

Spectrum	Name of Protein	Unique Peptide Counts	Fold Enrichment
LOCR_L_HPA028794_Band A	Heterogeneous nuclear ribonucleoprotein U-like protein 1 (Fragment) OS=Homo sapiens GN=HNRNPUL1 PE=1 SV=1 - [MOR3F1_HUMAN]	9	Not in Control IP
LOCR_L_HPA028794_Band A	60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2 - [CH60_HUMAN]	3	Not in Control IP
LOCR_L_HPA028794_Band A	Isoform 2 of ATP-dependent RNA helicase DDX1 OS=Homo sapiens GN=DDX1 - [DDX1_HUMAN]	2	Not in Control IP
LOCR_L_HPA028794_Band A	Isoform 2 of Dihydropyrimidinase-related protein 2 OS=Homo sapiens GN=DPYSL2 - [DPYL2_HUMAN]	2	Not in Control IP
LOCR_L_HPA028794_Band A	Ran GTPase-activating protein 1 OS=Homo sapiens GN=RANGAP1 PE=1 SV=1 - [RAGP1_HUMAN]	2	Not in Control IP
LOCR_L_HPA028794_Band A	TATA-binding protein-associated factor 2N OS=Homo sapiens GN=TAF15 PE=1 SV=1 - [A0A075B7D9_HUMAN]	2	Not in Control IP
LOCR_L_HPA028794_Band A	Ligand-dependent nuclear receptor corepressor-like protein OS=Homo sapiens GN=LCORL PE=1 SV=1 - [D6RA10_HUMAN]	1	Not in Control IP
LOCR_L_HPA028794_Band A	Pre-mRNA-splicing factor ATP-dependent RNA helicase DHX15 OS=Homo sapiens GN=DHX15 PE=1 SV=2 - [DHX15_HUMAN]	1	Not in Control IP
LOCR_L_HPA028794_Band A	Ligand-dependent corepressor OS=Homo sapiens GN=LCOR PE=1 SV=2 - [LCOR_HUMAN]	1	Not in Control IP
LOCR_L_HPA028794_Band A	Isoform 7 of Sarcolemmal membrane-associated protein OS=Homo sapiens GN=SLMAP - [SLMAP_HUMAN]	1	Not in Control IP
LOCR_L_HPA028794_Band A	Protein-glutamine gamma-glutamyltransferase 2 OS=Homo sapiens GN=TGM2 PE=1 SV=2 - [TGM2_HUMAN]	1	Not in Control IP
LOCR_L_HPA028794_Band A	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 2 (Fragment) OS=Homo sapiens GN=RPN2 PE=1 SV=5 - [QSJYR7_HUMAN]	1	Not in Control IP
LOCR_L_HPA028794_Band A	Phosphoenolpyruvate carboxykinase (GTP), mitochondrial OS=Homo sapiens GN=PCK2 PE=1 SV=1 - [HOYML5_HUMAN]	1	Not in Control IP
LOCR_L_HPA028794_Band A	Cleavage and polyadenylation-specificity factor subunit 2 (Fragment) OS=Homo sapiens GN=CPF2 PE=1 SV=1 - [HOYJF4_HUMAN]	1	Not in Control IP
LOCR_L_HPA028794_Band A	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1 OS=Homo sapiens GN=RPN1 PE=1 SV=1 - [F8WF32_HUMAN]	1	Not in Control IP
LOCR_L_HPA028794_Band A	Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM PE=1 SV=1 - [A0A087X0X3_HUMAN]	21	2.38
LOCR_L_HPA028794_Band A	Isoform 6 of RNA-binding protein EWS OS=Homo sapiens GN=EWSR1 - [EWS_HUMAN]	2	2.00
LOCR_L_HPA028794_Band A	Isoform Short of RNA-binding protein FUS OS=Homo sapiens GN=FUS - [FUS_HUMAN]	6	1.67
LOCR_L_HPA028794_Band A	Isoform 4 of Heterogeneous nuclear ribonucleoprotein R OS=Homo sapiens GN=HNRNPR - [HNRPR_HUMAN]	9	1.33
LOCR_L_HPA028794_Band A	Heterogeneous nuclear ribonucleoprotein Q OS=Homo sapiens GN=SYNCRIP PE=1 SV=1 - [B7Z645_HUMAN]	4	1.33
LOCR_L_HPA028794_Band A	Interleukin enhancer-binding factor 3 OS=Homo sapiens GN=ILF3 PE=1 SV=3 - [ILF3_HUMAN]	4	1.00
LOCR_L_HPA028794_Band A	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4 - [HS90B_HUMAN]	2	1.00
LOCR_L_HPA028794_Band A	Isoform 2 of Paternally-expressed gene 3 protein OS=Homo sapiens GN=PEG3 - [PEG3_HUMAN]	2	1.00
LOCR_L_HPA028794_Band A	Albumin, isoform CRA_k OS=Homo sapiens GN=ALB PE=1 SV=1 - [C9JKR2_HUMAN]	2	1.00
LOCR_L_HPA028794_Band A	Ig gamma-1 chain C region (Fragment) OS=Homo sapiens GN=IGHG1 PE=1 SV=1 - [A0A0A0MS07_HUMAN]	2	1.00
LOCR_L_HPA028794_Band A	Junction plakoglobin OS=Homo sapiens GN=JUP PE=1 SV=3 - [PLAK_HUMAN]	1	1.00
LOCR_L_HPA028794_Band A	Nuclear pore complex protein Nup93 (Fragment) OS=Homo sapiens GN=NUP93 PE=1 SV=1 - [H3BMX0_HUMAN]	1	1.00
LOCR_L_HPA028794_Band A	HCG2044799 OS=Homo sapiens GN=HNRNPUL2-BSCL2 PE=4 SV=1 - [H3BQZ7_HUMAN]	3	0.50
LOCR_L_HPA028794_Band A	Isoform Short of Splicing factor, proline- and glutamine-rich OS=Homo sapiens GN=SFPO - [SFPO_HUMAN]	10	0.40
LOCR_L_HPA028794_Band A	Matrin-3 OS=Homo sapiens GN=MATR3 PE=1 SV=1 - [D6REM6_HUMAN]	6	0.40
LOCR_L_HPA028794_Band A	Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2 - [GRP75_HUMAN]	19	0.38
LOCR_L_HPA028794_Band A	Isoform Short of Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU - [HNRPU_HUMAN]	24	0.21
LOCR_L_HPA028794_Band A	Isoform 2 of Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 - [HSP7C_HUMAN]	5	0.20
LOCR_L_HPA028794_Band A	Isoform 2 of Probable ATP-dependent RNA helicase DDX5 OS=Homo sapiens GN=DDX5 - [DDX5_HUMAN]	3	#REF!
LOCR_L_HPA028794_Band A	Probable ATP-dependent RNA helicase DDX17 OS=Homo sapiens GN=DDX17 PE=1 SV=2 - [DDX17_HUMAN]	5	#REF!
LOCR_L_HPA028794_Band A	78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2 - [GRP78_HUMAN]	19	#REF!