

Spectrum	Name of Protein	Unique Peptide Counts	Fold Enrichment
ATF7_HPA003384_Band A	Golgin subfamily A member 5 OS=Homo sapiens GN=GOLGA5 PE=1 SV=3 - [GOGA5_HUMAN]	14	Not in Control IP
ATF7_HPA003384_Band A	ATP-dependent DNA helicase Q1 OS=Homo sapiens GN=RECLQ PE=1 SV=3 - [RECLQ1_HUMAN]	4	Not in Control IP
ATF7_HPA003384_Band A	Nucleolar protein 58 OS=Homo sapiens GN=NOP58 PE=1 SV=1 - [NOP58_HUMAN]	3	Not in Control IP
ATF7_HPA003384_Band A	Putative uncharacterized protein DKFZp686H13261 OS=Homo sapiens GN=DKFZp686H13261 PE=1 SV=1 - [Q6N039_HUMAN]	3	Not in Control IP
ATF7_HPA003384_Band A	ATP-dependent RNA helicase DDX3X OS=Homo sapiens GN=DDX3X PE=1 SV=1 - [A0A0D95FB3_HUMAN]	2	Not in Control IP
ATF7_HPA003384_Band A	Pescadillo homolog OS=Homo sapiens GN=PE51 PE=1 SV=1 - [B5MCF9_HUMAN]	2	Not in Control IP
ATF7_HPA003384_Band A	Isoform 5 of Cyclic AMP-dependent transcription factor ATF-7 OS=Homo sapiens GN=ATF7 - [ATF7_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	Protein AATF OS=Homo sapiens GN=AATF PE=1 SV=1 - [AATF_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	Isoform 11 of ARF GTPase-activating protein GIT2 OS=Homo sapiens GN=GIT2 - [GIT2_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	Isoform 2 of GRAM domain-containing protein 1A OS=Homo sapiens GN=GRAMD1A - [GRM1A_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	UPF0606 protein KIAA1549L OS=Homo sapiens GN=KIAA1549L PE=2 SV=2 - [K154L_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	Lysine-specific demethylase 4A OS=Homo sapiens GN=KDM4A PE=1 SV=2 - [KDM4A_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	Something about silencing protein 10 OS=Homo sapiens GN=UTP3 PE=1 SV=1 - [SAS10_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	U3 small nucleolar RNA-associated protein 6 homolog OS=Homo sapiens GN=UTP6 PE=2 SV=2 - [UTP6_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	Isoform 3 of DBIRD complex subunit ZNF326 OS=Homo sapiens GN=ZNF326 - [ZN326_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	TATA-binding protein-associated factor 2N (Fragment) OS=Homo sapiens GN=TAF15 PE=1 SV=1 - [A0A075B7F4_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	U3 small nucleolar RNA-associated protein 14 homolog A (Fragment) OS=Homo sapiens GN=UTP14A PE=1 SV=1 - [X6RJO_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	Heterochromatin protein 1-binding protein 3 (Fragment) OS=Homo sapiens GN=HP1BP3 PE=1 SV=1 - [Q55WC8_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	Ribosomal L1 domain-containing protein 1 (Fragment) OS=Homo sapiens GN=RS1D1 PE=1 SV=1 - [J3Q5V6_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	WD repeat-containing protein 43 (Fragment) OS=Homo sapiens GN=WDR43 PE=1 SV=1 - [C9I2K7_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band A	Rho guanine nucleotide exchange factor 7 OS=Homo sapiens GN=ARHGEF7 PE=1 SV=1 - [E9PDQ5_HUMAN]	5	4.00
ATF7_HPA003384_Band A	Nucleolar protein 56 OS=Homo sapiens GN=NOP56 PE=1 SV=4 - [NOP56_HUMAN]	4	1.50
ATF7_HPA003384_Band A	Granulins (Fragment) OS=Homo sapiens GN=GRN PE=1 SV=1 - [K7EKL3_HUMAN]	4	1.50
ATF7_HPA003384_Band A	Isoform 6 of RNA-binding protein EWS OS=Homo sapiens GN=EWSR1 - [EWS_HUMAN]	3	1.00
ATF7_HPA003384_Band A	Isoform 2 of 60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 - [CH60_HUMAN]	2	1.00
ATF7_HPA003384_Band A	ATP-dependent RNA helicase DDX18 OS=Homo sapiens GN=DDX18 PE=1 SV=2 - [DDX18_HUMAN]	2	1.00
ATF7_HPA003384_Band A	Trifunctional enzyme subunit alpha, mitochondrial OS=Homo sapiens GN=HADHA PE=1 SV=2 - [ECHA_HUMAN]	2	1.00
ATF7_HPA003384_Band A	U3 small nucleolar RNA-associated protein 18 homolog OS=Homo sapiens GN=UTP18 PE=1 SV=3 - [UTP18_HUMAN]	2	1.00
ATF7_HPA003384_Band A	Protein TRAJ56 (Fragment) OS=Homo sapiens GN=TRAJ56 PE=4 SV=1 - [A0A075B622_HUMAN]	1	1.00
