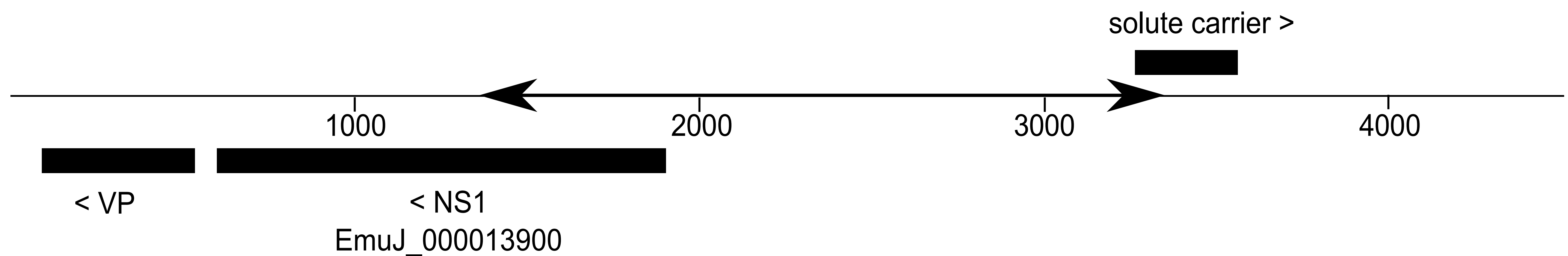
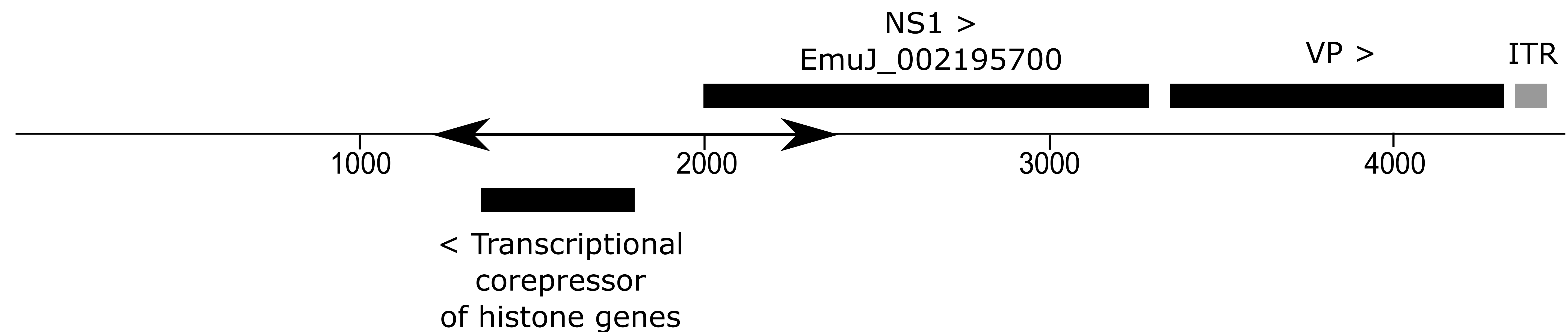


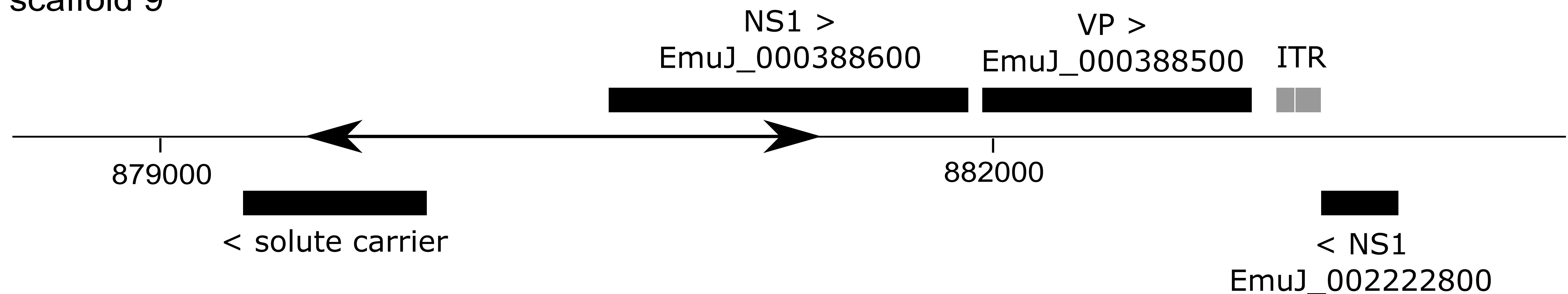
contig 0226



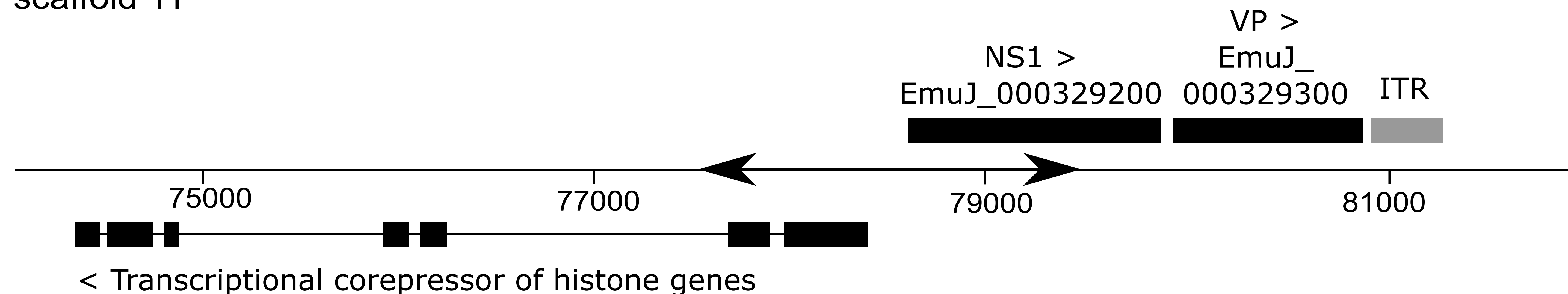
contig 0303



scaffold 9



scaffold 11



Additional file 4: Figure S3. Dengovirus integration sites in the *E. multilocularis* genome. Shown is a schematic representation of the complete contigs 0226 and 0303 and parts of scaffold 9 and 11. Exons and ITRs are represented by black and grey boxes, introns by lines. Arrowheads indicate gene orientation. Double headed arrows show sequenced genome regions. NS1: non-capsid protein 1; VP: capsid protein; ITR: inverted terminal repeat.