

Table S2. RNA-seq raw reads and alignment statistics		
	Total quality reads	Mapped reads to Spongospora
G1	74,130,035	4,299,542
G2	59,956,347	2,326,306
G3	71,378,260	6,509,697
<b>Average Gladiator</b>	<b>68,488,214</b>	<b>4,378,515</b>
I1	60,499,120	3,623,897
I2	62,383,322	4,123,537
I3	71,612,245	4,253,767
<b>Average Iwa</b>	<b>64,831,562</b>	<b>4,000,400</b>