

Spectrum	Name of Protein	Unique Peptide Counts	Fold Enrichment
ZNF217_A303-266A_Band A	Zinc finger protein 217 OS=Homo sapiens GN=ZNF217 PE=1 SV=1 - [ZNF217_HUMAN]	62	Not in Control IP
ZNF217_A303-266A_Band A	Biorientation of chromosomes in cell division protein 1-like 1 OS=Homo sapiens GN=BOD1L1 PE=1 SV=2 - [BD1L1_HUMAN]	59	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 5 of Pogo transposable element with ZNF domain OS=Homo sapiens GN=POGZ - [POGZ_HUMAN]	26	Not in Control IP
ZNF217_A303-266A_Band A	Protein Shroom2 OS=Homo sapiens GN=SHROOM2 PE=1 SV=1 - [SHRM2_HUMAN]	26	Not in Control IP
ZNF217_A303-266A_Band A	Structural maintenance of chromosomes protein OS=Homo sapiens GN=SMC1A PE=1 SV=2 - [G8JLG1_HUMAN]	22	Not in Control IP
ZNF217_A303-266A_Band A	Histone-lysine N-methyltransferase SETD1A OS=Homo sapiens GN=SETD1A PE=1 SV=3 - [SET1A_HUMAN]	15	Not in Control IP
ZNF217_A303-266A_Band A	Structural maintenance of chromosomes protein 3 OS=Homo sapiens GN=SMC3 PE=1 SV=2 - [SMC3_HUMAN]	14	Not in Control IP
ZNF217_A303-266A_Band A	Pyruvate carboxylase, mitochondrial OS=Homo sapiens GN=PC PE=1 SV=2 - [PYC_HUMAN]	12	Not in Control IP
ZNF217_A303-266A_Band A	Transcription intermediary factor 1-beta OS=Homo sapiens GN=TRIM28 PE=1 SV=5 - [TIF1B_HUMAN]	11	Not in Control IP
ZNF217_A303-266A_Band A	Cohesin subunit SA-2 OS=Homo sapiens GN=STAG2 PE=1 SV=3 - [STAG2_HUMAN]	9	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 2 of Heterogeneous nuclear ribonucleoprotein U-like protein 1 OS=Homo sapiens GN=HNRNPUL1 - [HNRL1_HUMAN]	8	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 2 of DNA-dependent protein kinase catalytic subunit OS=Homo sapiens GN=PRKDC - [PRKDC_HUMAN]	8	Not in Control IP
ZNF217_A303-266A_Band A	Splicing factor, proline- and glutamine-rich OS=Homo sapiens GN=SFQ PE=1 SV=2 - [SFQ_HUMAN]	8	Not in Control IP
ZNF217_A303-266A_Band A	Acyl-CoA dehydrogenase family member 11 OS=Homo sapiens GN=ACAD11 PE=1 SV=2 - [ACD11_HUMAN]	6	Not in Control IP
ZNF217_A303-266A_Band A	Hornerin OS=Homo sapiens GN=HRNR PE=1 SV=2 - [HORN_HUMAN]	5	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 2 of Protein Wiz OS=Homo sapiens GN=WIZ - [WIZ_HUMAN]	5	Not in Control IP
ZNF217_A303-266A_Band A	Transient receptor potential cation channel subfamily M member 7 OS=Homo sapiens GN=TRPM7 PE=1 SV=1 - [HOYLN8_HUMAN]	5	Not in Control IP
ZNF217_A303-266A_Band A	Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3 - [NUCL_HUMAN]	4	Not in Control IP
ZNF217_A303-266A_Band A	Histone-lysine N-methyltransferase EHMT2 OS=Homo sapiens GN=EHMT2 PE=4 SV=1 - [AOA0G2JIS2_HUMAN]	4	Not in Control IP
ZNF217_A303-266A_Band A	Ribosome-binding protein 1 OS=Homo sapiens GN=RRBP1 PE=1 SV=1 - [R7W755_HUMAN]	4	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 2 of Putative Polycomb group protein ASXL2 OS=Homo sapiens GN=ASXL2 - [ASXL2_HUMAN]	3	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 2 of Chromodomain-helicase-DNA-binding protein 3 OS=Homo sapiens GN=CHD3 - [CHD3_HUMAN]	3	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 4 of Histone-lysine N-methyltransferase EHMT1 OS=Homo sapiens GN=EHMT1 - [EHMT1_HUMAN]	3	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 2 of POZ-, AT hook-, and zinc finger-containing protein 1 OS=Homo sapiens GN=PATZ1 - [PATZ1_HUMAN]	3	Not in Control IP
