

Spectrum	Name of Protein	Unique Peptide Counts	Fold Enrichment
MCM3_9/61719_Band A	DNA replication licensing factor MCM3 OS=Homo sapiens GN=MCM3 PE=1 SV=3 - [MCM3_HUMAN]	51	Not in Control IP
MCM3_9/61719_Band A	DNA replication licensing factor MCM5 OS=Homo sapiens GN=MCM5 PE=1 SV=5 - [MCM5_HUMAN]	19	Not in Control IP
MCM3_9/61719_Band A	Cleavage and polyadenylation-specificity factor subunit 6 OS=Homo sapiens GN=CPSF6 PE=1 SV=1 - [F8WJN3_HUMAN]	5	Not in Control IP
MCM3_9/61719_Band A	Splicing factor 3B subunit 3 OS=Homo sapiens GN=SF3B3 PE=1 SV=4 - [SF3B3_HUMAN]	4	Not in Control IP
MCM3_9/61719_Band A	Pre-mRNA-splicing factor ATP-dependent RNA helicase DHX15 OS=Homo sapiens GN=DHX15 PE=1 SV=2 - [DHX15_HUMAN]	2	Not in Control IP
MCM3_9/61719_Band A	Isoform 5 of Histone H2A.V OS=Homo sapiens GN=H2AFV - [H2AFV_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	Poly [ADP-ribose] polymerase 1 OS=Homo sapiens GN=PARP1 PE=1 SV=4 - [PARP1_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	Isoform 2 of 116 kDa U5 small nuclear ribonucleoprotein component OS=Homo sapiens GN=EFTUD2 - [U5S1_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	Polyubiquitin-B (Fragment) OS=Homo sapiens GN=UBB PE=1 SV=1 - [J3QSA3_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	Isoform 2 of Probable ATP-dependent RNA helicase DDX5 OS=Homo sapiens GN=DDX5 - [DDX5_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	Isoform Short of RNA-binding protein FUS OS=Homo sapiens GN=FUS - [FUS_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	DNA replication licensing factor MCM6 OS=Homo sapiens GN=MCM6 PE=1 SV=1 - [MCM6_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	Isoform 2 of Ribosomal RNA processing protein 1 homolog B OS=Homo sapiens GN=RRP1B - [RRP1B_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	Isoform 4 of Splicing factor 1 OS=Homo sapiens GN=SF1 - [SF01_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	Isoform 2 of Splicing factor 3A subunit 1 OS=Homo sapiens GN=SF3A1 - [SF3A1_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	Complement C4-B OS=Homo sapiens GN=C4B_2 PE=4 SV=1 - [AOA0G2L54_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1 OS=Homo sapiens GN=RPN1 PE=1 SV=1 - [B7Z4L4_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	Nucleolar protein 56 (Fragment) OS=Homo sapiens GN=NOP56 PE=1 SV=1 - [HOY653_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	Nucleolar transcription factor 1 (Fragment) OS=Homo sapiens GN=UBTF PE=1 SV=1 - [E9PLT2_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band A	Nuclear pore complex protein Nup93 (Fragment) OS=Homo sapiens GN=NUP93 PE=1 SV=1 - [H3BM93_HUMAN]	3	3.00
MCM3_9/61719_Band A	DNA replication licensing factor MCM7 OS=Homo sapiens GN=MCM7 PE=1 SV=4 - [MCM7_HUMAN]	8	2.67
MCM3_9/61719_Band A	RNA-binding protein 14 OS=Homo sapiens GN=RBM14 PE=1 SV=2 - [RBM14_HUMAN]	2	2.00
MCM3_9/61719_Band A	DNA replication licensing factor MCM4 (Fragment) OS=Homo sapiens GN=MCM4 PE=1 SV=5 - [E5RFJ8_HUMAN]	2	2.00
MCM3_9/61719_Band A	Isoform 5 of Prelamin-A/C OS=Homo sapiens GN=LMNA - [LMNA_HUMAN]	9	1.80
MCM3_9/61719_Band A	Isoform 2 of Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM - [HNRPM_HUMAN]	5	1.67
