

Spectrum	Name of Protein	Unique Peptide Counts	Fold Enrichment
IKZF2_A303-618A_Band A	Zinc finger protein Helios OS=Homo sapiens GN=IKZF2 PE=1 SV=2 - [IKZF2_HUMAN]	6	Not in Control IP
IKZF2_A303-618A_Band A	Desmoglein-1 OS=Homo sapiens GN=DSG1 PE=1 SV=2 - [DSG1_HUMAN]	6	Not in Control IP
IKZF2_A303-618A_Band A	Kinesin-like protein KIFC1 OS=Homo sapiens GN=KIFC1 PE=1 SV=2 - [KIFC1_HUMAN]	6	Not in Control IP
IKZF2_A303-618A_Band A	Junction plakoglobin OS=Homo sapiens GN=LUP PE=1 SV=3 - [PLAK_HUMAN]	6	Not in Control IP
IKZF2_A303-618A_Band A	Isoform 7 of Zinc finger protein Aiolos OS=Homo sapiens GN=IKZF3 - [IKZF3_HUMAN]	5	Not in Control IP
IKZF2_A303-618A_Band A	Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2 - [H4_HUMAN]	4	Not in Control IP
IKZF2_A303-618A_Band A	Isoform 3 of ATP-dependent zinc metalloprotease YME1L1 OS=Homo sapiens GN=YME1L1 - [YME1_HUMAN]	4	Not in Control IP
IKZF2_A303-618A_Band A	Isoform DPII of Desmoplakin OS=Homo sapiens GN=DSP - [DSP_HUMAN]	3	Not in Control IP
IKZF2_A303-618A_Band A	Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=2 - [TBB5_HUMAN]	2	Not in Control IP
IKZF2_A303-618A_Band A	Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1 - [EF1A1_HUMAN]	2	Not in Control IP
IKZF2_A303-618A_Band A	Isoform 2 of Tubulin alpha-1A chain OS=Homo sapiens GN=TUBA1A - [TBA1A_HUMAN]	2	Not in Control IP
IKZF2_A303-618A_Band A	Tubulin beta-4B chain OS=Homo sapiens GN=TUBB4B PE=1 SV=1 - [TBB4B_HUMAN]	2	Not in Control IP
IKZF2_A303-618A_Band A	Cystatin-A OS=Homo sapiens GN=CSTA PE=1 SV=1 - [CYTA_HUMAN]	2	Not in Control IP
IKZF2_A303-618A_Band A	Isoform Ik4 of DNA-binding protein Ikaros OS=Homo sapiens GN=IKZF1 - [IKZF1_HUMAN]	2	Not in Control IP
IKZF2_A303-618A_Band A	Enolase (Fragment) OS=Homo sapiens GN=ENO1 PE=1 SV=1 - [K7EM90_HUMAN]	2	Not in Control IP
IKZF2_A303-618A_Band A	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1 OS=Homo sapiens GN=RPNI PE=1 SV=1 - [F8WF32_HUMAN]	2	Not in Control IP
IKZF2_A303-618A_Band A	Isoform 3 of Arginase-1 OS=Homo sapiens GN=ARG1 - [ARG1_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Dermcidin OS=Homo sapiens GN=DCD PE=1 SV=2 - [DCD_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Filaggrin-2 OS=Homo sapiens GN=FLG2 PE=1 SV=1 - [FLA2_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Filaggrin OS=Homo sapiens GN=FLG PE=1 SV=3 - [FLA_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Forkhead box protein S1 OS=Homo sapiens GN=FOXS1 PE=2 SV=2 - [FOX51_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Histone H1.2 OS=Homo sapiens GN=HIST1H1C PE=1 SV=2 - [H12_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	La-related protein 4B OS=Homo sapiens GN=LARP4B PE=1 SV=3 - [LAR4B_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Isoform 3 of Protein-methionine sulfoxide oxidase MICAL1 OS=Homo sapiens GN=MICAL1 - [MICA1_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Isoform 3 of Prolyl 4-hydroxylase subunit alpha-1 OS=Homo sapiens GN=P4HA1 - [P4HA1_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Isoform 2 of Phosphoglycerate kinase 1 OS=Homo sapiens GN=PGK1 - [PGK1_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2 - [PROF1_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Isoform 4 of Poly(U)-binding-splicing factor PUF60 OS=Homo sapiens GN=PUF60 - [PUF60_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Isoform 2 of Serpin B3 OS=Homo sapiens GN=SERPINB3 - [SPB3_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Isoform 2 of Zinc finger X-chromosomal protein OS=Homo sapiens GN=ZFX - [ZFX_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Peroxiredoxin 2, isoform CRA_a OS=Homo sapiens GN=PRDX2 PE=1 SV=2 - [A6NIV5_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Unconventional myosin-IXa (Fragment) OS=Homo sapiens GN=MYO9A PE=1 SV=1 - [H3BU05_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Salivary acidic proline-rich phosphoprotein 1/2 (Fragment) OS=Homo sapiens GN=PRH1 PE=4 SV=1 - [A0A087WYF5_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=1 - [E7EUT5_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Pyruvate kinase OS=Homo sapiens GN=PKM PE=1 SV=1 - [H3BR70_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Fructose-bisphosphate aldolase (Fragment) OS=Homo sapiens GN=ALDOA PE=1 SV=1 - [H3BR04_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1 - [A0A0A0MRQ5_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Keratin, type I cuticular Ha3-II OS=Homo sapiens GN=KRT33B PE=1 SV=1 - [A0A087X216_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Histone H2A OS=Homo sapiens GN=H2AFV PE=3 SV=1 - [C91386_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	Proliferation-associated protein 2G4 OS=Homo sapiens GN=PA2G4 PE=1 SV=1 - [F8VR77_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band A	78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2 - [GRP78_HUMAN]	16	3.25
IKZF2_A303-618A_Band A	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1 - [ACTB_HUMAN]	19	3.00
IKZF2_A303-618A_Band A	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1 - [A0A087X010_HUMAN]	6	3.00
IKZF2_A303-618A_Band A	Isoform Short of Splicing factor, proline- and glutamine-rich OS=Homo sapiens GN=SFPO - [SFPO_HUMAN]	3	2.00
IKZF2_A303-618A_Band A	Albumin, isoform CRA_k OS=Homo sapiens GN=ALB PE=1 SV=1 - [C91KR2_HUMAN]	2	2.00
IKZF2_A303-618A_Band A	Polyubiquitin-C (Fragment) OS=Homo sapiens GN=UBC PE=1 SV=1 - [F5H265_HUMAN]	5	1.67
IKZF2_A303-618A_Band A	Isoform 5 of Prelamin-A/C OS=Homo sapiens GN=LMNA - [LMNA_HUMAN]	2	1.00
IKZF2_A303-618A_Band A	Histone H3 OS=Homo sapiens GN=H3F3B PE=1 SV=1 - [K7EMV3_HUMAN]	2	1.00
IKZF2_A303-618A_Band A	DNA replication licensing factor MCM7 (Fragment) OS=Homo sapiens GN=MCM7 PE=1 SV=1 - [C9J8M6_HUMAN]	2	1.00
IKZF2_A303-618A_Band A	Histone H2B type 1-K OS=Homo sapiens GN=HIST1H2BK PE=1 SV=3 - [H2B1K_HUMAN]	1	1.00
IKZF2_A303-618A_Band A	Isoform 2 of Interferon-induced GTP-binding protein Mx2 OS=Homo sapiens GN=Mx2 - [MX2_HUMAN]	1	1.00
IKZF2_A303-618A_Band A	Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2 - [GRP75_HUMAN]	17	0.92
IKZF2_A303-618A_Band A	Isoform 2 of Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 - [HSP7C_HUMAN]	11	0.88
IKZF2_A303-618A_Band A	Lamin-B1 OS=Homo sapiens GN=LMBN1 PE=1 SV=2 - [LMBN1_HUMAN]	9	0.80
IKZF2_A303-618A_Band A	Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5 - [H590A_HUMAN]	3	0.11
IKZF2_A303-618A_Band A	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4 - [H590B_HUMAN]	7	0.08
IKZF2_A303-618A_Band B	cAMP-dependent protein kinase type I-alpha regulatory subunit OS=Homo sapiens GN=PRKARIA PE=1 SV=1 - [KAP0_HUMAN]	15	Not in Control IP
IKZF2_A303-618A_Band B	Zinc finger protein Helios OS=Homo sapiens GN=IKZF2 PE=1 SV=2 - [IKZF2_HUMAN]	5	Not in Control IP
IKZF2_A303-618A_Band B	Isoform 7 of Zinc finger protein Aiolos OS=Homo sapiens GN=IKZF3 - [IKZF3_HUMAN]	4	Not in Control IP
IKZF2_A303-618A_Band B	Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2 - [H4_HUMAN]	2	Not in Control IP
IKZF2_A303-618A_Band B	Ig gamma-4 chain C region OS=Homo sapiens GN=IGHG4 PE=1 SV=1 - [IGHG4_HUMAN]	2	Not in Control IP
IKZF2_A303-618A_Band B	Histone-binding protein RBBP4 (Fragment) OS=Homo sapiens GN=RBBP4 PE=1 SV=1 - [HOYEU5_HUMAN]	2	Not in Control IP
IKZF2_A303-618A_Band B	Desmoglein-1 OS=Homo sapiens GN=DSG1 PE=1 SV=2 - [DSG1_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band B	Isoform 2 of Noelin-3 OS=Homo sapiens GN=OLFM3 - [NOE3_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band B	Isoform 2 of Tubulin alpha-4A chain OS=Homo sapiens GN=TUBA4A - [TBA4A_HUMAN]	1	Not in Control IP
IKZF2_A303-618A_Band B	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1 - [A0A087X010_HUMAN]	6	3.58
IKZF2_A303-618A_Band B	Isoform 2 of ATP synthase subunit alpha, mitochondrial OS=Homo sapiens GN=ATP5A1 - [ATPA_HUMAN]	7	3.00
IKZF2_A303-618A_Band B	Vimentin OS=Homo sapiens GN=VIM PE=1 SV=4 - [VIME_HUMAN]	7	3.00
IKZF2_A303-618A_Band B	Isoform 2 of Tubulin alpha-1A chain OS=Homo sapiens GN=TUBA1A - [TBA1A_HUMAN]	6	2.25
IKZF2_A303-618A_Band B	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1 - [ACTB_HUMAN]	19	1.18
IKZF2_A303-618A_Band B	Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=2 - [TBB5_HUMAN]	10	1.00
IKZF2_A303-618A_Band B	Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1 - [EF1A1_HUMAN]	6	0.67