

Spectrum	Name of Protein	Unique Peptide Counts	Fold Enrichment
ZC3H8_A303-090A	Interleukin enhancer-binding factor 2 OS=Homo sapiens GN=ILF2 PE=1 SV=1 - [B4DY09_HUMAN]	14	Not in Control IP
ZC3H8_A303-090A	RNA-binding motif protein, X chromosome OS=Homo sapiens GN=RBMX PE=1 SV=3 - [RBMX_HUMAN]	13	Not in Control IP
ZC3H8_A303-090A	Isoform 3 of Core histone macro-H2A.1 OS=Homo sapiens GN=H2AFY - [H2AY_HUMAN]	12	Not in Control IP
ZC3H8_A303-090A	Serpine B4 OS=Homo sapiens GN=SERPINB4 PE=1 SV=2 - [SPB4_HUMAN]	9	Not in Control IP
ZC3H8_A303-090A	Histone H1.4 OS=Homo sapiens GN=HIST1H1E PE=1 SV=2 - [H14_HUMAN]	7	Not in Control IP
ZC3H8_A303-090A	Isoform 1 of RNA-binding protein Raly OS=Homo sapiens GN=RALY - [RALY_HUMAN]	7	Not in Control IP
ZC3H8_A303-090A	rRNA 2'-O-methyltransferase fibrillarin (Fragment) OS=Homo sapiens GN=FBL PE=1 SV=1 - [MOROP1_HUMAN]	6	Not in Control IP
ZC3H8_A303-090A	ELAV-like protein 1 OS=Homo sapiens GN=ELAVL1 PE=1 SV=2 - [ELAV1_HUMAN]	5	Not in Control IP
ZC3H8_A303-090A	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1 - [A0A087X010_HUMAN]	4	Not in Control IP
ZC3H8_A303-090A	14-3-3 protein sigma OS=Homo sapiens GN=SFN PE=1 SV=1 - [14335_HUMAN]	4	Not in Control IP
ZC3H8_A303-090A	Isoform 3 of Transformer-2 protein homolog beta OS=Homo sapiens GN=TRA2B - [TRA2B_HUMAN]	4	Not in Control IP
ZC3H8_A303-090A	Zinc finger CCH domain-containing protein 8 OS=Homo sapiens GN=ZC3H8 PE=1 SV=2 - [ZC3H8_HUMAN]	3	Not in Control IP
ZC3H8_A303-090A	Histone H2B type 1-K OS=Homo sapiens GN=HIST1H2BK PE=1 SV=3 - [H2BK_HUMAN]	3	Not in Control IP
ZC3H8_A303-090A	Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2 - [H4_HUMAN]	3	Not in Control IP
ZC3H8_A303-090A	Heterogeneous nuclear ribonucleoprotein D0 (Fragment) OS=Homo sapiens GN=HNRNPD PE=1 SV=5 - [D6RF44_HUMAN]	3	Not in Control IP
ZC3H8_A303-090A	Ribosome biogenesis protein BRX1 homolog OS=Homo sapiens GN=BRX1 PE=1 SV=2 - [BRX1_HUMAN]	3	Not in Control IP
ZC3H8_A303-090A	Isoform 3 of Centromere protein V OS=Homo sapiens GN=CENPV - [CENPV_HUMAN]	3	Not in Control IP
ZC3H8_A303-090A	Histone H1.5 OS=Homo sapiens GN=HIST1H1B PE=1 SV=3 - [H15_HUMAN]	3	Not in Control IP
ZC3H8_A303-090A	60S ribosomal protein L7a OS=Homo sapiens GN=RPL7A PE=1 SV=2 - [RL7A_HUMAN]	3	Not in Control IP
ZC3H8_A303-090A	Isoform 2 of Serpin B3 OS=Homo sapiens GN=SERPINB3 - [SPB3_HUMAN]	3	Not in Control IP
ZC3H8_A303-090A	Isoform 3 of Serine/arginine-rich splicing factor 7 OS=Homo sapiens GN=SRSF7 - [SRSF7_HUMAN]	3	Not in Control IP
ZC3H8_A303-090A	60S ribosomal protein L7 OS=Homo sapiens GN=RPL7 PE=1 SV=1 - [A8MUD9_HUMAN]	3	Not in Control IP
ZC3H8_A303-090A	Annexin (Fragment) OS=Homo sapiens GN=ANXA2 PE=1 SV=1 - [H0YK54_HUMAN]	3	Not in Control IP
ZC3H8_A303-090A	Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1 - [A0A087WV09_HUMAN]	3	Not in Control IP
ZC3H8_A303-090A	Histone H2A type 1-H OS=Homo sapiens GN=HIST1H2AH PE=1 SV=3 - [H2A1H_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	Isoform 2 of Tubulin alpha-1A chain OS=Homo sapiens GN=TUBA1A - [TBA1A_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	Histone H3 OS=Homo sapiens GN=H3F3B PE=1 SV=1 - [K7EMV3_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	Isoform 3 of Chromatin target of PRMT1 protein OS=Homo sapiens GN=CHTOP - [CHTOP_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	Probable rRNA-processing protein EBP2 OS=Homo sapiens GN=EBNA1BP2 PE=1 SV=2 - [EBP2_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	