

Sequence length distribution

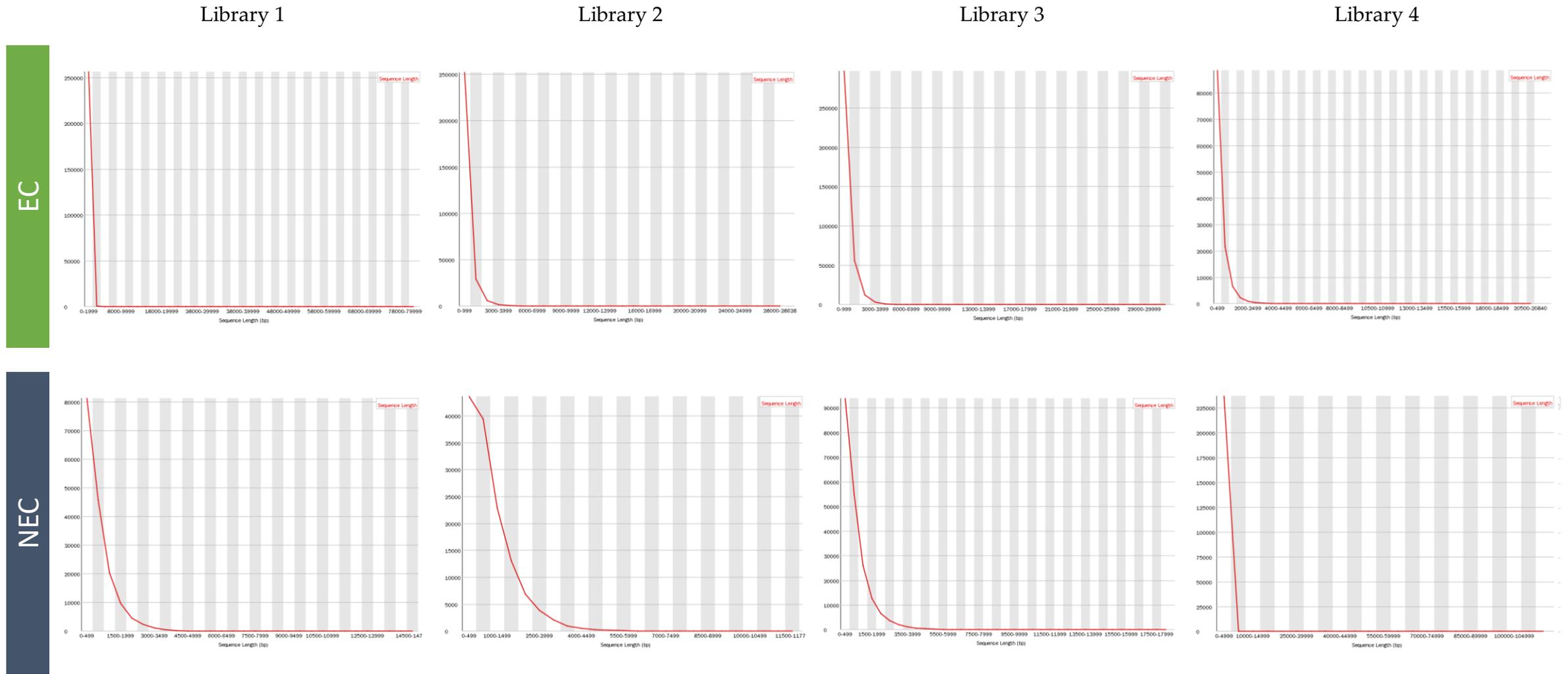


Figure S1. Distribution of sequence lengths, over all sequences of long-reads, generated from barcoded cDNA libraries, of *S. betaceum* compact embryogenic callus (EC) and friable non-embryogenic callus (NEC), by MinION sequencing from Oxford Nanopore Technologies®.

