

Supplemental table S1: Raw and clean reads of 9 RNA samples of *Fraxinus mandshurica* stigma

<b>Sample</b>	<b>Total Raw Reads (M)</b>	<b>Total Clean Reads (M)</b>	<b>Clean Base(G)</b>	<b>Q20(%)</b>	<b>Q30(%)</b>	<b>GC Content(%)</b>
<b>S1-1</b>	80.49	77.05	11.56	98.84	95.22	43.51
<b>S1-2</b>	90.44	86.85	13.03	99.03	95.95	43.56
<b>S1-3</b>	82.23	78.87	11.83	98.91	95.48	43.47
<b>S2-1</b>	80.77	77.71	11.66	98.86	95.29	43.22
<b>S2-2</b>	88.62	82.25	12.34	98.88	95.38	43.36
<b>S2-3</b>	74.23	70.94	10.64	98.83	95.21	43.23
<b>S3-1</b>	80.57	77.02	11.55	98.91	95.5	43.4
<b>S3-2</b>	80.92	76.84	11.53	98.94	95.66	43.38
<b>S3-3</b>	78.26	74.24	11.14	98.81	95.1	43.44