ZNF217_A303-266A_Band A	Pre-mRNA-processing-splicing factor 8 OS=Homo sapiens GN=PRPF8 PE=1 SV=2 - [PRP8_HUMAN]	3	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 7 of Shootin-1 OS=Homo sapiens GN=KIAA1598 - [SHOT1_HUMAN]	3	Not in Control IP
ZNF217_A303-266A_Band A	Zinc finger protein 316 OS=Homo sapiens GN=ZNF316 PE=1 SV=1 - [ZN316_HUMAN]	3	Not in Control IP
ZNF217_A303-266A_Band A	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1 - [AOA087X010_HUMAN]	3	Not in Control IP
ZNF217_A303-266A_Band A	Isoform EWS-B of RNA-binding protein EWS OS=Homo sapiens GN=EWSR1 - [EWS_HUMAN]	2	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 2 of Hypoxia up-regulated protein 1 OS=Homo sapiens GN=HYOU1 - [HYOU1_HUMAN]	2	Not in Control IP
ZNF217_A303-266A_Band A	Nuclear pore complex protein Nup205 OS=Homo sapiens GN=NUP205 PE=1 SV=3 - [NU205_HUMAN]	2	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 2 of Pre-mRNA-processing factor 6 OS=Homo sapiens GN=PRPF6 - [PRP6_HUMAN]	2	Not in Control IP
ZNF217_A303-266A_Band A	Double-strand-break repair protein rad21 homolog OS=Homo sapiens GN=RAD21 PE=1 SV=2 - [RAD21_HUMAN]	2	Not in Control IP
ZNF217_A303-266A_Band A	Albumin, isoform CRA_k OS=Homo sapiens GN=ALB PE=1 SV=1 - [C9JKR2_HUMAN]	2	Not in Control IP
ZNF217_A303-266A_Band A	DNA-directed RNA polymerase OS=Homo sapiens GN=POLR2A PE=1 SV=1 - [AOA0CADGZ0_HUMAN]	2	Not in Control IP
ZNF217_A303-266A_Band A	Filamin-A OS=Homo sapiens GN=FLNA PE=1 SV=1 - [Q5HY54_HUMAN]	2	Not in Control IP
ZNF217_A303-266A_Band A	Nuclear mitotic apparatus protein 1 OS=Homo sapiens GN=NUMA1 PE=1 SV=1 - [AOA087WY61_HUMAN]	2	Not in Control IP
ZNF217_A303-266A_Band A	Matrin-3 (Fragment) OS=Homo sapiens GN=MATR3 PE=1 SV=1 - [D6R991_HUMAN]	2	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 2 of AT-rich interactive domain-containing protein 1A OS=Homo sapiens GN=ARID1A - [ARI1A_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 5 of Dynamin-3 OS=Homo sapiens GN=DNM3 - [DYN3_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Filaggrin OS=Homo sapiens GN=FLG PE=1 SV=3 - [FILA_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 5 of Filamin-B OS=Homo sapiens GN=FLNB - [FLNB_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 2 of Golgin subfamily A member 5 OS=Homo sapiens GN=GOLGA5 - [GOGA5_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 2 of Host cell factor 1 OS=Homo sapiens GN=HCFC1 - [HCFC1_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 4 of Host cell factor 1 OS=Homo sapiens GN=HCFC1 - [HCFC1_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 4 of DNA mismatch repair protein Msh6 OS=Homo sapiens GN=MSH6 - [MSH6_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 2 of UDP-N-acetylglucosamine-peptide N-acetylglucosaminyltransferase 110 kDa subunit OS=Homo sapiens GN=OGT - [OGT1_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Poly (ADP-ribose) polymerase 1 OS=Homo sapiens GN=PARP1 PE=1 SV=4 - [PARP1_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	FACT complex subunit SPT16 OS=Homo sapiens GN=SPT16H PE=1 SV=1 - [SP16H_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Isoleucine--tRNA ligase, mitochondrial OS=Homo sapiens GN=IARS2 PE=1 SV=2 - [SYIM_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 2 of U5 small nuclear ribonucleoprotein 200 kDa helicase OS=Homo sapiens GN=SNRNP200 - [U520_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Isoform 3 of Zinc finger protein ZFAT