Table S1: Full RIP-ChIP dataset. PB ID: P. berghei gene names; Product Description: P. berghei product description; FC DOZI/WT: fold change expression from the DOZI-KO vs. wild type [Mair et al., 2006]; Confidence: corresponds to the false discovery rate (expected percent of false predictions in the set of predictions (see www.plasmodb.org); PF ID: P. falciparum gene names; PF Product Description: P. falciparum product description; Manual Annotation: obtained from literature scan, BLASTs etc; CITH: CITH immunoprecipitation, 0, not enriched, 1, enriched; DOZI: DOZI immunoprecipitation, 0, not enriched, 1, enriched; DOZI/CITH: 0, not enriched, 1, enriched in one IP, 2, enriched in both IPs; Gene Deletion Phenotype (RMgmDB, www.pberghei.eu); PB and PF annotated Gene Ontology (GO) Molecular Function, Biological Process and Cellular Component.

| | | FC | | | | | | | Gene Deletion Phenotype | Phenotype P. berghei annotated GO | | 0 | | syntenic P. falciparum orthologs annotated GO | |
|---------------|--|---------|------------|-----------------|---|---------------------|-----|----------|--|--|--|--|--------------------|---|--------------------|
| PBANKA | Product Description | DOZI/WT | Confidence | PF ID | PF Product Description | Manual Annotation | DOZ | DOZI/CIT | in mosquito and liver stages | Molecular Function | Biological Process | Cellular Component | Molecular Function | Biological Process | Cellular Component |
| PBANKA_134780 | 20 kDa chaperonin, putative (CPN20) | -2.73 | 0.75 | PF3D7_1333000 | 20 kDa chaperonin (CPN20) | chaperone | 1 1 | 2 | na | ATP binding, chaperone binding | protein folding | apicoplast | | | |
| PBANKA_071420 | ClpB protein, putative (ClpB1) | -1.63 | 0.7 | PF3D7_0816600 | ClpB protein, putative (ClpB1) | chaperone | 1 1 | 2 | na | ATP binding, nucleoside-triphosphatase activity, protein bind | ing protein metabolic process | apicoplast | | | |
| PBANKA_051500 | 25 kDa ookinete surface antigen precursor (P25) | -5.59 | 0.75 | PF3D7_1031000 | 25 kDa ookinete surface antigen precursor (Pfs25 | 5) adhesin | 1 1 | 2 | different in fertilization, ookinete and oocys | st null | null | cell surface, membrane | | | |
| PBANKA_051490 | 28 kDa ookinete surface protein (P28) | -2.62 | 0.75 | PF3D7_1030900 | 28 kDa ookinete surface protein (Pfs28) | adhesin | 1 1 | 2 | different in fertilization, ookinete and oocys | st null | null | cell surface, membrane | | | |
| PBANKA_051170 | topoisomerase, putative | | | PF3D7_1027600 | topoisomerase, putative | DNA replication | 1 1 | 2 | na | ATP binding, DNA binding, catalytic activity | DNA metabolic process | chromosome | | | |
| PBANKA_081680 | glycolipid transfer protein, putative | -5.27 | 0.75 | PF3D7_0915800 | glycolipid transfer protein, putative | lipid metabolism | 1 1 | 2 | na | glycolipid binding, glycolipid transporter activity | glycolipid transport | cytoplasm | | | |
| PBANKA_102870 | conserved Plasmodium protein, unknown function | | | PF3D7_1414000 | conserved protein, unknown function | unknown function | 1 1 | 2 | na | null | null | cytoplasm, nucleus | | | |
| PBANKA_071290 | high mobility group protein, putative (HMGB2) | -3.31 | 0.71 | PF3D7_0817900 | high mobility group protein (HMGB2) | chromatin | 1 1 | 2 | na | DNA binding | null | cytoplasm, nucleus | | | |
| PBANKA_092000 | multiprotein bridging factor type 1, putative | | | PF3D7_1128200 | multiprotein bridging factor type 1, putative | transcription | 1 1 | 2 | na | sequence-specific DNA binding | null | cytosol, nuclear periphery | | | |
| PBANKA_103840 | selenoprotein, putative (Sel1) | | | PF3D7_1403600 | selenoprotein (Sel1) | redoxmeth | 1 1 | 2 | na | null | null | endoplasmic reticulum | | | |
| PBANKA_060930 | GPI mannosyltransferase I, putative | | | PF3D7_1210900 | GPI mannosyltransferase I (PIG-M) | GPI | 1 1 | 2 | na | transferase activity, transferring hexosyl groups | GPI anchor biosynthetic process | endoplasmic reticulum membrane, integral to membrane | | | |
| PBANKA_083680 | Pb-fam-1 protein | | | PF3D7_1147800.1 | /2 merozoite adhesive erythrocytic binding protein (MAEB | BL) contaminant | 1 1 | 2 | na | null | null | host cell membrane | | | |
| PBANKA_133890 | glideosome associated protein with multiple membrane spans 1, putative | -4.14 | 0.75 | PF3D7_1323700 | glideosome associated protein with multiple membrane spans 1 (GAPM1 | 1) gliding motility | 1 1 | 2 | KO not successful | null | null | inner membrane complex | | | |
| PBANKA_090710 | inner membrane complex protein 1b (IMC1b) | -3.1 | 0.73 | PF3D7_1141900 | inner membrane complex protein 1b, putative (IMC1b | b) gliding motility | 1 1 | 2 | different in fertilization, ookinete and oocys | st null | null | inner membrane complex | | | |
| PBANKA_124060 | inner membrane complex protein 1g, putative (IMC1G) | -4.13 | 0.75 | PF3D7_0525800 | membrane skeletal protein IMC1-related | gliding motility | 1 1 | 2 | na | null | null | inner membrane complex | | | |
| PBANKA_143660 | inner membrane complex protein 1h (IMC1h) | -2.99 | 0.73 | PF3D7_1221400 | membrane skeletal protein, putative (ALV3) | gliding motility | 1 1 | 2 | different in fertilization, ookinete, oocyst, sporozoite and liver stage | null | null | inner membrane complex | | | |
| PBANKA_130290 | copper transporter, putative | | | PF3D7_1439000 | copper transporter putative | transporter | 1 1 | 2 | na | copper ion transmembrane transporter activity | copper ion transport | integral to membrane | | | |
| PBANKA_123130 | metabolite/drug transporter, putative | -3.99 | 0.75 | PF3D7_0516500 | metabolite/drug transporter, putative | transporter | 1 1 | 2 | not different from wild type | tetracycline:hydrogen antiporter activity | response to antibiotic, tetracycline transport | integral to membrane | | | |
| PBANKA_123780 | multidrug resistance protein, putative (MDR1) | -2.4 | 0.7 | PF3D7_0523000 | multidrug resistance protein (MDR1) | transporter | 1 1 | 2 | na | ATP binding, ATPase activity, coupled to transmembrane movement of substan | ces transport | integral to membrane | | | |

| | | | | | | • | 1000 | | | | | | |
|---------------|--|-------|------|-------------------|---|-----------------------|------|---|------|--|--|---|---|
| PBANKA_082140 | ADP-ribosylation factor-like protein | -4.56 | 0.75 | PF3D7_0920500 | ADP-ribosylation factor, putative | trafficking | 1 | 1 | 2 n | na | GTP binding | small GTPase mediated signal transduction | intracellular |
| PBANKA_051850 | ADP-ribosylation factor, putative | -4.45 | 0.75 | PF3D7_1034700 | ADP-ribosylation factor, putative | trafficking | 1 | 1 | 2 n | na | GTP binding | intracellular protein transport, small GTPase mediated signal transduction | intracellular |
| PBANKA_100830 | conserved Plasmodium protein, unknown function | | | PF3D7_1436300 | translocon component PTEX150 (PTEX150) | trafficking | 1 | 1 | 2 n | na | GTP binding | null | intracellular |
| PBANKA_142370 | GDP-mannose 4,6-dehydratase, putative | -2.74 | 0.75 | PF3D7_0813800 | GDP-mannose 4,6-dehydratase, putative | mannose metabolism | 1 | 1 | 2 n | na | GDP-mannose 4,6-dehydratase activity, coenzyme binding | GDP-mannose metabolic process | intracellular |
| PBANKA_122330 | GTPase, Rab18, putative | | | PF3D7_0807300 | Rab GTPase 18 (RAB18) | trafficking | 1 | 1 | 2 n | na | GTP binding, GTPase activity, protein binding | intracellular protein transport, nucleocytoplasmic transport, small GTPase mediated signal transduction | intracellular |
| PBANKA_141890 | Rab GTPase 11a (Rab11a) | -4.21 | 0.75 | PF3D7_1320600 | Rab GTPase 11a (RAB11a) | trafficking | 1 | 1 | 2 k | KO not successful | GTP binding, GTPase activity, protein binding | intracellular protein transport, nucleocytoplasmic transport, small GTPase mediated signal transduction | intracellular |
| PBANKA_030800 | Rab5a, GTPase, putative | | | PF3D7_0211200 | Rab GTPase 5a (RAB5a) | trafficking | 1 | 1 | 2 n | na | GTP binding, GTPase activity, protein binding | intracellular protein transport, nucleocytoplasmic transport, small GTPase mediated signal transduction | intracellular |
| PBANKA_041820 | Rab7, putative | | | PF3D7_0903200 | Rab GTPase 7 (Rab7) | trafficking | 1 | 1 | 2 n | na | GTP binding, GTPase activity, protein binding | intracellular protein transport, nucleocytoplasmic transport, small GTPase mediated signal transduction | intracellular |
| PBANKA_114350 | ribosome biogenesis protein MRT4, putative | | | PF3D7_1367600 | ribosome biogenesis protein MRT4, putative | ribosome assembly | 1 | 1 | 2 n | na | null | ribosome biogenesis | intracellular |
| PBANKA_144580 | small GTPase Rab2, putative | | | PF3D7_1231100 | Rab GTPase 2 (RAB2) | trafficking | 1 | 1 | 2 n | na | GTP binding, GTPase activity, protein binding | intracellular protein transport, nucleocytoplasmic transport, small GTPase mediated signal transduction | intracellular |
| PBANKA_071400 | meiotic recombination protein DMC1-like protein, putative | | | PF3D7_0816800 | meiotic recombination protein dmc1-like protein | DNA repair | 1 | 1 | 2 di | different in fertilization, ookinete, oocyst, sporozoite and liver stage | ATP binding, DNA binding, DNA-dependent ATPase activity, protein binding | reciprocal meiotic recombination | intracellular, nucleus |
| PBANKA_070910 | 60S ribosomal protein L22, putative | | | PF3D7_0821700 | 60S ribosomal protein L22, putative | ribosome | 1 | 1 | 2 n | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_134670 | 60S ribosomal protein L23, putative | | | PF3D7_1331800 | 60S ribosomal protein L23, putative | ribosome | 1 | 1 | 2 n | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_090490 | conserved Plasmodium protein, unknown function | | | PF3D7_1144100 | conserved protein, unknown function | unknown function | 1 | 1 | 2 n | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_124560 | mitochondrial ribosomal protein S16 precursor, putative | | | PF3D7_0531200 | mitochondrial ribosomal protein S16 precursor, putative | e ribosome mito | 1 | 1 | 2 n | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_061600 | mitochondrial ribosomal protein S8 precursor, putative | | | PF3D7_0718400 | mitochondrial ribosomal protein S8 precursor, putative | ribosome mito | 1 | 1 | 2 n | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_101810 | ABC transporter, putative | | | PF3D7_1426500 | ABC transporter, (EPP family), putative | transporter | 1 | 1 | 2 n | na | ATP binding, ATPase activity | null | membrane |
| PBANKA_070670 | dicarboxylate/tricarboxylate carrier, putative (DTC) | -5.45 | 0.75 | PF3D7_0823900 | dicarboxylate/tricarboxylate carrier (DTC) | transporter | 1 | 1 | 2 n | na | binding | transport | membrane |
| PBANKA_010210 | dihydroorotate dehydrogenase, putative (DHODH) | -2.58 | 0.75 | PF3D7_0603300 | dihydroorotate dehydrogenase, mitochondrial precursor (DHODH) | pyrimidine metabolism | 1 | 1 | 2 n | na | dihydroorotate dehydrogenase activity, dihydroorotate oxidase activity | 'de novo' pyrimidine nucleobase biosynthetic process, UMP biosynthetic process | membrane |
| PBANKA_135900 | sec61 alpha subunit, putative | | | PF3D7_1346100 | Sec61 alpha subunit, PfSec61 (SEC61) | trafficking | 1 | 1 | 2 n | na | P-P-bond-hydrolysis-driven protein transmembrane transporter activity | protein transport | membrane |
| PBANKA_030690 | Sec61-gamma subunit of protein translocation complex, putative | -3.5 | 0.75 | PF3D7_0210000 | secretory complex protein 61 gamma subunit (Sec61-gamma) | trafficking | 1 | 1 | 2 n | na | P-P-bond-hydrolysis-driven protein transmembrane transporter activity | protein targeting | membrane |
| PBANKA_052350 | clathrin assembly protein, putative | | | PF3D7_0423100 | clathrin assembly protein, putative | trafficking | 1 | 1 | 2 n | na | protein binding, protein transporter activity | intracellular protein transport, vesicle-mediated transport | membrane coat |
| PBANKA_143230 | cell traversal protein for ookinetes and sporozoites (CelTOS) | | | PF3D7_1216600 | cell traversal protein for ookinetes and sporozoites (CelTOS) | adhesin | 1 | 1 | 2 di | different in fertilization, ookinete, oocyst, sporozoite and liver stage | null | null | microneme |
| PBANKA_060710 | tim10 homologue, putative | | | PF3D7_1208600 | tim10 homologue, putative | trafficking | 1 | 1 | 2 n | na | null | protein import into mitochondrial inner membrane | mitochondrial intermembrane space protein transporter complex |
| PBANKA_110260 | zinc binding protein, putative | | | PF3D7_0502900 | mitochondrial inner membrane TIM10 associated protein, putative | trafficking | 1 | 1 | 2 n | na | null | protein import into mitochondrial inner membrane | mitochondrial intermembrane space protein transporter complex |
| PBANKA_082880 | cytochrome c oxidase, putative | | | PF3D7_0928000.1/2 | cytochrome c oxidase, putative | redoxmeth | 1 | 1 | 2 r | na | cytochrome-c oxidase activity | null | mitochondrion |
| PBANKA_135570 | myosin A (MyoA) | -4.46 | 0.75 | PF3D7_1342600 | myosin A (MyoA) | gliding motility | 1 | 1 | 2 (n | (mutated) different in fertilization, ookinete, oocyst and sporozoite | a ATP binding, motor activity | null | myosin complex |
| PBANKA_144940 | Splicing factor 3B subunit 3, putative (SF3B3) | | | PF3D7_1234800 | Splicing factor 3B subunit 3, putative (SF3B3) | mRNA | 1 | 1 | 2 n | na | nucleic acid binding | null | nucleus |
| PBANKA_090590 | transcription factor with AP2 domain(s) (AP2-O) | -3.76 | 0.75 | PF3D7_1143100 | transcription factor with AP2 domain(s) (AP2-O) | transcription | 1 | 1 | 2 d | different in fertilization, ookinete, oocyst and sporozoite | sequence-specific DNA binding transcription factor activity | regulation of transcription, DNA-dependent | nucleus |
| PBANKA_092880 | U2 snRNP auxiliary factor, small subunit, putative | | | PF3D7_1119300 | U2 snRNP auxiliary factor, small subunit, putative | mRNA | 1 | 1 | 2 n | na | RNA binding, zinc ion binding | null | nucleus |
| PBANKA_130820 | 1-acyl-sn-glycerol-3-phosphate acyltransferase, putative (LPAAT) | | | PF3D7_1444300 | apicoplast 1-acyl-sn-glycerol-3-phosphate acyltransferase, putative | fatty acids synth | 1 | 1 | 2 k | KO not successful | transferase activity, transferring acyl groups | metabolic process | null |
| | | | | | | | | | | | | | |

| | | | | | | _ | | | | | | |
|--|-------|------|--|--------------------------------|-----|---|---|---|------|--|---------------------------------------|--------------------------|
| PBANKA_100210 6-cysteine protein (P36) | -2.55 | 0.73 | PF3D7_0404400 6-cysteine protein (P36) | adhesin | 1 1 | 2 different in liver stage | null | null | null | null | null | null |
| PBANKA_100260 6-cysteine protein (P41) | | | PF3D7_0404900 6-cysteine protein (P41) | adhesin | 1 1 | 2 not different from wild type | null | null | null | null | null | null |
| PBANKA_145330 acyl-CoA synthetase, putative | -2.63 | 0.75 | PF3D7_1238800 acyl-CoA synthetase, PfACS11 (ACS11) | lipid metabolism | 1 1 | 2 na | catalytic activity | metabolic process | null | | | |
| PBANKA_122560 alpha/beta hydrolase, putative | -1.96 | 0.73 | PF3D7_0805000 alpha/beta hydrolase, putative | chaperone | 1 1 | 2 na | null | null | null | null | null | apicoplast |
| PBANKA_121690 ATP-dependent RNA helicase, putative | | | PF3D7_0321600 ATP-dependent RNA helicase, putative | mRNA | 1 1 | 2 na | ATP binding, ATP-dependent helicase activity, helicase activity, nucleic acid binding | g null | null | | | |
| PBANKA_041570 autophagy-related protein 3, putative (ATG3) | -1.86 | 0.72 | PF3D7_0905700.1/2/3 autophagy-related protein 3, putative (ATG3) | autophagy | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_090040 BIR protein | | | | BIR protein | 1 1 | 2 na | null | null | null | na | na | na |
| PBANKA_146520 BIR protein | | | | BIR protein | 1 1 | 2 na | null | null | null | | | |
| PBANKA_021500 BIR protein, pseudogene | | | | BIR protein | 1 1 | 2 na | null | null | null | na | na | na |
| PBANKA_060620 blood stage antigen 41-3 precursor, putative | -5.88 | 0.75 | PF3D7_1207700 blood stage antigen 41-3 precursor | | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_141110 branched-chain alpha keto-acid dehydrogenase, putative | | | PF3D7_1312600 branched-chain alpha keto-acid dehydrogenase, put | tative TCA | 1 1 | 2 na | oxidoreductase activity, acting on the aldehyde or oxo group of donors, disulfide as acceptor | metabolic process | null | | | |
| PBANKA_081590 BSD domain, putative | | | PF3D7_0914900 BSD domain, putative | transcription | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_102440 calmodulin, putative | -2.14 | 0.75 | PF3D7_1418300 calmodulin, putative | Ca2+ binding | 1 1 | 2 na | null | null | null | calcium ion binding | null | null |
| PBANKA_113320 cdc2-related kinase 2 (CRK2) | | | PF3D7_1356900 protein kinase 5 (PK5) | kinase | 1 1 | 2 different in liver stage | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | y protein phosphorylation | null | | | |
| PBANKA_124400 cell cycle regulator protein, putative | | | PF3D7_0529500 cell cycle regulator protein, putative | RNA binding | 1 1 | 2 na | RNA binding | null | null | | | |
| PBANKA_112190 chorismate synthase, putative (CS) | | | PF3D7_0623000 chorismate synthase (CS) | shikimate | 1 1 | 2 na | chorismate synthase activity | aromatic amino acid family biosynthetic process | null | | | |
| PBANKA_041290 circumsporozoite- and TRAP-related protein (CTRP) | | | PF3D7_0315200 circumsporozoite- and TRAP-related protein (CTI | RP) adhesin | 1 1 | different in fertilization, ookinete and oocy | st null | null | null | null | null | null |
| PBANKA_123520 conserved Plasmodium protein, unknown function | | | PF3D7_0520400 conserved Plasmodium protein, unknown funct | tion unknown function | 1 1 | 2 na | DNA binding | regulation of transcription, DNA-dependent | null | | | |
| PBANKA_030580 conserved Plasmodium protein, unknown function | | | PF3D7_0208700.1/2 conserved Plasmodium protein, unknown funct | tion unknown function | 1 1 | 2 na | DNA-directed RNA polymerase activity | transcription, DNA-dependent | null | | | |
| PBANKA_020360 conserved Plasmodium protein, unknown function | -6.26 | 0.75 | PF3D7_0110000 conserved Plasmodium protein, unknown funct | tion unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_030160 conserved Plasmodium protein, unknown function | -5.31 | 0.75 | PF3D7_0203800 conserved Plasmodium protein, unknown funct | tion phototransduction pathway | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_030660 conserved Plasmodium protein, unknown function | -2.39 | 0.74 | PF3D7_0209500 conserved Plasmodium protein, unknown funct | tion unknown function | 1 1 | 2 na | null | null | null | GTP binding, GTPase activity, translation initiation factor activity | translation, translational initiation | cytoplasm, intracellular |
| PBANKA_031070 conserved Plasmodium protein, unknown function | | | PF3D7_0213800 conserved Plasmodium protein, unknown funct | tion unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_031210 conserved Plasmodium protein, unknown function | | | PF3D7_0215400 conserved Plasmodium protein, unknown funct | tion unknown function | 1 1 | 2 na | null | null | null | null | null | membrane |
| PBANKA_031480 conserved Plasmodium protein, unknown function | | | PF3D7_0218100 conserved Plasmodium membrane protein, unknown fur | nction unknown function | 1 1 | 2 na | null | null | null | hydrolase activity, metal ion binding, nucleotide binding | transport | integral to membrane |
| PBANKA_031490 conserved Plasmodium protein, unknown function | | | PF3D7_0218200 conserved Plasmodium protein, unknown funct | tion unknown function | 1 1 | 2 na | null | null | null | rRNA (guanine-N1-)-methyltransferase activity | ribosomal large subunit assembly | apicoplast |
| PBANKA_040250 conserved Plasmodium protein, unknown function | -4.21 | 0.75 | PF3D7_0303900 conserved Plasmodium protein, unknown funct | tion unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_041720 conserved Plasmodium protein, unknown function | -3.16 | 0.75 | PF3D7_0904200 conserved Plasmodium protein, unknown funct | tion unknown function | 1 1 | 2 na | null | null | null | null | null | apicoplast, membrane |
| PBANKA_050520 conserved Plasmodium protein, unknown function | | | PF3D7_1021000 conserved Plasmodium protein, unknown funct | tion unknown function | 1 1 | 2 na | null | null | null | glutaminyl-tRNA synthase (glutamine-hydrolyzing) activity | translation | null |
| PBANKA_050610 conserved Plasmodium protein, unknown function | | | | proteasome | 1 1 | 2 na | null | null | null | na | na | na |
| PBANKA_050750 conserved Plasmodium protein, unknown function | -1.77 | 0.7 | PF3D7_1023300 conserved Plasmodium protein, unknown funct | tion unknown function | 1 1 | 2 na | null | null | null | null | null | membrane |

| PBANKA_051300 | conserved Plasmodium protein, unknown function | | | PF3D7_1028900 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | membrane |
|---------------|--|-------|------|---------------|--|-------|---|------|------|------|---|--|--|
| PBANKA_052110 | conserved Plasmodium protein, unknown function | | | PF3D7_1038200 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_060330 | conserved Plasmodium protein, unknown function | | | PF3D7_1204400 | conserved Plasmodium membrane protein, unknown function unknown function | 1 1 2 | na | null | null | null | na | na | na |
| PBANKA_061240 | conserved Plasmodium protein, unknown function | -4.07 | 0.75 | PF3D7_0411000 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | apicoplast, integral to membrane, membrane |
| PBANKA_061780 | conserved Plasmodium protein, unknown function | -2.09 | 0.75 | PF3D7_0720300 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_062030 | conserved Plasmodium protein, unknown function | -4.34 | 0.75 | PF3D7_0722800 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_062040 | conserved Plasmodium protein, unknown function | -3.41 | 0.75 | PF3D7_0722900 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_070330 | conserved Plasmodium protein, unknown function | -3.25 | 0.75 | PF3D7_0827400 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | cytoplasm |
| PBANKA_070490 | conserved Plasmodium protein, unknown function | | | PF3D7_0825700 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_071220 | conserved Plasmodium protein, unknown function | -2.64 | 0.75 | PF3D7_0818600 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | different in oocyst, sporozoite and liver stage | null | null | null | hydrolase activity | null | membrane |
| PBANKA_072090 | conserved Plasmodium protein, unknown function | -4.34 | 0.75 | PF3D7_0418800 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | cytoplasm, membrane |
| PBANKA_080290 | conserved Plasmodium protein, unknown function | | | PF3D7_0705200 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_080380 | conserved Plasmodium protein, unknown function | | | PF3D7_0706200 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_081330 | conserved Plasmodium protein, unknown function | -3.35 | 0.75 | PF3D7_0912200 | conserved Plasmodium membrane protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | integral to membrane, membrane |
| PBANKA_081540 | conserved Plasmodium protein, unknown function | -5.12 | 0.75 | PF3D7_0914400 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | nucleotide binding | null | cytoplasm |
| PBANKA_082120 | conserved Plasmodium protein, unknown function | | | PF3D7_0920300 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | cytoplasm |
| PBANKA_082320 | conserved Plasmodium protein, unknown function | -3.1 | 0.73 | PF3D7_0922300 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_082590 | conserved Plasmodium protein, unknown function | | | PF3D7_0925100 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | integral to membrane, membrane |
| PBANKA_082740 | conserved Plasmodium protein, unknown function | -4.21 | 0.75 | PF3D7_0926600 | conserved Plasmodium membrane protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | integral to membrane, membrane |
| PBANKA_083280 | conserved Plasmodium protein, unknown function | | | PF3D7_0932000 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | integral to membrane, membrane |
| PBANKA_083470 | conserved Plasmodium protein, unknown function | | | PF3D7_0933900 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | AT looks \$50 detail (TO) privaces aline, SELENNE, aparts speen superplaces aline, responses to looking, excluding scalaring periods aline, resources a looking scalaring periods aline, resources a looking scalaring scalaring periods aline, resources a looking scalaring periods aline, resources a looking scalaring periods aline, resources a looking scalaring periods aline, resources and resources are resourced as a looking scalaring period of the resources and resources are resourced as a looking scalaring periods are resourced as a looking scalaring period of the resource and resources are resourced as a looking scalaring period of the resource and resources are resourced as a looking scalaring period of the resource and resource are resourced as a looking scalaring period of the resource and resource are resourced as a looking scalaring period of the resource and resource are resourced as a looking scalaring period of the resource and resourced as a looking scalaring period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced as a looking scalar period of the resource are resourced a | DNA integration, DNA recombination, proteolysis, transposition | cytoplasm, nucleus |
| PBANKA_090170 | conserved Plasmodium protein, unknown function | | | PF3D7_1147400 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | protein farnesyltransferase activity, zinc ion binding | protein farnesylation | protein farnesyltransferase complex |
| PBANKA_090300 | conserved Plasmodium protein, unknown function | | | PF3D7_1146100 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | membrane |
| PBANKA_090620 | conserved Plasmodium protein, unknown function | | | PF3D7_1142800 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_090780 | conserved Plasmodium protein, unknown function | -1.93 | 0.73 | PF3D7_1141200 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | transferase activity | null | cytoplasm, membrane |
| PBANKA_091320 | conserved Plasmodium protein, unknown function | | | PF3D7_1135300 | conserved Plasmodium membrane protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | integral to membrane |
| PBANKA_091670 | conserved Plasmodium protein, unknown function | -3.42 | 0.75 | PF3D7_1131500 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | ATP binding, hydrolase activity | null | cytoplasm |
| PBANKA_093260 | conserved Plasmodium protein, unknown function | | | PF3D7_1115100 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_100170 | conserved Plasmodium protein, unknown function | | | PF3D7_0404200 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_100190 | conserved Plasmodium protein, unknown function | | | | unknown function | 1 1 2 | na | null | null | null | null | null | membrane |
| PBANKA_102680 | conserved Plasmodium protein, unknown function | | | PF3D7_1415900 | conserved Plasmodium protein, unknown function nucleic acid binding | 1 1 2 | na | null | null | null | zinc ion binding | null | nucleus |
| | | | | | | | | | | | | | |

| PBANKA_103300 conserved Plasmodium protein, unknown function | | PF3D7_1409400 | conserved Plasmodium membrane protein, unknown function unknown function | 1 1 | 2 na | null | null | null | na | na | na |
|--|---------|-------------------|--|-----|------|------|------|------|--|--|---|
| PBANKA_103880 conserved Plasmodium protein, unknown function | | PF3D7_1403200 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | extracellular region, membrane |
| PBANKA_110410 conserved Plasmodium protein, unknown function | -8.06 0 | .75 PF3D7_0504500 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | attachment of GPI anchor to protein | cytoplasm |
| PBANKA_110600 conserved Plasmodium protein, unknown function | | PF3D7_0506400 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_110930 conserved Plasmodium protein, unknown function | -3.59 0 | .75 PF3D7_0509700 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | apicoplast |
| PBANKA_110990 conserved Plasmodium protein, unknown function | -2.15 0 | .73 PF3D7_0510300 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | cytoplasm |
| PBANKA_111020 conserved Plasmodium protein, unknown function | -4.6 0 | .75 PF3D7_0510700 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_111270 conserved Plasmodium protein, unknown function | | PF3D7_0513000 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_111500 conserved Plasmodium protein, unknown function | -3.83 0 | .75 PF3D7_0515400 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_112590 conserved Plasmodium protein, unknown function | -2.28 0 | .75 | unknown function | 1 1 | 2 na | null | null | null | na | na | na |
| PBANKA_113230 conserved Plasmodium protein, unknown function | | PF3D7_1356000 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | ATP binding, hydrolase activity | null | null |
| PBANKA_113580 conserved Plasmodium protein, unknown function | | PF3D7_1359500 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_113740 conserved Plasmodium protein, unknown function | | PF3D7_1361300 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | membrane |
| PBANKA_113860 conserved Plasmodium protein, unknown function | -4.57 0 | .73 PF3D7_1362600 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_114380 conserved Plasmodium protein, unknown function | -4.19 0 | .75 PF3D7_1367900 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | apicoplast, membrane |
| PBANKA_120450 conserved Plasmodium protein, unknown function | | PF3D7_1006300 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_120930 conserved Plasmodium protein, unknown function | | PF3D7_1010900 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | ATP binding, protein kinase activity | protein phosphorylation | Golgi apparatus, cytoplasm, cytoplasmic vesicle, membrane |
| PBANKA_121440 conserved Plasmodium protein, unknown function | -3.47 0 | .75 PF3D7_1016000 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | ATP binding, protein binding | null | cytoplasm, plasma membrane |
| PBANKA_122530 conserved Plasmodium protein, unknown function | -2.93 0 | .72 PF3D7_0805300 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | $N-acetylgalactosaminyl-proteoglycan\ 3-beta-glucuronosyltransferase\ activity,\ zinconstruction and the proteoglycan activity activities and the proteoglycan activities and the proteoglycan activities activity and the proteoglycan activities activity activities and the proteoglycan activities activity activities ac$ | on binding null | intracellular, plasma membrane |
| PBANKA_122550 conserved Plasmodium protein, unknown function | -2.7 0 | .73 PF3D7_0805100 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_122750 conserved Plasmodium protein, unknown function | | PF3D7_0802700 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_122990 conserved Plasmodium protein, unknown function | -2.38 0 | .72 PF3D7_0615200 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_123900 conserved Plasmodium protein, unknown function | | PF3D7_0524200 | conserved Plasmodium membrane protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | integral to membrane, membrane |
| PBANKA_124010 conserved Plasmodium protein, unknown function | -3.8 0 | .75 PF3D7_0525300 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_124240 conserved Plasmodium protein, unknown function | | PF3D7_0527600 | conserved Plasmodium protein, unknown function DNA replication | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_124260 conserved Plasmodium protein, unknown function | | PF3D7_0528000 | conserved Plasmodium protein, unknown function proteasome | 1 1 | 2 na | null | null | null | null | null | cytoplasm |
| PBANKA_124520 conserved Plasmodium protein, unknown function | -2.94 0 | .73 PF3D7_0530800 | CPW-WPC family protein adhesin | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_130980 conserved Plasmodium protein, unknown function | | PF3D7_1446100 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | cytoplasm |
| PBANKA_131130 conserved Plasmodium protein, unknown function | -2.11 0 | .75 PF3D7_1447500 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | cytoplasm |
| PBANKA_131140 conserved Plasmodium protein, unknown function | | PF3D7_1447600 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | null | null | null |
| PBANKA_131260 conserved Plasmodium protein, unknown function | | PF3D7_1448900 | conserved Plasmodium protein, unknown function unknown function | 1 1 | 2 na | null | null | null | DNA binding, metal ion binding | regulation of transcription, DNA-dependent, transcription, | DNA-dependent nucleus |
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|---------------|--|-------|------|-----------------|--|-------|------------------------------|------|------|------|--|--|--|
| PBANKA_131850 | conserved Plasmodium protein, unknown function | -5.23 | 0.75 | PF3D7_1454800 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_131860 | conserved Plasmodium protein, unknown function | -3.25 | 0.73 | PF3D7_1454900 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | not different from wild type | null | null | null | null | null | membrane |
| PBANKA_131900 | conserved Plasmodium protein, unknown function | | | PF3D7_1455300 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | KO not successful | null | null | null | DNA binding, flap endonuclease activity, single-stranded DNA specific 5'-3' exodeoxyribonuclease activ | only DNA recombination, mismatch repair, response to DNA damage stimulus | nucleus |
| PBANKA_132080 | conserved Plasmodium protein, unknown function | -5.48 | 0.75 | PF3D7_1457100 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | membrane |
| PBANKA_132180 | conserved Plasmodium protein, unknown function | | | PF3D7_1458100 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | nucleotide binding, transferase activity | null | membrane |
| PBANKA_132390 | conserved Plasmodium protein, unknown function | | | PF3D7_1460200 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | membrane |
| PBANKA_132540 | conserved Plasmodium protein, unknown function | | | PF3D7_1461700 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | ATP binding, protein binding, zinc ion binding | $intracellular\ protein\ transmembrane\ transport,\ protein\ import,\ protein\ target in$ | ng cytoplasm, plasma membrane |
| PBANKA_132940 | conserved Plasmodium protein, unknown function | -4.02 | 0.75 | PF3D7_1466000 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_132990 | conserved Plasmodium protein, unknown function | | | PF3D7_1466500 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_133400 | conserved Plasmodium protein, unknown function | | | PF3D7_1470800 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_133470 | conserved Plasmodium protein, unknown function | -3.7 | 0.75 | PF3D7_1471500 | conserved Plasmodium membrane protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | integral to membrane, membrane |
| PBANKA_133620 | conserved Plasmodium protein, unknown function | | | PF3D7_1473000 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | membrane |
| PBANKA_133810 | conserved Plasmodium protein, unknown function | | | PF3D7_1322900 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_134230 | conserved Plasmodium protein, unknown function | -3.45 | 0.75 | PF3D7_1327100 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | beta-galactosidase activity, carbohydrate binding, cation binding | carbohydrate metabolic process | beta-galactosidase complex, extracellular region |
| PBANKA_134370 | conserved Plasmodium protein, unknown function | | | PF3D7_1328600 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | cytoplasm |
| PBANKA_135380 | conserved Plasmodium protein, unknown function | -7.77 | 0.75 | PF3D7_1340400 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_135490 | conserved Plasmodium protein, unknown function | -4.62 | 0.75 | PF3D7_1341800 | conserved Plasmodium protein, unknown function gliding motility | 1 1 2 | na | null | null | null | nucleotide binding | null | cytoplasm |
| PBANKA_135610 | conserved Plasmodium protein, unknown function | -3.07 | 0.75 | PF3D7_1343100 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_136040 | conserved Plasmodium protein, unknown function | | | PF3D7_1347600 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_136140 | conserved Plasmodium protein, unknown function | | | PF3D7_1348600 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | cytoplasm |
| PBANKA_136460 | conserved Plasmodium protein, unknown function | | | PF3D7_1351900 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | ATP binding, hydrolase activity | null | integral to membrane |
| PBANKA_140090 | conserved Plasmodium protein, unknown function | | | PF3D7_1302400 | conserved Plasmodium membrane protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | endoplasmic reticulum, integral to membrane |
| PBANKA_141480 | conserved Plasmodium protein, unknown function | -3.46 | 0.75 | PF3D7_1316300 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | null |
| PBANKA_141520 | conserved Plasmodium protein, unknown function | -3.5 | 0.75 | PF3D7_1316700 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | nucleotide binding | null | null |
| PBANKA_141540 | conserved Plasmodium protein, unknown function | -2.34 | 0.73 | PF3D7_1316900 | conserved Plasmodium protein, unknown function chromatin | 1 1 2 | na | null | null | null | protein binding | DNA repair, regulation of transcription, DNA-dependent, response to DNA damage stimulus, transcription, DNA-depend | nucleus |
| PBANKA_141740 | conserved Plasmodium protein, unknown function | -5.81 | 0.75 | PF3D7_1318900 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | ATP binding | null | null |
| PBANKA_141930 | conserved Plasmodium protein, unknown function | -2.85 | 0.75 | PF3D7_1321000 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | transport | integral to membrane, plasma membrane |
| PBANKA_142200 | conserved Plasmodium protein, unknown function | -1.89 | 0.73 | PF3D7_0715700 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | DNA binding, nucleotidyltransferase activity | null | null |
| PBANKA_142290 | conserved Plasmodium protein, unknown function | -5.15 | 0.75 | PF3D7_0814600 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | ATP binding, transferase activity | null | null |
| PBANKA_142860 | conserved Plasmodium protein, unknown function | | | PF3D7_1212600.1 | conserved Plasmodium membrane protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | integral to membrane, membrane |
| PBANKA_143030 | conserved Plasmodium protein, unknown function | | | PF3D7_1214500 | conserved Plasmodium protein, unknown function unknown function | 1 1 2 | na | null | null | null | null | null | apicoplast |
| | | | | | | | | | | | | | |

