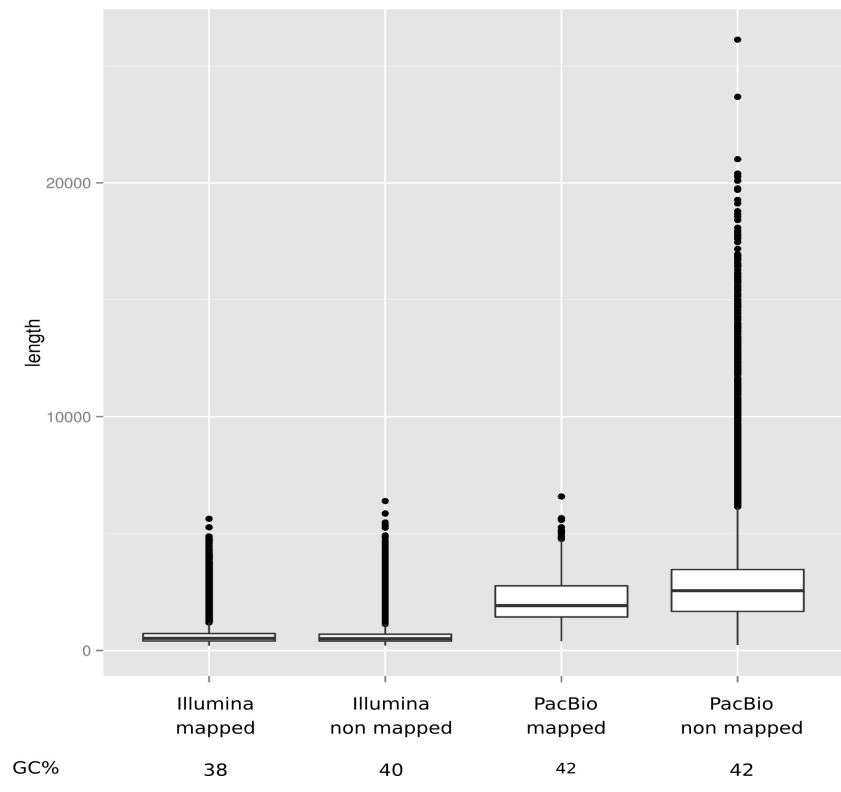


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**Figure S2.** Transcript length distribution of assembled transcripts where no CDS was identified. Data are shown for Illumina or PacBio libraries, and for transcripts that could be aligned (or not aligned) on the *P. lambertiana* genome. Average GC content of transcripts is shown in the bottom of the figure.