0	1.273E8	2.897E8	2.660E8