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|---------------|---|-------|------|------------------|---|-----------------------------|---|---|---|----|---|--|--------|
| PBANKA_143070 | conserved Plasmodium protein, unknown function | | | PF3D7_1214900 | conserved Plasmodium membrane protein, unknown function | n unknown function | 1 | 1 | 2 | na | null | null | null |
| PBANKA_143500 | conserved Plasmodium protein, unknown function | | | PF3D7_1219800 | conserved Plasmodium protein, unknown function | unknown function | 1 | 1 | 2 | na | null | null | null |
| PBANKA_143630 | conserved Plasmodium protein, unknown function | -2.46 | 0.73 | PF3D7_1221100 | conserved Plasmodium protein, unknown function | unknown function | 1 | 1 | 2 | na | null | null | null |
| PBANKA_143640 | conserved Plasmodium protein, unknown function | | | PF3D7_1221200 | conserved Plasmodium protein, unknown function | unknown function | 1 | 1 | 2 | na | null | null | null |
| PBANKA_143770 | conserved Plasmodium protein, unknown function | | | PF3D7_1222800 | conserved Plasmodium protein, unknown function | unknown function | 1 | 1 | 2 | na | null | null | null |
| PBANKA_144420 | conserved Plasmodium protein, unknown function | | | PF3D7_1229600 | conserved Plasmodium protein, unknown function | unknown function | 1 | 1 | 2 | na | null | null | null |
| PBANKA_144480 | conserved Plasmodium protein, unknown function | -4.42 | 0.75 | PF3D7_1230100 | conserved Plasmodium protein, unknown function | unknown function | 1 | 1 | 2 | na | null | null | null |
| PBANKA_144640 | conserved Plasmodium protein, unknown function | | | PF3D7_1231700 | conserved Plasmodium protein, unknown function | unknown function | 1 | 1 | 2 | na | null | null | null |
| PBANKA_145210 | conserved Plasmodium protein, unknown function | | | PF3D7_1237500 | conserved Plasmodium protein, unknown function | unknown function | 1 | 1 | 2 | na | null | null | null |
| PBANKA_145390 | conserved Plasmodium protein, unknown function | -1.76 | 0.7 | PF3D7_1239400 | conserved Plasmodium protein, unknown function | unknown function | 1 | 1 | 2 | na | null | null | null |
| PBANKA_146130 | conserved Plasmodium protein, unknown function | -5.22 | 0.75 | PF3D7_1248400 | conserved Plasmodium protein, unknown function | unknown function | 1 | 1 | 2 | na | null | null | null |
| PBANKA_146330 | conserved Plasmodium protein, unknown function | -5.38 | 0.75 | PF3D7_1250400 | conserved Plasmodium protein, unknown function | unknown function | 1 | 1 | 2 | na | null | null | null |
| PBANKA_100440 | conserved Plasmodium protein, unknown function | -1.75 | 0.73 | PF3D7_0406800 | conserved Plasmodium protein, unknown function | unknown function | 1 | 1 | 2 | na | null | translation | null |
| PBANKA_132380 | conserved Plasmodium protein, unknown function | -5.5 | 0.75 | | | unknown function | 1 | 1 | 2 | na | protein binding | null | null |
| PBANKA_052190 | conserved Plasmodium protein, unknown function | | | PF3D7_0420500 | conserved Plasmodium protein, unknown function | phosphatase | 1 | 1 | 2 | na | protein tyrosine/serine/threonine phosphatase activity | protein dephosphorylation | null |
| PBANKA_093040 | conserved Plasmodium protein, unknown function | | | PF3D7_1117600.1/ | 2 conserved Plasmodium protein, unknown function | translation | 1 | 1 | 2 | na | translation release factor activity | translational termination | null |
| PBANKA_101540 | CPW-WPC family protein | -2.61 | 0.73 | PF3D7_1429300 | CPW-WPC family protein | adhesin | 1 | 1 | 2 | na | null | null | null |
| PBANKA_112320 | CPW-WPC family protein | | | PF3D7_0624300 | CPW-WPC family protein | adhesin | 1 | 1 | 2 | na | null | null | null |
| PBANKA_135250 | CPW-WPC family protein | -3.01 | 0.75 | PF3D7_1338800 | CPW-WPC family protein | adhesin | 1 | 1 | 2 | na | null | null | null |
| PBANKA_134630 | CPW-WPC family protein (UIS19) | -3.77 | 0.73 | PF3D7_1331400 | CPW-WPC family protein | adhesin | 1 | 1 | 2 | na | null | null | null |
| PBANKA_094340 | CPW-WPC family protein, putative | -2.54 | 0.75 | PF3D7_1103500 | CPW-WPC family protein | adhesin | 1 | 1 | 2 | na | null | null | null |
| PBANKA_144930 | CPW-WPC family protein, putative | -2.93 | 0.74 | PF3D7_1234700 | CPW-WPC family protein | adhesin | 1 | 1 | 2 | na | null | null | null |
| PBANKA_132170 | cysteine proteinase, putative | | | PF3D7_1458000 | cysteine proteinase falcipain 1 | protease | 1 | 1 | 2 | na | cysteine-type peptidase activity | proteolysis | null |
| PBANKA_081910 | cytochrome b5, putative | | | PF3D7_0918100 | cytochrome b5-like heme/steroid binding protein, putative | utilization phospholipids | 1 | 1 | 2 | na | heme binding | null | null |
| PBANKA_141020 | cytochrome c2 precursor,putative | | | PF3D7_1311700 | cytochrome c2 precursor, putative | redoxmeth | 1 | 1 | 2 | na | electron carrier activity, heme binding, iron ion binding | null | null |
| PBANKA_051200 | DHHC-type zinc finger protein, putative | -3.85 | 0.75 | PF3D7_1027900 | DHHC-type zinc finger protein, putative | palmytoylation; prenylation | 1 | 1 | 2 | na | zinc ion binding | null | null |
| PBANKA_133460 | diacylglycerol kinase, putative | | | PF3D7_1471400 | diacylglycerol kinase, putative | utilization phospholipids | 1 | 1 | 2 | na | diacylglycerol kinase activity | protein kinase C-activating G-protein coupled receptor signaling pathway | y null |
| PBANKA_080620 | e3 ubiquitin-protein ligase rbx1, putative (RBX1) | | | PF3D7_0319100 | ubiquitin-protein ligase, putative | proteasome | 1 | 1 | 2 | na | protein binding, zinc ion binding | null | null |
| PBANKA_122950 | endonuclease III homologue, putative | | | PF3D7_0614800 | endonuclease III homologue, putative | DNA repair | 1 | 1 | 2 | na | 4 iron, 4 sulfur cluster binding, endonuclease activity | base-excision repair | null |
| PBANKA_124280 | eukaryotic translation initiation factor 3, subunit 6, putative | | | PF3D7_0528200 | eukaryotic translation initiation factor 3, subunit 6, putative | translation | 1 | 1 | 2 | na | null | null | null |
| PBANKA_062250 | exonuclease I, putative | | | PF3D7_0725000 | exonuclease I, putative | DNA repair | 1 | 1 | 2 | na | DNA binding, nuclease activity | DNA repair | null |
| | | | | | | | | | | | | | |

| te dehydrogenase activity, binding | glycolysis, malate metabolic process, tricarboxylic acid cycle | integral to membrane |
|---|--|---------------------------------------|
| | null | integral to membrane, plasma membrane |
| | null | null |
| ATPase activity, coupled to transmembrane movement of substances, chloride channel activity, chloride ion binding | transport | integral to membrane |
| | null | cytoplasm |
| binding | null | apicoplast, cytoplasm |
| | null | null |
| | null | null |
| | null | apicoplast, cytoplasm, membrane |
| nding, ligase activity | translation | cytoplasm |
| | null | apicoplast |
| nding, actin binding, calmodulin binding, motor activity | null | cytoplasm, myosin complex |
| | | |
| | | |
| | | |
| | | |
| | null | null |
| | null | membrane |
| | null | null |
| | null | apicoplast |
| | null | apicoplast |

null

| PBANKA_052390 glideosome associated protein with multiple membrane spans 2, putative (GAPM2) PF3D7_0423500 glideosome associated | stein with multiple membrane spans 2 (GAPM2) gliding motility 1 1 2 na | null | null | null | metal ion binding, protein binding | transport | actomyosin, actin part, inner membrane complex, integral to membrane, membrane |
|--|--|---|--|---------|------------------------------------|--|--|
| PBANKA_103540 glideosome associated protein with multiple membrane spans 3, putative -2.56 0.7 PF3D7_1406800 glideosome associated | stein with multiple membrane spans 3 (GAPM3) gliding motility 1 1 2 na | null | null | null | null | null | inner membrane complex, integral to membrane, membrane |
| PBANKA_111530 glideosome-associated protein 40, putative (GAP40) -4.69 0.75 PF3D7_0515700 glideosome-associated protein 40, putative (GAP40) | ated protein 40, putative (GAP40) gliding motility 1 1 2 na | null | null | null | null | null | null |
| PBANKA_143760 glideosome-associated protein 45, putative -4.44 0.74 PF3D7_1222700 glideosome-associated protein 45, putative | ciated protein 45 (GAP45) gliding motility 1 1 2 different in fertilization and | ookinete null | null | null | null | null | extrinsic to membrane, host cell membrane, inner membrane complex |
| PBANKA_145730 glucose inhibited division protein a homologue, putative PF3D7_1244000 glucose inhibited | vision protein a homologue, putative translation 1 1 2 na | flavin adenine dinucleotide binding | tRNA processing | null | | | |
| PBANKA_136430 glycerol kinase, putative -2.86 0.73 PF3D7_1351600 glycerol kinase | SK) glycolysis 1 1 2 KO not successful | glycerol kinase activity | carbohydrate metabolic process, glycerol-3-phosphate metabolic process | ss null | | | |
| PBANKA_136090 GTPase, putative PF3D7_1348100 Rab14 GTPase | outative trafficking 1 1 2 na | GTP binding | protein transport, small GTPase mediated signal transduction | on null | | | |
| PBANKA_102520 h/aca ribonucleoprotein complex subunit 4, putative (CBF5) PF3D7_1417500 pseudouridine | nthase, putative RNA 1 1 2 na | RNA binding, pseudouridine synthase activity | RNA processing, pseudouridine synthesis | null | | | |
| PBANKA_110190 HCNGP-like protein, putative PF3D7_0502100 HCNGP-like p | ein transcription 1 1 2 na | null | null | null | null | null | null |
| PBANKA_031000 heat shock 40 kDa protein, putative PF3D7_0213100 heat shock pro | n 40, putative chaperone 1 1 2 na | heat shock protein binding, unfolded protein binding | protein folding | null | | | |
| PBANKA_021480 heat shock protein, putative -3.38 0.75 | chaperone 1 1 2 KO not successful | heat shock protein binding, unfolded protein binding | protein folding | null | | | |
| PBANKA_120200 inner membrane complex protein 1c, putative (IMC1C, ALV5) -4.82 0.75 PF3D7_1003600 membrane ske | al protein IMC1-related (ALV5) gliding motility 1 1 2 na | null | null | null | null | cytoskeleton organization | inner membrane complex, subpellicular network |
| PBANKA_121910 inner membrane complex protein 1d, putative (IMC1D) -10.18 0.75 PF3D7_0708600 heat shock pro | n 86 family protein gliding motility 1 1 2 na | null | null | null | null | response to heat, response to unfolded protein | null |
| PBANKA_040270 inner membrane complex protein 1e, putative (IMC1E, ALV2) -6.85 0.75 PF3D7_0304100 membrane ske | al protein IMC1-related (ALV2) gliding motility 1 1 2 na | null | null | null | null | null | inner membrane complex |
| PBANKA_041760 LCCL domain-containing protein (CCp4) -5.57 0.75 PF3D7_0903800 LCCL domain- | staining protein (CCp4) adhesin 1 2 offerent in occypt and sportcoite; (tagged) different in fe | erifization and colonete null | null | null | null | null | null |
| PBANKA_131950 LCCL domain-containing protein CCP2 (CCP2) -7.76 0.75 PF3D7_1455800 LCCL domain- | ntaining protein (CCp2) adhesin 1 1 2 different in fertilization, ookinete, oocyst | and sporozoite receptor binding | cell adhesion, signal transduction | null | | | |
| PBANKA_111770 malate dehydrogenase, putative (MDH) PF3D7_0618500 malate dehydrogenase | enase (MDH) TCA 1 1 2 na | binding, oxidoreductase activity, acting on the CH-OH group of donors, NAD or NADP as acceptance. | optor glycolysis | null | | | |
| PBANKA_070710 membrane skeletal protein, putative -5.38 0.75 PF3D7_0823500 membrane skeletal | al protein IMC1-related gliding motility 1 1 2 na | null | null | null | null | null | null |
| PBANKA_081570 membrane transporter, putative -4.44 0.75 PF3D7_0914700 transporter, put | ve transporter 1 1 2 na | null | null | null | null | null | null |
| PBANKA_124210 methionine aminopeptidase, putative -1.99 0.74 PF3D7_0527300 methionine am | peptidase 1a, putative (MetAP1a) post-translational modification 1 1 2 na | aminopeptidase activity, metalloexopeptidase activity | cellular process, proteolysis | null | | | |
| PBANKA_082600 methyltransferase, putative PF3D7_0925200 rRNA processing an | omere maintaining methyltransferase, putative RNA 1 1 2 na | null | null | null | null | null | null |
| PBANKA_092110 methyltransferase, putative -1.61 0.7 PF3D7_1127300 methyltransfer | , putative RNA 1 1 2 na | tRNA (guanine-N7-)-methyltransferase activity | tRNA modification | null | | | |
| PBANKA_051520 MORN repeat-containing protein 1, putative (MORN1) PF3D7_1031200 MORN repeat | ntaining protein 1 (MORN1) lipid metabolism 1 1 2 na | null | null | null | null | null | null |
| PBANKA_141430 Myb1 protein, putative PF3D7_1315800 Myb1-related | scription factor (MYB1) transcription 1 1 2 na | DNA binding | null | null | | | |
| PBANKA_145950 myosin light chain 1, putative,myosin A tail domain interacting protein M -3.21 0.74 PF3D7_1246400 myosin light chain 1 | sin A tail domain interacting protein (MTIP) gliding motility 1 1 2 (mutated) different in fertilization a | and ookinete null | null | null | null | null | inner membrane complex |
| PBANKA_114340 NADH-cytochrome b5 reductase, putative PF3D7_1367500 NADH-cytochrome b5 reductase, putative | e b5 reductase, putative fatty acids synth 1 1 2 na | electron carrier activity, oxidoreductase activity | null | null | | | |
| PBANKA_124110 nucleolar Jumonji domain interacting protein, putative -3.72 0.75 PF3D7_0526300 nucleolar Jumonji | domain interacting protein, putative ribosome 1 1 2 na | null | null | null | null | null | null |
| PBANKA_090310 nucleolar preribosomal assembly protein, putative -2.02 0.74 PF3D7_1146000 nucleolar prer | somal assembly protein, putative ribosome assembly 1 1 2 na | null | null | null | null | null | null |
| PBANKA_103470 nucleolar preribosomal GTPase, putative PF3D7_1407500 nucleolar prer | somal GTPase, putative ribosome assembly 1 1 2 na | null | null | null | null | null | null |
| PBANKA_103340 oxidoreductase, putative PF3D7_1409100 aldo-keto redu | se, putative redoxmeth 1 1 2 na | oxidoreductase activity | oxidation-reduction process | null | | | |
| PBANKA_020060 Pb-fam-1 protein | Pb-fam-1 protein 1 1 2 na | null | null | null | na | na | na |
| | | | | | | | |

| PBANKA_021490 | Pb-fam-1 protein | | | | | Pb-fam-1 protein | 1 1 : | ! na | null | null | null | na | na | na |
|---------------|---|-----------|------|-----------------|--|---------------------------------|-------|---|---|------------------------------------|------|--|--|--|
| PBANKA_110090 | Pb-fam-1 protein | -1.64 | 0.71 | | | Pb-fam-1 protein | 1 1 : | ! na | null | null | null | | | |
| PBANKA_093220 | peptidyl-prolyl cis-trans isomerase, putative | | | PF3D7_1115600 | peptidyl-prolyl cis-trans isomerase (CYP19B) | chaperone | 1 1 : | not different from wild type | peptidyl-prolyl cis-trans isomerase activity | null | null | | | |
| PBANKA_124330 | peptidyl-prolyl cis-trans isomerase, putative (CYP23) | | | PF3D7_0528700 | peptidyl-prolyl cis-trans isomerase (CYP23) | chaperone | 1 1 : | ! na | peptidyl-prolyl cis-trans isomerase activity | null | null | | | |
| PBANKA_112040 | Pfs77 homologue, putative | -4.93 | 0.75 | PF3D7_0621400 | Pf77 protein (ALV7) | gliding motility | 1 1 : | ! na | null | null | null | null | null | inner membrane complex |
| PBANKA_050270 | PHF5-like protein, putative | | | PF3D7_1018500 | PHF5-like protein, putative | mRNA | 1 1 : | ! na | null | null | null | null | null | null |
| PBANKA_143490 | phosphatidylethanolamine-binding protein, putative | | | PF3D7_1219700 | raf kinase inhibitor (RKIP) | | 1 1 : | ! na | null | null | null | phosphatidylethanolamine binding, protein kinase regulator activ | ity null | null |
| PBANKA_133370 | phosphodiesterase delta (PDEdelta) | -2.25 | 0.74 | PF3D7_1470500 | phosphodiesterase delta, putative (PDEdelta) | purine metabolism | 1 1 : | different in fertilization, ookinete and oocy | yst 3',5'-cyclic-nucleotide phosphodiesterase activity | signal transduction | null | | | |
| PBANKA_135590 | phosphoenolpyruvate carboxykinase, putative (PEPCK) | -2.49 | 0.71 | PF3D7_1342800 | phosphoenolpyruvate carboxykinase (PEPCK) | TCA | 1 1 : | ! na | ATP binding, phosphoenolpyruvate carboxykinase (ATP) activity | ity gluconeogenesis | null | | | |
| PBANKA_050690 | phospholipid scramblase, putative | | | PF3D7_1022700 | phospholipid scramblase 1, putative | transporter | 1 1 : | ! na | null | null | null | null | null | null |
| PBANKA_040970 | plasmepsin VI | -2.73 | 0.73 | PF3D7_0311700 | plasmepsin VI | plasmepsin, protease | 1 1 : | different in oocyst and sporozoite | aspartic-type endopeptidase activity | proteolysis | null | | | |
| PBANKA_051760 | plasmepsin VII, putative | -5.22 | 0.75 | PF3D7_1033800 | plasmepsin VII | plasmepsin, protease | 1 1 : | ! na | aspartic-type endopeptidase activity | proteolysis | null | | | |
| PBANKA_132910 | plasmepsin VIII, putative | -3.59 | 0.75 | PF3D7_1465700 | plasmepsin VIII | plasmepsin, protease | 1 1 : | ! na | aspartic-type endopeptidase activity | proteolysis | null | | | |
| PBANKA_020040 | Plasmodium exported protein, unknown function | | | | | unknown function | 1 1 : | ! na | null | null | null | na | na | na |
| PBANKA_072260 | Plasmodium exported protein, unknown function | -3.76 | 0.75 | | | unknown function | 1 1 : | ! na | null | null | null | | | |
| PBANKA_146510 | Plasmodium exported protein, unknown function | | | | | unknown function | 1 1 : | ! na | null | null | null | | | |
| PBANKA_120510 | PPPDE peptidase, putative | -6.22 | 0.75 | PF3D7_1006900 | PPPDE peptidase, putative | protease | 1 1 : | ! na | null | null | null | null | null | null |
| PBANKA_051410 | pre-mRNA-splicing factor ATP-dependent RNA helicase PRP22, putative | e (PRP22) | | PF3D7_1030100 | pre-mRNA-splicing factor ATP-dependent RNA helicase PRP22, putative (PRF | P22) mRNA | 1 1 : | ! na | ATP binding, RNA binding, helicase activity | null | null | | | |
| PBANKA_083070 | procollagen lysine 5-dioxygenase, putative | -3.61 | 0.75 | PF3D7_0930000 | procollagen lysine 5-dioxygenase, putative | post-translational modification | 1 1 : | ! na | null | null | null | null | null | null |
| PBANKA_141450 | protein kinase, putative | -3.93 | 0.75 | PF3D7_1316000 | protein kinase, putative | kinase | 1 1 : | KO not successful | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | y protein phosphorylation | null | | | |
| PBANKA_131880 | protein phosphatase 7, putative | -2.1 | 0.75 | PF3D7_1455100 | protein phosphatase, putative | phosphatase | 1 1 : | na na | protein tyrosine/serine/threonine phosphatase activity | protein dephosphorylation | null | | | |
| PBANKA_142700 | protein phosphatase, putative | | | PF3D7_0810500 | protein phosphatase, putative | phosphatase | 1 1 : | na na | catalytic activity | null | null | | | |
| PBANKA_031100 | protein transport protein Sec31, putative (SEC31) | | | PF3D7_0214100 | protein transport protein sec31 (SEC31) | trafficking | 1 1 : | ! na | null | null | null | null | ER to Golgi vesicle-mediated transport, translocation of peptides or proteins into host cell cytop | asm COPII vesicle coat, ER to Golgi transport vesicle, Maurer's cleft, host cell cytoplasm |
| PBANKA_070900 | protein-transport protein sec61 beta 1 subunit, putative | | | PF3D7_0821800 | secretory complex protein 61 beta subunit (Sec61-be | eta) trafficking | 1 1 : | ! na | null | null | null | protein binding, protein transporter activity | intracellular protein transport, protein transport by the Sec comp | lex Sec61 translocon complex, endoplasmic reticulum membrane |
| PBANKA_146000 | rac-beta serine/threonine protein kinase, putative (PKB) | | | PF3D7_1246900 | RAC-beta serine/threonine protein kinase (PKB | 3) kinase | 1 1 : | KO not successful | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | y protein phosphorylation | null | | | |
| PBANKA_093950 | Rad51 homolog, putative | -7.81 | 0.75 | PF3D7_1107400 | Rad51 homolog (RAD51) | DNA repair | 1 1 : | na na | ATP binding, DNA-dependent ATPase activity, damaged DNA binding, protein binding | g DNA repair | null | | | |
| PBANKA_051380 | RAP protein, putative | | | PF3D7_1029800 | RAP protein, putative | mRNA | 1 1 : | na na | null | null | null | null | null | null |
| PBANKA_140730 | recombinase, putative | | | PF3D7_1308800 | recombinase, putative | | 1 1 : | ! na | DNA binding | DNA integration, DNA recombination | null | | | |
| PBANKA_123010 | ribonuclease, putative, fragment | | | | | | 1 1 : | na na | null | null | null | RNA binding, ribonuclease activity | RNA processing | null |
| PBANKA_031580 | ribosome associated membrane protein RAMP4, putative | | | PF3D7_0219400.1 | /2 ribosome associated membrane protein RAMP4, putati | tive translation | 1 1 : | ! na | null | null | null | null | null | membrane |
| PBANKA_112140 | RNA methyltransferase, putative | | | PF3D7_0622500 | RNA methyltransferase, putative | RNA | 1 1 : | na na | methyltransferase activity | rRNA processing | null | | | |
| | | | | | | | | | | | | | | |

| | null | inner membrane complex null |
|---|--|-----------------------------|
| ohatidylethanolamine binding, protein kinase regulator activity | | null |
| | | |
| | na | na |
| | | null |
| | | |
| | ER to Golgi vesicle-mediated transport, translocation of peptides or proteins into host cell cytoplasm intracellular protein transport, protein transport by the Sec complex | |
| | null | null |
| binding, ribonuclease activity | | null membrane |

| PBANKA_051470 | RNA methyltransferase, putative | | | PF3D7_1030700 | RNA methyltransferase, putative | RNA | 1 | 1 | 2 | na | RNA binding, RNA methyltransferase activity | RNA processing | null |
|---------------|--|-------|------|-------------------|--|----------------------|---|---|---|------------------------------------|---|--|------|
| PBANKA_142950 | RNA polymerase subunit 8c, putative | | | PF3D7_1213700 | RNA polymerase subunit 8c, putative | mRNA | 1 | 1 | 2 | na | null | transcription, DNA-dependent | null |
| PBANKA_071790 | RNA-binding protein, putative | | | PF3D7_0416000 | RNA binding protein, putative | mRNA | 1 | 1 | 2 | na | nucleic acid binding | null | null |
| PBANKA_120080 | RNA-binding protein, putative | -1.66 | 0.7 | PF3D7_1002400.1/2 | RNA binding protein, putative | mRNA | 1 | 1 | 2 | na | nucleic acid binding | null | null |
| PBANKA_070750 | RWD domain-containing protein, putative | | | PF3D7_0823100 | RWD domain-containing protein, putative | proteasome | 1 | 1 | 2 | na | null | null | null |
| PBANKA_082310 | S-adenosylmethionine synthetase, putative | -4.07 | 0.75 | PF3D7_0922200 | S-adenosylmethionine synthetase (SAMS) | | 1 | 1 | 2 | na | ATP binding, methionine adenosyltransferase activity | one-carbon metabolic process | null |
| PBANKA_123360 | secreted ookinete protein, putative (PSOP13) | | | PF3D7_0518800 | secreted ookinete protein, putative (PSOP13) | adhesin | 1 | 1 | 2 | different in oocyst and sporozoite | null | null | null |
| PBANKA_142170 | secreted ookinete protein, putative (PSOP20) | -5.61 | 0.75 | PF3D7_0715400 | secreted ookinete protein, putative (PSOP20) | adhesin | 1 | 1 | 2 | not different from wild type | null | null | null |
| PBANKA_112900 | secreted ookinete protein, putative (PSOP6) | -2.92 | 0.74 | PF3D7_0630200 | secreted ookinete protein, putative (PSOP6) | adhesin | 1 | 1 | 2 | KO not successful | null | null | null |
| PBANKA_102700 | serine C-palmitoyltransferase, putative | -3.3 | 0.75 | PF3D7_1415700 | serine C-palmitoyltransferase, putative | lipid metabolism | 1 | 1 | 2 | na | pyridoxal phosphate binding, transferase activity, transferring nitrogenous groups | biosynthetic process | null |
| PBANKA_030500 | serine repeat antigen 2 (SERA2) | | | | | | 1 | 1 | 2 | not different from wild type | cysteine-type peptidase activity | proteolysis | null |
| PBANKA_031140 | serine/threonine protein kinase, putative | -4.17 | 0.75 | PF3D7_0214600 | serine/threonine protein kinase, putative | kinase | 1 | 1 | 2 | KO not successful | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| PBANKA_082850 | serine/threonine protein phosphatase, putative | | | PF3D7_0927700 | serine/threonine protein phosphatase, putative | phosphatase | 1 | 1 | 2 | na | hydrolase activity | null | null |
| PBANKA_145310 | sphingomyelin phosphodiesterase, putative | -5.54 | 0.75 | PF3D7_1238600 | sphingomyelin phosphodiesterase, putative | lipid metabolism | 1 | 1 | 2 | na | null | null | null |
| PBANKA_112400 | sphingomyelin synthase 2, putative (SMS2) | -7.33 | 0.75 | PF3D7_0625100 | sphingomyelin synthase, putative | lipid metabolism | 1 | 1 | 2 | na | null | null | null |
| PBANKA_081070 | subpellicular microtubule protein 1, putative (SPM1) | | | PF3D7_0909500 | subpellicular microtubule protein 1, putative (SPM1) | gliding motility | 1 | 1 | 2 | na | null | null | null |
| PBANKA_082570 | telomeric repeat binding factor 1, putative | | | PF3D7_0924800 | telomeric repeat binding factor 1, putative (TRF1) | telomerase | 1 | 1 | 2 | na | DNA binding | null | null |
| PBANKA_132090 | thioredoxin, putative (TRX1) | | | PF3D7_1457200 | thioredoxin 1 (TRX1) | redoxmeth | 1 | 1 | 2 | na | electron carrier activity, protein disulfide oxidoreductase activity | cell redox homeostasis, glycerol ether metabolic process | null |
| PBANKA_131320 | transcription factor with AP2 domain(s), putative (ApiAP2) | -2.49 | 0.75 | PF3D7_1449500 | transcription factor with AP2 domain(s), putative (ApiAP2) | transcription | 1 | 1 | 2 | na | sequence-specific DNA binding transcription factor activity | regulation of transcription, DNA-dependent | null |
| PBANKA_134760 | translation initiation factor 6, putative | | | PF3D7_1332800 | translation initiation factor 6, putative | translation | 1 | 1 | 2 | na | ribosome binding | mature ribosome assembly | null |
| PBANKA_124460 | triose phosphate transporter, putative | | | PF3D7_0530200 | triose phosphate transporter (iTPT) | transporter | 1 | 1 | 2 | na | null | null | null |
| PBANKA_080650 | triosephophate isomerase, putative | | | PF3D7_0318800 | triosephophate isomerase, putative | glycolysis | 1 | 1 | 2 | na | triose-phosphate isomerase activity | metabolic process | null |
| PBANKA_103360 | tRNA-dihydrouridine synthase, putative | -1.87 | 0.73 | PF3D7_1408900 | tRNA-dihydrouridine synthase, putative | translation | 1 | 1 | 2 | na | flavin adenine dinucleotide binding, tRNA dihydrouridine synthase activity | oxidation-reduction process, tRNA processing | null |
| PBANKA_071200 | U3 small nucleolar ribonucleoprotein, putative | | | PF3D7_0818800 | U3 small nucleolar ribonucleoprotein protein, putative | mRNA | 1 | 1 | 2 | na | null | null | null |
| PBANKA_081610 | ubiquitin conjugating enzyme, putative | | | PF3D7_0915100 | ubiquitin conjugating enzyme, putative | proteasome | 1 | 1 | 2 | na | small conjugating protein ligase activity | post-translational protein modification, regulation of protein metabolic process | null |
| PBANKA_102970 | ubiquitin conjugating enzyme, putative | | | PF3D7_1412900.1 | ubiquitin conjugating enzyme, putative | proteasome | 1 | 1 | 2 | na | small conjugating protein ligase activity | post-translational protein modification, regulation of protein metabolic process | null |
| PBANKA_040420 | ubiquitin-conjugating enzyme e2, putative | | | PF3D7_0305700 | ubiquitin conjugating enzyme E2, putative | proteasome | 1 | 1 | 2 | na | small conjugating protein ligase activity | post-translational protein modification, regulation of protein metabolic process | null |
| PBANKA_122890 | von willebrand factor a-domain-related protein (WARP) | -5.64 | 0.75 | PF3D7_0801300 | von Willebrand factor A-domain related protein (WARP) | adhesin | 1 | 1 | 2 | not different from wild type | null | null | null |
| PBANKA_060860 | zinc finger protein, putative | -4.67 | 0.75 | PF3D7_1210200 | zinc finger protein, putative | nucleic acid binding | 1 | 1 | 2 | na | nucleic acid binding, zinc ion binding | null | null |
| PBANKA_113490 | zinc finger protein, putative | -6.57 | 0.75 | PF3D7_1358600 | zinc finger protein, putative | RNA binding | 1 | 1 | 2 | na | nucleic acid binding, zinc ion binding | null | null |
| PBANKA_143340 | zinc finger protein, putative | | | PF3D7_1217800 | zinc finger protein, putative | Ca2+ binding | 1 | 1 | 2 | na | null | null | null |
| | | | | | | | | | | | | | |

| null | null | null |
|---|-----------------|--|
| protein binding | null | nucleus |
| null | null | null |
| nucleotide binding | null | cytoplasm |
| nacconce smang | | Сугоризт |
| | | |
| | | |
| | | |
| null | null | null |
| null | null | null |
| null | null | null |
| | | |
| | | |
| | | |
| | | |
| $glucose\ 6-phosphate: phosphate\ antiporter\ activity,\ triose-phosphate: phosphate\ antiporter\ activity$ | glucose import | apicoplast, integral to plastid inner membrane, membrane |
| | | |
| | | |
| null | RNA processing | small nucleolar ribonucleoprotein complex |
| | | |
| | | |
| LBD domain binding, cell adhesion molecule binding, heparin binding | g call adhesion | microneme lumen |
| LED domain unumg, cen adhesion molecule unumg, nepami umum | g cen aunesion | This office fuller |
| | | |
| zinc ion binding | null | cytoplasm, nucleus |
| | | |

| PBANKA_14 | zinc finger protein, putative | | | PF3D7_1244800 | cytoplasmic translation machinery associated protein, putative | e translation | 1 | 1 2 | 2 na | à | null | null | null | null | null |
|-----------|---|-------|------|-------------------|---|-----------------------------|---|-----|--------------|--|--|--|--|------|------|
| PBANKA_01 | zinc finger protein, putative | -3.31 | 0.75 | PF3D7_0609800 | zinc finger protein, putative | nucleic acid binding | 1 | 1 2 | 2 na | a | zinc ion binding | null | null | | |
| PBANKA_09 | zinc finger, DHHC-type, putative | -2.88 | 0.75 | PF3D7_1121000 | zinc finger, DHHC-type, putative | palmytoylation; prenylation | 1 | 1 2 | 2 na | a | zinc ion binding | null | null | | |
| PBANKA_11 | conserved protein, unknown function | | | PF3D7_0626700 | conserved protein, unknown function | unknown function | 1 | 1 2 | 2 na | a | null | null | null | null | null |
| PBANKA_09 | conserved rodent malaria protein, unknown function | | | | | unknown function | 1 | 1 2 | 2 na | a | null | null | null | na | na |
| PBANKA_09 | folate transporter 2, putative (FT2) | -4.21 | 0.75 | PF3D7_1116500 | folate transporter 2 (FT2) | transporter | 1 | 1 2 | 2 na | a | folic acid transporter activity | null | plasma membrane | | |
| PBANKA_14 | ATP-dependent protease subunit ClpQ, putative (ClpQ) | | | PF3D7_1230400 | ATP-dependent protease subunit ClpQ (ClpQ) | protease | 1 | 1 2 | 2 na | a | threonine-type endopeptidase activity | proteolysis involved in cellular protein catabolic process | proteasome core complex | | |
| PBANKA_06 | ribonucleoside-diphosphate reductase, large subunit, putative | | | PF3D7_1437200 | ribonucleoside-diphosphate reductase, large subunit | t pyrimidine metabolism | 1 | 1 2 | 2 na | a | ATP binding, protein binding, ribonucleoside-diphosphate reductase activity, thioredoxin disuffide as acceptor | DNA replication, oxidation-reduction process | ribonucleoside-diphosphate reductase complex | | |
| PBANKA_14 | h/aca ribonucleoprotein complex subunit 1, putative (GAR1) | | | PF3D7_1309500 | small nucleolar ribonucleoprotein, putative | RNA | 1 | 1 2 | 2 na | a | rRNA binding | rRNA processing | small nucleolar ribonucleoprotein complex | | |
| PBANKA_10 | pre-mRNA-processing-splicing factor 8, putative (PRPF8) | | | PF3D7_0405400 | pre-mRNA-processing-splicing factor 8, putative (PRPF8 |) mRNA | 1 | 1 2 | 2 na | a | null | nuclear mRNA splicing, via spliceosome | spliceosomal complex | | |
| PBANKA_08 | profilin, putative (PFN) | | | PF3D7_0932200 | profilin, putative (PFN) | gliding motility | 1 | 0 1 | 1 (mı | utated) different in fertilization and ookinete | actin binding | cytoskeleton organization | actin cytoskeleton | | |
| PBANKA_07 | 690 conserved Plasmodium protein, unknown function | | | PF3D7_0414900 | conserved ARM repeats protein, unknown function | n centrosome, chromatin | 0 | 1 1 | 1 na | a | binding | null | anchored to membrane, rhoptry | | |
| PBANKA_11 | 790 triose phosphate transporter, putative | | | PF3D7_0508300 | triose phosphate transporter (oTPT) | transporter, apicoplast | 1 | 0 1 | 1 na | a | null | null | apicoplast, membrane | | |
| PBANKA_14 | conserved Plasmodium protein, unknown function | | | PF3D7_1217100.1/2 | conserved Plasmodium protein, unknown function | meiosis | 0 | 1 1 | 1 na | a | ATP binding, DNA binding, catalytic activity | DNA metabolic process | chromosome | | |
| PBANKA_10 | DNA topoisomerase II, putative | | | PF3D7_1433500 | DNA topoisomerase II, putative | DNA replication | 0 | 1 1 | 1 na | a | ATP binding, DNA topoisomerase (ATP-hydrolyzing) activity | DNA topological change | chromosome | | |
| PBANKA_13 | DNA topoisomerase III, putative | | | PF3D7_1347100 | DNA topoisomerase III, putative | DNA replication | 0 | 1 1 | 1 na | a | DNA topoisomerase type I activity | DNA topological change | chromosome | | |
| PBANKA_11 | 720 Sec24 subunit, putative | | | PF3D7_1361100 | Sec24 subunit a (SEC24a) | trafficking | 0 | 1 1 | 1 na | a | protein binding, zinc ion binding | ER to Golgi vesicle-mediated transport, intracellular protein transport | t COPII vesicle coat | | |
| PBANKA_10 | 280 Sec24-like protein, putative | | | PF3D7_0405100 | Sec24 subunit b (SEC24b) | trafficking | 1 | 0 1 | 1 na | a | protein binding | ER to Golgi vesicle-mediated transport, intracellular protein transport | t COPII vesicle coat | | |
| PBANKA_03 | peptide chain release factor subunit 1, putative | | | PF3D7_0212300 | peptide chain release factor subunit 1, putative | translation | 0 | 1 1 | 1 na | a | translation release factor activity, codon specific | translational termination | cytoplasm | | |
| PBANKA_11 | translationally controlled tumor protein homolog, putative | | | PF3D7_0511000 | translationally controlled tumor protein homolog, putative | e trafficking | 1 | 0 1 | 1 na | a | null | null | cytoplasm | | |
| PBANKA_06 | ubiquitin-conjugating enzyme e2, putative | | | PF3D7_1203900 | ubiquitin conjugating enzyme E2, putative (UBC) | proteasome | 1 | 0 1 | 1 na | a | ubiquitin-protein ligase activity | post-translational protein modification, regulation of protein metabolic process | cytoplasm | | |
| PBANKA_13 | 990 leucine aminopeptidase, putative | | | PF3D7_1446200 | M17 leucyl aminopeptidase (LAP) | protease | 0 | 1 1 | 1 na | a | aminopeptidase activity, manganese ion binding, metalloexopeptidase activity | protein metabolic process, proteolysis | cytoplasm, intracellular | | |
| PBANKA_12 | pre-mRNA-splicing factor, putative (SR1) | | | PF3D7_0517300 | pre-mRNA-splicing factor (SR1) | mRNA | 0 | 1 1 | 1 na | a | nucleic acid binding | RNA splicing | cytoplasm, nucleus | | |
| PBANKA_10 | RNA-binding protein, putative | | | PF3D7_1406000 | RNA binding protein, putative | mRNA | 0 | 1 1 | 1 na | a | RNA binding | RNA processing | cytoplasm, nucleus | | |
| PBANKA_13 | protein serine/threonine phosphatase, putative | | | PF3D7_1466100 | protein serine/threonine phosphatase | phosphatase | 1 | 0 1 | 1 diffe | erent in fertilization, ookinete, oocyst and sporozoite | iron ion binding, manganese ion binding, phosphoprotein phosphatase activity | null | cytoplasm, nucleus | | |
| PBANKA_08 | phosphoglycerate kinase, putative (PGK) | | | PF3D7_0922500 | phosphoglycerate kinase (PGK) | glycolysis | 0 | 1 1 | 1 na | a | phosphoglycerate kinase activity | glycolysis | cytosol | | |
| PBANKA_11 | SNAP protein, putative | | | PF3D7_0509000 | SNAP protein (soluble N-ethylmaleimide-sensitive factor Attachment Protein), putative | re trafficking | 0 | 1 1 | 1 na | a | null | intracellular protein transport | Golgi apparatus, endoplasmic reticulum | | |
| PBANKA_12 | conserved Plasmodium protein, unknown function | | | PF3D7_1011000 | conserved protein, unknown function | gliding motility | 0 | 1 1 | 1 (tagg | ged) different in fertilization, ookinete, oocyst, sprorozoite and liver stage | null | null | inner membrane complex | | |
| PBANKA_13 | conserved Plasmodium protein, unknown function | | | PF3D7_1345600 | conserved Plasmodium protein, unknown function | gliding motility | 1 | 0 1 | 1 na | à | null | null | inner membrane complex | | |
| PBANKA_11 | 780 longevity-assurance (LAG1) protein, putative | | | PF3D7_0508200 | longevity-assurance (LAG1) domain protein, putative | e ceramide biosynthesis | 0 | 1 1 | 1 na | à | null | null | integral to membrane | | |
| PBANKA_07 | 270 rhomboid protease, putative | | | PF3D7_0828000 | rhomboid protease ROM3 (ROM3) | rhomboid | 0 | 1 1 | 1 differe | rent in cocyst and sporozoite; (tagged) also different in fertilization and cokinete | null | null | integral to membrane | | |
| | | | | | | | | | | | | | | | |

