

Spectrum	Name of Protein	Unique Peptide Counts	Fold Enrichment
ZNF592_A301-531A_Band A	Zinc finger protein 592 OS=Homo sapiens GN=ZNF592 PE=1 SV=2 - [ZNF592_HUMAN]	9	Not in Control IP
ZNF592_A301-531A_Band A	Isoform 3 of Protein kinase C-binding protein 1 OS=Homo sapiens GN=ZMYND8 - [PKCB1_HUMAN]	9	Not in Control IP
ZNF592_A301-531A_Band A	Zinc finger protein 318 OS=Homo sapiens GN=ZNF318 PE=1 SV=2 - [ZN318_HUMAN]	7	Not in Control IP
ZNF592_A301-531A_Band A	Isoform 2 of ADAMTS-like protein 4 OS=Homo sapiens GN=ADAMTSL4 - [ATL4_HUMAN]	2	Not in Control IP
ZNF592_A301-531A_Band A	Isoform 2 of Nucleolar RNA helicase 2 OS=Homo sapiens GN=DDX21 - [DDX21_HUMAN]	2	Not in Control IP
ZNF592_A301-531A_Band A	DNA topoisomerase 2-alpha OS=Homo sapiens GN=TOP2A PE=1 SV=3 - [TOP2A_HUMAN]	2	Not in Control IP
ZNF592_A301-531A_Band A	Matrin-3 OS=Homo sapiens GN=MATR3 PE=1 SV=1 - [B3KM87_HUMAN]	2	Not in Control IP
ZNF592_A301-531A_Band A	Isoform 5 of Double-stranded RNA-specific adenosine deaminase OS=Homo sapiens GN=ADAR - [DSRAD_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	Isoform MBP-1 of Alpha-enolase OS=Homo sapiens GN=ENO1 - [ENOA_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	Msx2-interacting protein OS=Homo sapiens GN=SPEN PE=1 SV=1 - [MINT_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	Myosin regulatory light chain 12A OS=Homo sapiens GN=MYL12A PE=1 SV=2 - [ML12A_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3 - [NUCL_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	Protein RRP5 homolog OS=Homo sapiens GN=PDCD11 PE=1 SV=3 - [RRP5_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	Splicing factor 3B subunit 3 OS=Homo sapiens GN=SF3B3 PE=1 SV=4 - [SF3B3_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	DNA topoisomerase 1 OS=Homo sapiens GN=TOP1 PE=1 SV=2 - [TOP1_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	DNA damage-binding protein 1 (Fragment) OS=Homo sapiens GN=DDB1 PE=1 SV=1 - [F5GWIO_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	DNA topoisomerase 2 (Fragment) OS=Homo sapiens GN=TOP2B PE=1 SV=1 - [E9PCY5_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	Nuclear mitotic apparatus protein 1 (Fragment) OS=Homo sapiens GN=NUMA1 PE=1 SV=1 - [HOYFY6_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	60S ribosomal protein L9 (Fragment) OS=Homo sapiens GN=RPL9 PE=1 SV=1 - [E7ESE0_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	HEAT repeat-containing protein 1 OS=Homo sapiens GN=HEATR1 PE=1 SV=1 - [Q5T3Q7_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	Katanin p60 ATPase-containing subunit A-like 2 (Fragment) OS=Homo sapiens GN=KATNAL2 PE=1 SV=1 - [K7EM02_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	Trypsin-1 (Fragment) OS=Homo sapiens GN=PRSS1 PE=1 SV=1 - [HOY8D1_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	Creatine kinase B-type (Fragment) OS=Homo sapiens GN=CKB PE=1 SV=1 - [G3V461_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	Zinc finger protein 687 (Fragment) OS=Homo sapiens GN=ZNF687 PE=1 SV=1 - [HOY5I5_HUMAN]	1	Not in Control IP
ZNF592_A301-531A_Band A	DNA replication licensing factor MCM4 OS=Homo sapiens GN=MCM4 PE=1 SV=5 - [MCM4_HUMAN]	11	10.00
ZNF592_A301-531A_Band A	ATP-dependent RNA helicase A OS=Homo sapiens GN=DHX9 PE=1 SV=4 - [DHX9_HUMAN]	3	2.00
ZNF592_A301-531A_Band A	DNA replication licensing factor MCM6 OS=Homo sapiens GN=MCM6 PE=1 SV=1 - [MCM6_HUMAN]	4	1.00
ZNF592_A301-531A_Band A	Piezo-type mechanosensitive ion channel component 1 OS=Homo sapiens GN=PIEZO1 PE=1 SV=4 - [PIEZ1_HUMAN]	2	1.00
ZNF592_A301-531A_Band A	Polyubiquitin-B (Fragment) OS=Homo sapiens GN=UBB PE=1 SV=1 - [J3QSA3_HUMAN]	2	1.00
ZNF592_A301-531A_Band A	Isoform Short of Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU - [HNRPU_HUMAN]	7	0.60
ZNF592_A301-531A_Band A	Histone H2B type 1-J OS=Homo sapiens GN=HIST1H2BJ PE=1 SV=3 - [H2B1J_HUMAN]	1	0.50
ZNF592_A301-531A_Band A	Histone H2B type 1-K OS=Homo sapiens GN=HIST1H2BK PE=1 SV=3 - [H2B1K_HUMAN]	1	0.50