

Proteins Only in P. falciparum Treatment Samples. Column one provides the protein name and column two the accession number for 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	Q8IOU7_PLAF7

<i>Protein</i>	<i>Accession Number</i>
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
Coatamer 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
Heat shock protein 90, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF11_0188 PE=3 SV=1	Q8III6_PLAF7
ORF 2 protein (Fragment) OS=Plasmodium falciparum GN=ORF 2 PE=2 SV=1	Q02602_PLAFA
Ring-exported protein 1 OS=Plasmodium falciparum (isolate 3D7) GN=REX1 PE=4 SV=1	Q8I2G1_PLAF7
40S ribosomal protein S24 OS=Plasmodium falciparum (isolate 3D7) GN=PFE0975c PE=3 SV=1	Q8I3R6_PLAF7
Glutamate--cysteine ligase (Gamma-glutamylcysteine synthetase) OS=Plasmodium falciparum GN=GCS PE=2 SV=1	Q9TY17_PLAFA
Activator of Hsp90 ATPase homolog 1-like protein, putative OS=Plasmodium falciparum (isolate 3D7) GN=PFC0360w PE=4 SV=2	O97256_PLAF7
Ubiquitin conjugating enzyme E2, putative OS=Plasmodium falciparum (isolate 3D7) GN=PFL0190w PE=3 SV=1	Q8I607_PLAF7
Coatamer protein, beta subunit, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF14_0277 PE=4 SV=2	Q8ILG6_PLAF7
Putative uncharacterized protein OS=Plasmodium falciparum (isolate 3D7) GN=MAL8P1.62 PE=4 SV=1	Q8IB31_PLAF7
60S ribosomal protein L36 OS=Plasmodium falciparum (isolate 3D7) GN=PF11_0106 PE=3 SV=2	Q8I713_PLAF7

<i>Protein</i>	<i>Accession Number</i>
Serine rich protein homologue OS=Plasmodium falciparum PE=3 SV=1	Q26015_PLAFA
6-cysteine protein, putative OS=Plasmodium falciparum (isolate 3D7) GN=Pf41 PE=4 SV=1	Q811Y0_PLAF7
DNA replication licensing factor mcm7 homologue, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF07_0023 PE=3 SV=1	Q81C16_PLAF7
Rab2, GTPase OS=Plasmodium falciparum (isolate 3D7) GN=Rab2 PE=3 SV=1	Q815A9_PLAF7
Cytidine triphosphate synthetase OS=Plasmodium falciparum (isolate 3D7) GN=PF14_0100 PE=4 SV=1	Q81LZ3_PLAF7
Ubiquitin carboxyl-terminal hydrolase OS=Plasmodium falciparum (isolate 3D7) GN=PFE1355c PE=3 SV=1	Q813J3_PLAF7
Glycogen synthase kinase 3 OS=Plasmodium falciparum (isolate 3D7) GN=PfGSK-3 PE=4 SV=2	O77344_PLAF7
Acetyl-CoA synthetase OS=Plasmodium falciparum (isolate 3D7) GN=PFF1350c PE=4 SV=1	C6KTB4_PLAF7
DNAJ protein, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF08_0032 PE=4 SV=1	Q81B72_PLAF7
DNA topoisomerase 2 OS=Plasmodium falciparum (isolate 3D7) GN=PF14_0316 PE=3 SV=1	Q81LC8_PLAF7
Conserved Plasmodium protein OS=Plasmodium falciparum (isolate 3D7) GN=PFL0895c PE=4 SV=1	Q815M3_PLAF7
Spermidine synthase OS=Plasmodium falciparum (isolate 3D7) GN=PF11_0301 PE=1 SV=1	Q81I73_PLAF7
Putative uncharacterized protein OS=Plasmodium falciparum (isolate 3D7) GN=PF08_0137 PE=4 SV=1	Q81AK9_PLAF7
PfSec23 protein OS=Plasmodium falciparum (isolate 3D7) GN=Pfsec23 PE=2 SV=1	Q81B60_PLAF7
Replication factor C subunit 1 OS=Plasmodium falciparum GN=rfc1 PE=4 SV=1	Q9GQW6_PLAFA
60S ribosomal protein L8, putative OS=Plasmodium falciparum (isolate 3D7) GN=PFE0845c PE=4 SV=1	Q813T9_PLAF7

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Ubiquitin carboxyl-terminal hydrolase a, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF0680c PE=4 SV=1	Q811U8_PLAF7
Orotate phosphoribosyltransferase OS=Plasmodium falciparum GN=opr1 PE=2 SV=1	Q8N0R1_PLAFA
Calmodulin OS=Plasmodium falciparum (isolate 3D7) GN=PF14_0323 PE=3 SV=2	CALM_PLAF7
Haloacid dehalogenase-like hydrolase, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF11_0190 PE=4 SV=1	Q8III4_PLAF7
Thymidylate kinase, putative OS=Plasmodium falciparum (isolate 3D7) GN=TMK PE=1 SV=1	Q8I4S1_PLAF7
Myosin-A OS=Plasmodium falciparum (isolate FCBR / Columbia) PE=2 SV=1	MYOA_PLAFB
Probable DNA-directed RNA polymerase II subunit RPB11 OS=Plasmodium falciparum (isolate 3D7) GN=PF13_0023 PE=3 SV=1	RPB11_PLAF7
Zinc finger protein, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF13_0313 PE=4 SV=1	Q8IDC0_PLAF7
60S ribosomal protein L28, putative OS=Plasmodium falciparum (isolate 3D7) GN=PF11_0437 PE=4 SV=1	Q8IHU0_PLAF7