

Table 1 Genes used for hierarchical clustering of mRNAs differently abundant depending on time p.i..

<i>Data description</i>	<i>E. falciformis</i> genes	Mouse genes
Sum of 1st infection NMRI sample differences (including oocysts and sporozoites if appl.)	4	8052
Used in hierarchical clustering (heatmap)	1618	1313