

Proteins in Only P. falciparum Control Samples. Column one provides the protein name and column two the accession number of each protein. Any identifications made from the reverse concatenated database are not represented.

<i>Protein</i>	<i>Accession Number</i>
Putative uncharacterized protein OS=Plasmodium falciparum (isolate 3D7) GN=MAL8P1.103 PE=4 SV=1	C0H4U0_PLAF7
Conserved Plasmodium protein OS=Plasmodium falciparum (isolate 3D7) GN=PFL2120w PE=4 SV=1	Q8I4Y9_PLAF7
Putative uncharacterized protein OS=Plasmodium falciparum (isolate 3D7) GN=PFE1465w PE=4 SV=1	C0H4G7_PLAF7
DNA repair protein, putative OS=Plasmodium falciparum (isolate 3D7) GN=PFE0270c PE=3 SV=1	Q8I447_PLAF7
Histidine--tRNA ligase, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF14_0428 PE=4 SV=1	Q8IL22_PLAF7
40S ribosomal protein S28e, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF14_0585 PE=4 SV=1	Q8IKL9_PLAF7
10b antigen, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF10_0213 PE=4 SV=1	Q8IJI4_PLAF7
Ubiquitination-mediated degradation component, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF08_0020 PE=4 SV=1	C0H4Y0_PLAF7
Putative uncharacterized protein OS=Plasmodium falciparum (isolate 3D7) GN=PF07_0016 PE=4 SV=1	Q8IC27_PLAF7
Vacuolar ATP synthase subunit E, putative OS=Plasmodium falciparum (isolate 3D7) GN=PFI1670c PE=3 SV=2	Q8I2H3_PLAF7
V-type ATPase, subunit C, putative OS=Plasmodium falciparum (isolate 3D7) GN=PFA_0300c PE=4 SV=1	Q8I280_PLAF7
Arginyl-tRNA synthetase, putative OS=Plasmodium falciparum (isolate 3D7) GN=PFL0900c PE=3 SV=1	Q8I5M2_PLAF7
Conserved Plasmodium protein OS=Plasmodium falciparum (isolate 3D7) GN=PF11_0332 PE=4 SV=1	Q8II42_PLAF7
Putative uncharacterized protein OS=Plasmodium falciparum (isolate 3D7) GN=PFE0990w PE=4 SV=1	C0H4F1_PLAF7
Proteasome, putative OS=Plasmodium falciparum (isolate 3D7) GN=PFI1545c PE=4 SV=1	Q8I0U7_PLAF7
DNA-directed RNA polymerase OS=Plasmodium falciparum (isolate 3D7) GN=PFC0805w PE=3 SV=1	O77375_PLAF7
P-type calcium transporting ATPase OS=Plasmodium falciparum GN=serca PE=3 SV=1	E1CC54_PLAFA
60S ribosomal protein L23a, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF13_0132 PE=3 SV=1	Q8IE82_PLAF7
Coatmer alpha subunit, putative OS=Plasmodium falciparum (isolate 3D7) GN=PFF0330w PE=4 SV=1	C6KSR5_PLAF7
Pfmdr2 protein OS=Plasmodium falciparum GN=pfmdr2 PE=3 SV=1	Q25693_PLAFA
60S ribosomal protein L14, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF14_0296 PE=4 SV=1	Q8ILE8_PLAF7

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60S ribosomal protein L14, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF14_0296 PE=4 SV=1	Q8ILE8_PLAF7