MCM3_9/61719_Band A	Lamin-B1 OS=Homo sapiens GN=LMNB1 PE=1 SV=2 - [LMNB1_HUMAN]	5	1.67
MCM3_9/61719_Band A	X-ray repair cross-complementing protein 6 OS=Homo sapiens GN=XRCC6 PE=1 SV=1 - [B1AH9_HUMAN]	3	1.50
MCM3_9/61719_Band A	Heterogeneous nuclear ribonucleoprotein R OS=Homo sapiens GN=HNRNPR PE=1 SV=1 - [B4DT28_HUMAN]	3	1.50
MCM3_9/61719_Band A	78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2 - [GRP78_HUMAN]	4	1.33
MCM3_9/61719_Band A	Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2 - [GRP75_HUMAN]	13	1.30
MCM3_9/61719_Band A	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90A1 PE=1 SV=4 - [HS90B_HUMAN]	3	1.00
MCM3_9/61719_Band A	Isoform 2 of Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 - [HSP7C_HUMAN]	2	1.00
MCM3_9/61719_Band A	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1 - [AOA087X010_HUMAN]	2	1.00
MCM3_9/61719_Band A	Heterogeneous nuclear ribonucleoprotein Q (Fragment) OS=Homo sapiens GN=SYNCRIP PE=1 SV=1 - [F6UXX1_HUMAN]	1	1.00
MCM3_9/61719_Band A	RNA-binding protein EWS (Fragment) OS=Homo sapiens GN=EWSR1 PE=1 SV=5 - [F8WC90_HUMAN]	1	1.00
MCM3_9/61719_Band A	X-ray repair cross-complementing protein 5 OS=Homo sapiens GN=XRCC5 PE=1 SV=3 - [XRCC5_HUMAN]	2	0.67
MCM3_9/61719_Band A	Isoform 2 of Nucleolar RNA helicase 2 OS=Homo sapiens GN=DDX21 - [DDX21_HUMAN]	1	0.50
MCM3_9/61719_Band A	Isoform Short of Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRPU - [HNRPU_HUMAN]	9	0.43
MCM3_9/61719_Band B	DNA replication licensing factor MCM3 OS=Homo sapiens GN=MCM3 PE=1 SV=3 - [MCM3_HUMAN]	13	Not in Control IP
MCM3_9/61719_Band B	Vimentin OS=Homo sapiens GN=VIM PE=1 SV=4 - [VIME_HUMAN]	12	Not in Control IP
MCM3_9/61719_Band B	Inosine-5'-monophosphate dehydrogenase 2 (Fragment) OS=Homo sapiens GN=IMPDH2 PE=1 SV=1 - [HOY4R1_HUMAN]	11	Not in Control IP
MCM3_9/61719_Band B	Isoform 2 of Tubulin alpha-1A chain OS=Homo sapiens GN=TUBA1A - [TBA1A_HUMAN]	4	Not in Control IP
MCM3_9/61719_Band B	Poly(U)-binding-splicing factor PUF60 (Fragment) OS=Homo sapiens GN=PUF60 PE=1 SV=5 - [E9PQ56_HUMAN]	3	Not in Control IP
MCM3_9/61719_Band B	ATP synthase subunit alpha, mitochondrial (Fragment) OS=Homo sapiens GN=ATP5A1 PE=1 SV=1 - [K7EK77_HUMAN]	3	Not in Control IP
MCM3_9/61719_Band B	DNA replication licensing factor MCM5 OS=Homo sapiens GN=MCM5 PE=1 SV=5 - [MCM5_HUMAN]	2	Not in Control IP
MCM3_9/61719_Band B	Isoform 5 of Prelamin-A/C OS=Homo sapiens GN=LMNA - [LMNA_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band B	Putative translationally-controlled tumor protein-like protein TPT1P8 OS=Homo sapiens GN=TPT1P8 PE=5 SV=2 - [TCTP8_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band B	ATP synthase subunit beta (Fragment) OS=Homo sapiens GN=ATP5B PE=1 SV=1 - [HOYH81_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band B	Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1 - [AOA087WVQ9_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band B	Uncharacterized protein (Fragment) OS=Homo sapiens PE=3 SV=1 - [HOYAE9_HUMAN]	1	Not in Control IP
MCM3_9/61719_Band B	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1 - [AOA087X010_HUMAN]	1	0.33