

Spectrum	Name of Protein	Unique Peptide Counts	Fold Enrichment
IKZF1_GTX129438_Band A	Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2 - [GRP75_HUMAN]	17	Not in Control IP
IKZF1_GTX129438_Band A	Prelamin-A/C OS=Homo sapiens GN=LMNA PE=1 SV=1 - [LMNA_HUMAN]	8	Not in Control IP
IKZF1_GTX129438_Band A	Isoform Ik2 of DNA-binding protein Ikaros OS=Homo sapiens GN=IKZF1 - [IKZF1_HUMAN]	6	Not in Control IP
IKZF1_GTX129438_Band A	Isoform Ik7 of DNA-binding protein Ikaros OS=Homo sapiens GN=IKZF1 - [IKZF1_HUMAN]	5	Not in Control IP
IKZF1_GTX129438_Band A	Isoform Short of Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU - [HNRPU_HUMAN]	5	Not in Control IP
IKZF1_GTX129438_Band A	Isoform 2 of ATPase family AAA domain-containing protein 3A OS=Homo sapiens GN=ATAD3A - [ATD3A_HUMAN]	4	Not in Control IP
IKZF1_GTX129438_Band A	RNA-binding protein 14 OS=Homo sapiens GN=RBM14 PE=1 SV=2 - [RBM14_HUMAN]	4	Not in Control IP
IKZF1_GTX129438_Band A	X-ray repair cross-complementing protein 6 OS=Homo sapiens GN=XRCC6 PE=1 SV=1 - [B1AHC9_HUMAN]	3	Not in Control IP
IKZF1_GTX129438_Band A	Isoform A2 of Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 - [ROA2_HUMAN]	3	Not in Control IP
IKZF1_GTX129438_Band A	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=1 - [A0A087WWT3_HUMAN]	2	Not in Control IP
IKZF1_GTX129438_Band A	Eukaryotic initiation factor 4A-II OS=Homo sapiens GN=EIF4A2 PE=1 SV=1 - [E7EQG2_HUMAN]	2	Not in Control IP
IKZF1_GTX129438_Band A	Heterogeneous nuclear ribonucleoprotein D0 (Fragment) OS=Homo sapiens GN=HNRNPD PE=1 SV=5 - [D6RF44_HUMAN]	2	Not in Control IP
IKZF1_GTX129438_Band A	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=1 - [E7EUT5_HUMAN]	1	Not in Control IP
IKZF1_GTX129438_Band A	Isoform 2 of Desmoglein-1 OS=Homo sapiens GN=DSG1 - [DSG1_HUMAN]	1	Not in Control IP
IKZF1_GTX129438_Band A	Isoform 3 of TRAF3-interacting JNK-activating modulator OS=Homo sapiens GN=TRAF3IP3 - [T3JAM_HUMAN]	1	Not in Control IP
IKZF1_GTX129438_Band A	Transmembrane protein 232 OS=Homo sapiens GN=TMEM232 PE=2 SV=2 - [TM232_HUMAN]	1	Not in Control IP
IKZF1_GTX129438_Band A	Isoform 4 of Zinc finger C3H1 domain-containing protein OS=Homo sapiens GN=ZFC3H1 - [ZC3H1_HUMAN]	1	Not in Control IP
IKZF1_GTX129438_Band A	Spliceosome RNA helicase DDX39B (Fragment) OS=Homo sapiens GN=DDX39B PE=1 SV=1 - [H0Y400_HUMAN]	1	Not in Control IP
IKZF1_GTX129438_Band A	Heterogeneous nuclear ribonucleoprotein H OS=Homo sapiens GN=HNRNPH1 PE=1 SV=1 - [D6R9D3_HUMAN]	1	Not in Control IP
IKZF1_GTX129438_Band A	RNA-binding motif protein, X chromosome OS=Homo sapiens GN=RBMX PE=1 SV=1 - [H3BUY5_HUMAN]	1	Not in Control IP
IKZF1_GTX129438_Band A	TAR DNA-binding protein 43 (Fragment) OS=Homo sapiens GN=TARDBP PE=1 SV=1 - [A0A087WX29_HUMAN]	1	Not in Control IP
IKZF1_GTX129438_Band A	Cytochrome b-c1 complex subunit 2, mitochondrial (Fragment) OS=Homo sapiens GN=UQCRC2 PE=1 SV=1 - [H3BP04_HUMAN]	1	Not in Control IP
IKZF1_GTX129438_Band A	T-complex protein 1 subunit alpha OS=Homo sapiens GN=TCP1 PE=1 SV=1 - [E7EQR6_HUMAN]	1	Not in Control IP
IKZF1_GTX129438_Band A	Heat shock 70 kDa protein 1A OS=Homo sapiens GN=HSPA1A PE=1 SV=1 - [HS71A_HUMAN]	6	6.00
IKZF1_GTX129438_Band A	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1 - [HSP7C_HUMAN]	8	4.00
IKZF1_GTX129438_Band A	Ig gamma-3 chain C region (Fragment) OS=Homo sapiens GN=IGHG3 PE=1 SV=1 - [A0A075B6N8_HUMAN]	4	4.00
IKZF1_GTX129438_Band A	Vimentin OS=Homo sapiens GN=VIM PE=1 SV=1 - [B0YJC4_HUMAN]	2	2.00
IKZF1_GTX129438_Band A	ATP synthase subunit beta (Fragment) OS=Homo sapiens GN=ATP5B PE=1 SV=1 - [H0YH81_HUMAN]	2	2.00
IKZF1_GTX129438_Band A	Isoform DP11 of Desmoplakin OS=Homo sapiens GN=DSP - [DESP_HUMAN]	9	1.00
IKZF1_GTX129438_Band A	Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B PE=1 SV=1 - [TBA1B_HUMAN]	5	1.00
IKZF1_GTX129438_Band A	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1 - [ACTB_HUMAN]	3	1.00
IKZF1_GTX129438_Band A	Histone H2A type 1-H OS=Homo sapiens GN=HIST1H2AH PE=1 SV=3 - [H2A1H_HUMAN]	2	1.00
IKZF1_GTX129438_Band A	Isoform 2 of ATP synthase subunit alpha, mitochondrial OS=Homo sapiens GN=ATP5A1 - [ATPA_HUMAN]	9	0.67
IKZF1_GTX129438_Band A	Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=1 - [Q5JP53_HUMAN]	4	0.67
IKZF1_GTX129438_Band A	Junction plakoglobin OS=Homo sapiens GN=JUP PE=1 SV=3 - [PLAK_HUMAN]	7	0.33