

Spectrum	Name of Protein	Unique Peptide Counts	Fold Enrichment
L3MBTL2_HPA000815_Band A	Isoform 2 of Lethal(3)malignant brain tumor-like protein 2 OS=Homo sapiens GN=L3MBTL2 - [LMBL2_HUMAN]	6	Not in Control IP
L3MBTL2_HPA000815_Band A	Matrin-3 OS=Homo sapiens GN=MATR3 PE=1 SV=1 - [D6REM6_HUMAN]	5	Not in Control IP
L3MBTL2_HPA000815_Band A	X-ray repair cross-complementing protein 5 OS=Homo sapiens GN=XRCC5 PE=1 SV=3 - [XRCC5_HUMAN]	4	Not in Control IP
L3MBTL2_HPA000815_Band A	Protein AATF OS=Homo sapiens GN=AATF PE=1 SV=1 - [AATF_HUMAN]	2	Not in Control IP
L3MBTL2_HPA000815_Band A	Isoform 2 of Nucleolar RNA helicase 2 OS=Homo sapiens GN=DDX21 - [DDX21_HUMAN]	2	Not in Control IP
L3MBTL2_HPA000815_Band A	FACT complex subunit SSRP1 OS=Homo sapiens GN=SSRP1 PE=1 SV=1 - [SSRP1_HUMAN]	2	Not in Control IP
L3MBTL2_HPA000815_Band A	Nucleolin (Fragment) OS=Homo sapiens GN=NCL PE=1 SV=5 - [H7BY16_HUMAN]	2	Not in Control IP
L3MBTL2_HPA000815_Band A	MAX gene-associated protein (Fragment) OS=Homo sapiens GN=MGA PE=1 SV=1 - [H3BP52_HUMAN]	2	Not in Control IP
L3MBTL2_HPA000815_Band A	Isoform 3 of Sarcoplasmic/endoplasmic reticulum calcium ATPase 1 OS=Homo sapiens GN=ATP2A1 - [AT2A1_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	Isoform 2 of Ribosome biogenesis protein BOP1 OS=Homo sapiens GN=BOP1 - [BOP1_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	Isoform 5 of Interleukin enhancer-binding factor 3 OS=Homo sapiens GN=ILF3 - [ILF3_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	U3 small nucleolar ribonucleoprotein protein MPP10 OS=Homo sapiens GN=MPPH10 PE=1 SV=2 - [MPP10_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	Nucleolar protein 58 OS=Homo sapiens GN=NOP58 PE=1 SV=1 - [NOP58_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	Isoform Short of Splicing factor, proline- and glutamine-rich OS=Homo sapiens GN=SFPO - [SFPO_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	Transducin beta-like protein 3 OS=Homo sapiens GN=TBL3 PE=1 SV=2 - [TBL3_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	Polypyrimidine tract binding protein 1, isoform CRA_e OS=Homo sapiens GN=PTBP1 PE=1 SV=1 - [A0A0D9SF20_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	HCG2044799 OS=Homo sapiens GN=HNRNPUL2-BSCL2 PE=4 SV=1 - [H3BQZ7_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	Heterochromatin protein 1-binding protein 3 (Fragment) OS=Homo sapiens GN=HP1BP3 PE=1 SV=1 - [B0QZK4_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	Nucleolar and coiled-body phosphoprotein 1 OS=Homo sapiens GN=NOLC1 PE=1 SV=1 - [S4R341_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	DNA replication licensing factor MCM4 (Fragment) OS=Homo sapiens GN=MCM4 PE=1 SV=5 - [E5RFJ8_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	116 kDa U5 small nuclear ribonucleoprotein component (Fragment) OS=Homo sapiens GN=EFTUD2 PE=1 SV=1 - [K7EP67_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	Nuclear pore complex protein Nup93 (Fragment) OS=Homo sapiens GN=NUP93 PE=1 SV=1 - [H3BMX0_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	RNA-binding protein 28 (Fragment) OS=Homo sapiens GN=RBM28 PE=1 SV=1 - [H7C5G8_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	Heterogeneous nuclear ribonucleoprotein Q (Fragment) OS=Homo sapiens GN=SYNCRIP PE=1 SV=1 - [F6UXX1_HUMAN]	1	Not in Control IP
L3MBTL2_HPA000815_Band A	Isoform Short of Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU - [HNRPU_HUMAN]	7	7.00
L3MBTL2_HPA000815_Band A	Isoform 2 of X-ray repair cross-complementing protein 6 OS=Homo sapiens GN=XRCC6 - [XRCC6_HUMAN]	3	3.00
L3MBTL2_HPA000815_Band A	Isoform 2 of Probable ATP-dependent RNA helicase DDX5 OS=Homo sapiens GN=DDX5 - [DDX5_HUMAN]	2	2.00
L3MBTL2_HPA000815_Band A	Lamin-B1 OS=Homo sapiens GN=LMNB1 PE=1 SV=2 - [LMNB1_HUMAN]	4	1.50
L3MBTL2_HPA000815_Band A	Isoform 4 of Heterogeneous nuclear ribonucleoprotein R OS=Homo sapiens GN=HNRNPR - [HNRPR_HUMAN]	7	1.33
L3MBTL2_HPA000815_Band A	Isoform 2 of Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM - [HNRPM_HUMAN]	2	1.00
L3MBTL2_HPA000815_Band A	Complement C4-B OS=Homo sapiens GN=C4B_2 PE=4 SV=1 - [A0A0G2JL54_HUMAN]	2	1.00
L3MBTL2_HPA000815_Band A	Heterogeneous nuclear ribonucleoprotein U-like protein 1 (Fragment) OS=Homo sapiens GN=HNRNPUL1 PE=1 SV=1 - [M0R3F1_HUMAN]	2	1.00
L3MBTL2_HPA000815_Band A	Isoform 2 of Probable ATP-dependent RNA helicase DDX17 OS=Homo sapiens GN=DDX17 - [DDX17_HUMAN]	1	1.00
L3MBTL2_HPA000815_Band A	RNA-binding protein 14 OS=Homo sapiens GN=RBM14 PE=1 SV=2 - [RBM14_HUMAN]	4	0.33
L3MBTL2_HPA000815_Band A	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1 - [HSP7C_HUMAN]	9	0.30
L3MBTL2_HPA000815_Band A	DNA replication licensing factor MCM7 OS=Homo sapiens GN=MCM7 PE=1 SV=4 - [MCM7_HUMAN]	13	0.18
L3MBTL2_HPA000815_Band A	78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2 - [GRP78_HUMAN]	8	0.14
L3MBTL2_HPA000815_Band A	Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2 - [GRP75_HUMAN]	17	0.06