| PBANKA_111780 | conserved Plasmodium protein, unknown function,rhomboid protease RC | OM10, putative | | PF3D7_0618600 | rhomboid protease ROM10 (ROM10) | rhomboid 1 | 0 | 1 | not different from wild type | null | null | integral to membrane |
|---------------|---|----------------|------|---------------|---|--|---|---|------------------------------|---|---|--|
| PBANKA_141300 | cop-coated vesicle membrane protein p24 precursor, putative | | | PF3D7_1314500 | cop-coated vesicle membrane protein p24 precursor, puta | tive trafficking 1 | 0 | 1 | na | null | transport | integral to membrane |
| PBANKA_031330 | MtN3-like protein | | | PF3D7_0216600 | MtN3-like protein | trafficking 1 | 0 | 1 | na | null | null | integral to membrane |
| PBANKA_110650 | rhomboid protease, putative (ROM4) | | | PF3D7_0506900 | rhomboid protease ROM4 (ROM4) | rhomboid 1 | 0 | 1 | KO not successful | serine-type endopeptidase activity | null | integral to membrane |
| PBANKA_131230 | SNARE protein, putative | | | PF3D7_1448600 | SNARE protein, putative (VTI1) | trafficking 1 | 0 | 1 | na | null | vesicle-mediated transport | integral to membrane |
| PBANKA_133990 | SNARE protein, putative | | | PF3D7_1324700 | SNARE protein, putative (YKT6.2) | trafficking 1 | 0 | 1 | na | null | vesicle-mediated transport | integral to membrane |
| PBANKA_020700 | calcium-transporting ATPase, putative (SERCA) | | | PF3D7_0106300 | calcium-transporting ATPase (ATP6) | transporter 0 | 1 | 1 | na | ATP binding, calcium-transporting ATPase activity | ATP biosynthetic process, calcium ion transport, metabolic process | is integral to membrane, membrane |
| PBANKA_110310 | actin-depolymerizing factor 1 (ADF1) | | | PF3D7_0503400 | actin-depolymerizing factor 1 (ADF1) | gliding motility 0 | 1 | 1 | KO not successful | actin binding | null | intracellular |
| PBANKA_131340 | exosome complex exonuclease RRP6, putative (RRP6) | | | PF3D7_1449700 | exosome complex exonuclease RRP6, putative (RRI | P6) mRNA 0 | 1 | 1 | na | 3'-5' exonuclease activity, nucleic acid binding | nucleobase-containing compound metabolic process | intracellular |
| PBANKA_100810 | GTPase, putative | | | PF3D7_1436500 | conserved protein, unknown function | translation, transporter 0 | 1 | 1 | na | GTP binding | null | intracellular |
| PBANKA_145990 | signal recognition particle, beta subunit, putative | | | PF3D7_1246800 | signal recognition particle receptor, beta subunit (SRPR-bet | ta) post-translational modification 0 | 1 | 1 | na | GTP binding | intracellular protein transport, small GTPase mediated signal transduction | intracellular |
| PBANKA_132410 | ubiquitin carboxyl-terminal hydrolase isozyme L3, putative (UCHL3) | | | PF3D7_1460400 | ubiquitin carboxyl-terminal hydrolase isozyme L3 (UCHL | 3) proteasome 0 | 1 | 1 | na | ubiquitin thiolesterase activity | ubiquitin-dependent protein catabolic process | intracellular |
| PBANKA_130430 | conserved Plasmodium protein, unknown function | | | PF3D7_1440400 | conserved Plasmodium protein, unknown function | on unknown function 1 | 0 | 1 | na | zinc ion binding | null | intracellular |
| PBANKA_135410 | Rab GTPase 11b | | | PF3D7_1340700 | Rab GTPase 11b (RAB11b) | trafficking 1 | 0 | 1 | na | GTP binding, GTPase activity, protein binding | intracellular protein transport, nucleocytoplasmic transport, small GTPase mediated signal transduction | intracellular |
| PBANKA_060770 | zinc finger transcription factor, putative | | | PF3D7_1209300 | zinc finger transcription factor (KROX1) | transcription 1 | 0 | 1 | na | zinc ion binding | null | intracellular |
| PBANKA_100590 | zinc-finger, RAN binding protein, putative | | | PF3D7_0408300 | zinc finger, RAN binding protein, putative | transcription 1 | 0 | 1 | na | zinc ion binding | null | intracellular |
| PBANKA_050710 | CDGSH iron-sulfur domain-containing protein, putative | | | PF3D7_1022900 | CDGSH iron-sulfur domain-containing protein, putat | ive intracellular Mb-bounded organelle | 0 | 1 | na | 2 iron, 2 sulfur cluster binding | null | intracellular membrane-bounded organelle |
| PBANKA_140130 | 40S ribosomal protein S7, putative | | | PF3D7_1302800 | 40S ribosomal protein S7, putative | ribosome 0 | 1 | 1 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_123390 | mitochondrial ribosomal protein L14 precursor, putative | -1.88 | 0.73 | PF3D7_0519100 | mitochondrial ribosomal protein L14 precursor, putat | ive ribosome mito 0 | 1 | 1 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_123720 | mitochondrial ribosomal protein L17 precursor, putative | -2.23 | 0.75 | PF3D7_0522500 | mitochondrial ribosomal protein L17 precursor, putat | ive ribosome mito 0 | 1 | 1 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_145610 | 40S ribosomal protein S17, putative | | | PF3D7_1242700 | 40S ribosomal protein S17, putative | ribosome 1 | 0 | 1 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_090640 | 60S ribosomal protein L35ae, putative | | | PF3D7_1142600 | 60S ribosomal protein L35ae, putative | ribosome 1 | 0 | 1 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_091810 | 60S ribosomal protein L38e, putative | | | PF3D7_1130100 | 60S ribosomal protein L38e, putative | ribosome 1 | 0 | 1 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_135190 | 60S ribosomal protein L6-2, putative | | | PF3D7_1338200 | 60S ribosomal protein L6-2, putative | ribosome 1 | 0 | 1 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_081920 | organelle ribosomal protein L3 precursor, putative | | | PF3D7_0918200 | organelle ribosomal protein L3 precursor, putativ | ve ribosome 1 | 0 | 1 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_120190 | 40S ribosomal protein S20e, putative | | | PF3D7_1003500 | 40S ribosomal protein S20e, putative | ribosome 1 | 0 | 1 | na | structural constituent of ribosome | translation | intracellular, ribosome, small ribosomal subunit |
| PBANKA_101500 | mitochondrial ribosomal protein L15 precursor, putative | -1.72 | 0.71 | PF3D7_1429700 | mitochondrial ribosomal protein L15 precursor, putat | ive ribosome mito 0 | 1 | 1 | na | structural constituent of ribosome | translation | large ribosomal subunit |
| PBANKA_071880 | small GTP-binding protein sar1, putative (SAR1) | | | PF3D7_0416800 | small GTP-binding protein sar1 (SAR1) | trafficking 0 | 1 | 1 | na | GTP binding | intracellular protein transport, small GTPase mediated signal transduction | Maurer's cleft, intracellular |
| PBANKA_120910 | dolichyl-phosphate-mannose protein mannosyltransferase, putative | | | PF3D7_1010700 | dolichyl-phosphate-mannose protein mannosyltransferase, putar | tive O-glycan biosynthesis 0 | 1 | 1 | na | null | null | membrane |
| PBANKA_145500 | mitochondrial carrier protein, putative | -2.26 | 0.75 | PF3D7_1241600 | mitochondrial carrier protein, putative | transporter 0 | 1 | 1 | na | binding | transport | membrane |
| PBANKA_060110 | mitochondrial phosphate carrier protein, putative | | | PF3D7_1202200 | mitochondrial phosphate carrier protein (MPC) | transporter 0 | 1 | 1 | na | binding | transport | membrane |
| | | | | | | | | | | | | |

| PBA | ANKA_010770 | permease, putative | -2.25 | 0.75 | PF3D7_0609100 | Zn2 or Fe2 permease | acidocalcisome | 0 | 1 | 1 r | na | metal ion transmembrane transporter activity | metal ion transport | membrane |
|-----|-------------|---|-------|------|---------------|---|-------------------------------|---|---|-----|---|--|--|--|
| PBA | ANKA_050650 | zinc transporter, putative | | | PF3D7_1022300 | zinc transporter, putative | acidocalcisome | 0 | 1 | 1 r | na | metal ion transmembrane transporter activity | metal ion transport | membrane |
| PBA | ANKA_061040 | ATPase, putative | | | PF3D7_1211900 | non-SERCA-type Ca2 -transporting P-ATPase (ATP4) |) acidocalcisome | 1 | 0 | 1 r | na | ATP binding, ATPase activity, coupled to transmembrane movement of ions, phosphorylative mechanism | ATP biosynthetic process, cation transport, metabolic proces | s membrane |
| PBA | ANKA_142710 | conserved Plasmodium protein, unknown function | -3.76 | 0.75 | PF3D7_0810400 | conserved Plasmodium membrane protein, unknown function | n unknown function | 1 | 0 | 1 r | na | transporter activity | transport | membrane |
| PBA | ANKA_101380 | cytochrome c oxidase subunit II precursor, putative | | | PF3D7_1430900 | cytochrome c oxidase subunit II precursor, putative | e redoxmeth | 1 | 0 | 1 r | na | copper ion binding, cytochrome-c oxidase activity | null | membrane |
| PBA | ANKA_093900 | ion channel protein, putative | | | PF3D7_1107900 | mechanosensitive ion channel protein | transporter | 1 | 0 | 1 r | na | null | null | membrane |
| PBA | ANKA_141410 | phosphatidylinositol synthase, putative (PIS) | | | PF3D7_1315600 | phosphatidylinositol synthase (PIS) | inositol phosphate metabolism | 1 | 0 | 1 r | na | phosphotransferase activity, for other substituted phosphate groups | s phospholipid biosynthetic process | membrane |
| PBA | ANKA_131620 | SNARE protein, putative | | | PF3D7_1452500 | SNARE protein, putative (PfBet1) | trafficking | 1 | 0 | 1 r | na | protein binding | Golgi vesicle transport | membrane |
| PBA | ANKA_146340 | adaptor-related protein complex 3, sigma 2 subunit, putative | | | PF3D7_1250500 | adaptor-related protein complex 3, sigma 2 subunit, putative | trafficking | 0 | 1 | 1 r | na | protein binding, protein transporter activity | intracellular protein transport, vesicle-mediated transport | membrane coat |
| PBA | ANKA_136550 | intra-erythrocytic P. berghei-induced structures protein 1 (IBIS1) | | | | | | 1 | 0 | 1 0 | different in liver stage | null | null | membrane, symbiont-containing vacuole membrane |
| PBA | ANKA_082420 | perforin like protein 3 (PPLP3) | -2.93 | 0.75 | PF3D7_0923300 | perforin like protein 3 (PPLP3) | perforin | 0 | 1 | 1 d | different in fertilization, ookinete and oocyst | null | null | microneme |
| PBA | ANKA_021390 | dynein light chain, putative | | | PF3D7_0729800 | dynein light chain, putative | dynein/kinesin | 0 | 1 | 1 r | na | microtubule motor activity | microtubule-based process | microtubule associated complex |
| PBA | ANKA_030220 | dynein light chain, putative | | | PF3D7_0204400 | dynein light chain, putative | dynein/kinesin | 1 | 0 | 1 r | na | microtubule motor activity | microtubule-based process | microtubule associated complex |
| PBA | ANKA_083430 | conserved Plasmodium protein, unknown function | | | PF3D7_0933500 | conserved Plasmodium protein, unknown function | chromatin | 0 | 1 | 1 r | na | null | microtubule cytoskeleton organization | microtubule organizing center, spindle pole |
| PBA | ANKA_101010 | mitochondrial import inner membrane translocase subunit tim17, putative | | | PF3D7_1434700 | mitochondrial import inner membrane translocase subunit tim17, putative | transporter | 0 | 1 | 1 r | na | protein transporter activity | protein transport | mitochondrial inner membrane |
| PBA | ANKA_145630 | mitochondrial import inner membrane translocase subunit, putative | | | PF3D7_1242900 | mitochondrial import inner membrane translocase subunit, putative | e transporter | 0 | 1 | 1 r | na | null | protein import into mitochondrial inner membrane | mitochondrial intermembrane space protein transporter complex |
| PBA | ANKA_111650 | mitochondrial import receptor subunit, putative | | | PF3D7_0617000 | mitochondrial import receptor subunit tom40, putative | e transporter | 1 | 0 | 1 r | na | voltage-gated anion channel activity | anion transport | mitochondrial outer membrane |
| PBA | ANKA_142180 | mitochondrial ATP synthase F1, epsilon subunit, putative | | | PF3D7_0715500 | mitochondrial ATP synthase F1, epsilon subunit, putativ | e transporter | 1 | 0 | 1 r | na | hydrogen ion transporting ATP synthase activity, rotational mechanism, proton-transporting ATPase activity, rotational mechanism | ATP synthesis coupled proton transport | mitochondrial proton-transporting ATP synthase complex, catalytic core F(1), mitochondrion |
| PBA | ANKA_041330 | mitochondrial ribosomal protein L29/L47 precursor, putative | | | PF3D7_0315500 | mitochondrial ribosomal protein L29/L47 precursor, putativ | re ribosome mito | 1 | 0 | 1 r | na | structural constituent of ribosome | translation | mitochondrial ribosome |
| PBA | ANKA_092360 | GrpE protein homolog, mitochondrial, putative (MGE1) | -2.7 | 0.75 | PF3D7_1124700 | co-chaperone GrpE, putative | chaperone, transporter | 1 | 0 | 1 r | na | adenyl-nucleotide exchange factor activity, chaperone binding, protein homodimerization activity | protein folding, protein import into mitochondrial matrix | mitochondrion |
| PBA | ANKA_102500 | conserved Plasmodium protein, unknown function | | | PF3D7_1417700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 1 | 1 r | na | null | null | nuclear periphery, nucleus |
| PBA | ANKA_111700 | histone H2A, putative (H2A) | | | PF3D7_0617800 | histone H2A (H2A) | chaperone, chromatin | 0 | 1 | 1 r | na | DNA binding | nucleosome assembly | nucleosome, nucleus |
| PBA | ANKA_142060 | histone H2B, putative | | | PF3D7_0714000 | histone H2B variant, putative (H2Bv) | chaperone | 1 | 0 | 1 (| tagged) different in liver stage | DNA binding | nucleosome assembly | nucleosome, nucleus |
| PBA | ANKA_143940 | chromatin assembly protein, putative | | | PF3D7_1224500 | chromatin assembly protein (ASF1), putative | chromatin | 0 | 1 | 1 r | na | null | chromatin assembly or disassembly | nucleus |
| PBA | ANKA_142770 | RuvB DNA helicase, putative | | | PF3D7_0809700 | RuvB DNA helicase, putative | DNA replication | 0 | 1 | 1 r | na | ATP binding, ATP-dependent 5'-3' DNA helicase activity, ATPase activity, DNA helicase activity | null | nucleus |
| PBA | ANKA_113440 | conserved Plasmodium protein, unknown function | | | PF3D7_1358100 | small subunit rRNA processing factor, putative | RNA | 1 | 0 | 1 r | na | null | gene silencing | nucleus |
| PBA | ANKA_113090 | methyltransferase, putative | -2.13 | 0.74 | PF3D7_1354300 | large subunit rRNA methyltransferase, putative | RNA | 1 | 0 | 1 r | na | methyltransferase activity, nucleic acid binding | rRNA methylation | nucleus |
| PBA | ANKA_051140 | peroxiredoxin, putative (nPrx, UIS16) | | | PF3D7_1027300 | peroxiredoxin (nPrx) | redoxmeth | 1 | 0 | 1 r | na | antioxidant activity, oxidoreductase activity | null | nucleus |
| PBA | ANKA_090720 | phd finger protein, putative | | | PF3D7_1141800 | phd finger protein, putative | chromatin | 1 | 0 | 1 r | na | protein binding, zinc ion binding | null | nucleus |
| PBA | ANKA_111910 | splicing factor 3A subunit 2, putative (SF3A2) | | | PF3D7_0619900 | splicing factor 3A subunit 2, putative (SF3A2) | mRNA | 1 | 0 | 1 r | na | nucleic acid binding, zinc ion binding | null | nucleus |
| PBA | ANKA_133200 | U2 snRNP auxiliary factor, putative | | | PF3D7_1468800 | U2 snRNP auxiliary factor, putative | mRNA | 1 | 0 | 1 r | na | RNA binding | mRNA processing | nucleus |
| | | | | | | | | | | | | | | |

| PBANKA_132740 yl1 nuclear protein, putative | PF3D7_1464000 YL1 nuclear protein, putative transcription 1 0 | 1 na s | sequence-specific DNA binding transcription factor activity re | egulation of transcription, DNA-dependent | nucleus | | | |
|---|--|---|---|--|---|-----------------|---|--------------------------------|
| PBANKA_020290 transcription initiation factor TFIIB, putative | PF3D7_0110800 transcription initiation factor TFIIB, putative transcription 1 0 | 1 na p | protein binding, translation initiation factor activity, zinc ion binding | IA-dependent transcription, initiation, regulation of transcription, DNA-dependent, translational initiation | nucleus, transcription factor complex | | | |
| PBANKA_110580 TATA-box binding protein, putative, transcription initiation factor tfiid, putative (TBP) | | | DNA binding reg | gulation of transcription, DNA-dependent, transcription initiation from RNA polymerase II promoter | nucleus, transcription factor TFIID complex | | | |
| PBANKA_101590 transcription initiation TFIID-like, putative | PF3D7_1428800 transcription initiation TFIID-like, putative transcription 0 1 | 1 na [| DNA binding reg | gulation of transcription, DNA-dependent, transcription initiation from RNA polymerase II promoter | nucleus, transcription factor TFIID complex | | | |
| PBANKA_114400 26s proteasome regulatory subunit rpn11, putative (RPN11) | PF3D7_1368100 proteasome regulatory subunit, putative proteasome 0 1 | 1 na r | null nu | ull | null | null | ubiquitin-dependent protein catabolic process | proteasome regulatory particle |
| PBANKA_112620 4-methyl-5(B-hydroxyethyl)-thiazol monophosphate biosynthesis enzyme, putative | PF3D7_0627500 4-methyl-5(B-hydroxyethyl)-thiazol monophosphate biosynthesis enzyme thiamine (vitB1) metabolism 0 1 | 1 na r | null nu | ull | null | null | thiamine biosynthetic process | null |
| PBANKA_130550 acid cluster protein 33 homologue, putative -2.34 0.75 | PF3D7_1441600 acid cluster protein 33 homologue, putative 0 1 | 1 na r | null nu | ull | null | null | null | null |
| PBANKA_112340 anaphase-promoting complex subunit, putative | PF3D7_0624500 anaphase promoting complex subunit, putative meiosis 0 1 | 1 na p | protein binding, zinc ion binding nu | ull | null | | | |
| PBANKA_031500 apicoplast RNA methyltransferase precursor, putative | PF3D7_0218300 apicoplast RNA methyltransferase precursor, putative RNA 0 1 | 1 na F | RNA binding, RNA methyltransferase activity RI | NA processing | null | | | |
| PBANKA_123260 apicortin, putative -1.89 0.7 | PF3D7_0517800 apicortin, putative apicoplast 0 1 | 1 na r | null int | tracellular signal transduction | null | | | |
| PBANKA_135770 aspartate carbamoyltransferase, putative -4.56 0.75 | PF3D7_1344800 aspartate carbamoyltransferase (atcasE) pyrimidine metabolism 0 1 | 1 na a | amino acid binding, aspartate carbamoyltransferase activity de | novo' pyrimidine nucleobase biosynthetic process, cellular amino acid metabolic process | null | | | |
| PBANKA_140020 BIR protein | BIR protein 0 1 | 1 na r | null nu | ull | null | na | na | na |
| PBANKA_031640 BIR protein, pseudogene | BIR protein 0 1 | 1 na r | null nu | ull | null | | | |
| PBANKA_050080 BIR protein, pseudogene | BIR protein 0 1 | 1 na r | null nu | ull | null | na | na | na |
| PBANKA_040820 calcium dependent protein kinase 3 (CDPK3) -1.68 0.71 | PF3D7_0310100 calcium dependent protein kinase 3 (CDPK3) kinase 0 1 | different in fertilization, ookinete and oocyst | ATP binding, calcium ion binding, protein serine/threonine kinase activity, protein tyrosine kinase activity PT | rotein phosphorylation | null | | | |
| PBANKA_101980 calcium-dependent protein kinase, putative -6.75 0.75 | PF3D7_1423600 serine/threonine protein kinase, putative, cdlk kinase 0 1 | different in oocyst, sporozoite and liver stage A | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | rotein phosphorylation | null | | | |
| PBANKA_091210 casein kinase 1 (CK1) | PF3D7_1136500.1/2 casein kinase 1 (CK1) kinase 0 1 | 1 KO not successful A | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | rotein phosphorylation | null | | | |
| PBANKA_051910 CCAT-binding transcription factor-like protein, putative | PF3D7_1036600.1/2 conserved Plasmodium protein, unknown function transcription 0 1 | 1 na r | null nu | ull | null | na | na | na |
| PBANKA_101600 conserved Plasmodium protein, unknown function -2.1 0.73 | PF3D7_1428700 conserved protein, unknown function redoxmeth 0 1 | 1 na h | neme binding nu | ull | null | | | |
| PBANKA_060600 conserved Plasmodium protein, unknown function | PF3D7_1207500 conserved Plasmodium protein, unknown function unknown function 0 1 | 1 na r | nucleic acid binding nu | ull | null | | | |
| PBANKA_132780 conserved Plasmodium protein, unknown function | PF3D7_1464400 zinc finger protein, putative nucleic acid binding 0 1 | 1 na r | nucleic acid binding, zinc ion binding nu | ull | null | | | |
| PBANKA_010510 conserved Plasmodium protein, unknown function | PF3D7_0606400 conserved Plasmodium protein, unknown function unknown function 0 1 | 1 na r | null nu | ull | null | null | null | cytoplasm, membrane |
| PBANKA_010700 conserved Plasmodium protein, unknown function | PF3D7_0608400 conserved Plasmodium protein, unknown function unknown function 0 1 | 1 na r | null nu | ull | null | null | null | null |
| PBANKA_010960 conserved Plasmodium protein, unknown function | PF3D7_0611300 conserved Plasmodium protein, unknown function unknown function 0 1 | 1 na r | null nu | ull | null | null | null | null |
| PBANKA_010990 conserved Plasmodium protein, unknown function -3.42 0.74 | PF3D7_0611600 conserved Plasmodium protein, unknown function unknown function 0 1 | 1 na r | null nu | ull | null | null | null | null |
| PBANKA_020790 conserved Plasmodium protein, unknown function | PF3D7_0105400.1/2 conserved Plasmodium protein, unknown function unknown function 0 1 | 1 na r | null nu | ull | null | null | null | integral to membrane, membrane |
| PBANKA_021160 conserved Plasmodium protein, unknown function | PF3D7_0727500 conserved Plasmodium protein, unknown function mitochondrial transcription 0 1 | 1 na r | null nu | ull | null | null | null | apicoplast |
| PBANKA_021260 conserved Plasmodium protein, unknown function | PF3D7_0728500 conserved Plasmodium protein, unknown function kinase 0 1 | 1 na r | null nu | ull | null | null | null | null |
| PBANKA_021450 conserved Plasmodium protein, unknown function | PF3D7_0730400 conserved Plasmodium protein, unknown function unknown function 0 1 | 1 na r | null nu | ull | null | null | null | null |
| PBANKA_030280 conserved Plasmodium protein, unknown function | PF3D7_0205100 conserved Plasmodium protein, unknown function DNA repair 0 1 | 1 na r | null nu | ull | null | protein binding | null | null |
| PBANKA_030350 conserved Plasmodium protein, unknown function -2.1 0.73 | PF3D7_0205800 conserved Plasmodium protein, unknown function phospholipid binding 0 1 | 1 na r | null nu | ull | null | null | null | null |
| | | | | | | | | |

| PBANKA_040880 conserved Plasmodium protein, unknown function | | PF3D7_031080 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | apicoplast, integral to membrane, membrane |
|--|-----------|-----------------|--|---|-----|----|------|------|------|---|---|--|
| PBANKA_041140 conserved Plasmodium protein, unknown function | | PF3D7_031370 | 0.1/2 conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | null |
| PBANKA_041560 conserved Plasmodium protein, unknown function | | PF3D7_090580 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | DNA binding, protein binding, translation elongation factor activity, zinc ion bi | inding DNA-dependent transcription, elongation, regulation of transcription from RNA polymerase II promoter, transcription, DNA-dependent | oendent nucleus |
| PBANKA_050780 conserved Plasmodium protein, unknown function | | PF3D7_102360 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | ATP binding, motor activity, protein binding | null | cytoplasm |
| PBANKA_051600 conserved Plasmodium protein, unknown function | | PF3D7_103220 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | null |
| PBANKA_052200 conserved Plasmodium protein, unknown function | | PF3D7_042060 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | null |
| PBANKA_062020 conserved Plasmodium protein, unknown function | | PF3D7_072270 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | integral to membrane |
| PBANKA_071300 conserved Plasmodium protein, unknown function | | PF3D7_081780 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | metal ion binding, nucleotide binding | null | null |
| PBANKA_080140 conserved Plasmodium protein, unknown function | | PF3D7_070370 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | null |
| PBANKA_081510 conserved Plasmodium protein, unknown function | -3.06 0.7 | 75 PF3D7_091410 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | ATP binding, motor activity, protein binding | null | cytoplasm |
| PBANKA_082930 conserved Plasmodium protein, unknown function | | PF3D7_092850 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | protein binding | null | null |
| PBANKA_092250 conserved Plasmodium protein, unknown function | | PF3D7_112580 | kelch protein, putative microtubule | 0 | 1 1 | na | null | null | null | null | null | null |
| PBANKA_092270 conserved Plasmodium protein, unknown function | | PF3D7_112560 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | null |
| PBANKA_092330 conserved Plasmodium protein, unknown function | -3.36 0.7 | 75 PF3D7_112500 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | ATP binding, ATPase activity | transport | apicoplast, plasma membrane |
| PBANKA_092760 conserved Plasmodium protein, unknown function | | PF3D7_112070 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | integral to membrane, membrane, nucleus |
| PBANKA_092820 conserved Plasmodium protein, unknown function | | PF3D7_112000 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | ATP binding | null | cytoplasm |
| PBANKA_093060 conserved Plasmodium protein, unknown function | | PF3D7_111740 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | membrane |
| PBANKA_100320 conserved Plasmodium protein, unknown function | | PF3D7_040550 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | null |
| PBANKA_102600 conserved Plasmodium protein, unknown function | | PF3D7_141670 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | cytoplasm |
| PBANKA_103140 conserved Plasmodium protein, unknown function | | PF3D7_141110 | 0.1/2 conserved Plasmodium membrane protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | integral to membrane, plasma membrane |
| PBANKA_110750 conserved Plasmodium protein, unknown function | | PF3D7_050790 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | transport | integral to membrane, plasma membrane |
| PBANKA_111280 conserved Plasmodium protein, unknown function | -2.43 0.7 | 75 PF3D7_051310 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | cytoplasm |
| PBANKA_111410 conserved Plasmodium protein, unknown function | -4.23 0.7 | 75 PF3D7_051450 | conserved Plasmodium membrane protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | membrane |
| PBANKA_111930 conserved Plasmodium protein, unknown function | | PF3D7_062010 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | cytoplasm |
| PBANKA_120530 conserved Plasmodium protein, unknown function | | PF3D7_100710 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | null |
| PBANKA_122170 conserved Plasmodium protein, unknown function | | PF3D7_071110 | conserved Plasmodium membrane protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | null |
| PBANKA_123370 conserved Plasmodium protein, unknown function | | PF3D7_051890 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | cytoplasm |
| PBANKA_123470 conserved Plasmodium protein, unknown function | | PF3D7_051990 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | apicoplast |
| PBANKA_123810 conserved Plasmodium protein, unknown function | | PF3D7_052330 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | null |
| PBANKA_123840 conserved Plasmodium protein, unknown function | | PF3D7_052360 | conserved Plasmodium protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | null | null | cytoplasm |
| PBANKA_123870 conserved Plasmodium protein, unknown function | | PF3D7_052390 | conserved Plasmodium membrane protein, unknown function unknown function | 0 | 1 1 | na | null | null | null | G-protein coupled receptor activity | G-protein coupled receptor signaling pathway | integral to membrane, membrane |
| | | | | | | | | | | | | |

| PBANKA_131290 | conserved Plasmodium protein, unknown function | | | | ι | unknown function | 0 | 1 | 1 | na | null | null | null |
|---------------|---|-------|------|---------------|---|-------------------------------------|---|---|---|-------------------|--|------------------------------|------|
| PBANKA_131610 | conserved Plasmodium protein, unknown function | | | PF3D7_1452400 | conserved Plasmodium protein, unknown function u | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_131940 | conserved Plasmodium protein, unknown function | | | PF3D7_1455700 | conserved Plasmodium protein, unknown function u | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_132190 | conserved Plasmodium protein, unknown function | | | PF3D7_1458200 | conserved Plasmodium protein, unknown function u | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_132240 | conserved Plasmodium protein, unknown function | | | PF3D7_1458700 | conserved Plasmodium protein, unknown function u | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_132880 | conserved Plasmodium protein, unknown function | | | PF3D7_1465400 | conserved Plasmodium protein, unknown function u | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_133910 | conserved Plasmodium protein, unknown function | | | PF3D7_1323900 | conserved Plasmodium protein, unknown function | Ca2+ binding, prot-prot interaction | 0 | 1 | 1 | na | null | null | null |
| PBANKA_134110 | conserved Plasmodium protein, unknown function | | | PF3D7_1326000 | conserved Plasmodium protein, unknown function u | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_134420 | conserved Plasmodium protein, unknown function | | | PF3D7_1329200 | conserved Plasmodium protein, unknown function u | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_135170 | conserved Plasmodium protein, unknown function | | | PF3D7_1338000 | conserved Plasmodium membrane protein, unknown function U | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_136020 | conserved Plasmodium protein, unknown function | -2.25 | 0.74 | PF3D7_1347400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_136360 | conserved Plasmodium protein, unknown function | | | PF3D7_1350800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_140290 | conserved Plasmodium protein, unknown function | | | PF3D7_1304400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_141010 | conserved Plasmodium protein, unknown function | -6.66 | 0.75 | PF3D7_1311600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_141350 | conserved Plasmodium protein, unknown function | | | PF3D7_1315000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_142920 | conserved Plasmodium protein, unknown function | -3.32 | 0.75 | PF3D7_1213400 | conserved Plasmodium protein, unknown function | galactose metabolism | 0 | 1 | 1 | na | null | null | null |
| PBANKA_143060 | conserved Plasmodium protein, unknown function | | | PF3D7_1214800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_143580 | conserved Plasmodium protein, unknown function | -1.96 | 0.73 | PF3D7_1220600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_144780 | conserved Plasmodium protein, unknown function | | | PF3D7_1233100 | conserved protein, unknown function | unknown function | 0 | 1 | 1 | KO not successful | null | null | null |
| PBANKA_145220 | conserved Plasmodium protein, unknown function | | | PF3D7_1237600 | rRNA processing WD-repeat protein, putative | RNA | 0 | 1 | 1 | na | null | null | null |
| PBANKA_145550 | conserved Plasmodium protein, unknown function | | | PF3D7_1242100 | conserved Plasmodium protein, unknown function | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_061530 | conserved Plasmodium protein, unknown function | -3.55 | 0.75 | PF3D7_0717600 | conserved Plasmodium protein, unknown function | unknown function | 1 | 0 | 1 | na | binding | null | null |
| PBANKA_101870 | conserved Plasmodium protein, unknown function | | | PF3D7_1425900 | conserved Plasmodium protein, unknown function t | transcription | 1 | 0 | 1 | na | binding | null | null |
| PBANKA_103890 | conserved Plasmodium protein, unknown function | | | PF3D7_1403100 | conserved Plasmodium protein, unknown function | chromatin | 1 | 0 | 1 | na | binding | null | null |
| PBANKA_134700 | conserved Plasmodium protein, unknown function | -1.99 | 0.72 | PF3D7_1332200 | conserved Plasmodium protein, unknown function | unknown function | 1 | 0 | 1 | na | DNA binding, alkylbase DNA N-glycosylase activity | base-excision repair | null |
| PBANKA_131310 | conserved Plasmodium protein, unknown function | -3.93 | 0.75 | PF3D7_1449400 | DNA replication related protein, putative | DNA replication | 1 | 0 | 1 | na | DNA binding, nuclease activity, protein binding | DNA metabolic process | null |
| PBANKA_146240 | cutA, putative | | | PF3D7_1249500 | cutA, putative | protein binding, response to ion | 0 | 1 | 1 | na | null | response to metal ion | null |
| PBANKA_021130 | cysteine desulfurase, putative (NFS) | | | PF3D7_0727200 | cysteine desulfurase, putative (NFS) | thiamine (vitB1) metabolism | 0 | 1 | 1 | na | null | metabolic process | null |
| PBANKA_146070 | dipeptidyl peptidase 2, putative (DPAP2) | | | PF3D7_1247800 | dipeptidyl peptidase 2, putative (DPAP2) | post-translational modification | 0 | 1 | 1 | na | cysteine-type peptidase activity | proteolysis | null |
| PBANKA_092300 | DNA-dependent RNA polymerase, putative | | | PF3D7_1125300 | DNA-dependent RNA polymerase t | transcription | 0 | 1 | 1 | na | DNA binding, DNA-directed RNA polymerase activity | transcription, DNA-dependent | null |
| PBANKA_040720 | dual-specificity protein phosphatase, putative (YVH1) | | | PF3D7_0309000 | dual specificity phosphatase (YVH1) | phosphatase | 0 | 1 | 1 | na | protein tyrosine/serine/threonine phosphatase activity | protein dephosphorylation | null |
| | | | | | | | | | | | | | |

cytoplasm, nucleus cytoplasm regulation of cell shape mitochondrion null protein transport cytoplasm, nucleus nucleotide binding integral to membrane, membrane integral to membrane, plasma membrane membrane integral to membrane, membrane protein binding cytoplasm, nucleus apicoplast apicoplast, membrane cytoplasm, nucleus null