Emerin OS=Homo sapiens GN=EMD PE=1 SV=1 - [EMD_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	MKI67 FHA domain-interacting nucleolar phosphoprotein OS=Homo sapiens GN=NIFK PE=1 SV=1 - [MK671_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	Isoform 2 of Neuroguinid OS=Homo sapiens GN=NGDN - [NGDN_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	Nucleolar protein 7 OS=Homo sapiens GN=NOL7 PE=1 SV=2 - [NOL7_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	60S ribosomal protein L5 OS=Homo sapiens GN=RPL5 PE=1 SV=3 - [RL5_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	U2 small nuclear ribonucleoprotein A' OS=Homo sapiens GN=SNRPA1 PE=1 SV=2 - [RU2A_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	THO complex subunit 4 OS=Homo sapiens GN=ALYREF PE=1 SV=3 - [THOC4_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	40S ribosomal protein S3a OS=Homo sapiens GN=RPS3A PE=1 SV=1 - [D6R9B6_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	SWI/SNF-related matrix-associated actin-dependent regulator of chromatin subfamily 8 member 1 (Fragment) OS=Homo sapiens GN=SMARCB1 PE=4 SV=1 - [A0A0G2J5E9_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	Telomere-associated protein RIF1 (Fragment) OS=Homo sapiens GN=RIF1 PE=1 SV=1 - [H7C2B5_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	40S ribosomal protein S3 OS=Homo sapiens GN=RPS3 PE=1 SV=1 - [E9PQ96_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	U3 small nucleolar ribonucleoprotein protein IMP4 OS=Homo sapiens GN=IMP4 PE=1 SV=1 - [B82477_HUMAN]	2	Not in Control IP
ZC3H8_A303-090A	Heat shock protein HSP90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4 - [H590B_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	WD repeat-containing protein 74 (Fragment) OS=Homo sapiens GN=WDR74 PE=1 SV=1 - [H0YFP2_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Heterogeneous nuclear ribonucleoprotein H (Fragment) OS=Homo sapiens GN=HNRNPH1 PE=1 SV=1 - [H0YB39_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Isoform 2 of DNA dC->du-editing enzyme APOBEC-3A OS=Homo sapiens GN=APOBEC3A - [ABC3A_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Beta-1,3-galactosyltransferase 2 OS=Homo sapiens GN=B3GALT2 PE=2 SV=1 - [B3GT2_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Desmoglein-1 OS=Homo sapiens GN=DSG1 PE=1 SV=2 - [DSG1_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Homo sapiens GN=HNRNCP PE=1 SV=4 - [HNRPC_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Isoform 3 of Putative RNA-binding protein Luc7-like 2 OS=Homo sapiens GN=LUC7L2 - [LC7L2_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Protein MAK16 homolog OS=Homo sapiens GN=MAK16 PE=1 SV=2 - [MAK16_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Olfactory receptor 2G3 OS=Homo sapiens GN=OR2G3 PE=2 SV=1 - [OR2G3_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Poly(rC)-binding protein 1 OS=Homo sapiens GN=PCBP1 PE=1 SV=2 - [PCBP1_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Heterogeneous nuclear ribonucleoprotein A0 OS=Homo sapiens GN=HNRNPA0 PE=1 SV=1 - [ROA0_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	40S ribosomal protein S4, X isoform OS=Homo sapiens GN=RPS4X PE=1 SV=2 - [R54X_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Protein S100-A8 OS=Homo sapiens GN=S100A8 PE=1 SV=1 - [S10A8_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Isoform 3 of SWI/SNF-related matrix-associated actin-dependent regulator of