

**Table S4: Summary of sequence reads with matches on *S. miltiorrhiza* isotigs.**

<b>Sample</b>	<b>Mapped reads</b>	<b>Total reads</b>
0 h	5,013,001 (72.84%)	6,882,388
12 h	4,621,801 (73.36%)	6,300,372
24 h	4,321,814 (75.40%)	5,731,519
36 h	4,140,869 (72.77%)	5,690,024