| PBANKA_052420 | early transcribed membrane protein (SEP2) | | | | | early transcribed membrane protein (ETRAMP) | 0 | 1 | 1 | | null | null | null |
|---------------|--|-------|------|---------------|--|---|---|---|---|--|---|---|------|
| PBANKA 103190 | eukaryotic translation initiation factor 2 gamma subunit, putative | | | PF3D7 1410600 | | translation | 0 | 1 | ' | KO not successful; (tagged) different in fertilization, oddinere, oocyst, sporozoite and liver stage | GTP binding, GTPase activity | null | null |
| _ | • | | | _ | J | | | 1 | ' | | , | | |
| PBANKA_133430 | exported protein 2, putative (EXP2) | | | PF3D7_1471100 | exported protein 2 (EXP2) | trafficking | 0 | ' | | (tagged) different in sporozoite and liver stage | | null | null |
| PBANKA_124710 | fam-b protein glutaminyl-peptide cyclotransferase, putative | 2.00 | 0.74 | | | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_131070 | | -2.08 | 0.74 | PF3D7_1446900 | glutaminyl-peptide cyclotransferase, putative | | 0 | 1 | 1 | na | null | null | null |
| PBANKA_130970 | helicase, putative | | | PF3D7_1445900 | DEAD/DEAH box ATP-dependent RNA helicase, putative | mRNA | 0 | 1 | 1 | na | ATP binding, ATP-dependent helicase activity, helicase activity, nucleic acid binding | null | null |
| PBANKA_071840 | histone acetyltransferase, putative (HAT1) | | | PF3D7_0416400 | histone acetyltransferase, putative (HAT1) | chromatin modifying proteins | 0 | 1 | 1 | na | histone acetyltransferase activity | chromatin modification | null |
| PBANKA_136440 | inner membrane complex protein 1f, putative (IMC1F, ALV6) | -2.42 | 0.73 | PF3D7_1351700 | alveolin, putative (ALV6) | gliding motility | 0 | 1 | 1 | na | null | null | null |
| PBANKA_093890 | IWS1-like protein, putative | | | PF3D7_1108000 | IWS1-like protein, putative | transcription | 0 | 1 | 1 | na | null | null | null |
| PBANKA_050420 | large subunit rRNA processing RRM protein, putative | | | PF3D7_1020000 | large subunit rRNA processing RRM protein, putative | RNA | 0 | 1 | 1 | na | nucleic acid binding | null | null |
| PBANKA_030440 | merozoite surface protein 4/5 (MSP4/5) | | | PF3D7_0207000 | merozoite surface protein 4 (MSP4) | contaminant | 0 | 1 | 1 | na | null | null | null |
| PBANKA_010300 | mitochondrial ribosomal protein L41 precursor, putative | | | PF3D7_0604200 | mitochondrial ribosomal protein L41 precursor, putative | ribosome mito | 0 | 1 | 1 | na | null | null | null |
| PBANKA_131830 | mitochondrial ribosomal protein S11 precursor, putative | | | PF3D7_1454600 | mitochondrial ribosomal protein S11 precursor, putative | ribosome mito | 0 | 1 | 1 | na | null | null | null |
| PBANKA_122690 | mitogen-activated protein kinase organizer 1, putative | | | PF3D7_0803300 | mitogen-activated protein kinase organizer 1, putative (PfMORG1) | kinase | 0 | 1 | 1 | na | null | null | null |
| PBANKA_030720 | monocarboxylate transporter, putative | | | PF3D7_0210300 | monocarboxylate transporter, putative | glycolysis | 0 | 1 | 1 | na | null | null | null |
| PBANKA_020410 | N-terminal acetyltransferase, putative | | | PF3D7_0109500 | N-acetyltransferase, putative | chromatin modifying proteins | 0 | 1 | 1 | na | N-acetyltransferase activity | metabolic process | null |
| PBANKA_135060 | nuclear movement protein, putative | | | PF3D7_1336800 | nuclear movement protein, putative | microtubule | 0 | 1 | 1 | na | null | null | null |
| PBANKA_041630 | nucleoporin NUP100/NSP100, putative | -1.63 | 0.7 | PF3D7_0905100 | nucleoporin NUP100/NSP100, putative (NUP100) | RNA | 0 | 1 | 1 | na | null | null | null |
| PBANKA_010740 | ornithine aminotransferase, putative (OAT) | | | PF3D7_0608800 | ornithine aminotransferase (OAT) | AA metabolism | 0 | 1 | 1 | na | pyridoxal phosphate binding, transaminase activity | null | null |
| PBANKA_093170 | peptidyl-prolyl cis-trans isomerase, putative (CYP19C) | | | PF3D7_1116300 | peptidyl-prolyl cis-trans isomerase (CYP19C) | chaperone | 0 | 1 | 1 | na | peptidyl-prolyl cis-trans isomerase activity | null | null |
| PBANKA_051620 | phosphatidyl inositol glycan, class A, putative | | | PF3D7_1032400 | N-acetylglucosaminyl-phosphatidylinositol biosynthetic protein, putative | GPI | 0 | 1 | 1 | na | null | GPI anchor biosynthetic process, biosynthetic process | null |
| PBANKA_112810 | phospholipase (PL) | | | PF3D7_0629300 | phosphatidylcholine-sterol acyltransferase precursor, putative (PL) | utilization phospholipids | 0 | 1 | 1 | different in sporozoite and liver stage | phosphatidylcholine-sterol O-acyltransferase activity | lipid metabolic process | null |
| PBANKA_020460 | photosensitized INA-labeled protein 1, putative | | | PF3D7_0109000 | photosensitized INA-labeled protein 1, PhIL1, putative | gliding motility | 0 | 1 | 1 | na | null | null | null |
| PBANKA_122250 | plasmepsin X | | | PF3D7_0808200 | plasmepsin X | plasmepsin, protease | 0 | 1 | 1 | KO not successful | aspartic-type endopeptidase activity | proteolysis | null |
| PBANKA_000110 | Plasmodium exported protein, unknown function | | | | | | 0 | 1 | 1 | na | null | null | null |
| PBANKA_011090 | PP-loop family protein, putative | | | PF3D7_0612600 | PP-loop family protein, putative | translation | 0 | 1 | 1 | na | ATP binding | tRNA processing | null |
| PBANKA_111420 | ribose 5-phosphate epimerase, putative | -4.29 | 0.75 | PF3D7_0514600 | ribose 5-phosphate epimerase, putative | glycolysis | 0 | 1 | 1 | na | ribose-5-phosphate isomerase activity | pentose-phosphate shunt, non-oxidative branch | null |
| PBANKA_113630 | RNA-binding protein, putative | -1.66 | 0.71 | PF3D7_1360100 | RNA binding protein, putative | mRNA | 0 | 1 | 1 | na | nucleic acid binding | null | null |
| PBANKA_051240 | rrna-processing protein ebp2, putative (EBP2) | | | PF3D7_1028300 | nucleolar preribosomal assembly protein, putative | ribosome assembly | 0 | 1 | 1 | na | null | null | null |
| PBANKA_040780 | SECIS-binding protein 2, putative (SBP2) | | | PF3D7_0309700 | SECIS-binding protein 2, putative (SBP2); ribosomal protein I7ae, putative | ribosome | 0 | 1 | 1 | na | null | null | null |
| PBANKA_103780 | secreted ookinete adhesive protein (SOAP) | -3.26 | 0.75 | PF3D7_1404300 | secreted ookinete adhesive protein (SOAP) | adhesin | 0 | 1 | 1 | different in fertilization, ookinete, oocyst, sporozoite and liver stage | null | null | null |
| | | | | | | | | | | | | | |

null

| | | | | | | | _ | | | | | | |
|---------------|--|------------------|-------|-------------------|--|------------------|---|---|---|---|---|--|------|
| PBANKA_122200 | sin3 associated polypeptide p18 protein, putative | | | PF3D7_0711400 | sin3 associated polypeptide p18-like protein | chromatin | 0 | 1 | 1 | na | null | null | null |
| PBANKA_141880 | SNARE protein, putative | | | PF3D7_1320500 | SNARE protein, putative (SNAP23) | trafficking | 0 | 1 | 1 | na | null | null | null |
| PBANKA_134980 | sporozoite surface protein 2 (SSP2),thrombospondin-related anonymous | s protein (TRAP) |), S8 | PF3D7_1335900 | sporozoite surface protein 2 (TRAP) | gliding motility | 0 | 1 | 1 | different in sporozoite and liver stage | null | null | null |
| PBANKA_144500 | subpellicular microtubule protein 2, putative (SPM2) | -6.63 | 0.75 | PF3D7_1230300 | subpellicular microtubule protein 2, putative (SPM2) | gliding motility | 0 | 1 | 1 | na | null | null | null |
| PBANKA_112980 | Tetratricopeptide repeat protein, putative | -2.65 | 0.75 | PF3D7_0631000 | Tetratricopeptide repeat protein, putative | chaperone | 0 | 1 | 1 | not different from wild type | null | null | null |
| PBANKA_093230 | transcription factor with AP2 domain(s), putative (ApiAP2) | | | PF3D7_1115500 | transcription factor with AP2 domain(s), putative (ApiAP2) | transcription | 0 | 1 | 1 | na | sequence-specific DNA binding transcription factor activity | regulation of transcription, DNA-dependent | null |
| PBANKA_142390 | translation initiation factor SUI1, putative | | | PF3D7_0813600 | translation initiation factor SUI1, putative | translation | 0 | 1 | 1 | na | translation initiation factor activity | translational initiation | null |
| PBANKA_130380 | triose-phosphate isomerase, putative (TIM) | | | PF3D7_1439900 | triosephosphate isomerase (TIM) | glycolysis | 0 | 1 | 1 | na | triose-phosphate isomerase activity | metabolic process | null |
| PBANKA_062170 | type 2A phosphatase-associated protein 42, putative (TAP42) | | | PF3D7_0724200 | immunoglobulin-binding protein 1-related, putative | | 0 | 1 | 1 | na | null | regulation of signal transduction, response to biotic stimulus | null |
| PBANKA_114570 | U2 small nuclear ribonucleoprotein A, putative | | | PF3D7_1369700 | U1 small nuclear ribonucleoprotein a, putative | mRNA | 0 | 1 | 1 | na | protein binding | null | null |
| PBANKA_062010 | U3 snoRNA-associated small subunit rRNA processing protein, putative | | | PF3D7_0722600 | nucleolar rRNA processing protein, putative | RNA | 0 | 1 | 1 | na | null | null | null |
| PBANKA_082190 | ubiquitin conjugating enzyme, putative | | | PF3D7_0921000.1 | ubiquitin conjugating enzyme, putative | proteasome | 0 | 1 | 1 | na | small conjugating protein ligase activity | post-translational protein modification, regulation of protein metabolic process | null |
| PBANKA_080600 | ubiquitin-conjugating enzyme, putative | | | PF3D7_0319300 | ubiquitin conjugating enzyme, putative | proteasome | 0 | 1 | 1 | na | small conjugating protein ligase activity | post-translational protein modification, regulation of protein metabolic process | null |
| PBANKA_051770 | ubiquitin-conjugating enzyme, putative | | | PF3D7_1033900 | ubiquitin conjugating enzyme, putative | proteasome | 0 | 1 | 1 | na | ubiquitin-protein ligase activity | post-translational protein modification, protein ubiquitination, regulation of protein metabolic process | null |
| PBANKA_141390 | zinc finger (CCCH type) protein, putative | | | PF3D7_1315400 | zinc finger (CCCH type) protein, putative | transcription | 0 | 1 | 1 | na | null | null | null |
| PBANKA_090950 | AAA family ATPase, putative | | | PF3D7_1139500.1/2 | AAA family ATPase, putative | transporter | 1 | 0 | 1 | na | ATP binding, nucleoside-triphosphatase activity | null | null |
| PBANKA_051980 | acetyl-CoA transporter, putative | | | PF3D7_1036800 | acetyl-CoA transporter, putative | TCA | 1 | 0 | 1 | na | null | null | null |
| PBANKA_123990 | acyl-CoA synthetase, putative | | | PF3D7_0525100 | acyl-CoA synthetase (ACS10) | lipid metabolism | 1 | 0 | 1 | na | catalytic activity | metabolic process | null |
| PBANKA_123560 | adenosylhomocysteinase, putative (SAHH) | | | PF3D7_0520900 | S-adenosyl-L-homocysteine hydrolase (SAHH) | AA metabolism | 1 | 0 | 1 | na | adenosylhomocysteinase activity, binding | one-carbon metabolic process | null |
| PBANKA_103240 | alpha/beta hydrolase, putative | | | PF3D7_1410100 | alpha/beta hydrolase, putative | protease | 1 | 0 | 1 | na | null | null | null |
| PBANKA_123330 | ATP-dependent RNA helicase DDX23, putative (DDX23) | | | PF3D7_0518500 | ATP-dependent RNA helicase DDX23, putative (DDX23) | mRNA | 1 | 0 | 1 | na | ATP binding, ATP-dependent helicase activity, helicase activity, nucleic acid binding | null | null |
| PBANKA_010050 | BIR protein | | | | | BIR protein | 1 | 0 | 1 | na | null | null | null |
| PBANKA_060030 | BIR protein | | | | | BIR protein | 1 | 0 | 1 | na | null | null | null |
| PBANKA_146530 | BIR protein, pseudogene | | | | | BIR protein | 1 | 0 | 1 | na | null | null | null |
| PBANKA_145260 | calcyclin binding protein, putative | | | PF3D7_1238100 | calcyclin binding protein, putative | proteasome | 1 | 0 | 1 | na | null | null | null |
| PBANKA_094140 | caltractin (centrin), putative (CEN4) | | | PF3D7_1105500 | centrin-4 (CEN4) | chromatin | 1 | 0 | 1 | na | calcium ion binding | null | null |
| PBANKA_123430 | carbon catabolite repressor protein 4, putative (CCR4) | | | PF3D7_0519500 | carbon catabolite repressor protein 4, putative (CCR4) | mRNA | 1 | 0 | 1 | na | null | null | null |
| PBANKA_134430 | chromatin assembly factor 1 subunit, putative | | | PF3D7_1329300 | chromatin assembly factor 1 subunit, putative | chromatin | 1 | 0 | 1 | na | null | null | null |
| PBANKA_111460 | conserved Plasmodium protein, unknown function | | | PF3D7_0515000 | RNA recognition motif, putative | mRNA | 1 | 0 | 1 | na | nucleic acid binding | null | null |
| PBANKA_010150 | conserved Plasmodium protein, unknown function | -4.87 | 0.75 | PF3D7_0602700 | conserved Plasmodium protein, unknown function | unknown function | 1 | 0 | 1 | na | null | null | null |
| PBANKA_010180 | conserved Plasmodium protein, unknown function | -2.17 | 0.73 | PF3D7_0603000 | conserved Plasmodium protein, unknown function | unknown function | 1 | 0 | 1 | na | null | null | null |
| | | | | | | | | | | | | | |

| transcription corepressor activity | regulation of transcription from RNA polymerase II promoter | histone deacetylase complex |
|--|---|--------------------------------|
| SNAP receptor activity, protein binding | intracellular protein transport, regulation of vesicle fusion, vesicle-mediated transport | SNARE complex |
| null | cellular component movement, entry into host | cytoplasm, membrane, microneme |
| transferase activity | null | null |
| null | null | null |
| | | |
| | | |
| | | |
| | | |
| | | |
| null | null | null |
| | | |
| | | |
| | | |
| null | null | null |
| null | null | membrane |
| nuii | nuii | membrane |
| | | |
| null | null | membrane |
| | | |
| | | |
| null | null | integral to membrane |
| | | |
| null | null | membrane |
| | | |
| null | null | null |
| chromatin binding, histone binding, unfolded protein binding | chromatin assembly or disassembly | chromatin assembly complex |
| | | |
| null | null | null |
| null | null | cytoplasm, membrane |

| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_0603100 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | nucleic acid binding | null | nucleus |
|----------|--|-------|------|-------------------|---|-----|---|----|------|------|------|--|--|--------------------------------------|
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_0605200 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | acylphosphatase activity | null | null |
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_0612300 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | null | integral to membrane, membrane |
| PBANKA_0 | 11080 conserved Plasmodium protein, unknown function | | | PF3D7_0612500 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | null | null |
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_0613100 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | transport | integral to membrane, membrane |
| PBANKA_0 | 20500 conserved Plasmodium protein, unknown function | | | PF3D7_0108500 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | null | integral to membrane, membrane |
| PBANKA_0 | 20820 conserved Plasmodium protein, unknown function | | | PF3D7_0105100 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | null | null |
| PBANKA_0 | 21370 conserved Plasmodium protein, unknown function | | | PF3D7_0729600 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | null | null |
| PBANKA_0 | 30070 conserved Plasmodium protein, unknown function | -1.56 | 0.7 | PF3D7_0202600.1/2 | 2 conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | calcium ion binding | cytolysis, hemolysis by symbiont of host erythrocytes, pathogenesi | s extracellular region, membrane |
| PBANKA_0 | 30100 conserved Plasmodium protein, unknown function | | | PF3D7_0216700.1/2 | 2 conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | ATP binding, actin binding, calmodulin binding, motor activity | null | cytoplasm, myosin complex |
| PBANKA_0 | 30290 conserved Plasmodium protein, unknown function | | | PF3D7_0205200 | conserved Plasmodium protein, unknown function kinase | 1 0 | 1 | na | null | null | null | null | null | null |
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_0903600.1 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | ATP binding, actin binding, calmodulin binding, motor activity | null | cytoplasm, myosin complex |
| PBANKA_0 | conserved Plasmodium protein, unknown function | -2.15 | 0.7 | PF3D7_1019100 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | protein binding | null | cytoplasm |
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_1023500 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | null | null |
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_1024100 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | null | cytoplasm, nucleus |
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_1025800 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | null | null |
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_1026200 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | methyltransferase activity, nucleic acid binding | methylation | null |
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_1028800 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | null | integral to membrane, membrane |
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_1030200 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | nucleotide binding | null | cytoplasm, membrane, nucleus |
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | | unknown function | 1 0 | 1 | na | null | null | null | na | na | na |
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_0423300 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | GTP binding, GTPase activity, translation initiation factor activity | translation, translational initiation | apicoplast, cytoplasm, intracellular |
| PBANKA_0 | 60580 conserved Plasmodium protein, unknown function | -4.98 | 0.75 | PF3D7_1207300 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | DNA binding, hydrolase activity, transferase activity | null | apicoplast, cytoplasm |
| PBANKA_0 | 60750 conserved Plasmodium protein, unknown function | | | PF3D7_1209000 | conserved Plasmodium protein, unknown function mRNA | 1 0 | 1 | na | null | null | null | null | null | cytoplasm, nuclear periphery |
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_0717200 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | null | membrane |
| PBANKA_0 | conserved Plasmodium protein, unknown function | | | PF3D7_0717900 | thioredoxin-like protein redoxmeth | 1 0 | 1 | na | null | null | null | null | null | null |
| PBANKA_0 | conserved Plasmodium protein, unknown function | -2.18 | 0.75 | PF3D7_0718800 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | null | integral to membrane, membrane |
| PBANKA_0 | 70340 conserved Plasmodium protein, unknown function | | | | unknown function | 1 0 | 1 | na | null | null | null | na | na | na |
| PBANKA_0 | 71280 conserved Plasmodium protein, unknown function | | | PF3D7_0818000 | conserved Plasmodium protein, unknown function RNA | 1 0 | 1 | na | null | null | null | null | null | null |
| PBANKA_0 | 71770 conserved Plasmodium protein, unknown function | -1.8 | 0.73 | PF3D7_0415700 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | null | apicoplast, cytoplasm |
| PBANKA_0 | 80040 conserved Plasmodium protein, unknown function | | | | unknown function | 1 0 | 1 | na | null | null | null | | | |
| PBANKA_0 | 80720 conserved Plasmodium protein, unknown function | -4.01 | 0.75 | PF3D7_0318000 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na | null | null | null | null | null | apicoplast |
| | | | | | | | | | | | | | | |

| | | | | · | | | | | | | | | |
|--|-------|------|-----------------|--|-----|---|--|---|------|------|---|--|--|
| PBANKA_081550 conserved Plasmodium protein, unknown function | | | PF3D7_0914500 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | ATP binding | null | cytoplasm |
| PBANKA_082260 conserved Plasmodium protein, unknown function | | | PF3D7_0921700 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | nucleotide binding | null | apicoplast, cytoplasm |
| PBANKA_082460 conserved Plasmodium protein, unknown function | | | PF3D7_0923700 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | cytoplasm |
| PBANKA_082760 conserved Plasmodium protein, unknown function | | | PF3D7_0926800 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | receptor binding | cell differentiation, multicellular organismal development, signal transduc | tion extracellular region |
| PBANKA_083040 conserved Plasmodium protein, unknown function | -7.37 | 0.75 | PF3D7_0929600 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | different in fertilization, collinate, sporozoite and liver stage; (tagged) also different in occyst null | r | null | null | null | null | null |
| PBANKA_083500 conserved Plasmodium protein, unknown function | | | PF3D7_0934200 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | structural constituent of ribosome | translation | intracellular, ribosome, small ribosomal subunit |
| PBANKA_083570 conserved Plasmodium protein, unknown function | | | PF3D7_0934900 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | membrane |
| PBANKA_090800 conserved Plasmodium protein, unknown function | | | PF3D7_1141000 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | cytoplasm |
| PBANKA_091290 conserved Plasmodium protein, unknown function | | | | unknown function | 1 0 | 1 | na null | r | null | null | null | null | null |
| PBANKA_093750 conserved Plasmodium protein, unknown function | -3.2 | 0.75 | PF3D7_1110100 | conserved Plasmodium protein, unknown function lipid binding | 1 0 | 1 | na null | r | null | null | calcium ion binding, diacylglycerol binding, protein binding, zinc ion bindi | ing intracellular signal transduction | integral to membrane, plasma membrane |
| PBANKA_100350 conserved Plasmodium protein, unknown function | | | PF3D7_0405800 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | null |
| PBANKA_100500 conserved Plasmodium protein, unknown function | | | PF3D7_0407400 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | ATP binding | null | null |
| PBANKA_100560 conserved Plasmodium protein, unknown function | -5.22 | 0.75 | PF3D7_0408000 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | null |
| PBANKA_100790 conserved Plasmodium protein, unknown function | -2.55 | 0.74 | PF3D7_0410700 | GTPase, putative translation | 1 0 | 1 | na null | r | null | null | ATP binding | null | cytoplasm |
| PBANKA_101270 conserved Plasmodium protein, unknown function | | | PF3D7_1432200 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | cytoplasm |
| PBANKA_101280 conserved Plasmodium protein, unknown function | | | PF3D7_1432100 | conserved protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | voltage-gated anion channel activity | anion transport | external side of mitochondrial outer membrane, mitochondrial outer membrane, mitochondrion |
| PBANKA_101730 conserved Plasmodium protein, unknown function | -2.02 | 0.71 | PF3D7_1427300 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | ATP binding, actin binding, calmodulin binding | null | cytoplasm, myosin complex |
| PBANKA_101920 conserved Plasmodium protein, unknown function | | | PF3D7_1425300 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | cytoplasm |
| PBANKA_102480 conserved Plasmodium protein, unknown function | | | PF3D7_1417900 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | null |
| PBANKA_103180 conserved Plasmodium protein, unknown function | | | PF3D7_1410700 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | apicoplast |
| PBANKA_103330 conserved Plasmodium protein, unknown function | | | PF3D7_1409200.2 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | apicoplast |
| PBANKA_103970 conserved Plasmodium protein, unknown function | | | PF3D7_1402200 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | ATP binding, glutamate-tRNA ligase activity | glutamyl-tRNA aminoacylation, translation | cytoplasm, membrane |
| PBANKA_110210 conserved Plasmodium protein, unknown function | -4.07 | 0.75 | PF3D7_0502300 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | apicoplast |
| PBANKA_110530 conserved Plasmodium protein, unknown function | | | PF3D7_0505700 | conserved Plasmodium membrane protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | ATP binding, actin binding, calmodulin binding, motor activity | null | cytoplasm, myosin complex |
| PBANKA_111690 conserved Plasmodium protein, unknown function | | | PF3D7_0617300 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | 4 iron, 4 sulfur cluster binding, electron carrier activity, iron ion binding, molybdenum ion binding, nitrate reductase activity | Mo-molybdopterin cofactor biosynthetic process, nitrate assimilation, oxidation-reduction process, trans | port periplasmic space |
| PBANKA_112460 conserved Plasmodium protein, unknown function | | | PF3D7_0625700 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | null |
| PBANKA_113070 conserved Plasmodium protein, unknown function | | | PF3D7_1354100 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | null |
| PBANKA_113130 conserved Plasmodium protein, unknown function | | | PF3D7_1354700 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | GTP binding, GTPase activity, translation initiation factor activity | translation, translational initiation | intracellular |
| PBANKA_113800 conserved Plasmodium protein, unknown function | | | PF3D7_1362000 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | null | null | null |
| PBANKA_113950 conserved Plasmodium protein, unknown function | | | PF3D7_1363500 | DNase I-like protein, putative DNA repair | 1 0 | 1 | na null | r | null | null | RNA binding, exonuclease activity, magnesium ion binding, nuclease activity, poly(A)-specific ribonuclease activity, protein bin | regulation of transcription, DNA-dependent, transcription, DNA-depend | ent cytoplasm, nucleus |
| PBANKA_120180 conserved Plasmodium protein, unknown function | -3.57 | 0.75 | PF3D7_1003400 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | na null | r | null | null | ATP binding, actin binding, calmodulin binding, motor activity | null | membrane, myosin complex |
| | | | | | | | | | | | | | |

| PBANKA_120350 | conserved Plasmodium protein, unknown function | | | PF3D7_1005300 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
|---------------|--|-------|------|---------------|--|-----|---|-----------|--|----------------|-------------------------------|-----|-----------------------------------|
| PBANKA_121060 | conserved Plasmodium protein, unknown function | | | PF3D7_1012200 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
| PBANKA_121330 | conserved Plasmodium protein, unknown function | -2.24 | 0.7 | PF3D7_1014900 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | ATP binding, actin binding, ca |
| PBANKA_121490 | conserved Plasmodium protein, unknown function | | | PF3D7_0323600 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
| PBANKA_122590 | conserved Plasmodium protein, unknown function | -2.99 | 0.75 | PF3D7_0804700 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
| PBANKA_123500 | conserved Plasmodium protein, unknown function | | | PF3D7_0520200 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | (+)-abscisic acid 8'-hydroxylas |
| PBANKA_123830 | conserved Plasmodium protein, unknown function | | | PF3D7_0523500 | outer arm dynein lc3, putative dynein/kinesin | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
| PBANKA_124440 | conserved Plasmodium protein, unknown function | | | PF3D7_0530000 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
| PBANKA_130700 | conserved Plasmodium protein, unknown function | | | PF3D7_1443100 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
| PBANKA_131530 | conserved Plasmodium protein, unknown function | -4.48 | 0.75 | PF3D7_1451600 | LCCL-like protein (FNPA) adhesin | 1 0 | 1 | 1 differe | erent in occyst and sporozoite; (tagged) different in fertilization and ookinete | null | nul | ıll | null |
| PBANKA_131590 | conserved Plasmodium protein, unknown function | | | PF3D7_1452200 | aminomethyltransferase, putative post-translational processing | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
| PBANKA_132250 | conserved Plasmodium protein, unknown function | | | PF3D7_1458800 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
| PBANKA_133110 | conserved Plasmodium protein, unknown function | | | PF3D7_1467800 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
| PBANKA_134250 | conserved Plasmodium protein, unknown function | -5.93 | 0.75 | PF3D7_1327300 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | ATP binding, actin binding, ca |
| PBANKA_135310 | conserved Plasmodium protein, unknown function | | | PF3D7_1339700 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | ATP binding, actin binding, ca |
| PBANKA_135360 | conserved Plasmodium protein, unknown function | | | PF3D7_1340200 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
| PBANKA_135950 | conserved Plasmodium protein, unknown function | -1.79 | 0.7 | PF3D7_1346600 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | structural molecule activity |
| PBANKA_140520 | conserved Plasmodium protein, unknown function | | | PF3D7_1306700 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | metal ion binding, nucleotide bin |
| PBANKA_141720 | conserved Plasmodium protein, unknown function | -6.69 | 0.75 | PF3D7_1318700 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | ATP binding |
| PBANKA_142080 | conserved Plasmodium protein, unknown function | -3.66 | 0.75 | PF3D7_0714200 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | ATP binding, protein binding |
| PBANKA_142140 | conserved Plasmodium protein, unknown function | | | PF3D7_0715100 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | protein binding |
| PBANKA_142150 | conserved Plasmodium protein, unknown function | | | | unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | na |
| PBANKA_142300 | conserved Plasmodium protein, unknown function | | | PF3D7_0814500 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
| PBANKA_144710 | conserved Plasmodium protein, unknown function | | | PF3D7_1232400 | conserved protein, unknown function protein-protein interaction | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
| PBANKA_145080 | conserved Plasmodium protein, unknown function | | | PF3D7_1236200 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | null |
| PBANKA_145110 | conserved Plasmodium protein, unknown function | -6.17 | 0.75 | PF3D7_1236500 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | null | nul | ıll | 3-dehydroquinate dehydratas |
| PBANKA_101760 | conserved Plasmodium protein, unknown function | | | PF3D7_1427000 | conserved Plasmodium protein, unknown function unknown function | 1 0 | 1 | 1 na | a null | protein fo | olding, protein transport nul | ıll | |
| PBANKA_130570 | conserved Plasmodium protein, unknown function | | | PF3D7_1441800 | SNF7 family protein, putative large tethering | 1 0 | 1 | 1 na | a null | protein tr | ransport nul | ıll | |
| PBANKA_092140 | conserved Plasmodium protein, unknown function | | | PF3D7_1127000 | protein phosphatase, putative phosphatase | 1 0 | 1 | 1 na | a phosphatase activity | dephosp | horylation nul | ıll | |
| PBANKA_140330 | conserved Plasmodium protein, unknown function | | | | unknown function | 1 0 | 1 | 1 na | a phosphatase activity | null | nul | ıll | |
| PBANKA_010580 | conserved Plasmodium protein, unknown function | | | PF3D7_0607200 | conserved Plasmodium membrane protein, unknown function unknown function | 1 0 | 1 | 1 na | a protein binding, zinc ion | n binding null | nul | ıll | |
| | | | | | | | | | | | | | |

| | ·· · | |
|--|---|-------------------------------------|
| | null | apicoplast, membrane, nucleus |
| ng, actin binding, calmodulin binding, motor activity | null | cytoplasm, membrane, myosin complex |
| | null | null |
| | null | null |
| ic acid 8'-hydroxylase activity | null | null |
| | null | null |
| | null | nucleus |
| | null | nucleus |
| | null | null |
| | null | null |
| | null | cytoplasm, membrane |
| | null | null |
| ng, actin binding, calmodulin binding, motor activity | null | cytoplasm, myosin complex |
| ng, actin binding, calmodulin binding, motor activity | null | cytoplasm, myosin complex |
| | null | null |
| I molecule activity | null | null |
| binding, nucleotide binding, protein binding, transferase activity | regulation of transcription, DNA-dependent | cytoplasm, membrane, nucleus |
| ing | null | cytoplasm |
| ng, protein binding | null | null |
| inding | $regulation\ of\ transcription,\ DNA-dependent,\ transcription,\ DNA-dependent$ | cytoplasm, nucleus |
| | na | na |
| | null | null |
| | null | null |
| | null | cytoplasm |
| oquinate dehydratase activity | aromatic amino acid family biosynthetic process | null |
| | | |

apicoplast, membrane

| PBANKA_092500 | conserved Plasmodium protein, unknown function | | PF3D7_1123300 | conserved Plasmodium protein, unknown function | transcription | 1 | 0 | 1 | na | protein binding, zinc ion binding | null | null |
|---------------|--|------------|---------------|---|-----------------------|---|---|---|----|---|---|------|
| PBANKA_093270 | conserved Plasmodium protein, unknown function (PCAS_091160:exon:5) | | PF3D7_1115000 | conserved Plasmodium protein, unknown function | unknown function | 1 | 0 | 1 | na | null | null | null |
| PBANKA_041540 | conserved Plasmodium protein, unknown function, pseudogene | | | | unknown function | 0 | 1 | 1 | na | null | null | null |
| PBANKA_121800 | conserved Plasmodium protein, unknown function,pyrazinamidase/nicoti | -3.63 0.75 | PF3D7_0320500 | conserved Plasmodium protein, unknown function | unknown function | 1 | 0 | 1 | na | catalytic activity | metabolic process | null |
| PBANKA_072080 | conserved Plasmodium protein, unknwon function | | PF3D7_0418700 | conserved Plasmodium protein, unknown function | RNA | 0 | 1 | 1 | na | null | null | null |
| PBANKA_060460 | conserved protein, unknown function | -1.84 0.7 | PF3D7_1205900 | conserved protein, unknown function | unknown function | 1 | 0 | 1 | na | null | null | null |
| PBANKA_000370 | conserved rodent malaria protein, unknown function | | | | | 0 | 1 | 1 | na | null | null | null |
| PBANKA_000520 | conserved rodent malaria protein, unknown function | | | | unknown function | 1 | 0 | 1 | na | null | null | null |
| PBANKA_122220 | conserved rodent malaria protein, unknown function | | | | unknown function | 1 | 0 | 1 | na | null | null | null |
| PBANKA_142440 | CS domain protein, putative | | PF3D7_0813200 | CS domain protein, putative | adhesin | 1 | 0 | 1 | na | null | null | null |
| PBANKA_103230 | cytidine triphosphate synthetase, putative | | PF3D7_1410200 | cytidine triphosphate synthetase | pyrimidine metabolism | 1 | 0 | 1 | na | CTP synthase activity | pyrimidine nucleotide biosynthetic process | null |
| PBANKA_060080 | cytochrome c oxidase assembly protein COX19, putative (COX19) | -1.74 0.71 | PF3D7_1201800 | cytochrome c oxidase assembly protein, putative | redoxmeth | 1 | 0 | 1 | na | null | null | null |
| PBANKA_132630 | cytochrome c1 precursor, putative | | PF3D7_1462700 | cytochrome c1 precursor, putative | redoxmeth | 1 | 0 | 1 | na | electron carrier activity, heme binding, iron ion binding | null | null |
| PBANKA_031270 | DEAD/DEAH helicase, putative | -3.02 0.75 | PF3D7_0216000 | DEAD/DEAH box helicase, putative | RNA metabolism | 1 | 0 | 1 | na | ATP binding, DNA binding, helicase activity | null | null |
| PBANKA_041520 | developmental protein, putative | | PF3D7_0906100 | developmental protein, putative | large tethering | 1 | 0 | 1 | na | null | protein transport | null |
| PBANKA_102890 | diphthamide synthesis protein, putative | | PF3D7_1413800 | diphthamide synthesis protein, putative | ribosome assembly | 1 | 0 | 1 | na | null | null | null |
| PBANKA_130190 | DNA repair metallo-beta-lactamase protein, putative | | PF3D7_1474300 | DNA repair metallo-beta-lactamase protein, putative | e DNA repair | 1 | 0 | 1 | na | null | null | null |
| PBANKA_070810 | dynein light chain 1 (DLC1) | | PF3D7_0822500 | dynein light chain 1 (DLC1) | dynein/kinesin | 1 | 0 | 1 | na | protein binding | null | null |
| PBANKA_093360 | dynein light chain 2, putative | | PF3D7_1114000 | dynein light chain type 2, putative | dynein/kinesin | 1 | 0 | 1 | na | null | null | null |
| PBANKA_120260 | e3 ubiquitin-protein ligase, putative | -4.42 0.75 | PF3D7_1004300 | zinc finger, C3HC4 type, putative | proteasome | 1 | 0 | 1 | na | protein binding, zinc ion binding | null | null |
| PBANKA_123250 | eukaryotic translation initiation factor 3 subunit, putative | | PF3D7_0517700 | eukaryotic translation initiation factor, putative | translation | 1 | 0 | 1 | na | RNA binding, translation initiation factor activity | translational initiation | null |
| PBANKA_050430 | flagellar outer arm dynein-associated protein, putative | | PF3D7_1020100 | flagellar outer arm dynein-associated protein, putative | e dynein/kinesin | 1 | 0 | 1 | na | null | null | null |
| PBANKA_040470 | glutaredoxin 1, putative (GRX1) | | PF3D7_0306300 | glutaredoxin 1 (GRX1) | redoxmeth | 1 | 0 | 1 | na | electron carrier activity, protein disulfide oxidoreductase activity | cell redox homeostasis | null |
| PBANKA_041040 | glycogen synthase kinase 3 (GSK3) | | PF3D7_0312400 | glycogen synthase kinase 3 (GSK3) | kinase | 1 | 0 | 1 | na | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| PBANKA_121030 | heme oxygenase, putative (HO) | | PF3D7_1011900 | heme oxygenase (HO) | redoxmeth | 1 | 0 | 1 | na | null | null | null |
| PBANKA_140790 | HORMA domain protein, putative | -5.85 0.75 | PF3D7_1309400 | HORMA domain protein, putative | microtubule | 1 | 0 | 1 | na | null | mitosis | null |
| PBANKA_061090 | HSP40, subfamily A, putative | | PF3D7_1437900 | HSP40, subfamily A, putative | chaperone | 1 | 0 | 1 | na | heat shock protein binding, unfolded protein binding | protein folding | null |
| PBANKA_093600 | inositol-5 phosphatase, putative | | PF3D7_1111800 | peptidyl-prolyl cis-trans isomerase, putative | chaperone | 1 | 0 | 1 | na | null | null | null |
| PBANKA_142880 | iron-sulfur subunit of succinate dehydrogenase, putative | | PF3D7_1212800 | iron-sulfur subunit of succinate dehydrogenase | TCA | 1 | 0 | 1 | na | electron carrier activity, iron-sulfur cluster binding, oxidoreductase activity | r tricarboxylic acid cycle | null |
| PBANKA_135860 | isocitrate dehydrogenase (NADP), mitochondrial precursor, putative (IDH) | | PF3D7_1345700 | isocitrate dehydrogenase (NADP), mitochondrial precursor (IDH |) TCA | 1 | 0 | 1 | na | NAD binding, isocitrate dehydrogenase (NADP+) activity, magnesium ion binding | isocitrate metabolic process, oxidation-reduction process | null |
| PBANKA_120820 | kinesin-4, putative | | | | dynein/kinesin | 1 | 0 | 1 | na | ATP binding, microtubule motor activity | microtubule-based movement | null |
| | | | | | | | | | | | | |

| anchor binding, heparan sulfate proteoglycan binding | B cell activation involved in immune response, cell-cell adhesion, entry into host cell | null |
|--|---|--|
| | null | nucleus integral to membrane, membrane |
| | | |
| | na null | na null |
| | Tiuli | Tion |
| | na | na |
| | | |
| | | |
| | null | null |
| | Tiuli | Tiuli |
| | null | null |
| | | |
| | null | membrane |
| | | |
| binding | null | apicoplast |
| | | |
| | null | null |
| | | |

