

Spectrum	Name of Protein	Unique Peptide Counts	Fold Enrichment
NBN_HPA001429-Band A	Nibrin OS=Homo sapiens GN=NBN PE=1 SV=1 - [NBN_HUMAN]	38	Not in Control IP
NBN_HPA001429-Band A	DNA repair protein RAD50 OS=Homo sapiens GN=RAD50 PE=1 SV=1 - [RAD50_HUMAN]	36	Not in Control IP
NBN_HPA001429-Band A	60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2 - [CH60_HUMAN]	8	Not in Control IP
NBN_HPA001429-Band A	Phosphoenolpyruvate carboxykinase [GTP], mitochondrial OS=Homo sapiens GN=PKC2 PE=1 SV=3 - [PKCKM_HUMAN]	7	Not in Control IP
NBN_HPA001429-Band A	Nuclear pore complex protein Nup88 (Fragment) OS=Homo sapiens GN=NUP88 PE=1 SV=1 - [J3KMX1_HUMAN]	3	Not in Control IP
NBN_HPA001429-Band A	Isoform 3 of Sodium/potassium-transporting ATPase subunit alpha-1 OS=Homo sapiens GN=ATP1A1 - [AT1A1_HUMAN]	2	Not in Control IP
NBN_HPA001429-Band A	Hornerin OS=Homo sapiens GN=HRNR PE=1 SV=2 - [HORN_HUMAN]	2	Not in Control IP
NBN_HPA001429-Band A	Lamin-B2 OS=Homo sapiens GN=LMB2 PE=1 SV=3 - [LMB2_HUMAN]	2	Not in Control IP
NBN_HPA001429-Band A	Double-strand break repair protein MRE11A OS=Homo sapiens GN=MRE11A PE=1 SV=3 - [MRE11_HUMAN]	2	Not in Control IP
NBN_HPA001429-Band A	Isoform 4 of Nuclear pore complex protein Nup98-Nup96 OS=Homo sapiens GN=NUP98 - [NUP98_HUMAN]	2	Not in Control IP
NBN_HPA001429-Band A	Isoform 2 of Pre-mRNA-processing factor 6 OS=Homo sapiens GN=PRPF6 - [PRPF6_HUMAN]	2	Not in Control IP
NBN_HPA001429-Band A	Double-strand break repair protein MRE11A OS=Homo sapiens GN=MRE11A PE=1 SV=1 - [F8W7U8_HUMAN]	2	Not in Control IP
NBN_HPA001429-Band A	Isoform 2 of ATP synthase subunit alpha, mitochondrial OS=Homo sapiens GN=ATP5A1 - [ATPA_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1 - [ACTB_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Alpha-aminoacidic semialdehyde synthase, mitochondrial OS=Homo sapiens GN=AASS PE=1 SV=1 - [AASS_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Long-chain-fatty-acid-CoA ligase 5 OS=Homo sapiens GN=ACSL5 PE=1 SV=1 - [ACSL5_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	AFG3-like protein 2 OS=Homo sapiens GN=AFG3L2 PE=1 SV=2 - [AFG3L2_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Isoform 2 of Delta-1-pyrroline-5-carboxylate dehydrogenase, mitochondrial OS=Homo sapiens GN=ALDH4A1 - [AL4A1_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Cytoskeleton-associated protein 4 OS=Homo sapiens GN=CKAP4 PE=1 SV=2 - [CKAP4_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Endoplasmic reticulum protein OS=Homo sapiens GN=HSP90B1 PE=1 SV=1 - [ENPL_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Isoform 2 of Heat shock 70 kDa protein 1A OS=Homo sapiens GN=HSPA1A - [HS71A_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Heat shock-related 70 kDa protein 2 OS=Homo sapiens GN=HSPA2 PE=1 SV=1 - [HSP72_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Prolyl 4-hydroxylase subunit alpha-1 OS=Homo sapiens GN=P4HA1 PE=1 SV=2 - [P4HA1_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Splicing regulatory glutamine/lysine-rich protein 1 OS=Homo sapiens GN=SREK1 PE=1 SV=1 - [SREK1_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Isoleucine--tRNA ligase, mitochondrial OS=Homo sapiens GN=IARS2 PE=1 SV=2 - [SYIM_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Transmembrane and coiled-coil domain family 3, isoform CRA_c OS=Homo sapiens GN=TMCC3 PE=1 SV=1 - [G3V207_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Heterogeneous nuclear ribonucleoprotein L OS=Homo sapiens GN=HNRNPL PE=1 SV=1 - [B4DV8_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Protein