ATF7_HPA003384_Band A	ARF GTPase-activating protein GIT1 OS=Homo sapiens GN=GIT1 PE=1 SV=1 - [J3QRU8_HUMAN]	1	1.00
ATF7_HPA003384_Band A	FACT complex subunit SSRP1 OS=Homo sapiens GN=SSRP1 PE=1 SV=1 - [SSRP1_HUMAN]	7	0.83
ATF7_HPA003384_Band A	Isoform 2 of Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM - [HNRPM_HUMAN]	10	0.67
ATF7_HPA003384_Band A	RNA-binding protein 14 OS=Homo sapiens GN=RBM14 PE=1 SV=2 - [RBM14_HUMAN]	4	0.67
ATF7_HPA003384_Band A	Isoform 2 of Probable ATP-dependent RNA helicase DDX5 OS=Homo sapiens GN=DDX5 - [DDX5_HUMAN]	2	0.67
ATF7_HPA003384_Band A	Isoform Short of RNA-binding protein FUS OS=Homo sapiens GN=FUS - [FUS_HUMAN]	8	0.60
ATF7_HPA003384_Band A	Isoform ADelta10 of Prelamin-A/C OS=Homo sapiens GN=LMNA - [LMNA_HUMAN]	8	0.60
ATF7_HPA003384_Band A	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1 - [ACTB_HUMAN]	9	0.50
ATF7_HPA003384_Band A	Isoform 2 of Probable ATP-dependent RNA helicase DDX17 OS=Homo sapiens GN=DDX17 - [DDX17_HUMAN]	3	0.50
ATF7_HPA003384_Band A	Insulin-like growth factor 2 mRNA-binding protein 1 OS=Homo sapiens GN=IGF2BP1 PE=1 SV=2 - [IF2B1_HUMAN]	3	0.50
ATF7_HPA003384_Band A	Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2 - [GRP75_HUMAN]	25	0.47
ATF7_HPA003384_Band A	Transducin beta-like protein 3 OS=Homo sapiens GN=TBL3 PE=1 SV=2 - [TBL3_HUMAN]	4	0.33
ATF7_HPA003384_Band A	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=1 - [A0A0C4DGB6_HUMAN]	4	0.33
ATF7_HPA003384_Band A	Isoform 2 of Heat shock 70 kDa protein 1A OS=Homo sapiens GN=HSPA1A - [HS71A_HUMAN]	5	0.29
ATF7_HPA003384_Band A	X-ray repair cross-complementing protein 5 OS=Homo sapiens GN=XRCC5 PE=1 SV=3 - [XRCC5_HUMAN]	11	0.22
ATF7_HPA003384_Band A	78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2 - [GRP78_HUMAN]	19	0.21
ATF7_HPA003384_Band A	Heterogeneous nuclear ribonucleoprotein U-like protein 1 (Fragment) OS=Homo sapiens GN=HNRNPUL1 PE=1 SV=1 - [MOR3F1_HUMAN]	5	0.20
ATF7_HPA003384_Band A	Isoform 2 of X-ray repair cross-complementing protein 6 OS=Homo sapiens GN=XRCC6 - [XRCC6_HUMAN]	6	0.17
ATF7_HPA003384_Band A	Lamin-B1 OS=Homo sapiens GN=LMNB1 PE=1 SV=2 - [LMNB1_HUMAN]	9	0.13
ATF7_HPA003384_Band A	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1 - [HSP7C_HUMAN]	8	0.10
ATF7_HPA003384_Band B	Heterogeneous nuclear ribonucleoprotein A/B OS=Homo sapiens GN=HNRNPAB PE=1 SV=1 - [D6R9P3_HUMAN]	4	Not in Control IP
ATF7_HPA003384_Band B	Isoform 2 of Heterogeneous nuclear ribonucleoprotein A3 OS=Homo sapiens GN=HNRNP3 - [ROA3_HUMAN]	3	Not in Control IP
ATF7_HPA003384_Band B	Isoform 1 of Core histone macro-H2A.1 OS=Homo sapiens GN=H2AFY - [H2AY_HUMAN]	2	Not in Control IP
ATF7_HPA003384_Band B	Putative keratin-87 protein OS=Homo sapiens GN=KRT87 PE=5 SV=4 - [KR87P_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band B	Isoform 2 of Zinc finger CCCH domain-containing protein 13 OS=Homo sapiens GN=ZC3H13 - [ZC3HD_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band B	Zinc finger and SCAN domain-containing protein 2 OS=Homo sapiens GN=ZSCAN2 PE=1 SV=1 - [F8W6Y9_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band B	RNA-binding motif protein, X chromosome OS=Homo sapiens GN=RBMX PE=1 SV=1 - [H3BU5_HUMAN]	1	Not in Control IP
ATF7_HPA003384_Band B	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1 - [ACTB_HUMAN]	9	1.00
ATF7_HPA003384_Band B	Heterogeneous nuclear ribonucleoprotein D0 (Fragment) OS=Homo sapiens GN=HNRNPD PE=1 SV=5 - [D6RF44_HUMAN]	3	1.00
ATF7_HPA003384_Band B	ATP synthase subunit beta (Fragment) OS=Homo sapiens GN=ATP5B PE=1 SV=1 - [HOYH81_HUMAN]	2	0.50
ATF7_HPA003384_Band B	Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=1 - [Q5JF53_HUMAN]	4	0.33
ATF7_HPA003384_Band B	ATP synthase subunit alpha, mitochondrial (Fragment) OS=Homo sapiens GN=ATP5A1 PE=1 SV=1 - [K7ERX7_HUMAN]	4	0.25