

PlasmoDB ID	Annotation	Time point ^a	Fold change to relative t ₀	
			Oligo array (±SD) ^b	Semi-qPCR (±SD) ^c
PFL1885c	Calcium/calmodulin-dependent protein kinase 2, putative	t ₃	2.43 (±0.18)	1.74 (±0.40)
PFD0285c ^d	Lysine decarboxylase	t ₃	2.78	1.71 (±0.19)
PFE0435w	Ornithine aminotransferase	t ₃	1.88	2.14 (±0.30)
PF08_0131	1-Cys-peroxidoxin	t ₃	0.35	0.42 (±0.14)
PFD0830w ^d	Dihydrofolate reductase/thymidylate synthase (DHFR/TS)	t ₁	0.50 (±0.05)	Not detected ^e
		t ₃	0.62 (±0.03)	0.35 (±0.04)
PFI1090w	S-adenosylmethionine synthase	t ₃	0.42	0.53 (±0.07)

a. Treated sample of maximum increase/decrease in transcript abundance

b. Data of transcripts represented by multiple oligonucleotides ($p < 0.05$) were averaged with the standard deviation indicated in parentheses.

c. Real-time PCR was performed in triplicate with the standard deviation indicated in parentheses.

d. Relatively low abundance transcripts

e. Not detected in sample of maximum decrease in abundance (t₁) and repeated for t₃

