

**S2 Table. Estimation of the Yap hadal snailfish genome size (Kmer = 17).**

<b>K-mer</b>	<b>K-mer number</b>	<b>K-mer Depth</b>	<b>Genome Size (Mb)</b>	<b>Heterozygous Ratio (%)</b>
17	39,380,990,928	47	815.59	0.61