chromatin subfamily A containing DEAD/H box 1 OS=Homo sapiens GN=SMARCA1 - [SMRCD_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Isoform 2 of A-kinase anchor protein SPHKAP OS=Homo sapiens GN=SPHKAP - [SPKAP_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	60S ribosomal protein L8 OS=Homo sapiens GN=RPL8 PE=1 SV=1 - [G3V1A1_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Histone H3 OS=Homo sapiens GN=HIST2H3P52 PE=1 SV=1 - [Q5TEC6_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	TATA-binding protein-associated factor 2N OS=Homo sapiens GN=TAF15 PE=1 SV=1 - [A0A075B7D9_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=1 - [E7EUT5_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Fructose-bisphosphate aldolase (Fragment) OS=Homo sapiens GN=ALDOA PE=1 SV=1 - [H3BR04_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Polyubiquitin-C (Fragment) OS=Homo sapiens GN=UBC PE=1 SV=1 - [F5H265_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Nuclease-sensitive element-binding protein 1 (Fragment) OS=Homo sapiens GN=YBX1 PE=1 SV=1 - [C9J5V9_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Peptidyl-prolyl cis-trans isomerase E (Fragment) OS=Homo sapiens GN=PIIE PE=1 SV=5 - [E9PIB0_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	L-lactate dehydrogenase A chain (Fragment) OS=Homo sapiens GN=LDHA PE=1 SV=1 - [F5GYU2_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Annexin (Fragment) OS=Homo sapiens GN=ANXA1 PE=1 SV=1 - [Q5T3N1_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Nucleolar protein 56 (Fragment) OS=Homo sapiens GN=NOP56 PE=1 SV=1 - [H0Y653_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Nucleolar complex protein 4 homolog (Fragment) OS=Homo sapiens GN=NOC41 PE=1 SV=1 - [F5H303_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Protein IGHV1-58 (Fragment) OS=Homo sapiens GN=IGHV1-58 PE=4 SV=1 - [A0A0C4D39_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Non-POU domain-containing octamer-binding protein (Fragment) OS=Homo sapiens GN=NONO PE=1 SV=1 - [C9IZL7_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	ATPase ASNA1 (Fragment) OS=Homo sapiens GN=ASNA1 PE=1 SV=1 - [K7ERW9_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	14-3-3 protein zeta/delta (Fragment) OS=Homo sapiens GN=YWHAZ PE=1 SV=1 - [E7ESK7_HUMAN]	1	Not in Control IP
ZC3H8_A303-090A	Heterogeneous nuclear ribonucleoprotein A/B OS=Homo sapiens GN=HNRNPAB PE=1 SV=1 - [D6R9P3_HUMAN]	7	6.00
ZC3H8_A303-090A	Isoform 2 of Heterogeneous nuclear ribonucleoprotein A3 OS=Homo sapiens GN=HNRNPA3 - [ROA3_HUMAN]	7	6.00
ZC3H8_A303-090A	Isoform 3 of Heterogeneous nuclear ribonucleoprotein D-like OS=Homo sapiens GN=HNRNPD3 - [HNRDL_HUMAN]	4	6.00
ZC3H8_A303-090A	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1 - [ACTB_HUMAN]	10	5.00
ZC3H8_A303-090A	Isoform 3 of Heterogeneous nuclear ribonucleoprotein H3 OS=Homo sapiens GN=HNRNPH3 - [HNRH3_HUMAN]	10	3.33
ZC3H8_A303-090A	Isoform 2 of Heterogeneous nuclear ribonucleoprotein A1 OS=Homo sapiens GN=HNRNPA1 - [ROA1_HUMAN]	5	2.50
ZC3H8_A303-090A	Heterogeneous nuclear ribonucleoproteins C1/C2 (Fragment) OS=Homo sapiens GN=HNRNCP PE=1 SV=1 - [G3V4W0_HUMAN]	3	2.22
ZC3H8_A303-090A	Isoform 2 of Nucleophosmin OS=Homo sapiens GN=NPM1 - [NPM_HUMAN]	6	2.00
ZC3H8_A303-090A	Isoform A2 of Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 - [ROA2_HUMAN]	13	1.43
ZC3H8_A303-090A	Heterogeneous nuclear ribonucleoprotein R OS=Homo sapiens GN=HNRNPR PE=1 SV=1 - [B4D128_HUMAN]	1	1.00