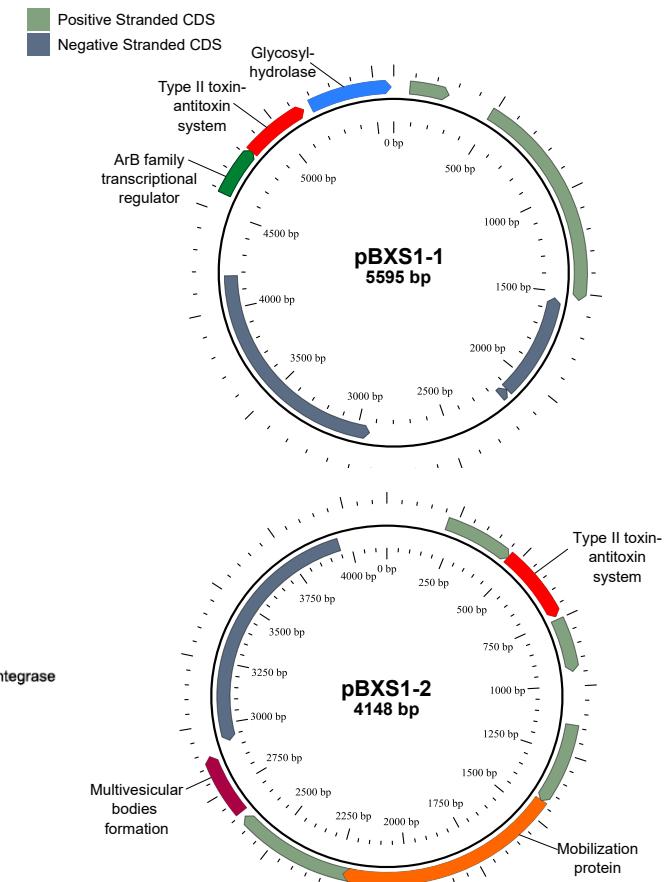
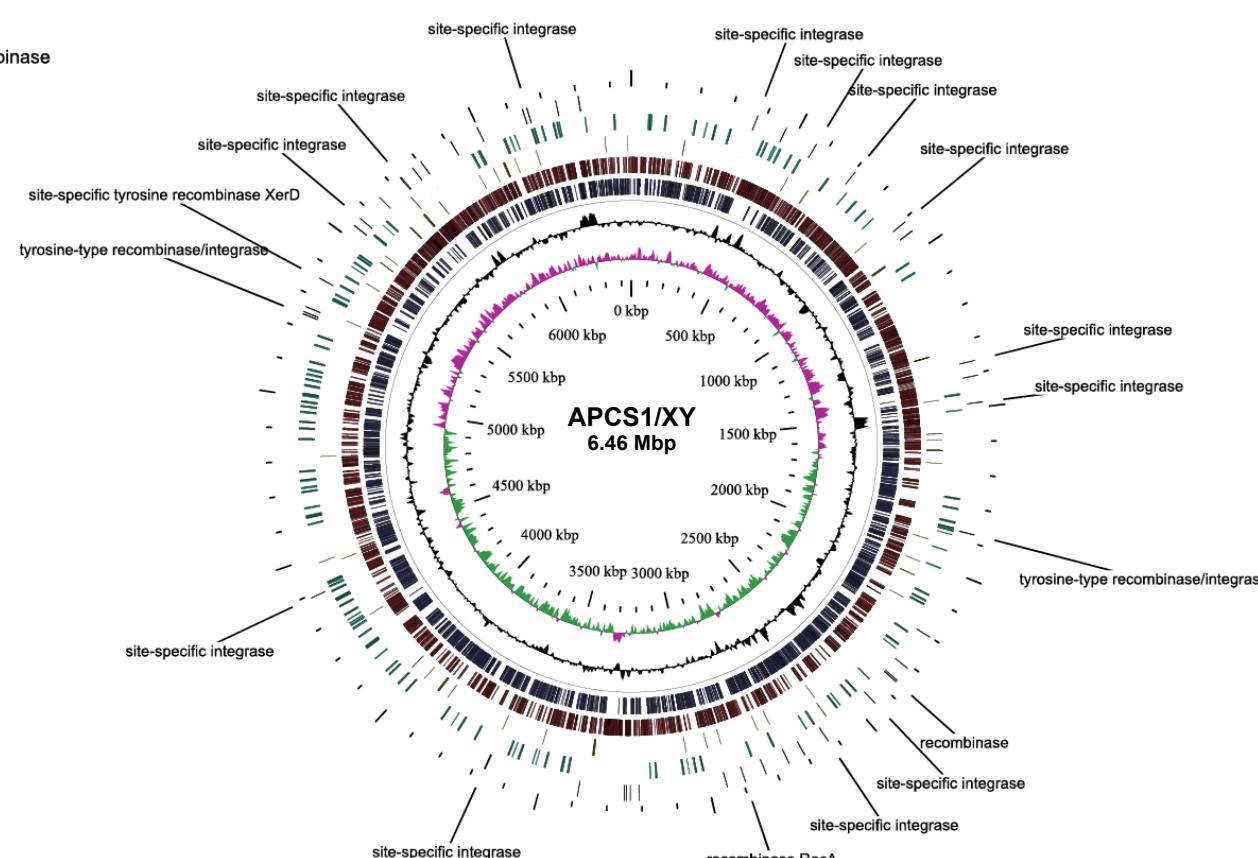
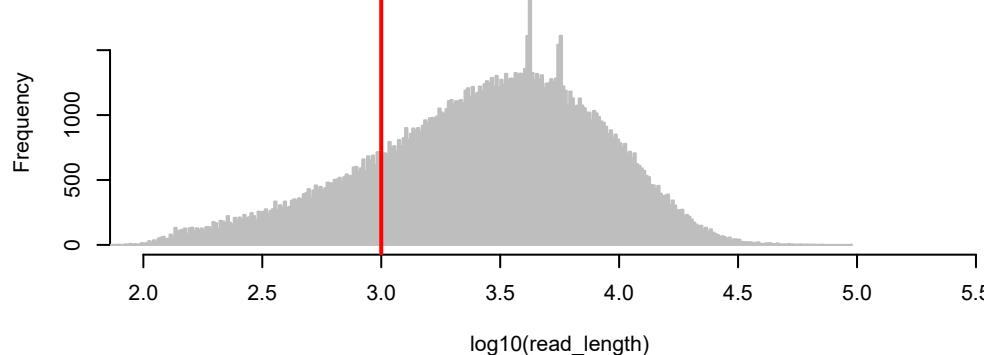
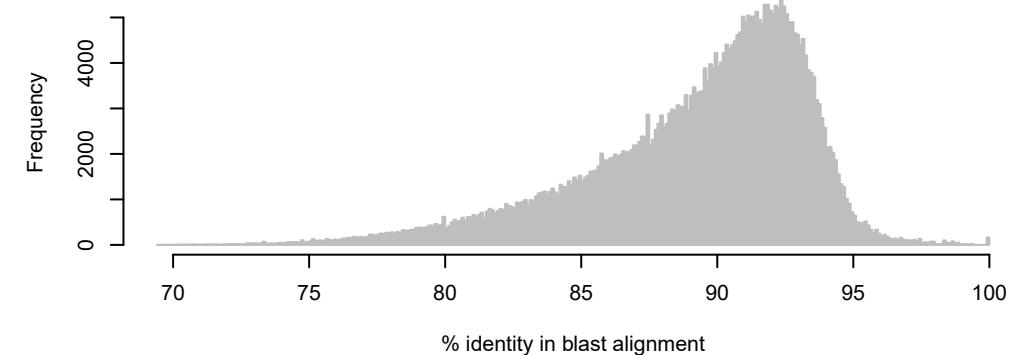


**a**

- █ Negative stranded CDS
- █ Positive stranded CDS
- █ tRNA and rRNA
- █ Sus gene family
- █ TonB dependent receptor
- █ Invertase/Integrase/Recombinase

**b****Read lengths distribution****c****BLAST alignment quality****d****Recombination hotpoints in individual ONT reads**