| PBANKA_011020 | | -1.89 | 0.73 | PF3D7_0611900 | Ism12, putative | mRNA | | na | null | null | null | null | null | null |
|---------------|---|--------|------|---------------|--|-------------------------------|-----|--|---|--|------|----------------|----------------------------|---|
| | lysophospholipase, putative | | | | | lipase | | na | null | null | null | na | na | na |
| PBANKA_090130 | merozoite adhesive erythrocytic binding protein (MAEBL) | | | | | unknown function | 0 1 | different in sporozoite and liver stag | ge null | null | null | binding | null | apical complex, membrane |
| PBANKA_120780 | metallopeptidase, putative | | | PF3D7_1009500 | metallopeptidase, putative | sumoylation | 0 1 | na | null | null | null | null | null | null |
| PBANKA_101020 | methionine aminopeptidase, type II, putative | | | PF3D7_1434600 | methionine aminopeptidase 2 (MetAP2) | post-translational processing | 0 1 | na | aminopeptidase activity, metalloexopeptidase activity | cellular process, proteolysis | null | | | |
| PBANKA_051580 | mitochondrial preribosomal assembly protein rimM precursor, putative | | | PF3D7_1032000 | mitochondrial preribosomal assembly protein rimM precursor, puta | tive ribosome mito | 0 1 | na | null | rRNA processing | null | | | |
| PBANKA_020440 | mRNA cleavage factor-like protein, putative | | | PF3D7_0109200 | mRNA cleavage factor-like protein, putative | mRNA | 0 1 | na | null | null | null | null | null | null |
| PBANKA_124070 | NIMA related kinase 2 (NEK2) | | | PF3D7_0525900 | NIMA related kinase 2 (NEK2) | kinase | 0 1 | different in fertilization, ookinete and ooc | yst ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | ty protein phosphorylation | null | | | |
| PBANKA_080560 | O-sialoglycoprotein endopeptidase, putative | | | PF3D7_0708300 | O-sialoglycoprotein endopeptidase, putative | protease | 0 1 | KO not successful | ATP binding, protein kinase activity | protein phosphorylation | null | | | |
| PBANKA_130030 | Pb-fam-1 protein, fragment | | | | | Pb-fam-1 protein | 0 1 | na | null | null | null | na | na | na |
| PBANKA_100730 | peptide chain release factor 2, putative | | | PF3D7_0409700 | peptide chain release factor 2, putative | translation | 0 1 | na | translation release factor activity | translational termination | null | | | |
| PBANKA_080930 | peptide deformylase, putative | | | PF3D7_0907900 | peptide deformylase (PDF) | protease | 0 1 | na | iron ion binding, peptide deformylase activity | translation | null | | | |
| PBANKA_093580 | peptidyl-prolyl cis-trans isomerase, putative | | | PF3D7_1113700 | glyoxalase I (GloI) | redoxmeth | 0 1 | na | null | protein folding | null | | | |
| PBANKA_060140 | peptidyl-prolyl cis-trans isomerase, putative (CYP26) | | | PF3D7_1202400 | peptidyl-prolyl cis-trans isomerase (CYP26) | chaperone | 0 1 | na | peptidyl-prolyl cis-trans isomerase activity | null | null | | | |
| PBANKA_100480 | peptidyl-tRNA hydrolase, putative | | | PF3D7_0407200 | peptidyl-tRNA hydrolase PTH2, putative | translation | 0 1 | na | aminoacyl-tRNA hydrolase activity | translation | null | | | |
| PBANKA_112870 | phosphatidylinositol/phosphatidylcholine transfer protein, putative | | | PF3D7_0629900 | sec14-like cytosolic factor or phosphatidylinositol/phosphatidylcholine transfer protein, pu | trafficking | 0 1 | na | null | null | null | binding | vesicle-mediated transport | Golgi apparatus |
| PBANKA_101450 | plasmepsin IX, putative | | | PF3D7_1430200 | plasmepsin IX | plasmepsin, protease | 0 1 | na | aspartic-type endopeptidase activity | proteolysis | null | | | |
| PBANKA_031610 | Plasmodium exported protein, unknown function | | | | | unknown function | 0 1 | na | null | null | null | | | |
| PBANKA_000590 | Plasmodium exported protein, unknown function, pseudogene | | | | | unknown function | 0 1 | na | null | null | null | | | |
| PBANKA_133720 | protein transport protein sft2, putative | | | PF3D7_1321800 | SFT2-like protein, putative | transporter, trafficking | 0 1 | na | null | null | null | null | null | membrane |
| PBANKA_131720 | pyridine nucleotide transhydrogenase, putative | -3.14 | 0.71 | PF3D7_1453500 | pyridine nucleotide transhydrogenase, putative | pyrimidine metabolism | 0 1 | na | NAD(P)+ transhydrogenase activity, binding, electron carrier activity | ity metabolic process, oxidation-reduction process | null | | | |
| PBANKA_121020 | QF122 antigen, putative | -3.15 | 0.75 | PF3D7_1011800 | QF122 antigen | mRNA | 0 1 | na | RNA binding | null | null | | | |
| PBANKA_092470 | RAP protein, putative | | | PF3D7_1123600 | RAP protein, putative | mRNA | 0 1 | na | null | null | null | null | null | null |
| PBANKA_040100 | regulatory protein, putative | | | PF3D7_0302000 | golgi organization and biogenesis factor, putativ | e | 0 1 | na | null | null | null | null | null | null |
| PBANKA_000160 | reticulocyte binding protein, putative, fragment | | | | | | 0 1 | na | null | null | null | | | |
| PBANKA_091460 | RNA (uracil-5-)methyltransferase, putative | -4.16 | 0.75 | PF3D7_1133800 | RNA (uracil-5-)methyltransferase, putative | RNA | 0 1 | na | RNA methyltransferase activity | RNA processing | null | | | |
| PBANKA_061370 | RNA binding protein, putative | | | PF3D7_0716000 | RNA binding protein, putative | mRNA | 0 1 | na | nucleic acid binding | null | null | | | |
| PBANKA_081770 | RNA-binding protein musashi, putative (HoMu) | -1.75 | 0.71 | PF3D7_0916700 | RNA-binding protein musashi, putative (HoMu) | mRNA | 0 1 | na | nucleic acid binding | null | null | | | |
| PBANKA_141920 | RNA-binding protein, putative | -1.95 | 0.73 | PF3D7_1320900 | RNA-binding protein, putative | mRNA | 0 1 | na | nucleic acid binding | null | null | | | |
| PBANKA_113570 | RNA-binding protein, putative | | | PF3D7_1359400 | rRNA associated RNA binding protein, putative | RNA | 0 1 | na | RNA binding, protein binding | null | null | | | |
| PBANKA_081900 | secreted acid phosphatase, putative,glideosome-associated protein 50, | , -4.4 | 0.75 | PF3D7_0918000 | glideosome-associated protein 50,secreted acid phosphatase (GAP) | 50) gliding motility | 0 1 | na | null | null | null | protein anchor | null | endoplasmic reticulum, inner membrane complex |
| | | | | | | | | | | | | | | |

| PBANKA_110300 | Ser/Arg-rich splicing factor, putative | | | PF3D7_0503300 | Ser/Arg-rich splicing factor, putative | mRNA | 1 | 0 | 1 r | na | nucleic acid binding | null | null |
|---------------|--|-------|------|---------------|--|------------------------------|---|---|-----|--|--|---|--|
| PBANKA_030510 | serine repeat antigen 1 (SERA1) | | | PF3D7_0207400 | serine repeat antigen 7 (SERA7) | protease | 1 | 0 | 1 r | not different from wild type | cysteine-type peptidase activity | proteolysis | null |
| PBANKA_131370 | serine/threonine protein kinase, putative | | | PF3D7_1450000 | serine/threonine protein kinase, putative | kinase | 1 | 0 | 1 F | KO not successful | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| PBANKA_141360 | serine/threonine protein kinase, putative | | | PF3D7_1315100 | serine/threonine protein kinase (PK9) | kinase | 1 | 0 | 1 F | KO not successful | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| PBANKA_093250 | SET domain protein, putative | -6.41 | 0.75 | PF3D7_1115200 | SET domain protein, putative (SET7) | chromatin modifying prtoeins | 1 | 0 | 1 r | na | null | null | null |
| PBANKA_136300 | small nuclear ribonucleoprotein E, putative (SNRPE) | | | PF3D7_1350200 | small nuclear ribonucleoprotein E, putative (SNRPE) | mRNA | 1 | 0 | 1 r | na | null | null | null |
| PBANKA_081100 | small nuclear ribonucleoprotein Sm D3, putative (SNRPD3) | | | PF3D7_0909800 | small nuclear ribonucleoprotein Sm D3, putative (SNRPD3 | B) mRNA | 1 | 0 | 1 r | na | null | null | null |
| PBANKA_102790 | small nuclear ribonucleoprotein-associated protein B, putative (SNRPB) | | | PF3D7_1414800 | small nuclear ribonucleoprotein-associated protein B, putative (SNRPB) | mRNA | 1 | 0 | 1 r | na | null | null | null |
| PBANKA_102320 | splicing factor 3B subunit 4, putative (SF3B4) | | | PF3D7_1420000 | splicing factor 3B subunit 4, putative (SF3B4) | mRNA | 1 | 0 | 1 r | na | nucleic acid binding | null | null |
| PBANKA_113680 | SUMO ligase, putative | | | PF3D7_1360700 | SUMO ligase, putative | sumoylation | 1 | 0 | 1 r | na | zinc ion binding | null | null |
| PBANKA_112240 | superoxide dismutase, putative | | | PF3D7_0623500 | superoxide dismutase (SOD2) | redoxmeth | 1 | 0 | 1 r | na | metal ion binding, superoxide dismutase activity | oxidation-reduction process, superoxide metabolic process | null |
| PBANKA_101850 | transcription factor 3b, putative | | | PF3D7_1426100 | basic transcription factor 3b, putative | transcription | 1 | 0 | 1 r | na | null | null | null |
| PBANKA_130580 | transcription factor TFIIH complex subunit Tfb5, putative | | | PF3D7_1441900 | transcription factor TFIIH complex subunit Tfb5, putative | e transcription | 1 | 0 | 1 r | na | DNA binding | nucleotide-excision repair | null |
| PBANKA_131970 | transcription factor with AP2 domain(s), putative (ApiAP2) | | | PF3D7_1456000 | transcription factor with AP2 domain(s) (ApiAP2) | transcription | 1 | 0 | 1 r | na | sequence-specific DNA binding transcription factor activity | regulation of transcription, DNA-dependent | null |
| PBANKA_123690 | transcription initiation factor tfiid subunit 10, putative (TAF10) | | | PF3D7_0522200 | conserved Plasmodium protein, unknown function | unknown function | 1 | 0 | 1 r | na | null | null | null |
| PBANKA_134160 | translation initiation factor EIF-2B gamma subunit, putative | | | PF3D7_1326400 | translation initiation factor EIF-2B gamma subunit, putative | e translation | 1 | 0 | 1 r | na | transferase activity | null | null |
| PBANKA_093990 | U6 snRNA-associated Sm-like protein LSm4, putative (LSM4) | | | PF3D7_1107000 | U6 snRNA-associated Sm-like protein LSm4, putative (LSM4) | mRNA | 1 | 0 | 1 r | na | null | null | null |
| PBANKA_060760 | U6 snRNA-associated Sm-like protein LSm7, putative (LSM7) | | | PF3D7_1209200 | U6 snRNA-associated Sm-like protein LSm7, putative (LSM7) | mRNA | 1 | 0 | 1 r | na | null | null | null |
| PBANKA_101770 | ubiquinol-cytochrome c reductase hinge protein, putative | | | PF3D7_1426900 | ubiquinol-cytochrome c reductase hinge protein, putative | e redoxmeth | 1 | 0 | 1 r | na | ubiquinol-cytochrome-c reductase activity | mitochondrial electron transport, ubiquinol to cytochrome c | null |
| PBANKA_124200 | ubiquitin carboxyl-terminal hydrolase, putative | | | PF3D7_0527200 | ubiquitin carboxyl-terminal hydrolase, putative | proteasome | 1 | 0 | 1 r | na | ubiquitin thiolesterase activity | ubiquitin-dependent protein catabolic process | null |
| PBANKA_091060 | ubiquitin-related modifier 1, putative (URM1) | | | PF3D7_1138100 | ubiquitin related modifier homologue | proteasome | 1 | 0 | 1 r | na | null | null | null |
| PBANKA_102290 | zinc finger protein, putative | | | PF3D7_1420300 | zinc finger protein, putative | transporter | 1 | 0 | 1 r | na | null | null | null |
| PBANKA_123760 | zinc finger protein, putative, fragment | -1.78 | 0.7 | PF3D7_0522900 | zinc finger protein, putative | RNA binding | 1 | 0 | 1 r | na | nucleic acid binding, zinc ion binding | null | null |
| PBANKA_071430 | small heat shock protein HSP20 (HSP20) | -4.76 | 0.75 | PF3D7_0816500 | small heat shock protein HSP20, putative (HSP20) |) chaperone | 0 | 1 | 1 a | ifferent in fertilization, ooklinete, sporozoite and liver stage; (tagged) different in oocyst too | null | null | plasma membrane |
| PBANKA_123400 | vacuolar ATP synthetase, putative | | | PF3D7_0519200 | vacuolar ATP synthetase | vacuolar ATP synthase | 0 | 1 | 1 r | na | hydrogen ion transmembrane transporter activity | ATP synthesis coupled proton transport | proton-transporting V-type ATPase, V0 domain |
| PBANKA_090890 | vacuolar ATP synthase subunit f, putative | | | PF3D7_1140100 | vacuolar ATP synthase subunit f, putative | vacuolar ATP synthase | 0 | 1 | 1 r | na | hydrogen ion transporting ATP synthase activity, rotational mechanism, proton-transporting ATPase activity, rotational mechanism | ATP synthesis coupled proton transport | proton-transporting V-type ATPase, V1 domain |
| PBANKA_093200 | rhoptry neck protein 4, putative (RON4) | | | PF3D7_1116000 | rhoptry neck protein 4 (RON4) | | 1 | 0 | 1 (| mutated) different in sporozoite and liver stage | null | null | rhoptry |
| PBANKA_133830 | 60S ribosomal protein L6, putative | | | PF3D7_1323100 | 60S ribosomal protein L6, putative | ribosome | 1 | 0 | 1 r | na | rRNA binding, structural constituent of ribosome | translation | ribosome |
| PBANKA_144620 | DIM1 protein homolog, putative | | | PF3D7_1231500 | DIM1 protein homolog, putative | ribosome assembly | 0 | 1 | 1 r | na | null | mitosis | spliceosomal complex |
| PBANKA_131780 | tRNA intron endonuclease, putative | | | PF3D7_1454100 | tRNA intron endonuclease, putative | translation | 0 | 1 | 1 r | na | tRNA-intron endonuclease activity | tRNA splicing, via endonucleolytic cleavage and ligation | tRNA-intron endonuclease complex |
| PBANKA_000330 | BIR protein | | | | | BIR protein | 0 | 0 | 0 r | na | null | null | null |
| | | | | | | | | | | | | | |

| PBANK | A_000340 | BIR protein | | | | BIR protein | 0 | 0 | 0 | na | null | null | null |
|-------|----------|---|------|------|--|---|---|---|---|--|--|---|--|
| PBANK | A_000380 | BIR protein | | | | BIR protein | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_000490 | conserved rodent malaria protein, unknown function | | | | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_000570 | BIR protein | | | | BIR protein | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_010060 | schizont membrane associated cytoadherence protein (SMAC) | | | | | 0 | 0 | 0 | na | null | cytoadherence to microvasculature, mediated by symbiont protein | host cell cytoplasm |
| PBANK | A_010070 | Plasmodium exported protein, unknown function | | | | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_010080 | conserved Plasmodium protein, unknown function | | | PF3D7_0602000 conserved Plasmodium protein, unknown for | unction microtubule | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_010270 | conserved Plasmodium protein, unknown function | | | PF3D7_0603900 conserved Plasmodium protein, unknown for | unction unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_010410 | aurora-related kinase 1, putative (ARK1) | | | PF3D7_0605300 serine/threonine protein kinase (ARK1) | kinase | 0 | 0 | 0 | KO not successful | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| PBANK | A_010470 | long chain polyunsaturated fatty acid elongation enzyme, putative | | | PF3D7_0605900 long chain polyunsaturated fatty acid elongation enzyme, | putative fatty acids synth | 0 | 0 | 0 | na | null | null | integral to membrane |
| PBANK | A_010560 | translation initiation factor IF-2, putative | | | PF3D7_0607000 translation initiation factor IF-2, putative | translation | 0 | 0 | 0 | na | GTP binding, GTPase activity | null | null |
| PBANK | A_010860 | Pre-mRNA-splicing factor SLU7, putative (SLU7) | | | PF3D7_0610100 Pre-mRNA-splicing factor SLU7, putative (| SLU7) mRNA | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_010870 | conserved Plasmodium protein, unknown function | -1.9 | 0.73 | PF3D7_0610200 conserved Plasmodium protein, unknown for | unction mRNA | 0 | 0 | 0 | na | nucleic acid binding | mRNA processing | null |
| PBANK | A_011000 | 60S ribosomal protein L39, putative | | | PF3D7_0611700 60S ribosomal protein L39, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANK | A_011030 | conserved Plasmodium protein, unknown function | | | PF3D7_0612000 conserved Plasmodium protein, unknown for | unction unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_011050 | leucine-rich repeat protein (LRR6) | -2.6 | 0.72 | PF3D7_0612200 leucine-rich repeat protein (LRR6) | kinase | 0 | 0 | 0 | na | protein binding | null | null |
| PBANK | A_011130 | conserved Plasmodium protein, unknown function | | | PF3D7_0613000.1/2 conserved Plasmodium protein, unknown for | unction unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_020020 | Pb-fam-1 protein | | | | Pb-fam-1 protein | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_020090 | conserved rodent malaria protein, unknown function | | | | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_020160 | early transcribed membrane protein (ETRAMP) | | | | early transcribed membrane protein (ETRAMP) | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_020190 | conserved Plasmodium protein, unknown function | | | PF3D7_0111900 conserved Plasmodium protein, unknown for | unction unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_020420 | tubulin-specific chaperone a, putative | | | PF3D7_0109400 tubulin-specific chaperone a, putative | chaperone | 0 | 0 | 0 | na | unfolded protein binding | tubulin complex assembly | microtubule |
| PBANK | A_020470 | conserved Plasmodium protein, unknown function | | | PF3D7_0108800 conserved Plasmodium protein, unknown for | unction transporter | 0 | 0 | 0 | na | binding | transport | membrane |
| PBANK | A_020510 | mitochondrial carrier protein, putative | | | PF3D7_0108400.1/2 mitochondrial carrier protein, putative | transporter | 0 | 0 | 0 | na | binding | transport | mitochondrial inner membrane |
| PBANK | A_020580 | serine/threonine protein kinase, putative (IK2) | | | PF3D7_0107600 serine/threonine protein kinase, putative, E | IK2 kinase | 0 | 0 | 0 | different in fertilization, ookinete, oocyst, sporozoite and liver stage (contradictory results) | ATP binding, protein serine/threonine kinase activity | protein phosphorylation | null |
| PBANK | A_020630 | centrin, putative | | | PF3D7_0107000 centrin-1 (CEN1) | chromatin | 0 | 0 | 0 | na | calcium ion binding | null | null |
| PBANK | A_020650 | Rab5c, GTPase, putative | | | PF3D7_0106800 Rab GTPase 5c (RAB5c) | trafficking | 0 | 0 | 0 | na | GTP binding, GTPase activity, protein binding | intracellular protein transport, nucleocytoplasmic transport, small GTPase mediated signal transduction | intracellular |
| PBANK | A_020720 | vacuolar ATP synthase subunit c, putative | | | PF3D7_0106100 vacuolar ATP synthase subunit c, putative | vacuolar ATP synthase | 0 | 0 | 0 | na | hydrolase activity, acting on acid anhydrides, catalyzing transmembrane movement of substances | ATP synthesis coupled proton transport | proton-transporting V-type ATPase, V1 domain |
| PBANK | A_020760 | asparagine-rich antigen, putative | | | | | 0 | 0 | 0 | na | null | null | null |
| PBANK | A_020930 | actin-related protein (ARP1) | | | PF3D7_0103800 actin-related protein (ARP1) | gliding motility | 0 | 0 | 0 | KO not successful | protein binding | null | null |
| PBANK | A_021120 | conserved Plasmodium protein, unknown function | | | PF3D7_0727100 conserved Plasmodium protein, unknown for | unction unknown function | 0 | 0 | 0 | na | null | null | null |
| | | | | | | | | | | | | | |

| PBANKA_021210 | eukaryotic translation initiation factor 2 alpha subunit, putative | PF3D7_0728000 | eukaryotic translation initiation factor 2 alpha subunit, putative | translation | 0 | 0 | 0 na | | RNA binding, translation initiation factor activity | translation | eukaryotic translation initiation factor 2 complex |
|---------------|---|---------------|---|---------------------------------|---|---|--|---|--|--|--|
| PBANKA_021300 | RNA-binding protein, putative | PF3D7_0728900 | RNA binding protein, putative | mRNA | 0 | 0 | 0 na | | nucleic acid binding | null | null |
| PBANKA_021310 | signal recognition particle subunit srp9, putative (SRP9) | PF3D7_0729000 | signal recognition particle SRP9 (SRP9) | post-translational modification | 0 | 0 | 0 na | | 7S RNA binding, protein binding | SRP-dependent cotranslational protein targeting to membrane, negative regulation of translational elongation | signal recognition particle |
| PBANKA_030060 | Plasmodium exported protein, unknown function | | | unknown function | 0 | 0 | 0 na | | null | null | null |
| PBANKA_030110 | conserved Plasmodium protein, unknown function -2.99 0.3 | PF3D7_0203200 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |
| PBANKA_030120 | ERCC1 nucleotide excision repair protein, putative | PF3D7_0203300 | ERCC1 nucleotide excision repair protein, putative | e DNA repair | 0 | 0 | 0 na | | damaged DNA binding, endonuclease activity | DNA repair | nucleus |
| PBANKA_030150 | nucleolar preribosomal assembly protein, putative | PF3D7_0203700 | nucleolar preribosomal assembly protein, putative | ribosome assembly | 0 | 0 | 0 na | | null | null | null |
| PBANKA_030250 | hexose transporter (HT) | PF3D7_0204700 | hexose transporter (HT) | glycolysis | 0 | 0 | O KID not auconsolut; (suggest) different in | threat in firstitusion, colones, coopst, sportscale and liver stage, (inustated) non-different from wild type | substrate-specific transmembrane transporter activity | transmembrane transport, transport | integral to membrane, membrane |
| PBANKA_030620 | phospholipase A2, putative | PF3D7_0209100 | phospholipase A2, putative | utilization phospholipids | 0 | 0 | 0 na | | null | lipid metabolic process | null |
| PBANKA_030680 | ATP-dependent RNA helicase UAP56, putative (UAP56) | PF3D7_0209800 | ATP-dependent RNA helicase UAP56 (UAP56) | mRNA | 0 | 0 | 0 na | | ATP binding, ATP-dependent helicase activity, helicase activity, nucleic acid binding | null | null |
| PBANKA_031020 | conserved Plasmodium protein, unknown function | PF3D7_0213300 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |
| PBANKA_031190 | RING zinc finger protein, putative | PF3D7_0215100 | RING zinc finger protein, putative | proteasome | 0 | 0 | 0 na | | protein binding, zinc ion binding | null | null |
| PBANKA_031240 | DNA-directed RNA polymerase II second largest subunit, putative | PF3D7_0215700 | DNA-directed RNA polymerase II second largest subunit, putative | e transcription | 0 | 0 | 0 na | | DNA binding, DNA-directed RNA polymerase activity, ribonucleoside binding | transcription, DNA-dependent | null |
| PBANKA_031250 | origin recognition complex subunit 5, putative (ORC5) | PF3D7_0215800 | origin recognition complex subunit 5 (ORC5) | DNA replication | 0 | 0 | 0 na | | null | null | null |
| PBANKA_031410 | conserved Plasmodium protein, unknown function | PF3D7_0217400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |
| PBANKA_031420 | calcium dependent protein kinase 1 (CDPK1) | PF3D7_0217500 | calcium dependent protein kinase 1 (CDPK1) | kinase | 0 | 0 | O different in oocyst; (muta | (mutated) fertilization and ookinete; (tagged) also in sporozoite | ATP binding, calcium ion binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| PBANKA_031450 | 40S ribosomal protein S26e, putative | PF3D7_0217800 | 40S ribosomal protein S26e, putative | ribosome | 0 | 0 | 0 na | | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_031560 | 40S ribosomal protein S30, putative | PF3D7_0219200 | 40S ribosomal protein S30, putative | ribosome | 0 | 0 | 0 na | | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_031600 | replication factor C subunit 1, putative | PF3D7_0219600 | replication factor C subunit 1, putative | DNA replication | 0 | 0 | 0 na | | ATP binding, DNA clamp loader activity | DNA replication | DNA replication factor C complex, intracellular |
| PBANKA_031620 | Plasmodium exported protein, unknown function, fragment | | | unknown function | 0 | 0 | 0 na | | null | null | null |
| PBANKA_040040 | BIR protein, pseudogene | | | BIR protein | 0 | 0 | 0 na | | null | null | null |
| PBANKA_040230 | dihydrolipoamide acyltransferase, putative | PF3D7_0303700 | dihydrolipoamide acyltransferase, putative | glycolysis, TCA | 0 | 0 | 0 na | | transferase activity, transferring acyl groups | metabolic process | null |
| PBANKA_040300 | 60S ribosomal protein L44, putative | PF3D7_0304400 | 60S ribosomal protein L44, putative | ribosome | 0 | 0 | 0 na | | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_040340 | conserved Plasmodium protein, unknown function | PF3D7_0304900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |
| PBANKA_040410 | AP endonuclease (DNA-(apurinic or apyrimidinic site) lyase), putative | PF3D7_0305600 | AP endonuclease (DNA-[apurinic or apyrimidinic site] lyase), putative | DNA repair | 0 | 0 | 0 na | | nuclease activity | DNA repair | null |
| PBANKA_040440 | conserved Plasmodium protein, unknown function | PF3D7_0306000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |
| PBANKA_040530 | 40S ribosomal protein S23, putative | PF3D7_0306900 | 40S ribosomal protein S23, putative | ribosome | 0 | 0 | 0 na | | structural constituent of ribosome | translation | intracellular, ribosome, small ribosomal subunit |
| PBANKA_040550 | 60S ribosomal protein L7, putative | PF3D7_0307200 | 60S ribosomal protein L7, putative | ribosome | 0 | 0 | 0 na | | structural constituent of ribosome | translation | intracellular, large ribosomal subunit, ribosome |
| PBANKA_040570 | ATP-dependent Clp protease proteolytic subunit, putative (ClpP) | PF3D7_0307400 | ATP-dependent Clp protease proteolytic subunit (ClpP | chaperone | 0 | 0 | 0 na | | serine-type endopeptidase activity | proteolysis | apicoplast |
| PBANKA_040600 | conserved Plasmodium protein, unknown function | PF3D7_0307700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | DNA binding | null | null |
| PBANKA_040650 | TCP-1/cpn60 chaperonin, putative | PF3D7_0308200 | TCP-1/cpn60 chaperonin family, putative | chaperone | 0 | 0 | 0 na | | ATP binding, unfolded protein binding | protein folding | null |
| | | | | | | | | | | | |

| PBANKA_040770 | 60S acidic ribosomal protein P2, putative | | | PF3D7_0309600 | 60S acidic ribosomal protein P2, putative | ribosome | 0 | 0 | 0 | KO not successful | structural constituent of ribosome | translational elongation | intracellular, ribosome |
|---------------|--|-------|------|---------------|--|---|---|---|---|--|--|--|---|
| PBANKA_040930 | phosphatidylinositol 3- and 4-kinase, putative | | | PF3D7_0311300 | phosphatidylinositol 3- and 4-kinase, putative | inositol phosphate metabolism | 0 | 0 | 0 | na | phosphotransferase activity, alcohol group as acceptor | null | null |
| PBANKA_040940 | protein kinase, putative (PKRP) | | | PF3D7_0311400 | protein kinase, putative | kinase | 0 | 0 | 0 | different in fertilization, ookinete, oocyst, sporozoite and liver stage | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| PBANKA_041000 | conserved Plasmodium protein, unknown function | | | PF3D7_0312000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_041130 | conserved Plasmodium protein, unknown function | | | PF3D7_0313600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_041310 | conserved Plasmodium protein, unknown function | | | | | RNA | 0 | 0 | 0 | na | null | microtubule cytoskeleton organization | microtubule organizing center, spindle pole |
| PBANKA_041460 | 40S ribosomal protein S15A, putative | | | PF3D7_0316800 | 40S ribosomal protein S15A, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_041480 | arginase | | | PF3D7_0906500 | arginase | AA metabolism | 0 | 0 | 0 | na | arginase activity, metal ion binding | arginine metabolic process | null |
| PBANKA_041660 | replication protein A1, small fragment | | | PF3D7_0904800 | replication protein A1, small fragment | DNA repair | 0 | 0 | 0 | na | nucleic acid binding | null | null |
| PBANKA_041740 | GTPase activating protein, putative | -1.86 | 0.73 | PF3D7_0904000 | GTPase activator, putative | trafficking | 0 | 0 | 0 | na | Rab GTPase activator activity | regulation of Rab GTPase activity | intracellular |
| PBANKA_041750 | 60S ribosomal protein L32, putative | | | PF3D7_0903900 | 60S ribosomal protein L32, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_041770 | alpha tubulin 1 | | | PF3D7_0903700 | alpha tubulin 1 | gliding motility | 0 | 0 | 0 | KO not successful | GTP binding, GTPase activity, structural molecule activity | microtubule-based movement, protein polymerization | microtubule, protein complex |
| PBANKA_050070 | Pb-fam-1 protein | | | | | Pb-fam-1 protein | 0 | 0 | 0 | na | null | null | null |
| PBANKA_050110 | early transcripted membrane protein (SEP3) | | | | | early transcribed membrane protein (ETRAMP) | 0 | 0 | 0 | KO not successful; (tagged) fertilization, ookinete and oocyst | null | null | null |
| PBANKA_050170 | phosphomannomutase, putative | | | PF3D7_1017400 | phosphomannomutase, putative | mannose metabolism | 0 | 0 | 0 | na | phosphomannomutase activity | mannose biosynthetic process | cytoplasm |
| PBANKA_050250 | serine/threonine protein phosphatase, putative | -3.02 | 0.74 | PF3D7_1018200 | serine/threonine protein phosphatase, putative | phosphatase | 0 | 0 | 0 | na | hydrolase activity | null | null |
| PBANKA_050280 | conserved Plasmodium protein, unknown function | | | PF3D7_1018600 | conserved Plasmodium protein, unknown function | translation | 0 | 0 | 0 | na | null | null | null |
| PBANKA_050350 | conserved Plasmodium protein, unknown function | -2.08 | 0.71 | PF3D7_1019300 | zinc finger protein, putative | nucleic acid binding | 0 | 0 | 0 | na | nucleic acid binding, zinc ion binding | null | null |
| PBANKA_050360 | 60S ribosomal protein L30e, putative | | | PF3D7_1019400 | 60S ribosomal protein L30e, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_050440 | conserved Plasmodium protein, unknown function | | | PF3D7_1020200 | kinesin, putative | dynein/kinesin | 0 | 0 | 0 | na | null | null | null |
| PBANKA_050510 | ADP-ribosylation factor, putative (ARF1) | | | PF3D7_1020900 | ADP-ribosylation factor (ARF1) | trafficking | 0 | 0 | 0 | na | GTP binding | intracellular protein transport, small GTPase mediated signal transduction | intracellular |
| PBANKA_050540 | conserved Plasmodium protein, unknown function | | | PF3D7_1021200 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_050580 | deoxyribose-phosphate aldolase, putative | | | PF3D7_1021600 | deoxyribose-phosphate aldolase, putative | gliding motility | 0 | 0 | 0 | na | deoxyribose-phosphate aldolase activity | deoxyribonucleotide catabolic process | cytoplasm |
| PBANKA_050630 | conserved Plasmodium protein, unknown function | | | PF3D7_1022100 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_050660 | pre-mRNA splicing factor, putative | -2.25 | 0.75 | PF3D7_1022400 | pre-mRNA splicing factor, putative | mRNA | 0 | 0 | 0 | na | nucleic acid binding | null | null |
| PBANKA_050670 | citrate synthase, mitochondrial precursor, putative | | | PF3D7_1022500 | citrate synthase, mitochondrial precursor, putative | TCA | 0 | 0 | 0 | (tagged) different in oocyst and liver stage | transferase activity, transferring acyl groups, acyl groups converted into alkyl on transfer | cellular carbohydrate metabolic process | null |
| PBANKA_050810 | chromodomain-helicase-DNA-binding protein 1, putative (CHD1) | | | PF3D7_1023900 | chromodomain-helicase-DNA-binding protein 1 homolog, putative (CHD1) | chromatin | 0 | 0 | 0 | KO not successful | ATP binding, DNA binding, chromatin binding, helicase activity | chromatin assembly or disassembly | chromatin, nucleus |
| PBANKA_050860 | conserved Plasmodium protein, unknown function | | | PF3D7_1024400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_050920 | formin 2, putative | | | PF3D7_1025000 | formin 2, putative | gliding motility | 0 | 0 | 0 | na | null | null | null |
| PBANKA_050930 | glucosaminefructose-6-phosphate aminotransferase, putative | | | PF3D7_1025100 | glucosamine-fructose-6-phosphate aminotransferase, putative | chitin precursor | 0 | 0 | 0 | na | carbohydrate binding | carbohydrate metabolic process | null |
| PBANKA_051010 | conserved Plasmodium protein, unknown function | | | PF3D7_1025900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| | | | | | | | | | | | | | |

| PBANKA_051020 | conserved Plasmodium protein, unknown function | PF3D7_1026000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
|---------------|--|-----------------|---|---|---|---|---|--|--|---|--|
| PBANKA_051060 | conserved Plasmodium protein, unknown function | PF3D7_1026400 | CDH1, WD-repeat protein, putative | kinetochore | 0 | 0 | 0 | different in fertilization and ookinete | null | cytokinesis | nucleus |
| PBANKA_051080 | conserved Plasmodium protein, unknown function | none | | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_051090 | 40S ribosomal protein S2B, putative | PF3D7_1026800 | 40S ribosomal protein S2B, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome, small ribosomal subunit |
| PBANKA_051120 | u3 small nucleolar ribonucleoprotein protein mpp10, putative (MPP10) | PF3D7_1027100 | small subunit rRNA processing stabilizing factor, putative | RNA | 0 | 0 | 0 | na | null | null | null |
| PBANKA_051180 | centrin-3, putative | PF3D7_1027700.2 | centrin-3 | chromatin | 0 | 0 | 0 | na | calcium ion binding | null | null |
| PBANKA_051190 | 60S ribosomal protein L3, putative | PF3D7_1027800 | 60S ribosomal protein L3, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_051270 | conserved Plasmodium protein, unknown function | PF3D7_1028600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_051280 | merozoite TRAP-like protein, putative (MTRAP) | PF3D7_1028700 | merozoite TRAP-like protein (MTRAP) | contaminant | 0 | 0 | 0 | na | null | null | null |
| PBANKA_051430 | conserved Plasmodium protein, unknown function -1.9 0.73 | PF3D7_1030300 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_051540 | conserved Plasmodium protein, unknown function | PF3D7_1031600 | conserved Plasmodium protein, unknown function (GEXP15 | i) unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_051630 | DER1-like protein, putative | PF3D7_1032500 | DER1-like protein, putative (Der1-2) | proteasome | 0 | 0 | 0 | na | null | null | null |
| PBANKA_051660 | leucine-rich repeat protein (LRR1) | PF3D7_1032800 | leucine-rich repeat protein (LRR1) | signal transduction | 0 | 0 | 0 | na | protein binding | null | null |
| PBANKA_051690 | S-adenosylmethionine decarboxylase, putative | PF3D7_1033100 | S-adenosylmethionine decarboxylase/ornithine decarboxylase (AdoMetDC/ODC) | polyamine biosynthesis | 0 | 0 | 0 | na | adenosylmethionine decarboxylase activity | spermidine biosynthetic process, spermine biosynthetic process | null |
| PBANKA_051700 | early transcribed membrane protein (ETRAMP) | | | early transcribed membrane protein (ETRAMP) | 0 | 0 | 0 | na | null | null | null |
| PBANKA_051740 | myb2 transcription factor, putative (Myb2) | PF3D7_1033600 | myb2 transcription factor, putative (Myb2) | transcription | 0 | 0 | 0 | na | DNA binding | null | null |
| PBANKA_051750 | bromodomain protein, putative | PF3D7_1033700 | bromodomain protein, putative | chromatin | 0 | 0 | 0 | na | null | null | nucleus |
| PBANKA_051820 | flavoprotein subunit of succinate dehydrogenase, putative | PF3D7_1034400 | flavoprotein subunit of succinate dehydrogenase (SDHA | A) TCA, redoxmeth | 0 | 0 | 0 | different in fertilization, ookinete and oocyst | electron carrier activity, flavin adenine dinucleotide binding, oxidoreductase activity, acting on the CH-CH group of donors | electron transport chain, oxidation-reduction process, tricarboxylic acid cycle | null |
| PBANKA_051900 | S-antigen, putative | | | | 0 | 0 | 0 | na | null | null | null |
| PBANKA_051960 | conserved Plasmodium protein, unknown function | | | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_052020 | ADP/ATP transporter on adenylate translocase, putative | PF3D7_1037300 | ADP/ATP transporter on adenylate translocase | transporter | 0 | 0 | 0 | na | binding, transporter activity | transport | mitochondrial inner membrane |
| PBANKA_052120 | conserved Plasmodium protein, unknown function | PF3D7_1038300 | conserved Plasmodium protein, unknown function | RNA | 0 | 0 | 0 | na | null | null | null |
| PBANKA_052210 | conserved Plasmodium protein, unknown function | PF3D7_0421700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_052270 | alpha tubulin 2 | PF3D7_0422300 | alpha tubulin 2 | gliding motility | 0 | 0 | 0 | na | GTP binding, GTPase activity, structural molecule activity | microtubule-based movement, protein polymerization | microtubule, protein complex |
| PBANKA_052360 | BSD-domain protein, putative | PF3D7_0423200 | BSD domain, putative | transcription | 0 | 0 | 0 | na | null | null | null |
| PBANKA_052430 | tryptophan/threonine-rich antigen, putative | | | | 0 | 0 | 0 | na | null | null | null |
| PBANKA_060060 | NIMA related kinase 3, putative (NEK3) | PF3D7_1201600 | NIMA related kinase 3 (NEK3) | kinase | 0 | 0 | 0 | KO not successful | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| PBANKA_060090 | conserved Plasmodium protein, unknown function | PF3D7_1201900 | conserved protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_060160 | conserved protein, unknown function | PF3D7_1202600 | conserved protein, unknown function | transcription | 0 | 0 | 0 | na | null | null | null |
| PBANKA_060190 | high mobility group protein, putative (HMGB1) | PF3D7_1202900 | high mobility group protein (HMGB1) | chromatin | 0 | 0 | 0 | KO not successful; (mutated) sprorozoite and liver stage | DNA binding | null | nucleus |
| PBANKA_060430 | conserved Plasmodium protein, unknown function | PF3D7_1205600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| | | | | | | | | | | | |

