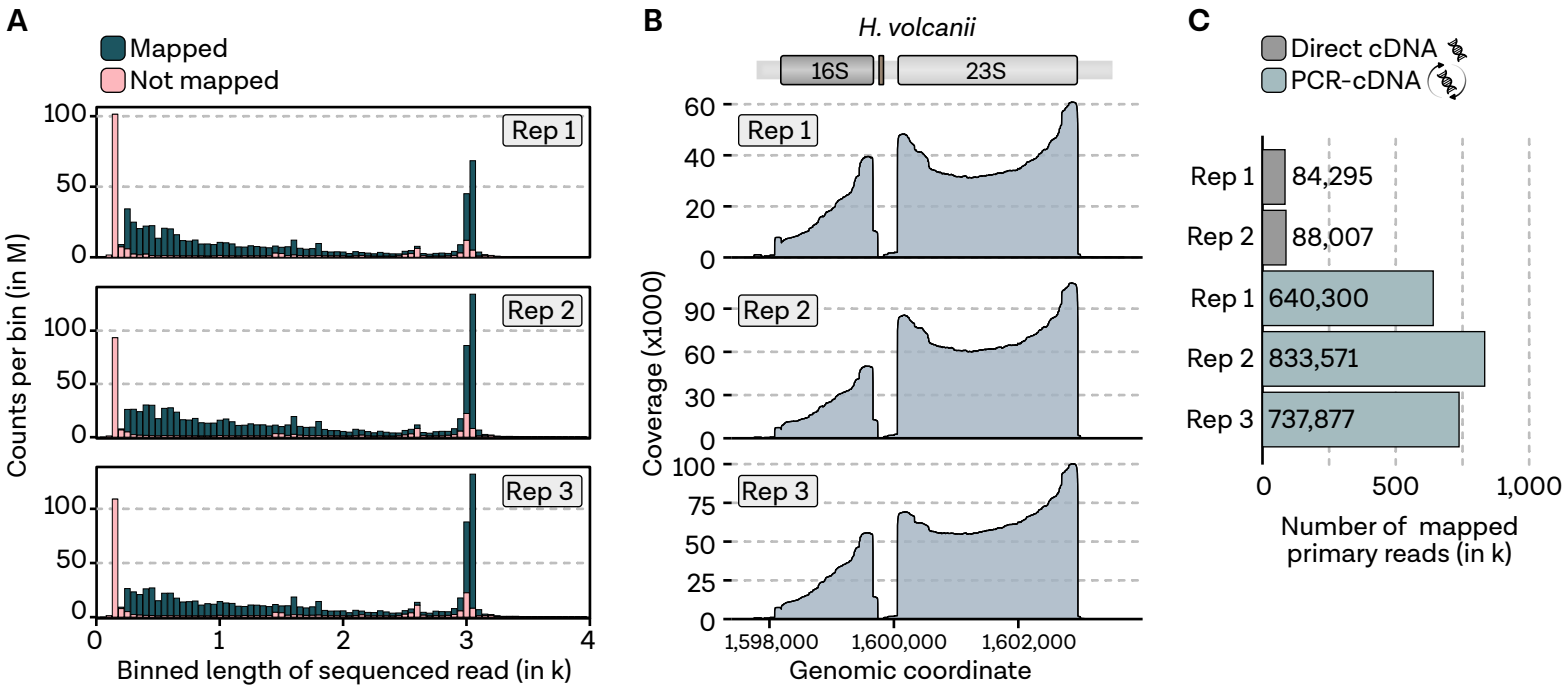




**Fig. S7** | Nanopore-based PCR-cDNA sequencing of *H. volcanii*.

**A**, Distribution of sequenced read length (bin size 50 nt), stacked by alignment group (blue: full-length mapped reads; pink: not mapped). **B**, Coverage profiles of pycloppe filtered full-length reads mapping to the rRNA region. **C**, Total number of primary mapped reads in direct cDNA (grey) and PCR-cDNA (blue) samples.