

PlasmoDB ID	Candidate	3D Structure
PF3D7_1202900	High mobility group protein B1 (HMGB1)	1CG7 (Predicted from <i>S. cerevisiae</i>)
PF3D7_1243600	Translation initiation factor SUI1, putative	2IF1 (Predicted from <i>H. sapiens</i>)
PF3D7_0818200	14-3-3 protein (14-3-3I)	1IB1 (Predicted from <i>O. aries</i>)
PF3D7_1444800	Fructose-biphosphate aldolase (FBPA)	1A5C*
PF3D7_1365900	Tetraubiquitin	1F9J (Predicted from <i>H. sapiens</i>)

S1 Table. Potential biomarkers for *Plasmodium falciparum* infection as proposed by the algorithm in Fig 1.