| P | PBANKA_060510 | Tat binding protein 1(TBP-1)-interacting protein, putative | | | PF3D7_1206500 | Tat binding protein 1(TBP-1)-interacting protein, putative | proteasome | 0 | 0 | 0 na | null | null | null |
|---|---------------|---|-------|------|---------------|---|------------------------------|---|---|--|---|--|--|
| P | PBANKA_060530 | eukaryotic translation initiation factor 5, putative | | | PF3D7_1206700 | eukaryotic translation initiation factor 5, putative | translation | 0 | 0 | 0 na | translation initiation factor activity | translational initiation | null |
| P | PBANKA_060560 | conserved Plasmodium protein, unknown function | 1.24 | 0.75 | PF3D7_1207000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_060630 | conserved Plasmodium protein, unknown function | | | PF3D7_1207800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_060810 | conserved Plasmodium protein, unknown function | | | PF3D7_1209700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_060950 | kinesin-7, putative | | | PF3D7_1211000 | kinesin-like protein, putative | dynein/kinesin | 0 | 0 | 0 na | ATP binding, microtubule motor activity | microtubule-based movement | null |
| P | PBANKA_060990 | heat shock protein DNAJ homologue Pfj4, putative | | | PF3D7_1211400 | heat shock protein DNAJ homologue Pfj4 (PfJ4) | chaperone | 0 | 0 | 0 na | heat shock protein binding | null | null |
| P | PBANKA_061020 | minchromosome maintenance (MCM) complex subunit, putative (MCM5) | | | PF3D7_1211700 | minchromosome maintenance (MCM) complex subunit, putative (MCM5) | DNA replication | 0 | 0 | 0 na | ATP binding, DNA binding | DNA replication | null |
| P | PBANKA_061070 | conserved Plasmodium protein, unknown function | | | | | unknown function | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_061180 | N-acetyltransferase, putative | | | PF3D7_1437000 | N-acetyltransferase, putative | chromatin modifying proteins | 0 | 0 | 0 na | N-acetyltransferase activity | metabolic process | null |
| P | PBANKA_061520 | calcium dependent protein kinase 4 (CDPK4) | | | PF3D7_0717500 | calcium dependent protein kinase 4 (CDPK4) | kinase | 0 | 0 | 0 different in fertilization, ookinete and oocys | st ATP binding, calcium ion binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| P | PBANKA_061550 | conserved Plasmodium protein, unknown function | | | PF3D7_0717800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_061670 | NIMA related kinase 4 (NEK4) | | | PF3D7_0719200 | NIMA related kinase 4 (NEK4) | kinase | 0 | 0 | 0 different in fertilization and ookinete | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| P | PBANKA_061710 | 60S ribosomal protein L11a, putative | | | PF3D7_0719600 | 60S ribosomal protein L11a, putative | ribosome | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, ribosome |
| P | PBANKA_061720 | 40S ribosomal protein S10, putative | | | PF3D7_0719700 | 40S ribosomal protein S10, putative | ribosome | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_061840 | conserved Plasmodium protein, unknown function | | | PF3D7_0720900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_061910 | 40S ribosomal protein S5, putative | | | PF3D7_0721600 | 40S ribosomal protein S5, putative | ribosome | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, ribosome, small ribosomal subunit |
| P | PBANKA_061920 | secreted ookinete protein, putative (PSOP1) | | | PF3D7_0721700 | secreted ookinete protein, putative (PSOP1) | adhesin | 0 | 0 | 0 not different from wild type | null | null | null |
| P | PBANKA_061930 | conserved Plasmodium protein, unknown function | | | PF3D7_0721800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_062000 | cell cycle control protein cwf15, putative | | | PF3D7_0722500 | cell cycle control protein cwf15, putative | mRNA | 0 | 0 | 0 na | null | nuclear mRNA splicing, via spliceosome | spliceosomal complex |
| P | PBANKA_062050 | tRNAHis guanylyltransferase, putative | | | PF3D7_0723000 | tRNAHis guanylyltransferase, putative | translation | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_062090 | conserved Plasmodium protein, unknown function | -1.81 | 0.73 | PF3D7_0723400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_062190 | mitochondrial import inner membrane translocase subunit tim14, putative | | | PF3D7_0724400 | mitochondrial import inner membrane translocase subunit tim14, putative | transporter | 0 | 0 | 0 na | heat shock protein binding | null | null |
| P | PBANKA_062240 | kinesin-19, putative | -4.93 | 0.75 | PF3D7_0724900 | kinesin-like protein, putative | dynein/kinesin | 0 | 0 | 0 na | ATP binding, microtubule motor activity | microtubule-based movement | null |
| P | PBANKA_062280 | conserved Plasmodium protein, unknown function | | | PF3D7_0725300 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_062300 | Plasmodium exported protein, unknown function | | | | | unknown function | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_062310 | conserved rodent malaria protein, unknown function | | | | | unknown function | 0 | 0 | 0 KO not successful | null | null | null |
| P | PBANKA_062330 | tryptophan-rich antigen, putative | | | | | | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_070020 | BIR protein | | | | | BIR protein | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_070110 | conserved Plasmodium protein, unknown function | | | | | unknown function | 0 | 0 | 0 na | null | null | null |
| P | PBANKA_070160 | conserved protein, unknown function | | | PF3D7_0829100 | conserved protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| | | | | | | | | | | | | | |

| PBANKA_070200 | conserved Plasmodium protein, unknown function | | | PF3D7_0828700 | conserved Plasmodium protein, unknown function | unknown function | 0 (| 0 (| 0 na | | null | null | null |
|---------------|--|-------|------|---------------|--|-----------------------------|-----|-----|-------------|---|---|---|-------------------------|
| PBANKA_070280 | protein disulfide isomerase | | | PF3D7_0827900 | protein disulfide isomerase (PDI8) | proteasome | 0 (| 0 (| 0 na | | isomerase activity | cell redox homeostasis | endoplasmic reticulum |
| PBANKA_070350 | conserved Plasmodium protein, unknown function | -2.88 | 0.75 | PF3D7_0827200 | conserved Plasmodium protein, unknown function | unknown function | 0 (| 0 (| 0 na | | null | null | null |
| PBANKA_070430 | SPRY domain, putative | | | PF3D7_0826300 | SPRY domain, putative | protein-protein interaction | 0 (| 0 (| 0 na | | null | null | null |
| PBANKA_070470 | conserved Plasmodium protein, unknown function | | | PF3D7_0825900 | conserved Plasmodium protein, unknown function | unknown function | 0 (| 0 (| 0 na | | null | null | null |
| PBANKA_070540 | translation initiation factor IF-3, putative | | | PF3D7_0825200 | translation initiation factor IF-3, putative | translation | 0 (| 0 (| 0 na | | translation initiation factor activity | translational initiation | null |
| PBANKA_070550 | conserved Plasmodium protein, unknown function | | | PF3D7_0825100 | conserved Plasmodium protein, unknown function | unknown function | 0 (| 0 (| 0 na | | null | null | null |
| PBANKA_070560 | conserved Plasmodium protein, unknown function | -4.04 | 0.75 | PF3D7_0825000 | conserved Plasmodium protein, unknown function | unknown function | 0 (| 0 (| 0 na | | null | null | null |
| PBANKA_070690 | conserved Plasmodium protein, unknown function | | | PF3D7_0823700 | conserved Plasmodium protein, unknown function | unknown function | 0 (| 0 (| 0 na | | null | null | null |
| PBANKA_071010 | 2-oxoglutarate dehydrogenase e1, putative | | | PF3D7_0820700 | 2-oxoglutarate dehydrogenase E1 component | TCA | 0 (| 0 (| 0 na | | oxoglutarate dehydrogenase (succinyl-transferring) activity, thiamine pyrophosphate binding | glycolysis | null |
| PBANKA_071120 | conserved Plasmodium protein, unknown function | | | PF3D7_0819600 | conserved Plasmodium protein, unknown function | unknown function | 0 (| 0 (| 0 na | | null | null | null |
| PBANKA_071190 | heat shock protein, putative (HSP70) | | | PF3D7_0818900 | heat shock protein 70 (Hsp70) | transporter | 0 (| 0 (| 0 na | | ATP binding | null | null |
| PBANKA_071230 | zinc finger protein, putative | | | PF3D7_0818500 | zinc finger protein, putative | proteasome | 0 (| 0 (| 0 na | | DNA binding, zinc ion binding | null | null |
| PBANKA_071260 | 14-3-3 protein, putative | | | PF3D7_0818200 | 14-3-3 protein, putative | | 0 (| 0 (| 0 na | | protein domain specific binding | null | cytoplasm, nucleus |
| PBANKA_071440 | conserved Plasmodium protein, unknown function | | | PF3D7_0816400 | conserved Plasmodium protein, unknown function | kinase | 0 (| 0 (| 0 na | | calcium ion binding | null | null |
| PBANKA_071550 | phosphoglucomutase-2 (PGM2) | | | PF3D7_0413500 | phosphoglucomutase-2 (PGM2) | glycolysis | 0 (| 0 (| 0 KO not su | successful; (tagged) fertilization and ookinete | null | null | cytoplasm |
| PBANKA_071560 | 26S proteasome AAA-ATPase subunit RPT3, putative | | | PF3D7_0413600 | 26S proteasome AAA-ATPase subunit RPT3, putative | proteasome | 0 (| 0 (| 0 na | | ATP binding, nucleoside-triphosphatase activity | protein catabolic process | cytoplasm, nucleus |
| PBANKA_071570 | lysine decarboxylase-like protein, putative | | | PF3D7_0413700 | lysine decarboxylase-like protein, putative | AA metabolism | 0 (| 0 (| 0 na | | null | null | null |
| PBANKA_071600 | chromosome associated protein, putative | | | PF3D7_0414000 | chromosome associated protein, putative | chromatin | 0 (| 0 (| 0 na | | ATP binding, protein binding | chromosome organization | chromosome |
| PBANKA_071660 | conserved Plasmodium protein, unknown function | | | PF3D7_0414600 | conserved protein, unknown function | unknown function | 0 (| 0 (| 0 na | | null | null | null |
| PBANKA_071780 | 60S ribosomal protein L15, putative | | | PF3D7_0415900 | 60S ribosomal protein L15, putative | ribosome | 0 (| 0 (| 0 na | | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_071970 | conserved Plasmodium protein, unknown function | | | PF3D7_0417600 | conserved Plasmodium protein, unknown function | unknown function | 0 (| 0 (| 0 na | | null | null | null |
| PBANKA_072040 | conserved Plasmodium protein, unknown function | | | PF3D7_0418300 | conserved Plasmodium protein, unknown function | unknown function | 0 (| 0 0 | 0 na | | null | null | null |
| PBANKA_072130 | CGI-141 protein homolog, putative | | | PF3D7_0419200 | CGI-141 protein homolog, putative | trafficking | 0 (| 0 0 | 0 na | | null | vesicle-mediated transport | null |
| PBANKA_072170 | ran binding protein 1, putative | | | PF3D7_0419600 | ran binding protein 1, putative | mRNA | 0 (| 0 0 | 0 na | | null | intracellular transport | null |
| PBANKA_080180 | conserved Plasmodium protein, unknown function | | | PF3D7_0704100 | conserved Plasmodium membrane protein, unknown function | unknown function | 0 (| 0 0 | 0 na | | null | null | null |
| PBANKA_080230 | e3 ubiquitin-protein ligase, putative | | | PF3D7_0704600 | ubiquitin transferase, putative | proteasome | 0 (| 0 0 | 0 na | | acid-amino acid ligase activity | cellular protein modification process | intracellular |
| PBANKA_080310 | minchromosome maintenance (MCM) complex subunit, putative (MCM7) | | | PF3D7_0705400 | minchromosome maintenance (MCM) complex subunit, putative (MCM7) | DNA replication | 0 (|) (| 0 na | | ATP binding, DNA binding, nucleoside-triphosphatase activity | DNA replication, DNA-dependent DNA replication initiation | nucleus |
| PBANKA_080340 | 40S ribosomal protein S29, putative | | | PF3D7_0705700 | 40S ribosomal protein S29, putative | ribosome | 0 (|) (| 0 na | | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_080400 | ribosomal protein, L37e, putative | | | PF3D7_0706400 | ribosomal protein, L37e, putative | ribosome | 0 (|) (| 0 na | | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_080410 | conserved Plasmodium protein, unknown function | | | PF3D7_0706500 | conserved Plasmodium protein, unknown function | unknown function | 0 (| 0 (| 0 na | | null | null | null |
| | | | | | | | | | | | | | |

| PBANKA_080570 | heat shock protein 90, putative (HSP90) | PF3D7_0708400 | heat shock protein 90 (HSP90) | chaperone | 0 | 0 | 0 | na | ATP binding, unfolded protein binding | protein folding, response to stress | null |
|---------------|--|---------------|--|-----------------------|---|---|---|---|---|--|--|
| PBANKA_080590 | kinesin, putative | PF3D7_0319400 | kinesin, putative | dynein/kinesin | 0 | 0 | 0 | na | ATP binding, microtubule motor activity | microtubule-based movement | null |
| PBANKA_080680 | conserved Plasmodium protein, unknown function | PF3D7_0318400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_080700 | dna-directed rna polymerase ii subunit rpb1, putative (RPB1) | PF3D7_0318200 | DNA-directed RNA polymerase II, putative | transcription | 0 | 0 | 0 | na | DNA binding, DNA-directed RNA polymerase activity | transcription from RNA polymerase II promoter | DNA-directed RNA polymerase II, core complex |
| PBANKA_080760 | 40S ribosomal protein S11, putative | PF3D7_0317600 | 40S ribosomal protein S11, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_080990 | tRNA pseudouridine synthase, putative | PF3D7_0908700 | tRNA pseudouridine synthase, putative | translation | 0 | 0 | 0 | na | RNA binding, pseudouridine synthase activity | pseudouridine synthesis | null |
| PBANKA_081050 | apoptosis-related protein, putative | PF3D7_0909300 | apoptosis-related protein, putative (ARP) | apoptosis | 0 | 0 | 0 | na | DNA binding | null | null |
| PBANKA_081300 | inhibitor of cysteine proteases (ICP) | PF3D7_0911900 | falstatin (ICP) | protease | 0 | 0 | 0 | KO not successful | null | null | null |
| PBANKA_081340 | conserved Plasmodium protein, unknown function | PF3D7_0912400 | conserved Plasmodium protein, unknown function | Ca2+ binding | 0 | 0 | 0 | na | null | null | null |
| PBANKA_081350 | conserved Plasmodium protein, unknown function | PF3D7_0912500 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | nucleic acid binding | null | nucleus |
| PBANKA_081480 | conserved Plasmodium protein, unknown function | PF3D7_0913800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_081690 | conserved Plasmodium protein, unknown function | PF3D7_0915900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_081700 | sugar transporter, putative | PF3D7_0916000 | sugar transporter, putative | transporter | 0 | 0 | 0 | na | null | null | null |
| PBANKA_081890 | heat shock protein 70, putative | PF3D7_0917900 | heat shock protein 70 (Hsp70-2) | chaperone | 0 | 0 | 0 | na | ATP binding | null | null |
| PBANKA_081990 | nucleosome assembly protein (NAPS) | PF3D7_0919000 | nucleosome assembly protein (NAPS) | chromatin | 0 | 0 | 0 | KO not successful; (tagged) different in fertilization, cokinete and oocyst | null | nucleosome assembly | nuclear periphery, nucleus |
| PBANKA_082110 | CS domain protein, putative | PF3D7_0920200 | CS-domain containing protein, conserved in Apicomplexa | adhesin | 0 | 0 | 0 | na | null | null | null |
| PBANKA_082170 | inosine-5'-monophosphate dehydrogenase, putative | PF3D7_0920800 | inosine-5'-monophosphate dehydrogenase | purine metabolism | 0 | 0 | 0 | na | IMP dehydrogenase activity | oxidation-reduction process | null |
| PBANKA_082200 | conserved Plasmodium protein, unknown function | PF3D7_0921100 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_082360 | pre-mRNA splicing factor, putative | PF3D7_0922700 | pre-mRNA splicing factor, putative | mRNA | 0 | 0 | 0 | na | null | RNA splicing | spliceosomal complex |
| PBANKA_082480 | RNA-binding protein, putative | PF3D7_0923900 | RNA binding protein, putative | mRNA | 0 | 0 | 0 | na | nucleic acid binding | null | null |
| PBANKA_082560 | splicing factor 3A subunit 3, putative (SF3A3) | PF3D7_0924700 | splicing factor 3A subunit 3, putative (SF3A3) | mRNA | 0 | 0 | 0 | na | nucleic acid binding, zinc ion binding | null | nucleus |
| PBANKA_082580 | conserved Plasmodium protein, unknown function | PF3D7_0925000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_082640 | zinc-binding protein (Yippee), putative | PF3D7_0925600 | zinc binding protein (Yippee), putative | senescence, apoptosis | 0 | 0 | 0 | na | null | null | null |
| PBANKA_082670 | conserved Plasmodium protein, unknown function | PF3D7_0925900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_082830 | dynein light chain, putative | PF3D7_0927500 | dynein light chain, putative | dynein/kinesin | 0 | 0 | 0 | na | null | null | null |
| PBANKA_082890 | conserved Plasmodium protein, unknown function | PF3D7_0928100 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_083030 | conserved Plasmodium protein, unknown function | PF3D7_0929500 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_083100 | merozoite surface protein 1 (MSP1) | PF3D7_0930300 | merozoite surface protein 1 (MSP1) | contaminant | 0 | 0 | 0 | KO not successful; (mutated) different in sporozoite and liver stage | null | pathogenesis | membrane |
| PBANKA_083170 | NifU-like protein, putative | PF3D7_0930900 | NifU-like protein, putative | apicoplast | 0 | 0 | 0 | na | iron ion binding, iron-sulfur cluster binding | iron-sulfur cluster assembly | null |
| PBANKA_083260 | proteasome subunit beta type-6, putative | PF3D7_0931800 | proteasome precursor, putative | proteasome | 0 | 0 | 0 | na | threonine-type endopeptidase activity | proteolysis involved in cellular protein catabolic process | proteasome core complex |
| PBANKA_083310 | aspartyl aminopeptidase, putative | PF3D7_0932300 | M18 aspartyl aminopeptidase (M18AAP) | protease | 0 | 0 | 0 | na | aminopeptidase activity, zinc ion binding | proteolysis | vacuole |
| | | | | | | | | | | | |

| PBANKA_083520 | transcription factor with AP2 domain(s), putative | | | PF3D7_0934400 | transcription factor with AP2 domain(s), putative (ApiAP2) | transcription | 0 | 0 | 0 | na | sequence-specific DNA binding transcription factor activity | regulation of transcription, DNA-dependent | null |
|---------------|--|-------|------|---------------|---|-------------------------------|---|---|---|---|--|---|--|
| PBANKA_083700 | BIR protein | | | | | BIR protein | 0 | 0 | 0 | na | null | null | null |
| PBANKA_090140 | mitochondrial ATP synthase delta subunit, putative | | | PF3D7_1147700 | mitochondrial ATP synthase delta subunit, putative | ATP synthase complex | 0 | 0 | 0 | na | hydrogen ion transporting ATP synthase activity, rotational mechanism, proton-transporting ATPase activity, rotational mechanism | m ATP synthesis coupled proton transport | mitochondrion, proton-transporting ATP synthase complex, catalytic core F(1) |
| PBANKA_090190 | tubulin-tyrosine ligase, putative | -3.79 | 0.75 | PF3D7_1147200 | tubulin-tyrosine ligase, putative | microtubule | 0 | 0 | 0 | na | tubulin-tyrosine ligase activity | cellular protein modification process | null |
| PBANKA_090260 | CCAAT-box DNA binding protein subunit B, putative | | | PF3D7_1146600 | CCAAT-box DNA binding protein subunit B | chromatin | 0 | 0 | 0 | na | sequence-specific DNA binding | null | intracellular, nucleus |
| PBANKA_090370 | conserved Plasmodium protein, unknown function | | | PF3D7_1145300 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_090420 | conserved Plasmodium protein, unknown function | -3.03 | 0.75 | PF3D7_1144800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_090460 | conserved Plasmodium protein, unknown function | | | PF3D7_1144400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_090470 | 60S ribosomal protein L41, putative | | | PF3D7_1144300 | 60S ribosomal protein L41, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | ribosome |
| PBANKA_090500 | 40S ribosomal protein S21e, putative | | | PF3D7_1144000 | 40S ribosomal protein S21e, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_090560 | translation initiation factor eIF-1A, putative | | | PF3D7_1143400 | translation initiation factor eIF-1A, putative | translation | 0 | 0 | 0 | na | RNA binding, translation initiation factor activity | translational initiation | null |
| PBANKA_090730 | conserved Plasmodium protein, unknown function | | | PF3D7_1141700 | conserved Plasmodium protein, unknown function | protease | 0 | 0 | 0 | na | null | null | null |
| PBANKA_090790 | conserved Plasmodium protein, unknown function | -6.19 | 0.75 | | | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_090870 | conserved Plasmodium protein, unknown function | | | PF3D7_1140300 | P-loop containing nucleoside triphosphate hydrolase, putative | purine metabolism | 0 | 0 | 0 | na | null | null | null |
| PBANKA_090880 | conserved Plasmodium protein, unknown function | | | PF3D7_1140200 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_090980 | conserved Plasmodium protein, unknown function | | | PF3D7_1139100 | conserved Plasmodium protein, unknown function | RNA binding | 0 | 0 | 0 | na | nucleic acid binding | null | null |
| PBANKA_091020 | protein phosphatase 2C, putative | | | PF3D7_1138500 | protein phosphatase 2c | phosphatase | 0 | 0 | 0 | na | catalytic activity | null | null |
| PBANKA_091090 | conserved Plasmodium protein, unknown function | | | PF3D7_1137800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_091170 | subtilisin-like protease 2 (SUB2) | | | PF3D7_1136900 | subtilisin-like protease 2 (SUB2) | protease | 0 | 0 | 0 | KO not successful | serine-type endopeptidase activity | proteolysis | null |
| PBANKA_091200 | conserved Plasmodium protein, unknown function | | | PF3D7_1136600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_091250 | conserved Plasmodium protein, unknown function | -3.34 | 0.75 | PF3D7_1136100 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_091260 | conserved Plasmodium protein, unknown function | | | PF3D7_1107300 | polyadenylate-binding protein-interacting protein 1, putative (PAIP1) | mRNA | 0 | 0 | 0 | na | null | null | null |
| PBANKA_091440 | heat shock protein hsp70 homologue, putative (UIS24) | | | PF3D7_1134000 | heat shock protein 70 (Hsp70-3) | transporter | 0 | 0 | 0 | na | ATP binding, unfolded protein binding | protein folding | null |
| PBANKA_091500 | apical membrane antigen 1 (AMA1) | | | PF3D7_1133400 | apical membrane antigen 1 (AMA1) | contaminant | 0 | 0 | 0 | KO not successful; (mutated) sporozoite and liver stage | null | pathogenesis | membrane |
| PBANKA_091540 | conserved Plasmodium protein, unknown function | | | PF3D7_1133000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_091560 | aquaglyceroporin (AQP) | | | PF3D7_1132800 | aquaglyceroporin (AQP) | glycolysis | 0 | 0 | 0 | not different from wild type | transporter activity | transport | membrane, perinuclear region of cytoplasm |
| PBANKA_091720 | RNA-binding protein s1, putative | -1.7 | 0.7 | | | mRNA | 0 | 0 | 0 | na | nucleic acid binding | null | null |
| PBANKA_091800 | 60S ribosomal protein, putative | | | PF3D7_1130200 | 60S ribosomal protein P0 | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | ribosome biogenesis, translational elongation | intracellular, ribosome |
| PBANKA_091830 | transporter, putative | | | PF3D7_1129900 | transporter, putative | transporter | 0 | 0 | 0 | na | null | null | null |
| PBANKA_091860 | phosphatidylinositol-4-phosphate 5-kinase, putative | | | PF3D7_1129600 | phosphatidylinositol-4-phosphate-5-kinase, putative | inositol phosphate metabolism | 0 | 0 | 0 | na | null | null | null |
| PBANKA_091910 | parasitophorous vacuolar protein 1, putative (PV1) | | | PF3D7_1129100 | parasitophorous vacuolar protein 1 (PV1) | PV membrane protein | 0 | 0 | 0 | na | null | null | null |
| | | | | | | | | | | | | | |

| PBANKA_092080 | CRAL/TRIO domain-containing protein, putative | PF3D7_1127600 | CRAL/TRIO domain-containing protein, putative | trafficking | 0 | 0 | 0 na | null | null | null |
|---------------|--|---------------|---|-------------------------------|---|---|---|--|--|-------------------------|
| PBANKA_092170 | conserved Plasmodium protein, unknown function | PF3D7_1126700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_092280 | small nuclear ribonucleoprotein D1, putative (SNRPD1) | PF3D7_1125500 | small nuclear ribonucleoprotein D1, putative (SNRPD1) | mRNA | 0 | 0 | 0 na | null | null | null |
| PBANKA_092340 | 60S ribosomal protein L35, putative | PF3D7_1124900 | 60S ribosomal protein L35, putative | ribosome | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_092490 | translation elongation factor EF-1, subunit alpha, putative | PF3D7_1123400 | translation elongation factor EF-1, subunit alpha, putative | translation | 0 | 0 | 0 na | GTP binding, GTPase activity | null | null |
| PBANKA_092550 | calcium dependent protein kinase 6 (CDPK6) | PF3D7_1122800 | calcium dependent protein kinase 6 (CDPK6) | kinase | 0 | 0 | 0 different in sporozoite and liver stage | ATP binding, calcium ion binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | y protein phosphorylation | null |
| PBANKA_092670 | circumsporozoite-related antigen | PF3D7_1121600 | circumsporozoite-related antigen (CRA) | adhesin | 0 | 0 | 0 different in liver stage | null | null | null |
| PBANKA_092750 | conserved Plasmodium protein, unknown function | PF3D7_1120800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_092800 | conserved Plasmodium protein, unknown function | PF3D7_1120200 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_092810 | phosphoglycerate mutase, putative (PGM1) | PF3D7_1120100 | phosphoglycerate mutase, putative (PGM1) | glycolysis | 0 | 0 | 0 na | intramolecular transferase activity, phosphotransferases | glycolysis | null |
| PBANKA_092920 | conserved Plasmodium protein, unknown function | PF3D7_1118900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_093090 | deubiquinating/deneddylating enzyme, putative (UCH54) | PF3D7_1117100 | deubiquinating/deneddylating enzyme (UCH54) | proteasome | 0 | 0 | 0 na | ubiquitin thiolesterase activity | ubiquitin-dependent protein catabolic process | intracellular |
| PBANKA_093120 | heat shock protein 101 (ClpB2) | PF3D7_1116800 | heat shock protein 101 (HSP101) | chaperone | 0 | 0 | KO not successful; (tagged) different in sporozoite and liver stage | $_{\mathrm{pe}}$ ATP binding, nucleoside-triphosphatase activity, protein binding | protein metabolic process | null |
| PBANKA_093130 | dipeptidyl aminopeptidase, putative | PF3D7_1116700 | cathepsin C, homolog,dipeptidyl peptidase 1 (DPAP1) | protease | 0 | 0 | 0 na | cysteine-type peptidase activity | proteolysis | null |
| PBANKA_093210 | conserved Plasmodium protein, unknown function | PF3D7_1115900 | conserved Plasmodium membrane protein, unknown function | unknown function | 0 | 0 | 0 na | zinc ion binding | null | null |
| PBANKA_093300 | serine/threonine protein kinase, puative | PF3D7_1114700 | serine/threonine protein kinase, putative (CLK3) (PRK4) | kinase | 0 | 0 | 0 KO not successful | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| PBANKA_093390 | glyoxalase I, putative | PF3D7_1147100 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_093720 | asparagine-rich antigen, putative | PF3D7_1111600 | endonuclease/exonuclease/phosphatase family protein, putative | inositol phosphate metabolism | 0 | 0 | 0 na | nucleic acid binding | null | null |
| PBANKA_093730 | conserved Plasmodium protein, unknown function | PF3D7_1110400 | asparagine-rich antigen | | 0 | 0 | 0 na | nucleic acid binding | null | intracellular |
| PBANKA_093770 | apicoplast ribosomal protein L36e precursor, putative | PF3D7_1109900 | apicoplast ribosomal protein L36e precursor, putative | ribosome apico | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_093840 | endoplasmic reticulum-resident calcium binding protein, putative | PF3D7_1108600 | endoplasmic reticulum-resident calcium binding protein (ERC) | HG ERG | 0 | 0 | 0 na | calcium ion binding | null | null |
| PBANKA_093880 | conserved Plasmodium protein, unknown function | PF3D7_1108100 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_094080 | apicoplast ribosomal protein S15 precursor, putative | PF3D7_1106100 | apicoplast ribosomal protein S15 precursor, putative | ribosome apico | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_094110 | conserved Plasmodium protein, unknown function | PF3D7_1105800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_094190 | histone H4, putative | PF3D7_1105000 | histone H4 (H4) | chromatin | 0 | 0 | 0 na | DNA binding | DNA-dependent transcription, initiation, nucleosome assemb | ly nucleosome, nucleus |
| PBANKA_094250 | conserved Plasmodium protein, unknown function | PF3D7_1104400 | conserved protein, unknown function | nucleic acid binding | 0 | 0 | 0 na | null | null | null |
| PBANKA_094270 | chromatin remodeling protein, putative (SNF2L) | PF3D7_1104200 | chromatin remodeling protein (SNF2L) | chromatin | 0 | 0 | 0 na | ATP binding, DNA binding, helicase activity | null | null |
| PBANKA_094360 | 60S acidic ribosomal protein, putative | PF3D7_1103100 | 60S ribosomal protein P1, putative | ribosome | 0 | 0 | 0 na | structural constituent of ribosome | translational elongation | intracellular, ribosome |
| PBANKA_100060 | erythrocyte membrane antigen 1 | PF3D7_0403700 | CGI-201 protein, short form | mRNA | 0 | 0 | 0 na | null | null | null |
| PBANKA_100140 | splicing factor, putative | PF3D7_0403800 | conserved Plasmodium protein, unknown function | mRNA | 0 | 0 | 0 na | null | RNA processing | intracellular |
| PBANKA_100150 | conserved Plasmodium protein, unknown function | PF3D7_0404000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| | | | | | | | | | | |

| PBANKA_100200 | conserved Plasmodium protein, unknown function | PF3D7_0404300 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | intracellular transport | null |
|---------------|--|-----------------|--|-----------------------------|---|---|--|--|--|---|
| PBANKA_100340 | lysine decarboxylase, putative (UIS14) | PF3D7_0405700 | lysine decarboxylase, putative | AA metabolism | 0 | 0 | 0 na | catalytic activity | null | null |
| PBANKA_100380 | vacuolar ATP synthase subunit b, putative | PF3D7_0406100 | vacuolar ATP synthase subunit b | vacuolar ATP synthase | 0 | 0 | 0 na | hydrogen ion transporting ATP synthase activity, rotational mechanism, proton-transporting ATPase activity, rotational mechanism | a ATP synthesis coupled proton transport | proton-transporting V-type ATPase, V1 domain, proton-transporting two-sector ATPase complex |
| PBANKA_100610 | flap endonuclease 1, putative | PF3D7_0408500 | flap endonuclease 1 (FEN1) | DNA replication, DNA repair | 0 | 0 | 0 na | 5'-3' exonuclease activity, DNA binding | DNA repair | null |
| PBANKA_100670 | U4/U6 small nuclear ribonucleoprotein PRP31, putative (PRPF31) | PF3D7_0409100 | U4/U6 small nuclear ribonucleoprotein PRP31, putative (PRPF31) | mRNA | 0 | 0 | 0 na | null | null | null |
| PBANKA_100720 | replication protein A1, large subunit, putative (RPA1) | PF3D7_0409600 | replication protein A1, large subunit (RPA1) | DNA repair | 0 | 0 | 0 na | nucleic acid binding | null | null |
| PBANKA_100750 | actin-like protein, putative | PF3D7_0409900 | actin-like protein, putative | gliding motility | 0 | 0 | 0 na | protein binding | null | null |
| PBANKA_100770 | protein phosphatase 2C, putative | PF3D7_0410500 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | catalytic activity | null | null |
| PBANKA_100850 | translocon component PTEX150 (PTEX150) | PF3D7_1436000 | glucose-6-phosphate isomerase (GPI) | glycolysis | 0 | 0 | 0 KO not successful; (tagged) sporozoite and liver stage | null | null | null |
| PBANKA_100880 | glucose-6-phosphate isomerase, putative | PF3D7_1435700.1 | conserved Plasmodium protein, unknown function | proteasome | 0 | 0 | 0 na | glucose-6-phosphate isomerase activity | gluconeogenesis, glycolysis | null |
| PBANKA_100910 | conserved Plasmodium protein, unknown function | PF3D7_1435700.2 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_100950 | NAD(P)H-dependent glutamate synthase, putative | PF3D7_1435300 | NAD(P)H-dependent glutamate synthase, putative | AA metabolism | 0 | 0 | 0 na | election confer activity, glutamate synthase activity, inon-water charter binding, cedioxeducines exclusity, acting on the CH-NPQ group of disnors, INIX or NACP as acceptor | glutamate biosynthetic process, nitrogen compound metabolic process, oxidation-reduction process | null |
| PBANKA_100990 | conserved Plasmodium protein, unknown function | PF3D7_1434900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_101000 | conserved Plasmodium protein, unknown function | PF3D7_1434800 | conserved protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | mitochondrial matrix |
| PBANKA_101050 | Hsp70/Hsp90 organizing protein, putative (HOP) | PF3D7_1434300 | Hsp70/Hsp90 organizing protein (HOP) | chaperone | 0 | 0 | 0 na | null | null | cytosol |
| PBANKA_101190 | h/aca ribonucleoprotein complex subunit 3, putative (NOP10) | PF3D7_1433000 | ribosome biogenesis protein, NOP10-like | ribosome | 0 | 0 | 0 na | null | null | null |
| PBANKA_101310 | 60S ribosomal protein L14, putative | PF3D7_1431700 | 60S ribosomal protein L14, putative | ribosome | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_101320 | ATP-specific succinyl-CoA synthetase beta subunit, putative | PF3D7_1431600 | ATP-specific succinyl-CoA synthetase beta subunit, putative | TCA | 0 | 0 | 0 na | catalytic activity | metabolic process | null |
| PBANKA_101400 | NADP-specific glutamate dehydrogenase, putative (GDH2) | PF3D7_1430700 | glutamate dehydrogenase, putative (GDHb) | nucleic acid binding | 0 | 0 | 0 na | binding, glutamate dehydrogenase (NADP+) activity | glutamate biosynthetic process, oxidation-reduction process | apicoplast |
| PBANKA_101560 | apicoplast ribosomal protein L15 precursor, putative | | apicoplast ribosomal protein L15 precursor, putative | ribosome apico | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, large ribosomal subunit, ribosome |
| PBANKA_101630 | proliferation-associated protein 2g4, putative | PF3D7_1428300 | proliferation-associated protein 2g4, putative | protease | 0 | 0 | 0 na | null | cellular process | null |
| PBANKA_101860 | 60S ribosomal protein L21e, putative | PF3D7_1426000 | 60S ribosomal protein L21e, putative | ribosome | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_101880 | conserved Plasmodium protein, unknown function | | | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_101940 | 60S ribosomal protein L7-3, putative | PF3D7_1424400 | 60S ribosomal protein L7-3, putative | ribosome | 0 | 0 | 0 na | null | null | null |
| PBANKA_101950 | 60S ribosomal protein L5, putative | PF3D7_1424100 | 60S ribosomal protein L5, putative | ribosome | 0 | 0 | 0 na | 5S rRNA binding, structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_102000 | conserved Plasmodium protein, unknown function | PF3D7_1423400 | conserved Plasmodium membrane protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_102070 | conserved Plasmodium protein, unknown function | PF3D7_1422700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_102110 | DnaJ protein, putative | PF3D7_1422300 | DnaJ protein, putative | chaperone | 0 | 0 | 0 na | heat shock protein binding | null | null |
| PBANKA_102180 | mitochondrial protein import protein TIM13, putative | PF3D7_1421500 | mitochondrial protein import protein TIM13, putative | transporter | 0 | 0 | 0 na | null | protein import into mitochondrial inner membrane | mitochondrial intermembrane space protein transporter complex |
| PBANKA_102200 | 40S ribosomal protein S25, putative | PF3D7_1421200 | 40S ribosomal protein S25, putative | ribosome | 0 | 0 | 0 na | null | null | null |
| PBANKA_102350 | conserved Plasmodium protein, unknown function | PF3D7_1419700 | conserved Plasmodium protein, unknown function | translation | 0 | 0 | 0 na | null | null | null |
| | | | | | | | | | | |

| PBANKA_102470 | ubiquitin fusion degradation protein UFD1, putative | PF3D7_1418000 | ubiquitin fusion degradation protein UFD1, putative (UFD1) | proteasome | 0 | 0 | 0 | na | null | ubiquitin-dependent protein catabolic process | null |
|---------------|--|-----------------|--|----------------------|---|---|---|---|--|---|-------------------------|
| PBANKA_102490 | minchromosome maintenance (MCM) complex subunit, putative (MCM2) | PF3D7_1417800 | minchromosome maintenance (MCM) complex subunit (MCM2) | DNA replication | 0 | 0 | 0 | na | ATP binding, DNA binding, nucleoside-triphosphatase activity | DNA replication, DNA-dependent DNA replication initiation | nucleus |
| PBANKA_102550 | NOT family protein, putative | PF3D7_1417200 | NOT family protein, putative | mRNA | 0 | 0 | 0 | na | null | null | null |
| PBANKA_102830 | serine/threonine protein phosphatase, putative | PF3D7_1414400 | serine/threonine protein phosphatase (PP1) | phosphatase | 0 | 0 | 0 | na | hydrolase activity | null | null |
| PBANKA_102840 | 60S ribosomal protein L10, putative | PF3D7_1414300 | 60S ribosomal protein L10, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_102940 | conserved Plasmodium protein, unknown function | PF3D7_1413200 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_103010 | actin II | PF3D7_1412500 | actin II (ACT2) | gliding motility | 0 | 0 | 0 | different in fertilization, ookinete, oocyst and sporozoite | protein binding | null | cytoplasm |
| PBANKA_103060 | conserved Plasmodium protein, unknown function | PF3D7_1412000 | p1/s1 nuclease, putative | DNA repair | 0 | 0 | 0 | na | hydrolase activity, acting on ester bonds | null | null |
| PBANKA_103160 | conserved Plasmodium protein, unknown function | PF3D7_1410900 | conserved Plasmodium protein, unknown function | transcription | 0 | 0 | 0 | na | null | null | null |
| PBANKA_103310 | conserved Plasmodium protein, unknown function | PF3D7_1409200.1 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_103370 | conserved Plasmodium protein, unknown function | PF3D7_1408800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_103390 | 40S ribosomal protein S8e, putative | PF3D7_1408600 | 40S ribosomal protein S8e, putative | ribosome | 0 | 0 | 0 | na | null | null | intracellular |
| PBANKA_103430 | transcription factor with AP2 domain(s) (API2-G2) | PF3D7_1408200 | transcription factor with AP2 domain(s) (ApiAP2) | transcription | 0 | 0 | 0 | different in fertilization and ookinete | sequence-specific DNA binding transcription factor activity | regulation of transcription, DNA-dependent | null |
| PBANKA_103440 | plasmepsin IV (PM4) | PF3D7_1407800 | plasmepsin IV (PM4) | plasmepsin, protease | 0 | 0 | 0 | na | aspartic-type endopeptidase activity | proteolysis | null |
| PBANKA_103490 | pre-mRNA splicing factor, putative | PF3D7_1407300 | pre-mRNA splicing factor, putative | mRNA | 0 | 0 | 0 | na | null | null | null |
| PBANKA_103510 | fibrillarin, putative (NOP1) | PF3D7_1407100 | fibrillarin, putative (NOP1) | ribosome assembly | 0 | 0 | 0 | na | RNA binding | rRNA processing | nucleus |
| PBANKA_103520 | LCCL domain-containing protein (CCp3) | PF3D7_1407000 | LCCL domain-containing protein (CCp3) | adhesin | 0 | 0 | 0 | different in occyst and sporozoite; (tagged) also in fertilization and ookinete | scavenger receptor activity | null | membrane |
| PBANKA_103630 | RNA-binding protein, putative | PF3D7_1405900 | RNA binding protein, putative | mRNA | 0 | 0 | 0 | na | nucleic acid binding | null | null |
| PBANKA_103760 | rrna biogenesis protein rrp5, putative (RRP5) | PF3D7_1404500 | U3 small nucleolar ribonucleoprotein, U3 snoRNP, putative | mRNA | 0 | 0 | 0 | na | null | RNA processing | intracellular |
| PBANKA_103800 | cytochrome c, putative | PF3D7_1404100 | cytochrome c, putative | redoxmeth | 0 | 0 | 0 | na | electron carrier activity, heme binding, iron ion binding | null | null |
| PBANKA_103860 | conserved Plasmodium protein, unknown function | PF3D7_1403400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_103940 | ubiquitin-40s ribosomal protein s27a, putative | PF3D7_1402500 | 40S ribosomal protein S31/UBI, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_104020 | BIR protein | | | BIR protein | 0 | 0 | 0 | na | null | null | null |
| PBANKA_110110 | Plasmodium exported protein, unknown function | | | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_110130 | conserved rodent malaria protein, unknown function | | | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_110230 | conserved Plasmodium protein, unknown function | PF3D7_0502600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_110340 | 60S ribosomal protein L31, putative | PF3D7_0503800 | 60S ribosomal protein L31, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_110390 | conserved Plasmodium protein, unknown function | PF3D7_0504300 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_110550 | conserved Plasmodium protein, unknown function | PF3D7_0505900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_110670 | 60S ribosomal protein L4, putative | PF3D7_0507100 | 60S ribosomal protein L4, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_110680 | subtilisin-like protease 3, putative (SUB3) | PF3D7_0507200 | subtilisin-like protease 3, putative (SUB3) | protease | 0 | 0 | 0 | na | serine-type endopeptidase activity | proteolysis | null |
| | | | | | | | | | | | |

