

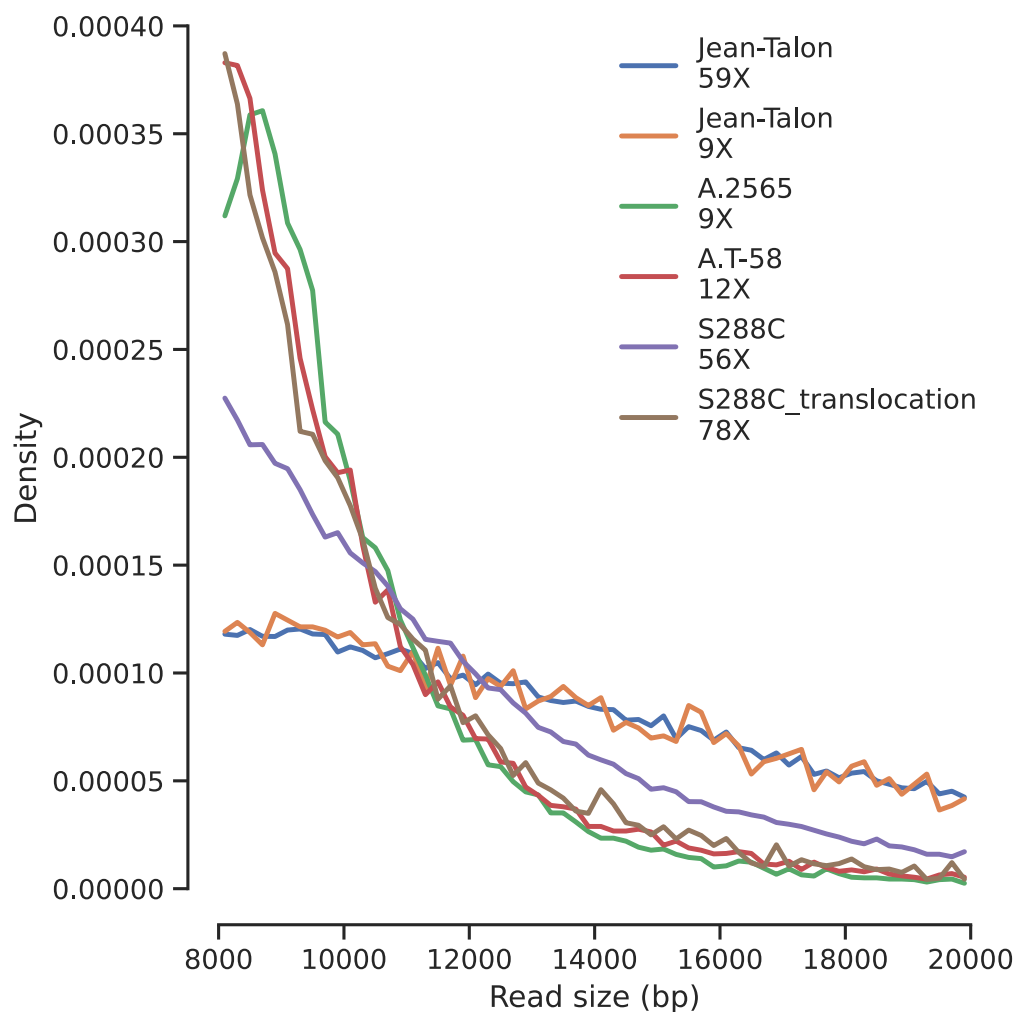


Figure S5. Read length distribution for the long-read datasets used in this study. Jean-Talon corresponds to the Oxford Nanopore dataset generated in this study; A.2565 and A.T-58 correspond to PacBio datasets from Fay *et al.* 2019; S288C corresponds to the PacBio dataset from Yue *et al.* 2017; S288c_translocation corresponds to the simulated PacBio dataset from the S288C assembly harboring translocations as shown in Figure S4.