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Supplemental Information

Profiling the Targets of Protective

CD8⁺ T Cell Responses to Infection

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Supplemental Information:

Table S1. Vector yields on various size plates

Plate Size	Adg. <i>Py</i> HEP17		Adg.CMVp65	
	ffu/ml	ffu total	ffu/ml	ffu total
60 mm	3.10E+08	1.30E+09	8.20E+08	3.44E+09
6 well	4.55E+08	9.55E+08	6.70E+08	1.41E+09
12 well	9.00E+08	7.56E+08	1.10E+09	9.24E+08
24 well	1.33E+09	5.59E+08	1.02E+09	4.28E+08
48 well	1.38E+09	5.80E+07	1.27E+09	5.30E+07
96 well	1.26E+09	2.26E+07	1.19E+09	2.10E+07

Table S2. Thirteen antigens that recalled CD8⁺ T cell responses more frequent than CSP from mice immunized with *P. yoelii* SPZ+CQ

Rank	Gene ID <i>P. yoelii</i>	Gene ID <i>P. falciparum</i>	% Identity ^a	<i>P. yoelii</i> Protein Name - Function	Liver stage active genes from Tarum et al., 2008	
					Transcriptome Clusters ^h	Proteome
1	PY02605	PF3D7_0932900	40	Hypothetical Protein	C3	-
2	PY02989	PF3D7_0212800	47	Hypothetical Protein ^b	C4	-
3	PY02326	PF3D7_1309500	74	snornp protein gar1 ^c	C3	+
4	PY02030	PF3D7_0624000	86	hexokinase		+
5	PY00629	PF3D7_0108300	33	asparagine-rich protein, putative		+
6	PY07062	PF3D7_1446400	82	pyruvate dehydrogenase E1 beta subunit	C3	+
7	PY00361	PF3D7_1357100	99	translation elongation factor EF-1, subunit alpha		+
8	PY04421	PF3D7_1121600	56	U43539 hepatocyte erythrocyte protein 17 kDa ^d	C3	+
9	PY05048	PF3D7_1108400	95	Protein kinase domain ^e		+
10	PY07825	PF3D7_0814200	86	Arabidopsis thaliana At1g20220/T20H2_3- related ^f	C3	+
11	PY05748	PF3D7_0930300	32	merozoite surface protein 1 precursor	C1	-
12	PY04475	PF3D7_0621200	86	ethylene-inducible protein hever ^g		+
13	PY02148	PF3D7_1334600	31	merozoite surface protein 7 precursor	C1	-

^a Percent identity between *P. falciparum* and *P. yoelii* orthologues established using blastpp suite-2sequences, National Center for Biotechnology Information.

^b *P. falciparum* protein function is multidrug efflux pump, putative.

^c *P. falciparum* protein function is H/ACA ribonucleoprotein complex subunit 1, putative.

^d *P. falciparum* protein function is exported protein 1.

^e *P. falciparum* protein function is casein kinase 2, alpha subunit.

^f *P. falciparum* protein function is DNA/RNA-binding protein Alba 1.

^g *P. falciparum* protein function pyridoxine biosynthesis protein PDX1.

^h Clusters from Tarum *et al.*, 2008 - C1 includes LS genes mainly up-regulated in LS24 compared with LS40 and LS50, but that were also expressed in the mosquito and erythrocytic stages. C2 encompassed genes that were mainly up-regulated at LS24; C3 showed most of the genes up-regulated in all three LS time points queried. C4 showed genes mainly up-regulated in LS40 and LS50.