TE: DmeI−L−B516−Map1−TRIM/nan
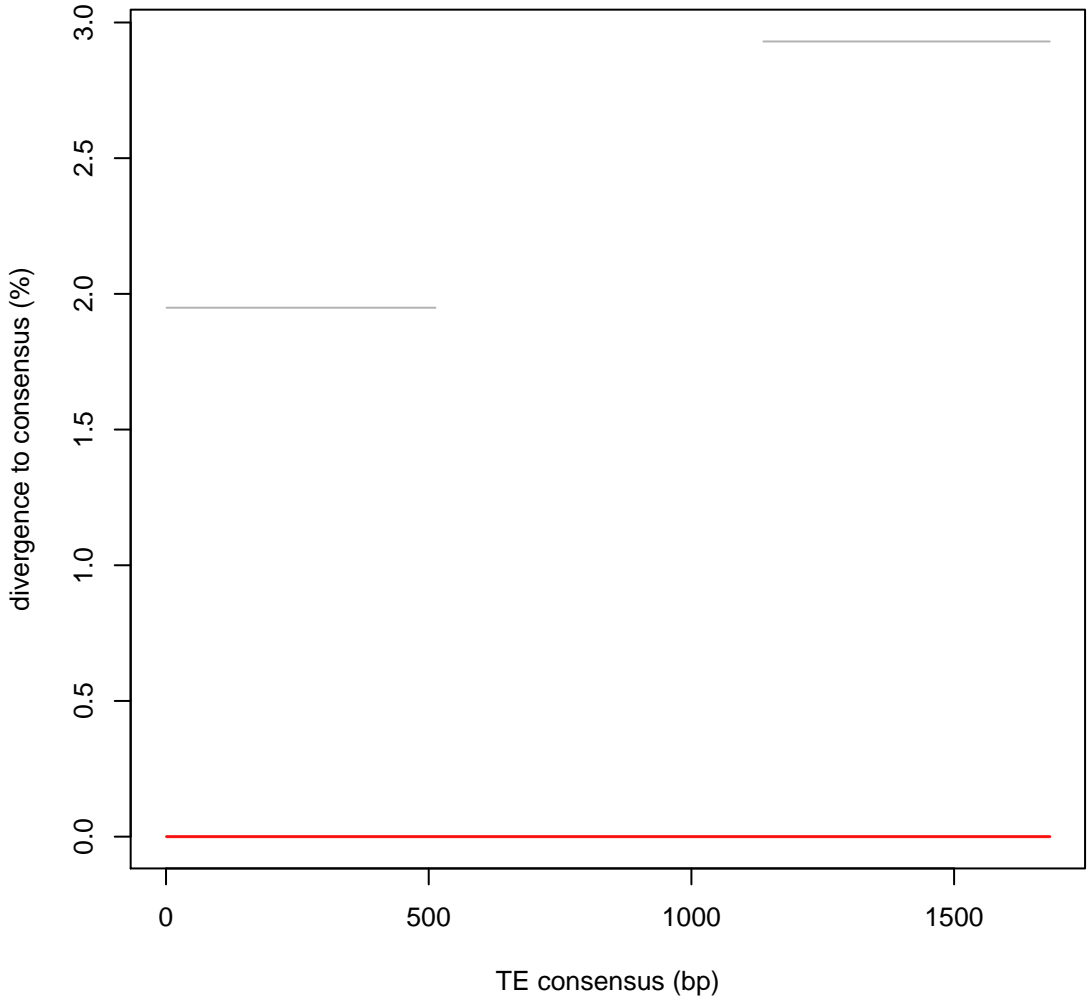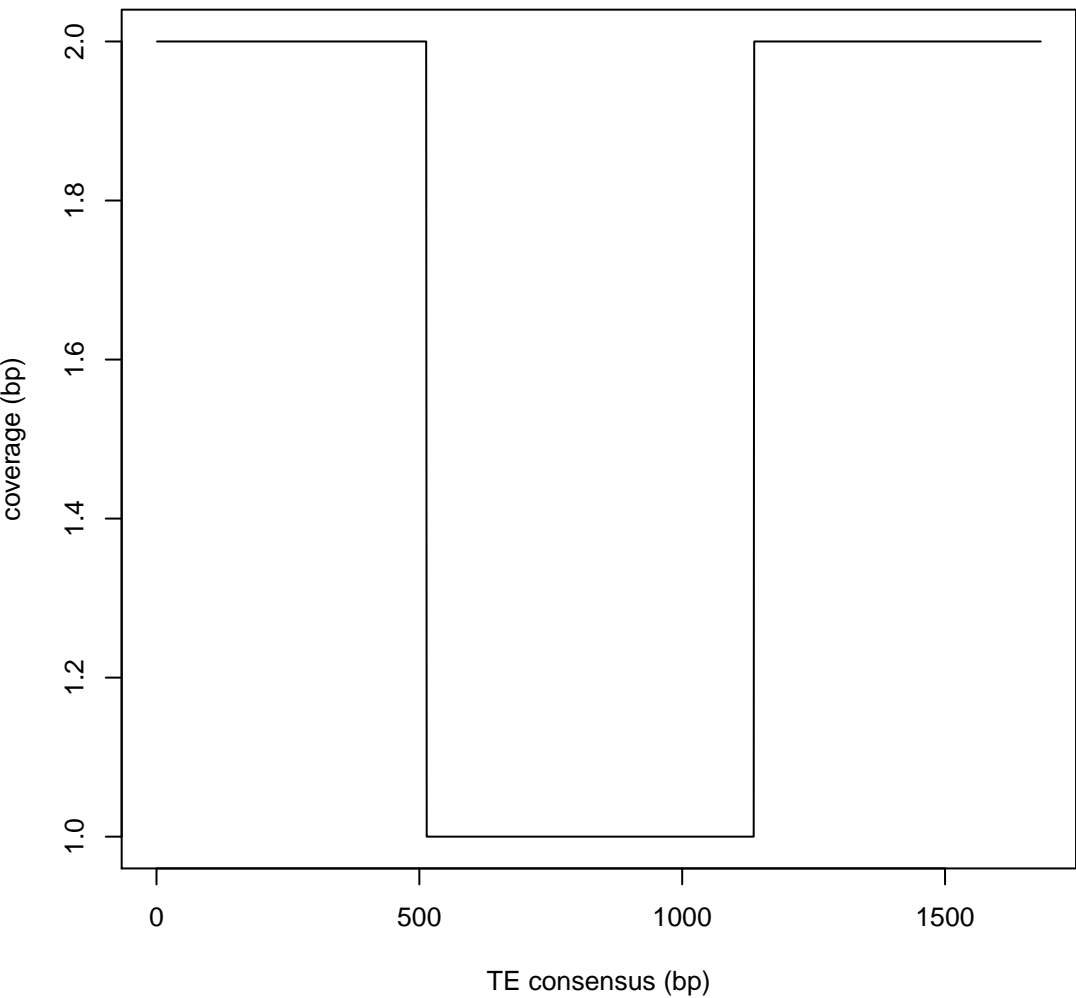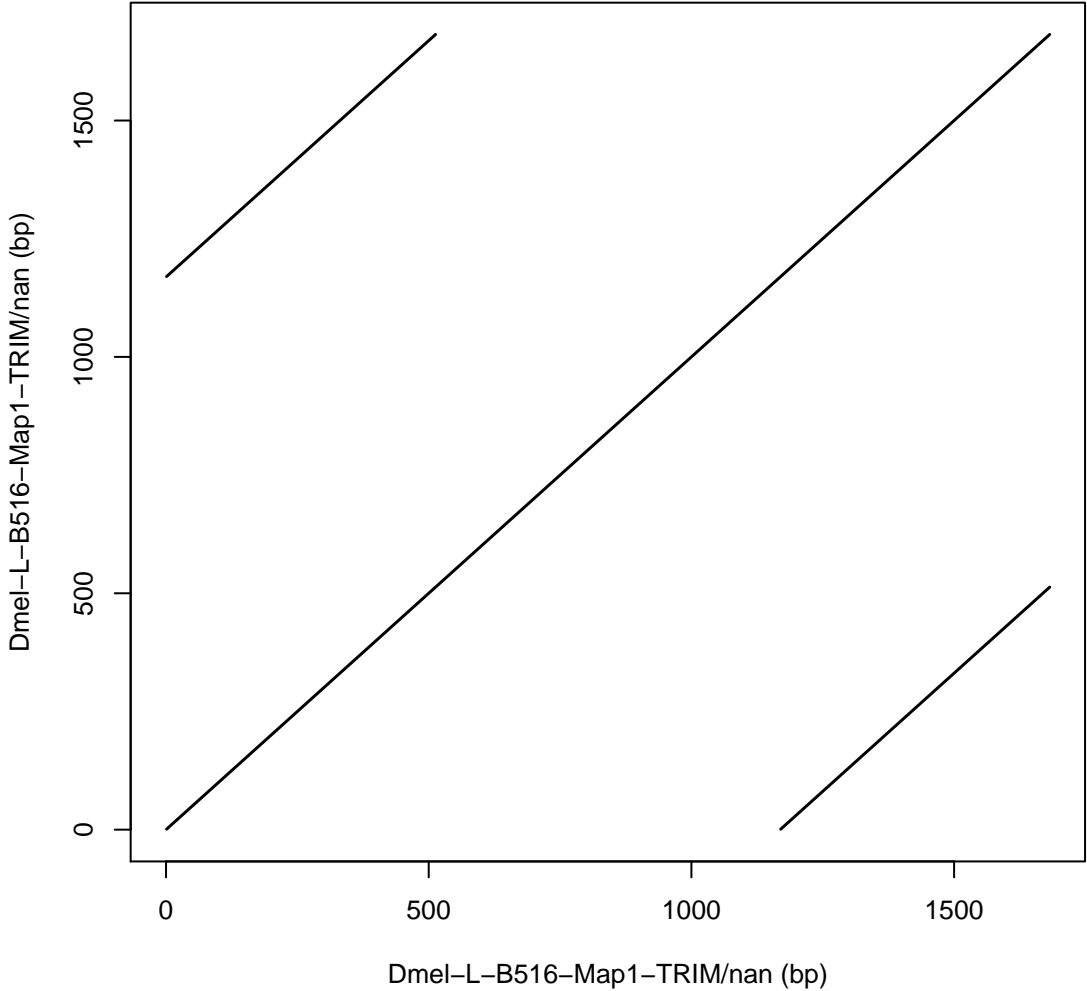consensus size: 1682bp; fragments: 3; full length: 1 (>=1513.8bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

Gypsy−22_DWil_pol LTR/Gypsy
UN−ISRm5_Sme TIR/MULE−MuDR
EX−1_CR_pol LINE/RTE−X
Gypsy−82_CQ_pol LTR/Gypsy
CR1−7_AMi_pol LINE/CR1