

Supplementary Material for

SMRT SEQUENCING OF *XANTHOMONAS* *ORYZAE* GENOMES REVEALS A DYNAMIC STRUCTURE AND COMPLEX TAL EFFECTOR GENE RELATIONSHIPS

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File S2. Traces of Sanger reads aligned to *tal2b* in the BLS256 reference genome assembly. In the first (top) and fourth trace the coordinate marker in the middle indicates the site of the frameshift call. The second and third traces appear to be corrupted and do not contain coordinates. The fourth trace is on the opposite strand from the rest.