| PBANKA_110710 | subtilisin-like protease 1, putative (SUB1) | PF3D7_0507500 | subtilisin-like protease 1 (SUB1) | protease | 0 | 0 | 0 | KO not successful; (mutated) different in liver stage | serine-type endopeptidase activity | proteolysis | null |
|---------------|---|---------------|---|--------------------------------|---|---|---|---|--|---|---------------------------------|
| PBANKA_110800 | transcription factor IIb, putative | PF3D7_0508400 | transcription factor IIb, putative | transcription | 0 | 0 | 0 | na | null | null | null |
| PBANKA_110820 | conserved Plasmodium protein, unknown function | PF3D7_0508600 | conserved Plasmodium protein, unknown function | nucleic acid binding | 0 | 0 | 0 | na | null | null | null |
| PBANKA_110850 | conserved Plasmodium protein, unknown function | PF3D7_0508900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_110890 | conserved Plasmodium protein, unknown function | PF3D7_0509300 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_111040 | conserved Plasmodium protein, unknown function | PF3D7_0510800 | conserved Plasmodium protein, unknown function | transcription | 0 | 0 | 0 | na | null | null | null |
| PBANKA_111130 | kinase binding protein CGI-121, putative | | | | 0 | 0 | 0 | na | null | null | null |
| PBANKA_111170 | conserved Plasmodium protein, unknown function | PF3D7_0512100 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_111230 | Rab1b, putative | PF3D7_0512600 | Rab GTPase 1b (Rab1b) | trafficking | 0 | 0 | 0 | na | GTP binding, GTPase activity, protein binding | intracellular protein transport, nucleocytoplasmic transport, small GTPase mediated signal transduction | intracellular |
| PBANKA_111300 | purine nucleotide phosphorylase, putative | PF3D7_0513300 | purine nucleoside phosphorylase (PNP) | purine metabolism | 0 | 0 | 0 | na | catalytic activity | nucleoside metabolic process | null |
| PBANKA_111330 | deoxyribodipyrimidine photolyase, putative | PF3D7_0513600 | deoxyribodipyrimidine photolyase (photoreactivating enzyme, DNA photolyase), putative | DNA replication | 0 | 0 | 0 | na | deoxyribodipyrimidine photo-lyase activity | DNA repair | null |
| PBANKA_111370 | tubulin-tyrosine ligase, putative | PF3D7_0514000 | tubulin-tyrosine ligase, putative (TTL) | microtubule | 0 | 0 | 0 | na | tubulin-tyrosine ligase activity | cellular protein modification process | null |
| PBANKA_111520 | conserved Plasmodium protein, unknown function | PF3D7_0515600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_111540 | conserved Plasmodium protein, unknown function | PF3D7_0515800 | BolA-like protein, putative | cell division | 0 | 0 | 0 | na | null | null | null |
| PBANKA_111610 | conserved Plasmodium protein, unknown function | PF3D7_0616600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_111740 | conserved Plasmodium protein, unknown function | PF3D7_0618200 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_111750 | ribosomal protein L27a, putative | PF3D7_0618300 | 60S ribosomal protein L27a, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_111860 | cell division cycle protein 48 homologue, putative | PF3D7_0619400 | cell division cycle protein 48 homologue, putative | cell division | 0 | 0 | 0 | na | ATP binding, nucleoside-triphosphatase activity | null | null |
| PBANKA_111920 | conserved Plasmodium protein, unknown function -3.28 0.75 | PF3D7_0620000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_111960 | merozoite surface protein 10, putative (MSP10) | PF3D7_0620400 | merozoite surface protein 10 (MSP10) | contaminant | 0 | 0 | 0 | na | null | null | null |
| PBANKA_111990 | DnaJ protein, putative | PF3D7_0620700 | DnaJ protein, putative | chaperone | 0 | 0 | 0 | na | heat shock protein binding | null | null |
| PBANKA_112100 | conserved Plasmodium protein, unknown function | PF3D7_0622100 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_112200 | coronin binding protein, putative | PF3D7_0623100 | coronin binding protein, putative | gliding motility | 0 | 0 | 0 | na | null | null | null |
| PBANKA_112350 | SNF2 helicase, putative (ISWI) | PF3D7_0624600 | SNF2 helicase, putative (ISWI) | chromatin | 0 | 0 | 0 | na | ATP binding, DNA binding, helicase activity, protein binding, zinc ion binding | null | nucleus |
| PBANKA_112380 | conserved Plasmodium protein, unknown function | PF3D7_0624900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_112470 | conserved Plasmodium protein, unknown function | PF3D7_0625900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_112560 | pyruvate kinase, putative | PF3D7_0626800 | pyruvate kinase (PyrK) | glycolysis | 0 | 0 | 0 | na | magnesium ion binding, potassium ion binding, pyruvate kinase activity | glycolysis | null |
| PBANKA_112580 | conserved Plasmodium protein, unknown function | PF3D7_0627000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_112700 | choline/ethanolaminephosphotransferase (CEPT) | PF3D7_0628300 | choline/ethanolaminephosphotransferase, putative (CEPT) | phosphatidylcholine metabolism | 0 | 0 | 0 | KO not successful | phosphotransferase activity, for other substituted phosphate group | s phospholipid biosynthetic process | endoplasmic reticulum, membrane |
| PBANKA_112780 | DnaJ protein, putative | PF3D7_0629200 | DnaJ protein, putative | chaperone | 0 | 0 | 0 | na | heat shock protein binding, unfolded protein binding | protein folding | null |
| PBANKA_112890 | conserved Plasmodium protein, unknown function | PF3D7_0630100 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| | | | | | | | | | | | |

| F | BANKA_112910 | DNA polymerase epsilon, putative | | | PF3D7_0630300 | DNA polymerase epsilon, catalytic subunit a, putative | DNA repair | 0 | 0 | 0 | na | DNA binding, DNA-directed DNA polymerase activity, nucleotide binding | J DNA replication | null |
|---|--------------|--|-------|------|---------------|--|------------------|---|---|---|-------------------|--|--|--|
| F | BANKA_112960 | conserved Plasmodium protein, unknown function | | | PF3D7_0630800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| F | BANKA_113050 | proteasome subunit alpha type-7, putative | | | PF3D7_1353900 | proteasome subunit, putative | proteasome | 0 | 0 | 0 | na | threonine-type endopeptidase activity | ubiquitin-dependent protein catabolic process | proteasome core complex |
| F | BANKA_113150 | conserved Plasmodium protein, unknown function | | | PF3D7_1354900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| F | BANKA_113160 | minchromosome maintenance (MCM) complex subunit, putative (MCM6) | | | PF3D7_1355100 | minchromosome maintenance (MCM) complex subunit (MCM6) | DNA replication | 0 | 0 | 0 | na | ATP binding, DNA binding | DNA replication, DNA-dependent DNA replication initiation | nucleus |
| F | BANKA_113260 | ubiquitin conjugating enzyme, putative | | | PF3D7_1356300 | ubiquitin conjugating enzyme, putative | proteasome | 0 | 0 | 0 | na | small conjugating protein ligase activity | post-translational protein modification, regulation of protein metabolic process | null |
| F | BANKA_113390 | conserved Plasmodium protein, unknown function | | | PF3D7_1357600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| F | BANKA_113460 | rhomboid protease ROM7, putative (ROM7) | | | PF3D7_1358300 | rhomboid protease ROM7 (ROM7) | rhomboid | 0 | 0 | 0 | KO not successful | null | null | integral to membrane |
| F | BANKA_113500 | conserved Plasmodium protein, unknown function | | | PF3D7_1358700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| F | BANKA_113700 | falcilysin, putative (FLN) | | | PF3D7_1360800 | falcilysin (FLN) | protease | 0 | 0 | 0 | na | catalytic activity, metalloendopeptidase activity, zinc ion binding | proteolysis | apicoplast, food vacuole |
| F | BANKA_113770 | cytochrome c oxidase subunit 2, putative | | | PF3D7_1361700 | cytochrome c oxidase subunit 2, putative | redoxmeth | 0 | 0 | 0 | na | copper ion binding, cytochrome-c oxidase activity | null | membrane |
| F | BANKA_113790 | proliferating cell nuclear antigen, putative (PCNA) | | | PF3D7_1361900 | proliferating cell nuclear antigen (PCNA) | DNA replication | 0 | 0 | 0 | na | DNA binding, DNA polymerase processivity factor activity | regulation of DNA replication | PCNA complex |
| F | BANKA_113820 | RuvB-like helicase, putative (RUVB3) | -2.45 | 0.73 | PF3D7_1362200 | ATP-dependent DNA helicase, putative | DNA replication | 0 | 0 | 0 | na | ATP binding, DNA helicase activity, nucleoside-triphosphatase activity | null | nucleus |
| F | BANKA_114060 | DNA-directed RNA polymerase II, putative | | | PF3D7_1364800 | DNA-directed RNA polymerase 2, putative | transcription | 0 | 0 | 0 | na | DNA binding, DNA-directed RNA polymerase activity | transcription, DNA-dependent | nucleus |
| F | BANKA_114170 | 60S ribosomal protein L40/UBI, putative | | | PF3D7_1365900 | 60S ribosomal protein L40/UBI, putative | proteasome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| F | BANKA_114280 | conserved Plasmodium protein, unknown function | | | PF3D7_1366900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| F | BANKA_114300 | U1 small nuclear ribonucleoprotein, putative | | | PF3D7_1367100 | U1 small nuclear ribonucleoprotein, putative | mRNA | 0 | 0 | 0 | na | nucleic acid binding | null | null |
| F | BANKA_114590 | conserved rodent malaria protein, unknown function | | | | | unknown function | 0 | 0 | 0 | na | null | null | null |
| F | BANKA_120010 | BIR protein | | | | | BIR protein | 0 | 0 | 0 | na | null | null | null |
| F | BANKA_120040 | BIR protein | | | | | BIR protein | 0 | 0 | 0 | na | null | null | null |
| F | BANKA_120060 | Plasmodium exported protein, unknown function | | | | | unknown function | 0 | 0 | 0 | KO not successful | null | null | null |
| F | BANKA_120150 | conserved Plasmodium protein, unknown function | | | PF3D7_1003100 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| F | BANKA_120220 | U5 small nuclear ribonuclear protein, putative | | | PF3D7_1003800 | U5 small nuclear ribonuclear protein, putative | mRNA | 0 | 0 | 0 | na | GTP binding, GTPase activity | null | null |
| F | BANKA_120240 | 60S ribosomal protein L13, putative | | | PF3D7_1004000 | 60S ribosomal protein L13, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, large ribosomal subunit, ribosome |
| F | BANKA_120270 | RNA-binding protein, putative | -2.26 | 0.75 | PF3D7_1004400 | RNA binding protein, putative | mRNA | 0 | 0 | 0 | na | nucleic acid binding | null | null |
| F | BANKA_120370 | regulator of nonsense transcripts, putative | | | PF3D7_1005500 | regulator of nonsense transcripts, putative | mRNA | 0 | 0 | 0 | na | ATP binding, DNA binding, helicase activity, zinc ion binding | nuclear-transcribed mRNA catabolic process, nonsense-mediated decay | y cytoplasm |
| F | BANKA_120420 | conserved Plasmodium protein, unknown function | | | PF3D7_1006000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | DNA binding, alkylbase DNA N-glycosylase activity | base-excision repair | null |
| F | BANKA_120440 | DNA/RNA-binding protein Alba 3, putative (ALBA3) | | | PF3D7_1006200 | DNA/RNA-binding protein Alba 3 (ALBA3) | mRNA | 0 | 0 | 0 | na | DNA binding, RNA binding | null | nucleus |
| F | BANKA_120480 | thioredoxin, putative | | | PF3D7_1006600 | thioredoxin, putative | redoxmeth | 0 | 0 | 0 | na | null | null | null |
| F | BANKA_120500 | RNA-binding protein, putative | | | PF3D7_1006800 | RNA binding protein, putative | mRNA | 0 | 0 | 0 | na | nucleic acid binding | null | null |
| F | BANKA_120540 | rhoGAP GTPase, putative | | | PF3D7_1007200 | rhoGAP GTPase, putative | gliding motility | 0 | 0 | 0 | na | null | signal transduction | intracellular |
| | | | | | | | | | | | | | | |

| PBANKA_120660 | 26S proteasome regulatory subunit 4, putative | | PF3D7_1008400 | 26S proteasome regulatory subunit 4, putative | proteasome | 0 | 0 | 0 | na | ATP binding, nucleoside-triphosphatase activity | protein catabolic process | cytoplasm, nucleus |
|---------------|--|------|-------------------|---|--------------------|---|---|---|--|---|---|--|
| PBANKA_120690 | tubulin beta chain, putative | | PF3D7_1008700 | tubulin beta chain | gliding motility | 0 | 0 | 0 | na | GTP binding, GTPase activity, structural molecule activity | microtubule-based movement, protein polymerization | microtubule, protein complex |
| PBANKA_120700 | small subunit rRNA processing protein, putative | | PF3D7_1008800 | small subunit rRNA processing protein, putative | RNA | 0 | 0 | 0 | na | null | null | null |
| PBANKA_120880 | conserved Plasmodium protein, unknown function | | PF3D7_1010400 | conserved Plasmodium protein, unknown function | kinase | 0 | 0 | 0 | na | null | null | null |
| PBANKA_120890 | conserved Plasmodium protein, unknown function | | | | proteasome | 0 | 0 | 0 | na | null | null | null |
| PBANKA_120900 | eukaryotic translation initiation factor 2, beta, putative | | PF3D7_1010600 | eukaryotic translation initiation factor 2 beta subunit, putative | translation | 0 | 0 | 0 | na | translation initiation factor activity | translational initiation | null |
| PBANKA_120990 | conserved Plasmodium protein, unknown function | | PF3D7_1011500 | conserved Plasmodium membrane protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_121070 | ubiquinol-cytochrome c reductase complex subunit, putative | | PF3D7_1012300 | ubiquinol-cytochrome c reductase complex subunit, putative | redoxmeth | 0 | 0 | 0 | na | ubiquinol-cytochrome-c reductase activity | mitochondrial electron transport, ubiquinol to cytochrome c | null |
| PBANKA_121090 | phosphoglucomutase, putative | | PF3D7_1012500 | phosphoglucomutase, putative | glycolysis | 0 | 0 | 0 | na | intramolecular transferase activity, phosphotransferases | carbohydrate metabolic process | null |
| PBANKA_121260 | male gamete fusion factor HAP2 (HAP2) | | PF3D7_1014200 | male gamete fusion factor HAP2, putative (HAP2) | sexual development | 0 | 0 | 0 | different in fertilization, ookinete, oocyst, sporozoite and liver stage | null | null | null |
| PBANKA_121270 | conserved Plasmodium protein, unknown function | | PF3D7_1014300 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | not different from wild type | null | null | null |
| PBANKA_121300 | transcriptional activator ADA2, putative | | PF3D7_1014600 | transcriptional coactivator ADA2 (ADA2) | transcription | 0 | 0 | 0 | na | DNA binding, zinc ion binding | null | null |
| PBANKA_121350 | conserved Plasmodium protein, unknown function | | PF3D7_1015100 | conserved protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_121360 | cysteinyl-tRNA synthetase, putative | | PF3D7_1015200.1/2 | cysteinyl-tRNA synthetase, putative | translation | 0 | 0 | 0 | na | ATP binding, cysteine-tRNA ligase activity | cysteinyl-tRNA aminoacylation, translation | cytoplasm |
| PBANKA_121400 | heat shock protein 60, putative | | PF3D7_1015600 | heat shock protein 60 (HSP60) | chaperone | 0 | 0 | 0 | na | ATP binding, protein binding | cellular protein metabolic process | null |
| PBANKA_121430 | enolase, putative (ENO) | | PF3D7_1015900 | enolase (ENO) | glycolysis | 0 | 0 | 0 | na | phosphopyruvate hydratase activity | glycolysis | phosphopyruvate hydratase complex |
| PBANKA_121470 | conserved Plasmodium protein, unknown function | | PF3D7_0323800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_121530 | conserved Plasmodium protein, unknown function | | PF3D7_0323200 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_121650 | peptidyl-prolyl cis-trans isomerase, putative (CYP19A) | | PF3D7_0322000 | peptidyl-prolyl cis-trans isomerase (CYP19A) | chaperone | 0 | 0 | 0 | na | peptidyl-prolyl cis-trans isomerase activity | null | null |
| PBANKA_121760 | histone H2A variant, putative (H2A.Z) | | PF3D7_0320900 | histone H2A variant, putative (H2A.Z) | chromatin | 0 | 0 | 0 | na | DNA binding | nucleosome assembly | nucleosome, nucleus |
| PBANKA_121770 | ATP-dependent RNA Helicase (DOZI) | | PF3D7_0320800 | ATP-dependent RNA helicase, putative (DOZI) | mRNA | 0 | 0 | 0 | different in fertilization and ookinete | ATP binding, ATP-dependent helicase activity, helicase activity, nucleic acid binding | null | null |
| PBANKA_121830 | CPW-WPC family protein, putative -2.75 | 0.75 | PF3D7_0320200 | CPW-WPC family protein | adhesin | 0 | 0 | 0 | na | null | null | null |
| PBANKA_121840 | SNARE protein, putative | | PF3D7_0320100 | SNARE protein (SEC22) | trafficking | 0 | 0 | 0 | na | null | vesicle-mediated transport | integral to membrane |
| PBANKA_121850 | conserved Plasmodium protein, unknown function | | PF3D7_0320000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_121860 | conserved Plasmodium protein, unknown function | | PF3D7_0319900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_121890 | elongation factor 1 (EF-1), putative | | | | translation | 0 | 0 | 0 | na | translation elongation factor activity | translational elongation | eukaryotic translation elongation factor 1 complex |
| PBANKA_121900 | heat shock protein 90, putative | | PF3D7_0708500 | heat shock protein 86 family protein | chaperone | 0 | 0 | 0 | na | null | null | null |
| PBANKA_121920 | Cg8 protein, putative | | PF3D7_0708700 | Cg8 protein | redoxmeth | 0 | 0 | 0 | na | null | null | null |
| PBANKA_122120 | 60S ribosomal protein L34a, putative | | PF3D7_0710600 | 60S ribosomal protein L34a, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_122130 | conserved Plasmodium protein, unknown function | | PF3D7_0710700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_122240 | ubiquitin regulatory protein, putative | | PF3D7_0808300 | ubiquitin regulatory protein, putative | proteasome | 0 | 0 | 0 | na | null | null | null |
| | | | | | | | | | | | | |

| PBANKA_122280 | tyrosyl-tRNA synthetase, putative | PF3D7_0807900 | tyrosyl-tRNA synthetase, putative | translation | 0 | 0 | 0 na | ATP binding, tyrosine-tRNA ligase activity | translation, tyrosyl-tRNA aminoacylation | cytoplasm |
|---------------|---|---------------|--|-----------------------------|---|---|--|--|--|--|
| PBANKA_122420 | DnaJ protein, putative | PF3D7_0806500 | DnaJ protein, putative | chaperone | 0 | 0 | 0 na | heat shock protein binding, unfolded protein binding | protein folding | null |
| PBANKA_122540 | conserved Plasmodium protein, unknown function | PF3D7_0805200 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_122660 | tubulin gamma chain, putative | PF3D7_0803700 | tubulin gamma chain (g-tub) | gliding motility | 0 | 0 | 0 na | GTP binding, GTPase activity | microtubule-based process, protein polymerization | microtubule, protein complex |
| PBANKA_122680 | DNA repair protein rad54, putative -1.65 0.7 | PF3D7_0803400 | DNA repair protein rad54, putative | DNA repair | 0 | 0 | 0 na | ATP binding, DNA binding, helicase activity | null | null |
| PBANKA_122720 | peptidyl-prolyl cis-trans isomerase, putative (CYP81) | PF3D7_0803000 | peptidyl-prolyl cis-trans isomerase (CYP81) | chaperone | 0 | 0 | 0 na | peptidyl-prolyl cis-trans isomerase activity | null | null |
| PBANKA_122740 | serine/threonine protein phosphatase 2B catalytic subunit A, putative (CNA) | PF3D7_0802800 | serine/threonine protein phosphatase, putative | phosphatase | 0 | 0 | 0 na | hydrolase activity | null | null |
| PBANKA_122780 | conserved Plasmodium protein, unknown function | PF3D7_0802400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_122830 | conserved Plasmodium protein, unknown function | PF3D7_0801900 | conserved Plasmodium protein, unknown function | redoxmeth | 0 | 0 | 0 na | electron carrier activity, oxidoreductase activity | null | null |
| PBANKA_122920 | 60S ribosomal protein L19, putative | PF3D7_0614500 | 60S ribosomal protein L19, putative | ribosome | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_123005 | ribonuclease, putative, fragment | PF3D7_0615400 | ribonuclease, putative | mRNA | 0 | 0 | 0 na | RNA binding, ribonuclease activity | null | null |
| PBANKA_123100 | 40S ribosomal protein S14, putative | PF3D7_0516200 | 40S ribosomal protein S14, putative | ribosome | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_123160 | transcription factor with AP2 domain(s), putative (ApiAP2) | PF3D7_0516800 | transcription factor with AP2 domain(s) (ApiAP2) | transcription | 0 | 0 | 0 na | sequence-specific DNA binding transcription factor activity | regulation of transcription, DNA-dependent | null |
| PBANKA_123170 | 60S ribosomal protein L8, putative | PF3D7_0516900 | 60S ribosomal protein L8, putative | ribosome | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_123180 | 60S ribosomal protein L12, putative | PF3D7_0517000 | 60S ribosomal protein L12, putative | ribosome | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_123350 | mRNA-binding protein PUF1 (PUF1) | PF3D7_0518700 | mRNA-binding protein PUF1 (PUF1) | mRNA | 0 | 0 | O different in fertilization, ookinete, oocyst, sporozoite and liver stage | RNA binding | null | null |
| PBANKA_123420 | 40S ribosomal protein S24, putative | PF3D7_0519400 | 40S ribosomal protein S24, putative | ribosome | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_123450 | conserved Plasmodium protein, unknown function | PF3D7_0519700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_123480 | 40S ribosomal protein S9, putative | PF3D7_0520000 | 40S ribosomal protein S9, putative | ribosome | 0 | 0 | 0 na | rRNA binding, structural constituent of ribosome | translation | intracellular, small ribosomal subunit |
| PBANKA_123510 | U6 snRNA-associated sm-like protein Ism2, putative (LSM2) | PF3D7_0520300 | U6 snRNA-associated sm-like protein lsm2, putative (LSM2) | mRNA | 0 | 0 | 0 na | null | null | null |
| PBANKA_123770 | zinc finger protein, putative, fragment | | | | 0 | 0 | 0 na | null | null | null |
| PBANKA_123880 | karyopherin beta, putative | PF3D7_0524000 | karyopherin beta (KASbeta) | nuclear pore | 0 | 0 | 0 na | binding | null | null |
| PBANKA_123920 | cytosolic preribosomal GTP-binding protein, putative | PF3D7_0524400 | cytosolic preribosomal GTP-binding protein, putative | ribosome assembly | 0 | 0 | 0 na | GTP binding | null | intracellular |
| PBANKA_123940 | organelle ribosomal protein L7/L12 precursor, putative | PF3D7_0524600 | organelle ribosomal protein L7/L12 precursor, putative | ribosome | 0 | 0 | 0 na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_123980 | zinc finger protein, putative | PF3D7_0525000 | zinc finger protein, putative | RNA binding, RNA processing | 0 | 0 | 0 na | nucleic acid binding, zinc ion binding | null | null |
| PBANKA_124180 | minchromosome maintenance (MCM) complex subunit, putative (MCM3) | PF3D7_0527000 | minchromosome maintenance (MCM) complex subunit, putative (MCM3) | DNA replication | 0 | 0 | 0 na | ATP binding, DNA binding, nucleoside-triphosphatase activity | / DNA replication, DNA-dependent DNA replication initiation | nucleus |
| PBANKA_124190 | ubiquitin-conjugating enzyme E2 N, putative (UBC13) | PF3D7_0527100 | ubiquitin-conjugating enzyme E2 N, putative (UBC13) | proteasome | 0 | 0 | 0 na | small conjugating protein ligase activity | post-translational protein modification, regulation of protein metabolic process | null |
| PBANKA_124220 | conserved Plasmodium protein, unknown function | PF3D7_0527400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| PBANKA_124230 | 58 kDa phosphoprotein,heat shock-related protein (HRP) | PF3D7_0527500 | hsp70 interacting protein, putative | trafficking | 0 | 0 | 0 na | null | chaperone cofactor-dependent protein refolding | null |
| PBANKA_124310 | F-actin capping protein, alpha subunit, putative | PF3D7_0528500 | f-actin capping protein alpha subunit, putative | gliding motility | 0 | 0 | 0 na | actin binding | actin cytoskeleton organization | F-actin capping protein complex |
| PBANKA_124320 | conserved Plasmodium protein, unknown function | PF3D7_0528600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | null | null | null |
| | | | | | | | | | | |

| PBANKA_124590 | Plasmodium exported protein, unknown function | | | | | unknown function | 0 | 0 | 0 | na | null | null | null |
|---------------|---|-------|------|-------------------|--|------------------|---|---|---|---|--|--|--|
| PBANKA_124640 | BIR protein | | | | | BIR protein | 0 | 0 | 0 | na | null | null | null |
| PBANKA_124690 | BIR protein, pseudogene | | | | | BIR protein | 0 | 0 | 0 | na | null | null | null |
| PBANKA_130050 | tubulin, putative | | | PF3D7_1475700 | tubulin, putative | chromatin | 0 | 0 | 0 | na | GTP binding, GTPase activity, structural molecule activity | microtubule-based movement, protein polymerization | microtubule, protein complex |
| PBANKA_130070 | LCCL domain-containing protein (CCp1) | | | PF3D7_1475500 | LCCL domain-containing protein (CCp1) | adhesin | 0 | 0 | 0 | different in occyst and sporozoite; (tagged) also in fertilization and ookinete | receptor binding | cell adhesion, signal transduction | null |
| PBANKA_130130 | trailer hitch homolog, putative (CITH) | | | PF3D7_1474900 | trailer hitch homolog, putative (CITH) | mRNA | 0 | 0 | 0 | different in fertilization and ookinete | null | null | null |
| PBANKA_130140 | proteosome subunit alpha type 1, putative | | | PF3D7_1474800 | proteosome subunit alpha type 1, putative | proteasome | 0 | 0 | 0 | na | threonine-type endopeptidase activity | ubiquitin-dependent protein catabolic process | proteasome core complex |
| PBANKA_130160 | conserved Plasmodium protein, unknown function | | | PF3D7_1474600 | conserved Plasmodium membrane protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_130260 | small subunit DNA primase, putative | | | PF3D7_1438700 | small subunit DNA primase | DNA replication | 0 | 0 | 0 | na | DNA primase activity | DNA replication, synthesis of RNA primer | null |
| PBANKA_130280 | thioredoxin peroxidase 1 (TPx1) | | | PF3D7_1438900 | thioredoxin peroxidase 1 (Trx-Px1) | redoxmeth | 0 | 0 | 0 | different in sporozoite and liver stage | antioxidant activity, oxidoreductase activity | cell redox homeostasis | null |
| PBANKA_130330 | ubiquinol-cytochrome c reductase, iron-sulfur subunit, putative | | | PF3D7_1439400 | ubiquinol-cytochrome c reductase iron-sulfur subunit, putative | redoxmeth | 0 | 0 | 0 | na | 2 iron, 2 sulfur cluster binding, electron carrier activity, ubiquinol-cytochrome-c reductase activity | oxidation-reduction process | membrane |
| PBANKA_130350 | conserved Plasmodium protein, unknown function | | | PF3D7_1439600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | mitochondrion |
| PBANKA_130390 | conserved Plasmodium protein, unknown function | -3.01 | 0.75 | PF3D7_1440000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | RNA-directed RNA polymerase activity | viral genome replication | null |
| PBANKA_130510 | 60S ribosomal protein L10a, putative | | | PF3D7_1441200 | 60S ribosomal protein L1, putative | ribosome | 0 | 0 | 0 | na | RNA binding | RNA processing | null |
| PBANKA_130590 | ADP-ribosylation-like factor, putative | | | PF3D7_1442000 | ADP-ribosylation factor, putative | trafficking | 0 | 0 | 0 | na | GTP binding | intracellular protein transport, small GTPase mediated signal transduction | intracellular |
| PBANKA_130600 | conserved Plasmodium protein, unknown function | | | PF3D7_1442100 | conserved Plasmodium protein, unknown function | DNA repair | 0 | 0 | 0 | na | null | null | null |
| PBANKA_130620 | tRNA binding protein, putative | | | PF3D7_1442300 | tRNA binding protein, putative | translation | 0 | 0 | 0 | na | tRNA binding | null | null |
| PBANKA_130770 | zinc finger protein, putative | | | PF3D7_1443800 | zinc finger protein, putative | mRNA | 0 | 0 | 0 | na | nucleic acid binding, protein binding, zinc ion binding | null | null |
| PBANKA_130810 | calmodulin-like protein | | | PF3D7_1444200 | calmodulin-like protein | kinase | 0 | 0 | 0 | na | null | null | null |
| PBANKA_130920 | serine/threonine kinase-1, putative | | | PF3D7_1445400 | protein serine/threonine kinase-1 (CLK1) | kinase | 0 | 0 | 0 | KO not successful | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| PBANKA_130930 | conserved Plasmodium protein, unknown function | | | PF3D7_1445500 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_131040 | centrin, putative | | | PF3D7_1446600 | centrin-2 (CEN2) | chromatin | 0 | 0 | 0 | na | calcium ion binding | null | null |
| PBANKA_131080 | 40S ribosomal protein S2, putative | | | PF3D7_1447000 | 40S ribosomal protein S2, putative | ribosome | 0 | 0 | 0 | na | RNA binding, structural constituent of ribosome | translation | intracellular, ribosome, small ribosomal subunit |
| PBANKA_131090 | conserved Plasmodium protein, unknown function | | | PF3D7_1447100 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_131440 | conserved Plasmodium protein, unknown function | | | PF3D7_1450700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_131480 | elongation factor 2, putative | | | PF3D7_1451100 | elongation factor 2 | translation | 0 | 0 | 0 | na | GTP binding, GTPase activity | null | null |
| PBANKA_131490 | conserved Plasmodium protein, unknown function | -2.62 | 0.75 | PF3D7_1451200 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_131550 | sortilin, putative | | | PF3D7_1451800 | sortilin, putative | large tethering | 0 | 0 | 0 | na | null | null | integral to membrane |
| PBANKA_131740 | co-chaperone p23, putative | | | PF3D7_1453700 | co-chaperone p23 (P23) | chaperone | 0 | 0 | 0 | na | null | null | null |
| PBANKA_131890 | conserved Plasmodium protein, unknown function | | | PF3D7_1455200.1/2 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | methyltransferase activity | metabolic process | null |
| PBANKA_132010 | conserved Plasmodium protein, unknown function (UIS25) | | | PF3D7_1456400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| | | | | | | | | | | | | | |

| PBA | NKA_132020 | conserved Plasmodium protein, unknown function | | | PF3D7_1456500 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
|-----|------------|---|-------|------|---------------|---|---------------------------------|---|---|---|------------------------------------|--|--|--|
| PBA | NKA_132070 | signal peptide peptidase, putative (SPP) | | | PF3D7_1457000 | signal peptide peptidase (SPP) | post-translational modification | 0 | 0 | 0 | na | aspartic-type endopeptidase activity | null | endoplasmic reticulum, integral to membrane |
| PBA | NKA_132150 | conserved Plasmodium protein, unknown function | | | PF3D7_1457800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_132400 | ribosomal protein L29, putative | | | PF3D7_1460300 | ribosomal protein L29, putative | ribosome | 0 | 0 | 0 | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBA | NKA_132510 | conserved Plasmodium protein, unknown function | | | PF3D7_1461400 | conserved Plasmodium protein, unknown function | kinase | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_132520 | conserved Plasmodium protein, unknown function | | | PF3D7_1461500 | conserved Plasmodium protein, unknown function | meiosis | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_132610 | conserved Plasmodium protein, unknown function | | | PF3D7_1462500 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_132640 | glyceraldehyde-3-phosphate dehydrogenase, putative | | | PF3D7_1462800 | glyceraldehyde-3-phosphate dehydrogenase (GAPDH |) glycolysis | 0 | 0 | 0 | na | NAD binding, glyceraldehyde-3-phosphate dehydrogenase (NAD+) (phosphorylating) activity | glucose metabolic process | null |
| PBA | NKA_132730 | conserved Plasmodium protein, unknown function | | | | | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_132830 | ATP-dependent protease la, putative | | | PF3D7_1464900 | ATP-dependent protease la, putative | protease | 0 | 0 | 0 | na | ATP binding, metalloendopeptidase activity, nucleoside-triphosphatase activity, zinc ion binding | protein catabolic process, proteolysis | integral to membrane, membrane |
| PBA | NKA_132840 | conserved Plasmodium protein, unknown function | | | PF3D7_1465000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_132900 | conserved Plasmodium protein, unknown function | | | PF3D7_1465600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_132930 | 40S ribosomal protein S3, putative | | | PF3D7_1465900 | 40S ribosomal protein S3, putative | ribosome | 0 | 0 | 0 | na | RNA binding, structural constituent of ribosome | translation | intracellular, ribosome, small ribosomal subunit |
| PBA | NKA_132980 | transcription factor with AP2 domain(s) (AP2-SP) | | | PF3D7_1466400 | transcription factor with AP2 domain(s) (ApiAP2) | transcription | 0 | 0 | 0 | different in oocyst and sporozoite | sequence-specific DNA binding transcription factor activity | regulation of transcription, DNA-dependent | null |
| PBA | NKA_133140 | conserved Plasmodium protein, unknown function | | | PF3D7_1468100 | conserved Plasmodium protein, unknown function | galactose metabolism | 0 | 0 | 0 | KO not successful | ATP binding | null | null |
| PBA | NKA_133210 | conserved Plasmodium protein, unknown function | | | PF3D7_1468900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_133300 | conserved Plasmodium protein, unknown function | | | PF3D7_1469800 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_133320 | conserved Plasmodium protein, unknown function | | | PF3D7_1470000 | ATG12 | autophagy | 0 | 0 | 0 | na | null | autophagic vacuole assembly | cytoplasm |
| PBA | NKA_133480 | conserved Plasmodium protein, unknown function | -2.16 | 0.72 | PF3D7_1471600 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_133490 | conserved Plasmodium protein, unknown function | -4.61 | 0.75 | | | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_133520 | pre-mRNA-splicing factor, putative | | | PF3D7_1472000 | pre-mRNA-splicing factor ISY1 homolog, putative | mRNA | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_133570 | conserved Plasmodium protein, unknown function | | | PF3D7_1472500 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_133660 | conserved Plasmodium protein, unknown function | | | PF3D7_1473400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_133690 | conserved Plasmodium protein, unknown function | | | | | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_133710 | splicing factor, putative | | | PF3D7_1321700 | splicing factor, putative | mRNA | 0 | 0 | 0 | na | RNA binding | mRNA processing | nucleus |
| PBA | NKA_133840 | vacuolar ATP synthase subunit g, putative | | | PF3D7_1323200 | vacuolar ATP synthase subunit g, putative | vacuolar ATP synthase | 0 | 0 | 0 | KO not successful | hydrolase activity, acting on acid anhydrides, catalyzing transmembrane movement of substances | proton transport | vacuolar proton-transporting V-type ATPase complex |
| PBA | NKA_133940 | microfibril-associated protein homologue, putative | | | PF3D7_1324200 | micro-fibrillar-associated protein, putative | gliding motility | 0 | 0 | 0 | na | null | null | extracellular region |
| PBA | NKA_134010 | L-lactate dehydrogenase (LDH) | | | PF3D7_1324900 | L-lactate dehydrogenase (LDH) | glycolysis | 0 | 0 | 0 | na | binding, oxidoreductase activity, acting on the CH-OH group of donors, NAD or NADP as acceptor | glycolysis | null |
| PBA | NKA_134020 | U6 snRNA-associated Sm-like protein LSm6, putative (LSM6) | -2.49 | 0.75 | PF3D7_1325000 | U6 snRNA-associated Sm-like protein LSm6, putative (LSM6) | mRNA | 0 | 0 | 0 | na | null | null | null |
| PBA | NKA_134040 | oxidoreductase, putative | | | PF3D7_1325200 | oxidoreductase, putative | glycolysis | 0 | 0 | 0 | na | binding, oxidoreductase activity, acting on the CH-OH group of donors, NAD or NADP as acceptor | glycolysis | null |
| PBA | NKA_134150 | RNA binding protein, putative | | | PF3D7_1326300 | RNA binding protein, putative | mRNA | 0 | 0 | 0 | na | nucleic acid binding | null | null |
| | | | | | | | | | | | | | | |