arginine N-methyltransferase 5 (Fragment) OS=Homo sapiens GN=PRMT5 PE=1 SV=1 - [HOY177_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Poly(U)-binding-splicing factor PUF60 (Fragment) OS=Homo sapiens GN=PUF60 PE=1 SV=1 - [E9PMU7_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Polypeptide N-acetylgalactosaminyltransferase 13 (Fragment) OS=Homo sapiens GN=GALNT13 PE=4 SV=1 - [H7BZG2_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Terminal uridylyltransferase 7 OS=Homo sapiens GN=ZCCHC6 PE=1 SV=1 - [A0A0C4DFW3_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Protein TRAJ56 (Fragment) OS=Homo sapiens GN=TRAJ56 PE=4 SV=1 - [A0A0758622_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Granulins (Fragment) OS=Homo sapiens GN=GRN PE=1 SV=1 - [K7EKL3_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Long-chain-fatty-acid-CoA ligase 3 (Fragment) OS=Homo sapiens GN=ACSL3 PE=1 SV=1 - [F5GW2_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band A	Albumin, isoform CRA_k OS=Homo sapiens GN=ALB PE=1 SV=1 - [C9JKR2_HUMAN]	5	5.00
NBN_HPA001429-Band A	Junction plakoglobin OS=Homo sapiens GN=JUP PE=1 SV=3 - [PLAK_HUMAN]	5	4.00
NBN_HPA001429-Band A	Nuclear pore complex protein Nup93 OS=Homo sapiens GN=NUP93 PE=1 SV=2 - [NUP93_HUMAN]	4	3.00
NBN_HPA001429-Band A	Isoform 2 of Complement C4-A OS=Homo sapiens GN=C4A - [C04A_HUMAN]	3	3.00
NBN_HPA001429-Band A	Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5 - [HS90A_HUMAN]	1	3.00
NBN_HPA001429-Band A	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4 - [HS90B_HUMAN]	1	3.00
NBN_HPA001429-Band A	Isoform 6 of RNA-binding protein EWS OS=Homo sapiens GN=EWSR1 - [EWS_HUMAN]	2	2.00
NBN_HPA001429-Band A	Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2 - [GRP75_HUMAN]	31	1.81
NBN_HPA001429-Band A	Isoform 2 of Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 - [HSP7C_HUMAN]	6	1.80
NBN_HPA001429-Band A	Calcium-binding mitochondrial carrier protein Aralar2 OS=Homo sapiens GN=SLC25A13 PE=1 SV=2 - [CMC2_HUMAN]	9	1.50
NBN_HPA001429-Band A	Ig gamma-1 chain C region (Fragment) OS=Homo sapiens GN=IGHG1 PE=1 SV=1 - [A0A0A0MS07_HUMAN]	3	1.50
NBN_HPA001429-Band A	Isoform 2 of ATPase family AAA domain-containing protein 3A OS=Homo sapiens GN=ATAD3A - [ATD3A_HUMAN]	6	1.00
NBN_HPA001429-Band A	78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2 - [GRP78_HUMAN]	26	0.95
NBN_HPA001429-Band A	Isoform C of Prelamin A/C OS=Homo sapiens GN=LMA - [LMA_HUMAN]	4	0.67
NBN_HPA001429-Band A	Trifunctional enzyme subunit alpha, mitochondrial OS=Homo sapiens GN=HADHA PE=1 SV=2 - [ECHA_HUMAN]	3	0.50
NBN_HPA001429-Band A	Lamin-B1 OS=Homo sapiens GN=LMB1 PE=1 SV=2 - [LMB1_HUMAN]	3	0.50
NBN_HPA001429-Band A	Nucleolin (Fragment) OS=Homo sapiens GN=NCL PE=1 SV=5 - [H7BY16_HUMAN]	2	0.50
NBN_HPA001429-Band A	Polyubiquitin-B (Fragment) OS=Homo sapiens GN=UBB PE=1 SV=1 - [J3QSA3_HUMAN]	2	0.50
NBN_HPA001429-Band A	Isoform Short of Delta-1-pyrroline-5-carboxylate synthase OS=Homo sapiens GN=ALDH18A1 - [P5CS_HUMAN]	4	0.33
NBN_HPA001429-Band A	Isoform 2 of Probable ATP-dependent RNA helicase DDX5 OS=Homo sapiens GN=DDX5 - [DDX5_HUMAN]	3	0.33
NBN_HPA001429-Band A	Isoform Short of RNA-binding protein FUS OS=Homo sapiens GN=FUS - [FUS_HUMAN]	3	0.33
NBN_HPA001429-Band A	Isoform 2 of Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM - [HNRPM_HUMAN]	10	0.25
NBN_HPA001429-Band A	Isoform Short of Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU - [HNRPU_HUMAN]	29	0.11
NBN_HPA001429-Band A	Splicing factor, proline- and glutamine-rich OS=Homo sapiens GN=SFQ PE=1 SV=2 - [SFQ_HUMAN]	11	0.10
