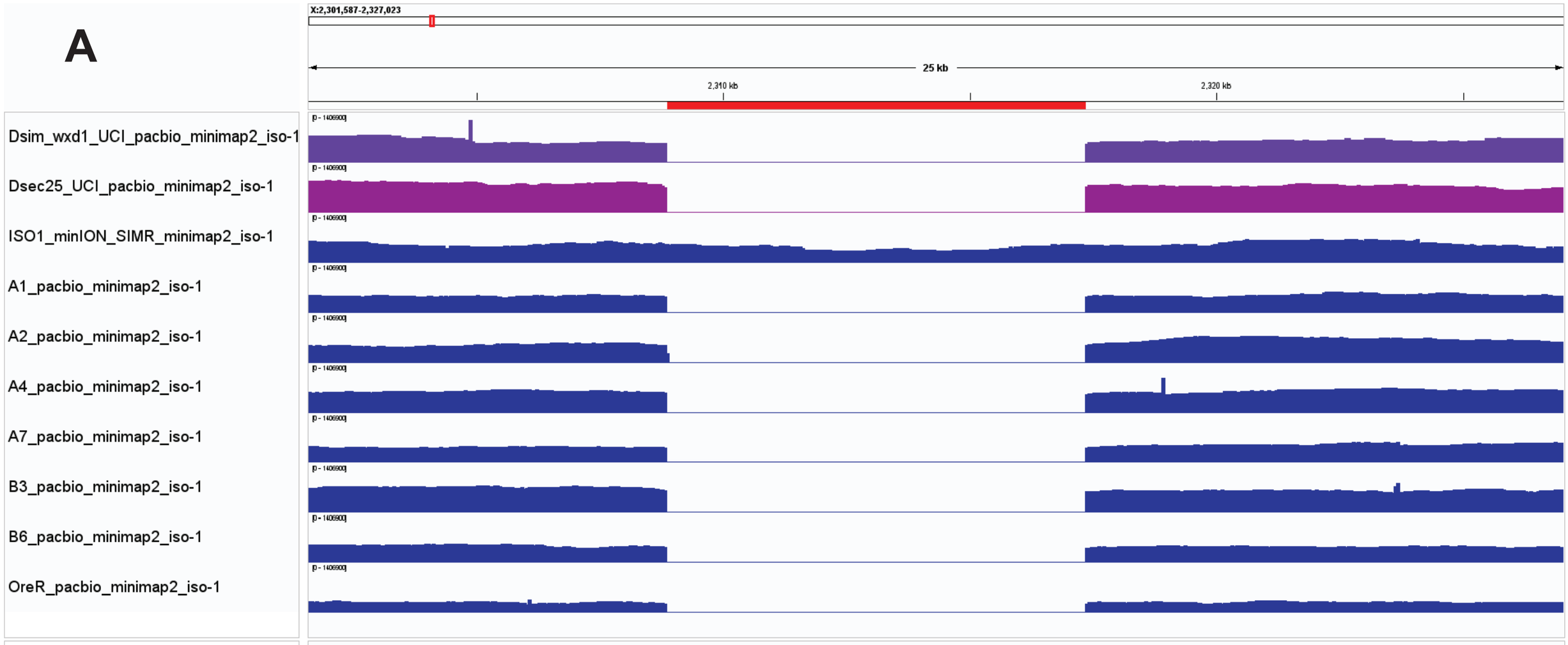

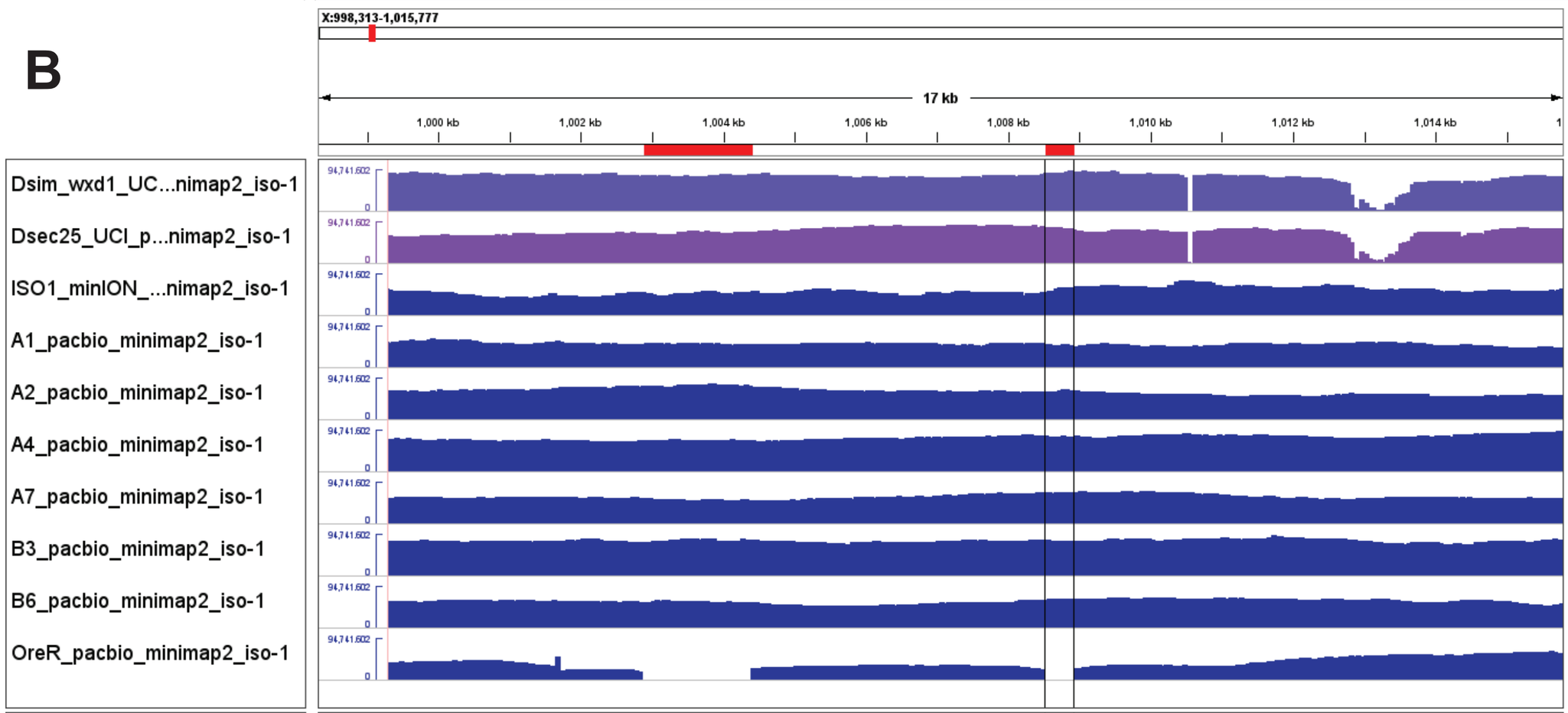

Supplementary Fig S10. IGV browsershots of INDELs supported by whole genome sequencing read coverage. *(A)* Absence of read coverage in ~8Kb region (highlighted in red) from all strains and *D. simulans* and *D. sechelia* except iso-1 aligned to iso-1 genome assembly supports a insertion specific to iso-1. *(B)* Absence of read coverage in 1.3kb and 0.4kb regions (highlighted in red) specifically in Oregon-R track supports two OreR-specific deletion events.