

**Table S5: Summary of *S. miltiorrhiza* isotigs with matched sequence reads.**

<b>Sample</b>	<b>Total isotigs</b>	<b>Mapped isotigs</b>
0 h		20,553 (79.68%)
12 h		20,195 (78.30%)
24 h	25,793	20,305 (78.72%)
36 h		20,367 (78.96%)
Total mapped cDNAs		22,464 (87.09%)