

Supplementary Table S4 Summary of raw sequencing data quality for five *Engelhardia* species

Sample Name	Clean Reads	Clean bases	Read length (bp)	Q20(%)	GC(%)
<i>E. spicata</i> var. <i>spicata</i>	920,933,333	138,128,887,500	150	98.40	38.25
<i>E. spicata</i> var. <i>colebrookeana</i>	784,466,667	117,670,000,050	150	98.50	36.28
<i>E. spicata</i> var. <i>aceriflora</i>	897,000,000	134,550,000,000	150	98.50	36.32
<i>E. roxburghiana</i>	1,107,733,333	166,160,000,000	150	98.30	36.48
<i>E. fenzelii</i>	791,400,000	118,710,000,000	150	97.60	39.90