

Supplement A

New paradigms for understanding and step changes in treating active and chronic, persistent apicomplexan infections

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Fibroblasts , MonoMacs, Neuronal stem cells

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Supplement "

Box. Co-crystallization data collection and refinement statistics

Cytochrome <i>bc1</i> –MJM170	
Data collection	
Space group	P65
Cell dimensions	
a, b, c (Å)	129.5, 129.5, 720.3
a, b, g (°)	90, 90, 120
Resolution (Å)	50-3.5(3.56-3.50)
Rmerge (%)	15.6(82)
I/sl	5.2(1.4)
Completeness (%)	99.6(99.7)
CC1/2	0.971(0.331)
Redundancy	9.2(9.2)
Refinement	
Resolution (Å)	50 – 3.5
No. reflections	80565
Rwork/ Rfree	24.13(29.13)
No. atoms	31811
Protein	30980
Ligand/ion	831
B-factors	
Protein	93.84
Ligand/ion	87.86
R.m.s deviations	
Bond lengths (Å)	0.010
Bond angles (°)	1.564
PDB access code	5HJ4

Supplemental Figures

Figure S1

Heat Maps for HFF infected with EGS in culture for 2, 18, and 48 hours for host and parasite cells

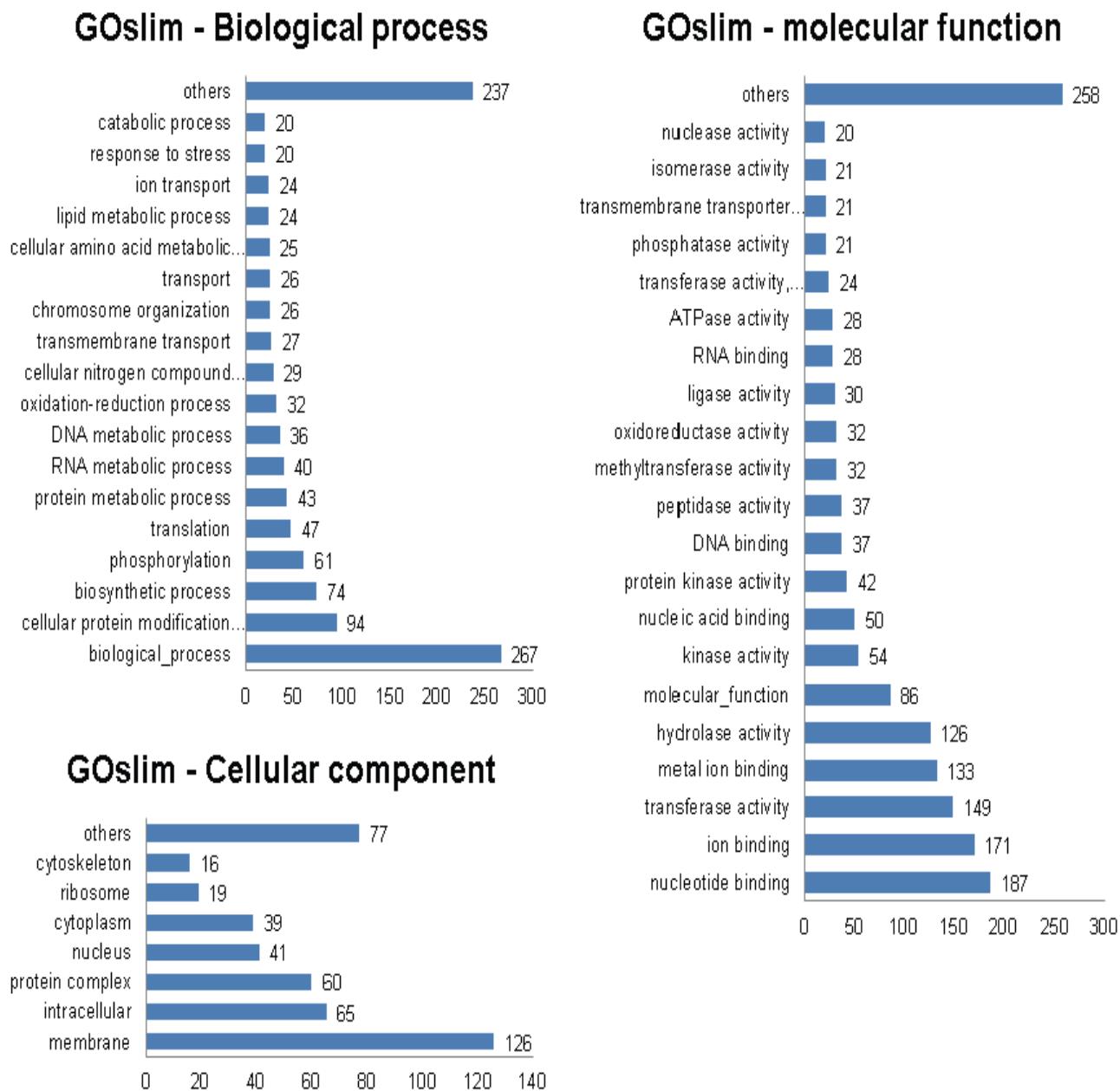


Figure S2

Log2 transformed normalized read count distributions were used to determine transcripts with low read count. The bars represent number of transcripts with different read count. Transcripts with read count below the 25 percentile (dark gray arrow) on the overall read count distribution were removed for further analysis. Yellow and green colors were used for presenting MM6 and NSC data.