| PBA | NKA_134320 | conserved Plasmodium protein, unknown function | PF3D7_1328000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | na | null | null | null |
|-----|------------|--|---------------|---|-------------------------------|---|---|------|-------------------|--|--|---|
| PBA | NKA_134560 | elongation factor Tu, putative | PF3D7_1330600 | elongation factor Tu, putative | translation | 0 | 0 | 0 na | na | GTP binding, GTPase activity, translation elongation factor activity | translational elongation | intracellular |
| PBA | NKA_134610 | conserved Plasmodium protein, unknown function | PF3D7_1331200 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_134840 | histone h3-like centromeric protein cse4, putative (CENPA) | PF3D7_1333700 | histone H3 variant, putative (CenH3) | chromatin | 0 | 0 | 0 na | na | DNA binding | nucleosome assembly | nucleosome, nucleus |
| PBA | NKA_135000 | conserved Plasmodium protein, unknown function | PF3D7_1336100 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_135030 | glucose-induced degradation protein 8, putative (GID8) | PF3D7_1336400 | glucose-induced degradation protein 8, putative (GID8) |) protein-protein interaction | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_135050 | conserved Plasmodium protein, unknown function | PF3D7_1336700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | na | calcium ion binding | null | null |
| PBA | NKA_135130 | conserved Plasmodium protein, unknown function | | | unknown function | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_135210 | conserved Plasmodium protein, unknown function | PF3D7_1338400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_135430 | sodium-dependent phosphate transporter, putative (PiT) | PF3D7_1340900 | sodium-dependent phosphate transporter (PiT) | acidocalcisome, transporter | 0 | 0 | 0 na | na | inorganic phosphate transmembrane transporter activity | phosphate ion transport | membrane |
| PBA | NKA_135440 | 60S ribosomal protein L18, putative | PF3D7_1341200 | 60S ribosomal protein L18, putative | ribosome | 0 | 0 | 0 na | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBA | NKA_135450 | 60S ribosomal protein L18-2, putative | PF3D7_1341300 | 60S ribosomal protein L18-2, putative | ribosome | 0 | 0 | 0 na | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBA | NKA_135500 | vacuolar ATP synthase subunit d, putative | PF3D7_1341900 | vacuolar ATP synthase subunit d, putative | vacuolar ATP synthase | 0 | 0 | 0 na | na | proton-transporting ATPase activity, rotational mechanism | ATP synthesis coupled proton transport | proton-transporting two-sector ATPase complex, catalytic domain |
| PBA | NKA_135510 | 40S ribosomal protein S6, putative | PF3D7_1342000 | 40S ribosomal protein S6, putative | ribosome | 0 | 0 | 0 na | na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBA | NKA_135550 | casein kinase II beta chain, putative | PF3D7_1342400 | casein kinase II beta chain (CK2beta2) | kinase | 0 | 0 | 0 na | na | protein kinase regulator activity | null | protein kinase CK2 complex |
| PBA | NKA_135580 | transcription activator, putative | PF3D7_1342700 | transcription activator, putative | transcription | 0 | 0 | 0 na | na | DNA binding, DNA-directed RNA polymerase activity | transcription, DNA-dependent | null |
| PBA | NKA_135630 | conserved Plasmodium protein, unknown function | PF3D7_1343300 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_135670 | kelch protein k13, putative | PF3D7_1343700 | kelch protein, putative | galactose metabolism | 0 | 0 | 0 na | na | protein binding, voltage-gated potassium channel activity | potassium ion transport | membrane, voltage-gated potassium channel complex |
| PBA | NKA_135780 | conserved Plasmodium protein, unknown function | PF3D7_1344900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_135830 | conserved Plasmodium protein, unknown function | PF3D7_1345400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_135980 | conserved Plasmodium protein, unknown function | PF3D7_1346900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_136030 | DNA/RNA-binding protein Alba 4, putative (ALBA4) | PF3D7_1347500 | DNA/RNA-binding protein Alba 4 (ALBA4) | mRNA | 0 | 0 | 0 na | na | DNA binding, RNA binding | null | cytoplasm, nucleus |
| PBA | NKA_136120 | conserved Plasmodium protein, unknown function | PF3D7_1348400 | conserved Plasmodium membrane protein, unknown function | n unknown function | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_136150 | conserved Plasmodium protein, unknown function | PF3D7_1348700 | conserved Plasmodium protein, unknown function | phototransduction | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_136200 | glutamate - tRNA ligase, putative | PF3D7_1349200 | glutamatetRNA ligase, putative | translation | 0 | 0 | 0 na | na | ATP binding, glutamate-tRNA ligase activity | glutamyl-tRNA aminoacylation, translation | cytoplasm |
| PBA | NKA_136420 | 60S ribosomal protein L17, putative | PF3D7_1351400 | 60S ribosomal protein L17, putative | ribosome | 0 | 0 | 0 na | na | structural constituent of ribosome | translation | intracellular, large ribosomal subunit, ribosome |
| PBA | NKA_140010 | BIR protein | | | BIR protein | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_140060 | cytoadherence linked asexual protein, putative | | | | 0 | 0 | 0 K | KO not successful | null | null | null |
| PBA | NKA_140070 | Plasmodium exported protein, unknown function | | | unknown function | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_140150 | conserved Plasmodium protein, unknown function | PF3D7_1303000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | na | null | null | null |
| PBA | NKA_140260 | DNA ligase I, putative | PF3D7_1304100 | DNA ligase I (LigI) | DNA repair | 0 | 0 | 0 K | KO not successful | ATP binding, DNA binding, DNA ligase (ATP) activity | DNA recombination, DNA repair, DNA replication | null |
| | | | | | | | | | | | | |

| PBANKA_140490 | 26s protease regulatory subunit 10b, putative (RPT4) | PF3D7_1306400 | 26S proteasome regulatory subunit, putative | proteasome | 0 (|) | 0 na | ATP binding, nucleoside-triphosphatase activity | protein catabolic process | cytoplasm, nucleus |
|---|---|---|---|--|---|--------|--|--|---|---|
| PBANKA_140610 | DNA-directed RNA polymerase alpha chain, putative | PF3D7_1307600 | DNA-directed RNA polymerase alpha chain, putative | transcription | 0 (|) | 0 na | DNA binding, DNA-directed RNA polymerase activity, protein dimerization activity | transcription, DNA-dependent | null |
| PBANKA_140700 | conserved Plasmodium protein, unknown function | PF3D7_1308500 | conserved Plasmodium protein, unknown function | glycolysis | 0 (|) | O na | null | null | null |
| PBANKA_140740 | mRNA-decapping enzyme 2, putative (DCP2) | PF3D7_1308900 | mRNA-decapping enzyme 2, putative (DCP2) | mRNA | 0 (|) | 0 na | hydrolase activity | null | null |
| PBANKA_140760 | 60S ribosomal protein L24, putative | PF3D7_1309100 | 60S ribosomal protein L24, putative | ribosome | 0 (|) | O na | structural constituent of ribosome | translation | intracellular, ribosome |
| PBANKA_140780 | U4/U6 small nuclear ribonucleoprotein PRP3, putative (PRPF3) | PF3D7_1309300 | U4/U6 small nuclear ribonucleoprotein PRP3, putative (PRPF3) | mRNA | 0 (|) | 0 na | null | null | null |
| PBANKA_140920 | rna-binding protein, putative | PF3D7_1310700 | conserved Plasmodium protein, unknown function | unknown function | 0 (|) | O na | nucleic acid binding | null | null |
| PBANKA_140960 | conserved Plasmodium protein, unknown function | PF3D7_1311100 | conserved Plasmodium protein, unknown function | unknown function | 0 (|) | O na | null | null | null |
| PBANKA_141000 | 26s protease regulatory subunit 7, putative (RPT1) | PF3D7_1311500 | 26S proteasome regulatory subunit 7, putative | proteasome | 0 (|) | O na | ATP binding, nucleoside-triphosphatase activity | protein catabolic process | cytoplasm, nucleus |
| PBANKA_141120 | conserved Plasmodium protein, unknown function | PF3D7_1312800 | conserved Plasmodium protein, unknown function | unknown function | 0 (|) | O na | null | null | null |
| PBANKA_141140 | ubiquitin-like protein nedd8 homologue, putative | PF3D7_1313000 | ubiquitin-like protein nedd8 homologue, putative (Nedd8) |) proteasome | 0 (|) | O na | null | null | null |
| PBANKA_141170 | peptidyl-prolyl cis-trans isomerase, putative | PF3D7_1313300 | peptidyl-prolyl cis-trans isomerase, putative | chaperone | 0 (|) | O na | null | protein folding | null |
| PBANKA_141320 | conserved Plasmodium protein, unknown function | PF3D7_1314700 | conserved Plasmodium protein, unknown function | unknown function | 0 (|) | O na | null | null | null |
| PBANKA_141400 | conserved Plasmodium protein, unknown function | PF3D7_1315500 | conserved Plasmodium protein, unknown function | unknown function | 0 (|) | O na | null | null | null |
| PBANKA_141470 | ADP-ribosylation factor, putative | PF3D7_1316200 | ADP-ribosylation factor, putative | trafficking | 0 (|) | 0 KO not successful | GTP binding | small GTPase mediated signal transduction | intracellular |
| PBANKA_141560 | minchromosome maintenance (MCM) complex subunit, putative (MCM4) | PF3D7_1317100 | minchromosome maintenance (MCM) complex subunit (MCM4) | DNA replication | 0 (|) | 0 na | ATP binding, DNA binding | DNA replication, DNA-dependent DNA replication initiation | nucleus |
| | | | | | | | | | | |
| PBANKA_141630 | 40S ribosomal protein S15/S19, putative | PF3D7_1317800 | 40S ribosomal protein S15/S19, putative | ribosome | 0 (|) | 0 na | structural constituent of ribosome | translation | intracellular, ribosome, small ribosomal subunit |
| PBANKA_141630 PBANKA_141690 | 40S ribosomal protein S15/S19, putative chromosome segregation protein, putative | PF3D7_1317800 PF3D7_1318400 | 40S ribosomal protein S15/S19, putative chromosome segregation protein, putative | ribosome | 0 0 | • | 0 na 0 na | structural constituent of ribosome ATP binding, protein binding | translation chromosome organization | intracellular, ribosome, small ribosomal subunit chromosome |
| | | | | | |) | | | | , , |
| PBANKA_141690 | chromosome segregation protein, putative | PF3D7_1318400 | chromosome segregation protein, putative | chromatin trafficking | 0 (|)) | 0 na | ATP binding, protein binding | chromosome organization | chromosome |
| PBANKA_141690 PBANKA_141730 | chromosome segregation protein, putative translocation protein sec63, putative | PF3D7_1318400 PF3D7_1318800 | chromosome segregation protein, putative secretory complex protein 63 (SEC63) | chromatin trafficking chaperone | 0 (|) | o na O na | ATP binding, protein binding heat shock protein binding, unfolded protein binding | chromosome organization protein folding | chromosome |
| PBANKA_141690 PBANKA_141730 PBANKA_141840 | chromosome segregation protein, putative translocation protein sec63, putative ATP-dependent Clp protease adaptor protein, putative | PF3D7_1318400 PF3D7_1318800 PF3D7_1320100 | chromosome segregation protein, putative secretory complex protein 63 (SEC63) ATP-dependent Clp protease adaptor protein ClpS, putative | chromatin trafficking chaperone unknown function | 0 0 |) | O na O na O na | ATP binding, protein binding heat shock protein binding, unfolded protein binding null | chromosome organization protein folding protein catabolic process | chromosome null |
| PBANKA_141690 PBANKA_141730 PBANKA_141840 PBANKA_142050 | chromosome segregation protein, putative translocation protein sec63, putative ATP-dependent Clp protease adaptor protein, putative conserved Plasmodium protein, unknown function | PF3D7_1318400 PF3D7_1318800 PF3D7_1320100 PF3D7_0713900 | chromosome segregation protein, putative secretory complex protein 63 (SEC63) ATP-dependent Clp protease adaptor protein ClpS, putative conserved Plasmodium protein, unknown function | chromatin trafficking chaperone unknown function | 0 0 | | 0 na 0 na 0 na 0 na | ATP binding, protein binding heat shock protein binding, unfolded protein binding null null | chromosome organization protein folding protein catabolic process null | chromosome null null |
| PBANKA_141690 PBANKA_141730 PBANKA_141840 PBANKA_142050 PBANKA_142070 | chromosome segregation protein, putative translocation protein sec63, putative ATP-dependent Clp protease adaptor protein, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function | PF3D7_1318400 PF3D7_1318800 PF3D7_1320100 PF3D7_0713900 PF3D7_0714100 | chromosome segregation protein, putative secretory complex protein 63 (SEC63) ATP-dependent Clp protease adaptor protein ClpS, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function | chromatin trafficking chaperone unknown function unknown function | 0 0 0 | | 0 na 0 na 0 na 0 na 0 na | ATP binding, protein binding heat shock protein binding, unfolded protein binding null null | chromosome organization protein folding protein catabolic process null null | chromosome null null null |
| PBANKA_141690 PBANKA_141730 PBANKA_141840 PBANKA_142050 PBANKA_142070 PBANKA_142100 | chromosome segregation protein, putative translocation protein sec63, putative ATP-dependent Clp protease adaptor protein, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function calmodulin, putative | PF3D7_1318400 PF3D7_1318800 PF3D7_1320100 PF3D7_0713900 PF3D7_0714100 PF3D7_0714400 | chromosome segregation protein, putative secretory complex protein 63 (SEC63) ATP-dependent Clp protease adaptor protein ClpS, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function calmodulin, putative | chromatin trafficking chaperone unknown function unknown function kinase | 0 0 0 | | 0 na | ATP binding, protein binding heat shock protein binding, unfolded protein binding null null null | chromosome organization protein folding protein catabolic process null null | chromosome null null null null |
| PBANKA_141690 PBANKA_141730 PBANKA_141840 PBANKA_142050 PBANKA_142070 PBANKA_142100 PBANKA_142350 | chromosome segregation protein, putative translocation protein sec63, putative ATP-dependent Clp protease adaptor protein, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function calmodulin, putative 60S ribosomal protein L13-2, putative | PF3D7_1318400 PF3D7_1318800 PF3D7_1320100 PF3D7_0713900 PF3D7_0714100 PF3D7_0714400 PF3D7_0814000 | chromosome segregation protein, putative secretory complex protein 63 (SEC63) ATP-dependent Clp protease adaptor protein ClpS, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function calmodulin, putative 60S ribosomal protein L13-2, putative | chromatin trafficking chaperone unknown function unknown function kinase ribosome | 0 | | 0 na | ATP binding, protein binding heat shock protein binding, unfolded protein binding null null null structural constituent of ribosome | chromosome organization protein folding protein catabolic process null null translation | chromosome null null null null intracellular, ribosome |
| PBANKA_141690 PBANKA_141730 PBANKA_141840 PBANKA_142050 PBANKA_142070 PBANKA_142100 PBANKA_142350 PBANKA_142360 | chromosome segregation protein, putative translocation protein sec63, putative ATP-dependent Clp protease adaptor protein, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function calmodulin, putative 60S ribosomal protein L13-2, putative 40S ribosomal protein S16, putative | PF3D7_1318400 PF3D7_1318800 PF3D7_1320100 PF3D7_0713900 PF3D7_0714100 PF3D7_0714400 PF3D7_0814000 PF3D7_0813900 | chromosome segregation protein, putative secretory complex protein 63 (SEC63) ATP-dependent Clp protease adaptor protein ClpS, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function calmodulin, putative 60S ribosomal protein L13-2, putative 40S ribosomal protein S16, putative | chromatin trafficking chaperone unknown function unknown function kinase ribosome | 0 | | 0 na | ATP binding, protein binding heat shock protein binding, unfolded protein binding null null null structural constituent of ribosome structural constituent of ribosome | chromosome organization protein folding protein catabolic process null null translation translation | chromosome null null null null intracellular, ribosome intracellular, ribosome |
| PBANKA_141690 PBANKA_141730 PBANKA_141840 PBANKA_142050 PBANKA_142070 PBANKA_142100 PBANKA_142350 PBANKA_142360 PBANKA_142430 | chromosome segregation protein, putative translocation protein sec63, putative ATP-dependent Clp protease adaptor protein, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function calmodulin, putative 60S ribosomal protein L13-2, putative 40S ribosomal protein S16, putative conserved Plasmodium protein, unknown function | PF3D7_1318400 PF3D7_1318800 PF3D7_1320100 PF3D7_0713900 PF3D7_0714100 PF3D7_0714400 PF3D7_0814000 PF3D7_0813900 PF3D7_0813300 | chromosome segregation protein, putative secretory complex protein 63 (SEC63) ATP-dependent Clp protease adaptor protein ClpS, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function calmodulin, putative 60S ribosomal protein L13-2, putative 40S ribosomal protein S16, putative conserved Plasmodium protein, unknown function | chromatin trafficking chaperone unknown function unknown function kinase ribosome ribosome unknown function | | | 0 na | ATP binding, protein binding heat shock protein binding, unfolded protein binding null null null structural constituent of ribosome structural constituent of ribosome | chromosome organization protein folding protein catabolic process null null translation translation | chromosome null null null null intracellular, ribosome intracellular, ribosome |
| PBANKA_141690 PBANKA_141730 PBANKA_141840 PBANKA_142050 PBANKA_142070 PBANKA_142100 PBANKA_142350 PBANKA_142360 PBANKA_142430 PBANKA_142430 | chromosome segregation protein, putative translocation protein sec63, putative ATP-dependent Clp protease adaptor protein, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function calmodulin, putative 60S ribosomal protein L13-2, putative 40S ribosomal protein S16, putative conserved Plasmodium protein, unknown function ubiquitin-conjugating enzyme, putative | PF3D7_1318400 PF3D7_1318800 PF3D7_1320100 PF3D7_0713900 PF3D7_0714100 PF3D7_0714400 PF3D7_0814000 PF3D7_0813900 PF3D7_0813300 PF3D7_0812600 | chromosome segregation protein, putative secretory complex protein 63 (SEC63) ATP-dependent Clp protease adaptor protein ClpS, putative conserved Plasmodium protein, unknown function calmodulin, putative 60S ribosomal protein L13-2, putative 40S ribosomal protein S16, putative conserved Plasmodium protein, unknown function ubiquitin conjugating enzyme, putative | chromatin trafficking chaperone unknown function unknown function kinase ribosome ribosome unknown function proteasome | | | 0 na | ATP binding, protein binding heat shock protein binding, unfolded protein binding null null null structural constituent of ribosome structural constituent of ribosome null small conjugating protein ligase activity | chromosome organization protein folding protein catabolic process null null translation translation null post-translational protein modification, regulation of protein metabolic process | chromosome null null null null intracellular, ribosome intracellular, null null |
| PBANKA_141690 PBANKA_141730 PBANKA_141840 PBANKA_142050 PBANKA_142070 PBANKA_142100 PBANKA_142350 PBANKA_142360 PBANKA_142430 PBANKA_142430 PBANKA_142490 PBANKA_142510 | chromosome segregation protein, putative translocation protein sec63, putative ATP-dependent Clp protease adaptor protein, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function calmodulin, putative 60S ribosomal protein L13-2, putative 40S ribosomal protein S16, putative conserved Plasmodium protein, unknown function ubiquitin-conjugating enzyme, putative karyopherin alpha, putative (KARalpha) | PF3D7_1318400 PF3D7_1318800 PF3D7_1320100 PF3D7_0713900 PF3D7_0714100 PF3D7_0714400 PF3D7_0814000 PF3D7_0813900 PF3D7_0812600 PF3D7_0812400 | chromosome segregation protein, putative secretory complex protein 63 (SEC63) ATP-dependent Clp protease adaptor protein ClpS, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function calmodulin, putative 60S ribosomal protein L13-2, putative 40S ribosomal protein S16, putative conserved Plasmodium protein, unknown function ubiquitin conjugating enzyme, putative karyopherin alpha (KARalpha) | chromatin trafficking chaperone unknown function unknown function kinase ribosome ribosome unknown function proteasome nuclear pore | | | 0 na | ATP binding, protein binding heat shock protein binding, unfolded protein binding null null null structural constituent of ribosome structural constituent of ribosome null small conjugating protein ligase activity binding, protein transporter activity | chromosome organization protein folding protein catabolic process null null translation translation null post-translational protein modification, regulation of protein metabolic process protein import into nucleus | chromosome null null null null intracellular, ribosome intracellular, ribosome null null cytoplasm, nuclear pore, nucleus |
| PBANKA_141690 PBANKA_141730 PBANKA_141840 PBANKA_142050 PBANKA_142070 PBANKA_142100 PBANKA_142350 PBANKA_142360 PBANKA_142430 PBANKA_142420 PBANKA_142490 PBANKA_142510 PBANKA_142620 | chromosome segregation protein, putative translocation protein sec63, putative ATP-dependent Clp protease adaptor protein, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function calmodulin, putative 60S ribosomal protein L13-2, putative 40S ribosomal protein S16, putative conserved Plasmodium protein, unknown function ubiquitin-conjugating enzyme, putative karyopherin alpha, putative (KARalpha) CCR4-associated factor 1, putative (CAF1) | PF3D7_1318400 PF3D7_1318800 PF3D7_1320100 PF3D7_0713900 PF3D7_0714100 PF3D7_0714400 PF3D7_0814000 PF3D7_0813900 PF3D7_0812600 PF3D7_0812600 PF3D7_0812400 PF3D7_0811300 | chromosome segregation protein, putative secretory complex protein 63 (SEC63) ATP-dependent Clp protease adaptor protein ClpS, putative conserved Plasmodium protein, unknown function conserved Plasmodium protein, unknown function calmodulin, putative 60S ribosomal protein L13-2, putative 40S ribosomal protein S16, putative conserved Plasmodium protein, unknown function ubiquitin conjugating enzyme, putative karyopherin alpha (KARalpha) CCR4-associated factor 1 (CAF1) | chromatin trafficking chaperone unknown function unknown function kinase ribosome ribosome unknown function proteasome nuclear pore mRNA | | | na n | ATP binding, protein binding heat shock protein binding, unfolded protein binding null null null structural constituent of ribosome structural constituent of ribosome null small conjugating protein ligase activity binding, protein transporter activity nucleic acid binding | chromosome organization protein folding protein catabolic process null null null translation translation null post-translational protein modification, regulation of protein metabolic process protein import into nucleus null | chromosome null null null null intracellular, ribosome intracellular, ribosome null null cytoplasm, nuclear pore, nucleus nucleus |

| PAMACL, 14-248 Pamachanin pretinin, publishe Pamachanin | 5544444 | 0-0 | | | DE0D7 4040700 | | | | | | | | | |
|--|-----------|---|-------|------|---------------|---|---------------------------------|---|---|------------|---|--|---|---------------------------------------|
| Part | _ | | | | PF3D7_1212700 | | | 0 | 0 | 0 na | | null | null | null |
| March Marc | PBANKA_14 | 890 bromodomain protein, putative | | | PF3D7_1212900 | bromodomain protein, putative | chromatin | 0 | 0 | 0 na | | null | null | nucleus |
| Ministry | PBANKA_14 | 940 dynein light chain 1, putative | | | PF3D7_1213600 | dynein light chain 1, putative | dynein/kinesin | 0 | 0 | 0 na | | microtubule motor activity | microtubule-based process | microtubule associated complex |
| Post | PBANKA_14 | 120 conserved Plasmodium protein, unknown function | | | PF3D7_1215500 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | cAMP-dependent protein kinase regulator activity | signal transduction | null |
| Pankle 14224 Pankle 14225 Pankle 14225 | PBANKA_14 | 140 conserved Plasmodium protein, unknown function | | | PF3D7_1215700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |
| Parameter Para | PBANKA_14 | signal recognition particle 19 kD protein, putative | | | PF3D7_1216300 | signal recognition particle SRP19 (SRP19) | post-translational modification | 0 | 0 | 0 na | | 7S RNA binding | SRP-dependent cotranslational protein targeting to membrane | e signal recognition particle |
| PRIVING 1982 PRIV | PBANKA_14 | male development gene 1 (MDV1) | | | PF3D7_1216500 | male development gene 1 (MDV1) | egress, sexual development | 0 | 0 | 0 differer | ent in fertilization, ookinete, oocyst and sporozoite | null | null | null |
| PARING 14500 145 | PBANKA_14 | perforin like protein 2 (PPLP2) | | | PF3D7_1216700 | perforin like protein 2 (PPLP2) | perforin | 0 | 0 | 0 differen | erent in fertilization, ookinete and oocyst | null | null | null |
| PAMPAL, 14-94 PAMPAL, 14-9 | PBANKA_14 | 300 conserved Plasmodium protein, unknown function | | | PF3D7_1217400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |
| PSMMA 1-13916 series of policy in pulsine PSOP17) PSMMA 1-13926 series of policy in pulsine production | PBANKA_14 | conserved Plasmodium protein, unknown function | | | PF3D7_1217700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |
| PRANK 14500 1450 | PBANKA_14 | 400 phosphate translocator, putative | | | PF3D7_1218400 | triose or hexose phosphate/phosphate translocator, putative | transporter | 0 | 0 | 0 na | | null | null | null |
| PRAVIAC, 143731 Somework Pleamodum protein, unknown function Fish 71,125000 Somework Pleamodum protein, unknown functi | PBANKA_14 | secreted ookinete protein, putative (PSOP17) | | | PF3D7_1218800 | conserved Plasmodium protein, unknown function (PSOP17) |) adhesin | 0 | 0 | 0 KO r | not successful | null | null | null |
| PRANK_148170 PRAN | PBANKA_14 | aminophospholipid-transporting P-ATPase, putative | | | PF3D7_1219600 | aminophospholipid-transporting P-ATPase (ATPase2) | transporter | 0 | 0 | 0 KO r | not successful | ATP binding, ATP-see activity, coupled to transmentarium reverent of lone, phosphoryletive mechanism, responsium ton binding, phospholytid-translocating ATP-see activity. | ATP biosynthetic process, phospholipid transport | integral to membrane, membrane |
| PBANKA_143870 Networkmain protein 1, putative (PH*1) 1.92 0.74 PF307_122190 conserved Plasmodium protein, unknown function 1.92 0.74 PF307_122290 conserved Plasmodium protein, unknown function 1.92 0.74 | PBANKA_14 | conserved Plasmodium protein, unknown function | | | PF3D7_1220000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |
| PBANK_14989 or desired Plasmodium protein, whoreon function 1.92 0.74 PSD7_122300 conserved Plasmodium protein, whoreon function unless on function of particular protein, whoreon function of particular protein, whoreon function or adoptasmin homolog precursor, putative or protein protein, putative protein, putative (PABP) and protein protein, | PBANKA_14 | ribosome biogenesis protein TSR3, putative (TSR3) | | | PF3D7_1220500 | ribosome biogenesis protein TSR3, putative (TSR3) | ribosome | 0 | 0 | 0 na | | null | null | null |
| PBANK_14370 Conserved Plasmodium protein, unknown function PF307_122100 Conserved Plasmodium protein, unknown function PF307_122200 PF | PBANKA_14 | heterochromatin protein 1, putative (HP1) | | | PF3D7_1220900 | heterochromatin protein 1 (HP1) | chromatin | 0 | 0 | 0 na | | chromatin binding | chromatin assembly or disassembly | chromatin, nucleus |
| PBANKA_143730 and protein prot | PBANKA_14 | conserved Plasmodium protein, unknown function | -1.92 | 0.74 | PF3D7_1221300 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |
| PBANKA_143700 CAMP-dependent protein, jutative (PKAr) | PBANKA_14 | 710 conserved Plasmodium protein, unknown function | | | PF3D7_1221900 | conserved Plasmodium membrane protein, unknown function | n unknown function | 0 | 0 | 0 na | | null | null | null |
| PBANK_143800 CAMP-dependent protein kinase regulatory subunit, putative (PKAP) FP57_1223100 CAMP-dependent protein kinase regulatory subunit, putative (PABP) regulation of protein protein kinase regulatory subunit, putative (PABP) read (AMP-dependent protein kinase regulator) activity regulation of protein phosphorylation cAMP-dependent protein kinase regulatory subunit, putative (PABP) read (AMP-dependent protein kinase regulator) activity regulation of protein phosphorylation cAMP-dependent protein kinase regulatory subunit, putative (PABP) read (AMP-dependent protein kinase subunit, putative (PABP) read (AMP-d | PBANKA_14 | endoplasmin homolog precursor, putative | | | PF3D7_1222300 | endoplasmin homolog precursor, putative | | 0 | 0 | 0 na | | ATP binding, unfolded protein binding | protein folding, response to stress | null |
| PBANKA_144392 plyadenylate-binding protein, putative (PABP) plyadenylate-binding protein, putative (PABP) protein, putati | PBANKA_14 | 740 ATP-binding protein, putative | | | PF3D7_1222500 | ATP-binding protein, putative | | 0 | 0 | 0 na | | null | null | null |
| PBANKA_144100 splicing factor 3b subunit, putative (SF3B14) splicing factor 3b subunit, putative (SF3B14) mRNA 0 0 0 0 0 na nucleic acid binding null nucleic acid binding | PBANKA_14 | 800 CAMP-dependent protein kinase regulatory subunit, putative (PKAr) | | | PF3D7_1223100 | cAMP-dependent protein kinase regulatory subunit (PKAr) |) kinase | 0 | 0 | 0 na | | cAMP-dependent protein kinase regulator activity | regulation of protein phosphorylation | cAMP-dependent protein kinase complex |
| PBANKA_144110 hydrolase, putative conserved Plasmodium protein, unknown function pBANKA_144180 conserved Plasmodium protein, unknown function pBANKA_144300 NIMA related kinase 1, putative (NEK1) PF3D7_122800 NIMA related kinase 1 (NEK1) PF3D7_122800 NIMA related kinase 1 (NEK1) NIMA related kinase 1, putative (NEK1) PF3D7_123070 protein transport protein Sec13, putative transport protein served Plasmodium protein, unknown function in unknown function pBANKA_144570 conserved Plasmodium protein, unknown function in PBANKA_144590 conserved Plasmodium protein, unknown function in | PBANKA_14 | polyadenylate-binding protein, putative (PABP) | | | PF3D7_1224300 | polyadenylate-binding protein, putative (PABP) | mRNA | 0 | 0 | 0 na | | RNA binding | null | null |
| PBANKA_144180 conserved Plasmodium protein, unknown function | PBANKA_14 | 980 splicing factor 3b subunit, putative (SF3B14) | | | PF3D7_1224900 | splicing factor 3b subunit, putative (SF3B14) | mRNA | 0 | 0 | 0 na | | nucleic acid binding | null | null |
| PBANKA_144300 NIMA related kinase 1, putative (NEK1) | PBANKA_14 | 110 hydrolase, putative | | | PF3D7_1226300 | cof-like hydrolase, had-superfamily, subfamily iib | hydrolase | 0 | 0 | 0 na | | hydrolase activity | metabolic process | null |
| PBANKA_144540 transport protein Sec13, putative transport protein Sec13, putative transport protein Sec13, putative trafficking 0 0 0 0 0 na null null null null null null null n | PBANKA_14 | 180 conserved Plasmodium protein, unknown function | | | PF3D7_1227000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |
| PBANKA_144570 conserved Plasmodium protein, unknown function -1.74 0.7 PF3D7_1231000 conserved Plasmodium protein, unknown function 0 0 0 na null PBANKA_144590 conserved Plasmodium protein, unknown function 0 0 0 na null PBANKA_144900 conserved Plasmodium protein, unknown function 0 0 0 na null PBANKA_144900 conserved Plasmodium protein, unknown function 0 0 0 na null PBANKA_144900 conserved Plasmodium protein, unknown function 0 0 0 na null PBANKA_144900 conserved Plasmodium protein, unknown function 0 0 0 na null PBANKA_144900 conserved Plasmodium protein, unknown function 0 0 0 na null PBANKA_144900 conserved Plasmodium protein, unknown function 0 0 0 na null PBANKA_144900 conserved Plasmodium protein, unknown function 0 0 0 na null PBANKA_144900 conserved Plasmodium protein, unknown function 0 0 0 na null PBANKA_144900 conserved Plasmodium protein, unknown function 0 0 0 na null | PBANKA_14 | NIMA related kinase 1, putative (NEK1) | | | PF3D7_1228300 | NIMA related kinase 1 (NEK1) | kinase | 0 | 0 | 0 KO r | not successful | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| PBANKA_144590 conserved Plasmodium protein, unknown function PF3D7_1231200 conserved Plasmodium protein, unknown function unk | PBANKA_14 | 540 transport protein Sec13, putative | | | PF3D7_1230700 | protein transport protein Sec13, putative | trafficking | 0 | 0 | 0 na | | null | null | null |
| PBANKA_144900 conserved Plasmodium protein, unknown function | PBANKA_14 | 570 conserved Plasmodium protein, unknown function | -1.74 | 0.7 | PF3D7_1231000 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |
| | PBANKA_14 | 590 conserved Plasmodium protein, unknown function | | | PF3D7_1231200 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |
| PBANKA_144950 conserved Plasmodium protein, unknown function PF3D7_1234900 conserved Plasmodium protein, unknown function unk | PBANKA_14 | 900 conserved Plasmodium protein, unknown function | | | PF3D7_1234400 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 KO r | not successful | aspartic-type endopeptidase activity | proteolysis | null |
| | PBANKA_14 | 950 conserved Plasmodium protein, unknown function | | | PF3D7_1234900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 na | | null | null | null |

| PBANKA_145020 | serine hydroxymethyltransferase, putative (SHMT) | | | PF3D7_1235600 | serine hydroxymethyltransferase (SHMT) | pyrimidine metabolism | 0 | 0 | 0 | na | glycine hydroxymethyltransferase activity | L-serine metabolic process, glycine metabolic process | null |
|---------------|---|------|------|---------------|---|--------------------------|---|---|---|------------------------------|---|--|----------------------------------|
| PBANKA_145070 | clustered-asparagine-rich protein, putative | | | PF3D7_1236100 | clustered-asparagine-rich protein | | 0 | 0 | 0 | na | nucleic acid binding, protein binding | null | null |
| PBANKA_145150 | conserved Plasmodium protein, unknown function | | | PF3D7_1236900 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_145240 | ubiquitin-like modifier hub1, putative (HUB1) | | | PF3D7_1237800 | ubiquitin-like protein, putative | proteasome | 0 | 0 | 0 | na | null | null | null |
| PBANKA_145250 | conserved Plasmodium protein, unknown function | | | PF3D7_1237900 | conserved Plasmodium protein, unknown function | RNA | 0 | 0 | 0 | na | null | null | null |
| PBANKA_145270 | conserved Plasmodium protein, unknown function | | | PF3D7_1238200 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_145340 | protein kinase 2 (PK2) | -2.9 | 0.75 | PF3D7_1238900 | protein kinase 2 (PK2) | kinase | 0 | 0 | 0 | KO not successful | ATP binding, protein serine/threonine kinase activity, protein tyrosine kinase activity | protein phosphorylation | null |
| PBANKA_145480 | RNA binding protein, putative | | | PF3D7_1241400 | RNA binding protein, putative | mRNA | 0 | 0 | 0 | na | nucleic acid binding | null | null |
| PBANKA_145510 | replication factor C subunit 4, putative | | | PF3D7_1241700 | replication factor C subunit 4, putative | DNA replication | 0 | 0 | 0 | na | ATP binding, DNA clamp loader activity | DNA replication | DNA replication factor C complex |
| PBANKA_145570 | conserved Plasmodium protein, unknown function | | | PF3D7_1242300 | conserved Plasmodium membrane protein, unknown function | n unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_145680 | endosome sorting protein (SNF7 homologue), putative | | | PF3D7_1243500 | endosome sorting protein (SNF7 homologue), putative | e large tethering | 0 | 0 | 0 | na | null | protein transport | null |
| PBANKA_145690 | translation initiation factor SUI1, putative | | | PF3D7_1243600 | translation initiation factor SUI1, putative | translation | 0 | 0 | 0 | na | translation initiation factor activity | translation, translational initiation | null |
| PBANKA_145720 | double C2-like domain-containing protein, putative (DOC2) | | | PF3D7_1243900 | double C2-like domain-containing protein (DOC2) | trafficking, transporter | 0 | 0 | 0 | na | null | entry into host | null |
| PBANKA_145780 | ADP-ribosylation factor, putative | | | PF3D7_1244600 | ADP-ribosylation factor GTPase-activating protein, putative | e trafficking | 0 | 0 | 0 | na | ARF GTPase activator activity, zinc ion binding | regulation of ARF GTPase activity | null |
| PBANKA_145790 | conserved Plasmodium protein, unknown function | | | PF3D7_1244700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_145850 | nedd8-conjugating enzyme ubc12, putative (UBC12) | | | PF3D7_1245300 | ubiquitin conjugating enzyme E2, putative | proteasome | 0 | 0 | 0 | na | ATP binding, small conjugating protein ligase activity | post-translational protein modification, regulation of protein metabolic process | null |
| PBANKA_145930 | actin I | | | PF3D7_1246200 | actin I (ACT1) | gliding motility | 0 | 0 | 0 | na | protein binding | null | cytoplasm, nucleus |
| PBANKA_146060 | conserved Plasmodium protein, unknown function | | | PF3D7_1247700 | conserved Plasmodium protein, unknown function | unknown function | 0 | 0 | 0 | na | uroporphyrinogen-III synthase activity | tetrapyrrole biosynthetic process | null |
| PBANKA_146120 | conserved Plasmodium protein, unknown function | | | PF3D7_1248300 | conserved Plasmodium membrane protein, unknown function | n unknown function | 0 | 0 | 0 | na | null | null | null |
| PBANKA_146140 | conserved Plasmodium protein, unknown function | | | PF3D7_1248500 | conserved protein, unknown function | apoptosis | 0 | 0 | 0 | na | null | null | null |
| PBANKA_146180 | tat-binding protein homolog, putative | | | PF3D7_1248900 | tat-binding protein homolog | proteasome | 0 | 0 | 0 | na | ATP binding, nucleoside-triphosphatase activity | protein catabolic process | cytoplasm, nucleus |
| PBANKA_146300 | osmiophilic body protein (G377) | | | PF3D7_1250100 | osmiophilic body protein (Pfg377) | adhesin | 0 | 0 | 0 | not different from wild type | null | null | osmiophilic body |