

**Table S1. The library information and data statistics**

<b>Platform</b>	<b>Number of libraries</b>	<b>Size</b>	<b>Average Read Length (bp)</b>	<b>Total Bases (Gb)</b>	<b>Sequence Depth</b>	<b>Application</b>
<b>PacBio</b>	<b>RS II and Sequel</b>	<b>Library Mean Size</b>	<b>7,707</b>	<b>8.753</b>	<b>213 ×</b>	<b>Genome assembly</b>
	Sequel	20 Kb	7,717	1.396		
	RS II	20 Kb	7,705	7.357		