OS=Homo sapiens GN=ZFAT - [ZFAT_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Zinc finger MYM-type protein 2 OS=Homo sapiens GN=ZMYM2 PE=1 SV=1 - [ZMYM2_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Adenomatous polyposis coli protein (Fragment) OS=Homo sapiens GN=APC PE=1 SV=1 - [B722B6_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Complement C4-B OS=Homo sapiens GN=C4B_2 PE=4 SV=1 - [AOA0G2JL54_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	40S ribosomal protein S15 OS=Homo sapiens GN=RPS15 PE=1 SV=2 - [K7EM56_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	HCG2044799 OS=Homo sapiens GN=HNRNPUL2-BSCL2 PE=4 SV=1 - [H3BQ27_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Biorientation of chromosomes in cell division protein 1-like 1 (Fragment) OS=Homo sapiens GN=BOD1L1 PE=1 SV=1 - [HOY9Y2_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Transformation/transcription domain-associated protein (Fragment) OS=Homo sapiens GN=TRRAP PE=1 SV=1 - [HOY4W2_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	60S ribosomal protein L26 OS=Homo sapiens GN=RPL26 PE=4 SV=1 - [J3K550_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	DNA topoisomerase 2 (Fragment) OS=Homo sapiens GN=TOP2B PE=1 SV=1 - [E9PCY5_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	DNA-directed RNA polymerase OS=Homo sapiens GN=POLR2B PE=1 SV=2 - [C9J4M6_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Lysine-specific histone demethylase 1A (Fragment) OS=Homo sapiens GN=KDM1A PE=1 SV=5 - [R4GMP9_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Polyubiquitin-B (Fragment) OS=Homo sapiens GN=UBB PE=1 SV=1 - [J3QSA3_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Bleomycin hydrolase OS=Homo sapiens GN=BLMH PE=1 SV=1 - [J3KS79_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Zinc finger protein 217 (Fragment) OS=Homo sapiens GN=ZNF217 PE=1 SV=1 - [Q9BZ31_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Replication factor C subunit 1 (Fragment) OS=Homo sapiens GN=RFC1 PE=1 SV=1 - [HOY8U4_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Importin subunit beta-1 (Fragment) OS=Homo sapiens GN=KPNA1 PE=1 SV=5 - [J3QR48_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	HEAT repeat-containing protein 1 OS=Homo sapiens GN=HEATR1 PE=1 SV=1 - [Q5T3Q7_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	DNA repair protein RAD50 OS=Homo sapiens GN=RAD50 PE=1 SV=1 - [E7ESD9_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Protein EMSY (Fragment) OS=Homo sapiens GN=C11orf30 PE=1 SV=1 - [HOYET8_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	HEAT repeat-containing protein 6 OS=Homo sapiens GN=HEATR6 PE=1 SV=1 - [K7ELR8_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Transferrin receptor protein 1 OS=Homo sapiens GN=TFRC PE=1 SV=1 - [F8WBES_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Zinc finger protein 687 (Fragment) OS=Homo sapiens GN=ZNF687 PE=1 SV=1 - [HOY515_HUMAN]	1	Not in Control IP
ZNF217_A303-266A_Band A	Desmoplakin OS=Homo sapiens GN=DSP PE=1 SV=3 - [DESP_HUMAN]	17	17.00
ZNF217_A303-266A_Band A	Isoform 7 of Plectin OS=Homo sapiens GN=PLEC - [PLEC_HUMAN]	8	8.00
ZNF217_A303-266A_Band A	Isoform Short of Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNP - [HNRPU_HUMAN]	14	4.67
ZNF217_A303-266A_Band A	ATP-dependent RNA helicase A OS=Homo sapiens GN=DHX9 PE=1 SV=4 - [DHX9_HUMAN]	4	4.00