NBN_HPA001429-Band A	Desmoplakin OS=Homo sapiens GN=DSP PE=1 SV=3 - [DESP_HUMAN]	14	0.07
NBN_HPA001429-Band A	X-ray repair cross-complementing protein 6 OS=Homo sapiens GN=XRCC6 PE=1 SV=2 - [XRCC6_HUMAN]	37	0.03
NBN_HPA001429-Band B	Nibrin OS=Homo sapiens GN=NBN PE=1 SV=1 - [NBN_HUMAN]	14	Not in Control IP
NBN_HPA001429-Band B	DNA repair protein RAD50 OS=Homo sapiens GN=RAD50 PE=1 SV=1 - [RAD50_HUMAN]	8	Not in Control IP
NBN_HPA001429-Band B	Sideroflexin OS=Homo sapiens GN=SFN3 PE=1 SV=1 - [A0A0A0MS41_HUMAN]	7	Not in Control IP
NBN_HPA001429-Band B	Golgin-45 (Fragment) OS=Homo sapiens GN=BLZF1 PE=1 SV=1 - [Q5T532_HUMAN]	4	Not in Control IP
NBN_HPA001429-Band B	Ig gamma-1 chain C region (Fragment) OS=Homo sapiens GN=IGHG1 PE=1 SV=1 - [A0A0A0MS07_HUMAN]	3	Not in Control IP
NBN_HPA001429-Band B	Protein PRRC1 OS=Homo sapiens GN=PRRC1 PE=1 SV=1 - [PRRC1_HUMAN]	3	Not in Control IP
NBN_HPA001429-Band B	Hornerin OS=Homo sapiens GN=HRNR PE=1 SV=2 - [HORN_HUMAN]	2	Not in Control IP
NBN_HPA001429-Band B	Double-strand break repair protein MRE11A OS=Homo sapiens GN=MRE11A PE=1 SV=3 - [MRE11_HUMAN]	2	Not in Control IP
NBN_HPA001429-Band B	Double-strand break repair protein MRE11A OS=Homo sapiens GN=MRE11A PE=1 SV=1 - [F8W7U8_HUMAN]	2	Not in Control IP
NBN_HPA001429-Band B	Asialoglycoprotein receptor 1 (Fragment) OS=Homo sapiens GN=ASGR1 PE=1 SV=1 - [K7EP55_HUMAN]	2	Not in Control IP
NBN_HPA001429-Band B	Centrosomal protein of 72 kDa OS=Homo sapiens GN=CEP72 PE=1 SV=2 - [CEP72_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band B	7-dehydrocholesterol reductase OS=Homo sapiens GN=DHCR7 PE=1 SV=1 - [DHCR7_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band B	Ig kappa chain V-1 region Roy OS=Homo sapiens GN=KV116 PE=1 SV=1 - [KV116_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band B	Probable serine carboxypeptidase CPVL (Fragment) OS=Homo sapiens GN=CPVL PE=1 SV=1 - [H7COX5_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band B	Sideroflexin-1 OS=Homo sapiens GN=SFN1 PE=1 SV=1 - [D6RAE9_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band B	High affinity cationic amino acid transporter 1 (Fragment) OS=Homo sapiens GN=SLC7A1 PE=1 SV=1 - [Q5JR49_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band B	Pyrroline-5-carboxylate reductase 1, mitochondrial (Fragment) OS=Homo sapiens GN=PYCR1 PE=1 SV=1 - [J3QL24_HUMAN]	1	Not in Control IP
NBN_HPA001429-Band B	Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2 - [GRP75_HUMAN]	31	4.00
NBN_HPA001429-Band B	Elongation factor Tu, mitochondrial OS=Homo sapiens GN=TUFM PE=1 SV=2 - [EFTU_HUMAN]	4	3.00
NBN_HPA001429-Band B	Heterogeneous nuclear ribonucleoproteins C1/C2 (Fragment) OS=Homo sapiens GN=HNRNPC PE=1 SV=1 - [G3V5X6_HUMAN]	4	3.00
NBN_HPA001429-Band B	Isoform 2 of ATP synthase subunit alpha, mitochondrial OS=Homo sapiens GN=ATP5A1 - [ATPA_HUMAN]	16	2.60
NBN_HPA001429-Band B	Albumin, isoform CRA_k OS=Homo sapiens GN=ALB PE=1 SV=1 - [C9JKR2_HUMAN]	5	1.00
NBN_HPA001429-Band B	Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1 - [EF1A1_HUMAN]	4	1.00
NBN_HPA001429-Band B	Isoform 2 of Heterogeneous nuclear ribonucleoprotein A1 OS=Homo sapiens GN=HNRNPA1 - [ROA1_HUMAN]	2	1.00
NBN_HPA001429-Band B	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1 - [ACTB_HUMAN]	13	0.78
NBN_HPA001429-Band B	ATP synthase subunit beta, mitochondrial OS=Homo sapiens GN=ATP5B PE=1 SV=3 - [ATPB_HUMAN]	7	0.75
NBN_HPA001429-Band B	Tubulin alpha-1C chain OS=Homo sapiens GN=TUBA1C PE=1 SV=1 - [TBA1C_HUMAN]	5	0.75
NBN_HPA001429-Band B	Isoform A2 of Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 - [ROA2_HUMAN]	5	0.67