

Single-molecule long-read sequencing of the full-length transcriptome of *Rhododendron lapponicum* L.

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Table S1 Summary of polymerase reads from PacBio single-molecule long read sequencing

Size	Post-filter polymerase reads	Post-filter total number of subread bases	Post-filter number of subread	Post-filter subreads N50	Post-filter mean subread length
1-2K	387,454	5.27G	2,861,654	1,782	1,659
2-3K	334,062	6.43G	3,573,056	2,531	2,440
3-6K	235,516	3.67G	1,825,478	3,406	3,211