Duty time grouped

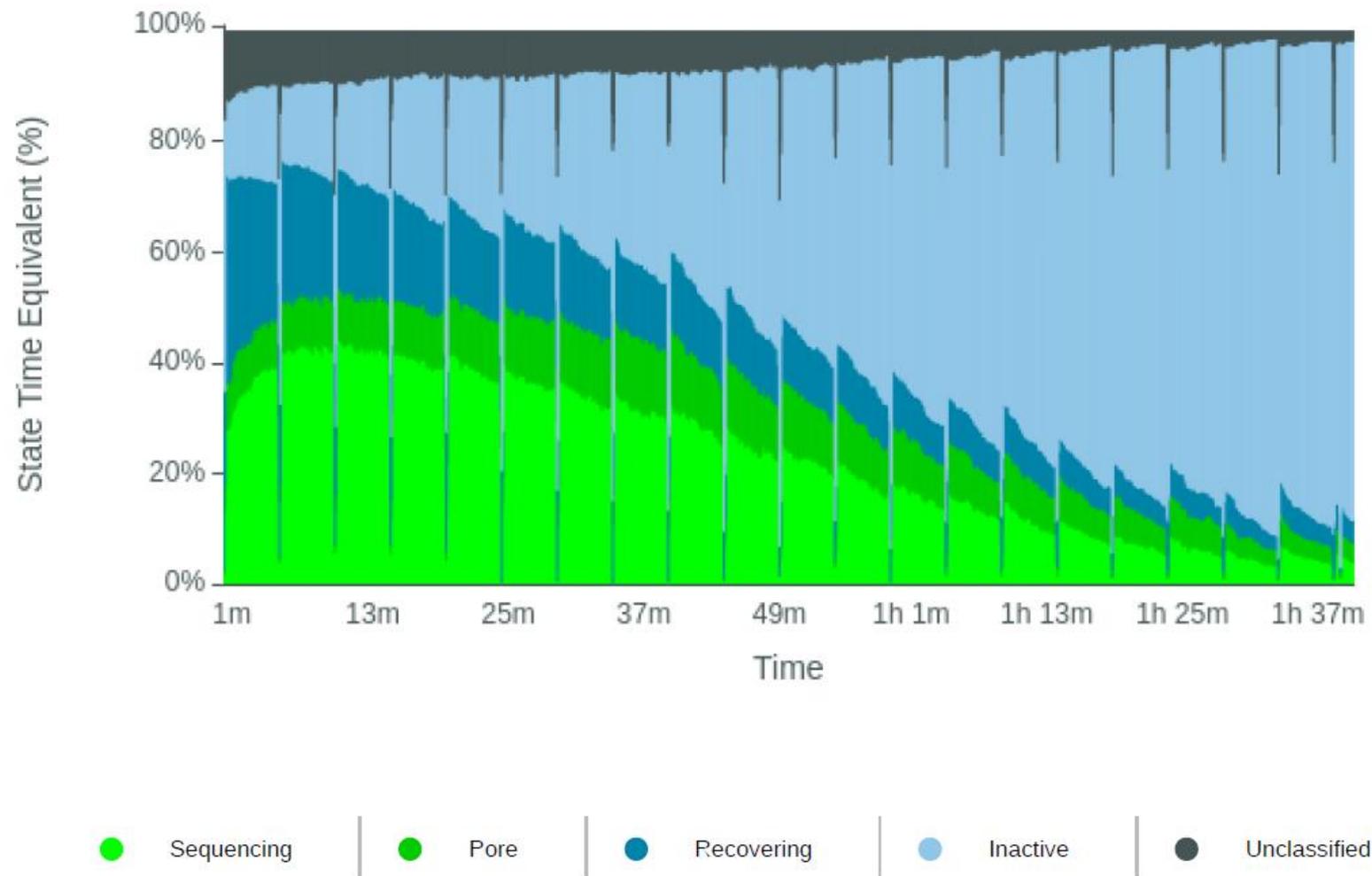


Figure S2. Duty time plot, obtained by MinKNOW to access the quality of run carried out by MinION sequencing from Oxford Nanopore Technologies®.