

**Sequence methods and library kits used in the hybrid assembly of snout otter clam (*L. rhynchaena*) genome**

	<b>Platform</b>	<b>Read length</b>	<b>Number of Reads</b>	<b>Total Bases</b>	<b>Estimated genome coverage</b>
Genomic	Illumina (NovaSeq) - PCR-free	2 × 150bp	367,711,652	55.16 Gbp	100×
	Illumina (NovaSeq) - PCR	2 × 150bp	445,111,088	66.77 Gbp	121×
	ONT (MinION)	variable	2,830,557	14.04 Gbp	25×
Transcriptomic	Illumina (NovaSeq)	2 × 150 bp	230,852,564	34.63 Gbp	N/A