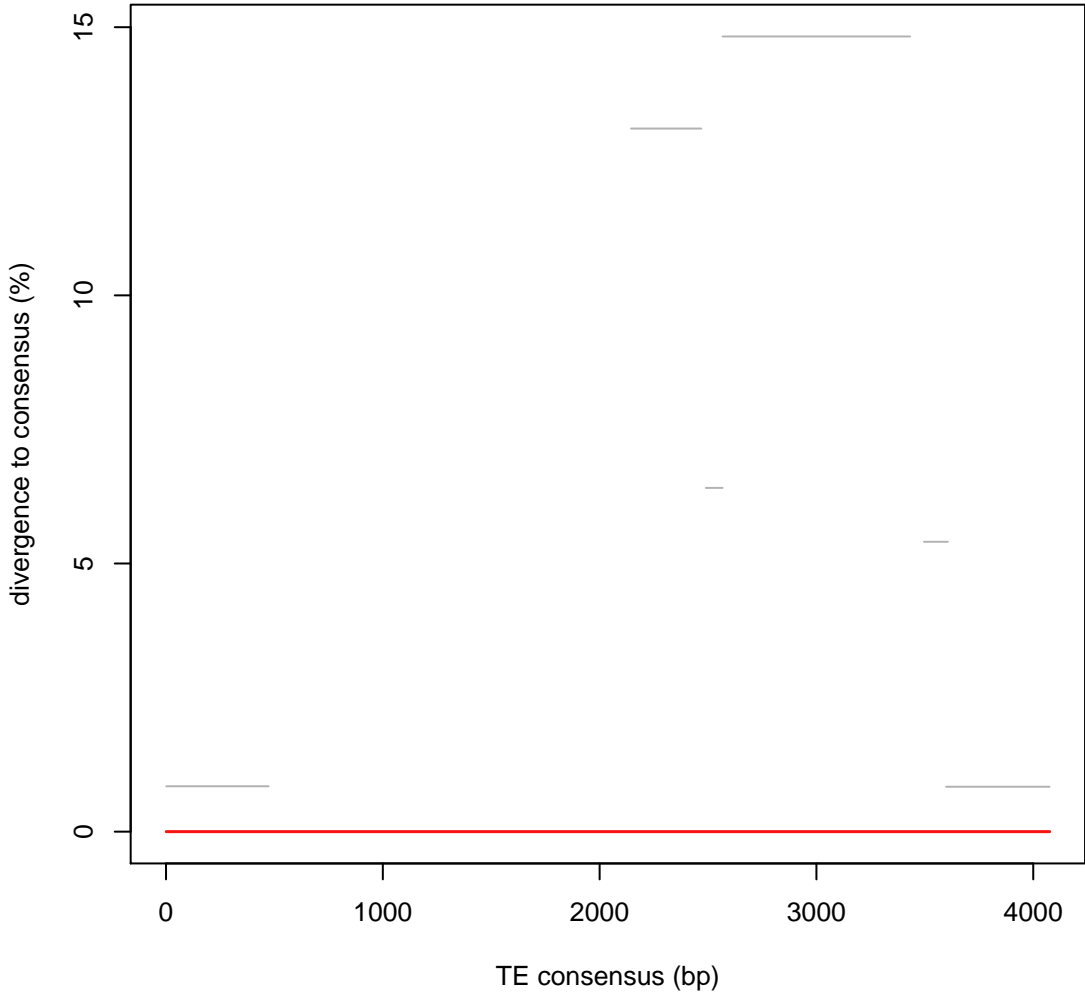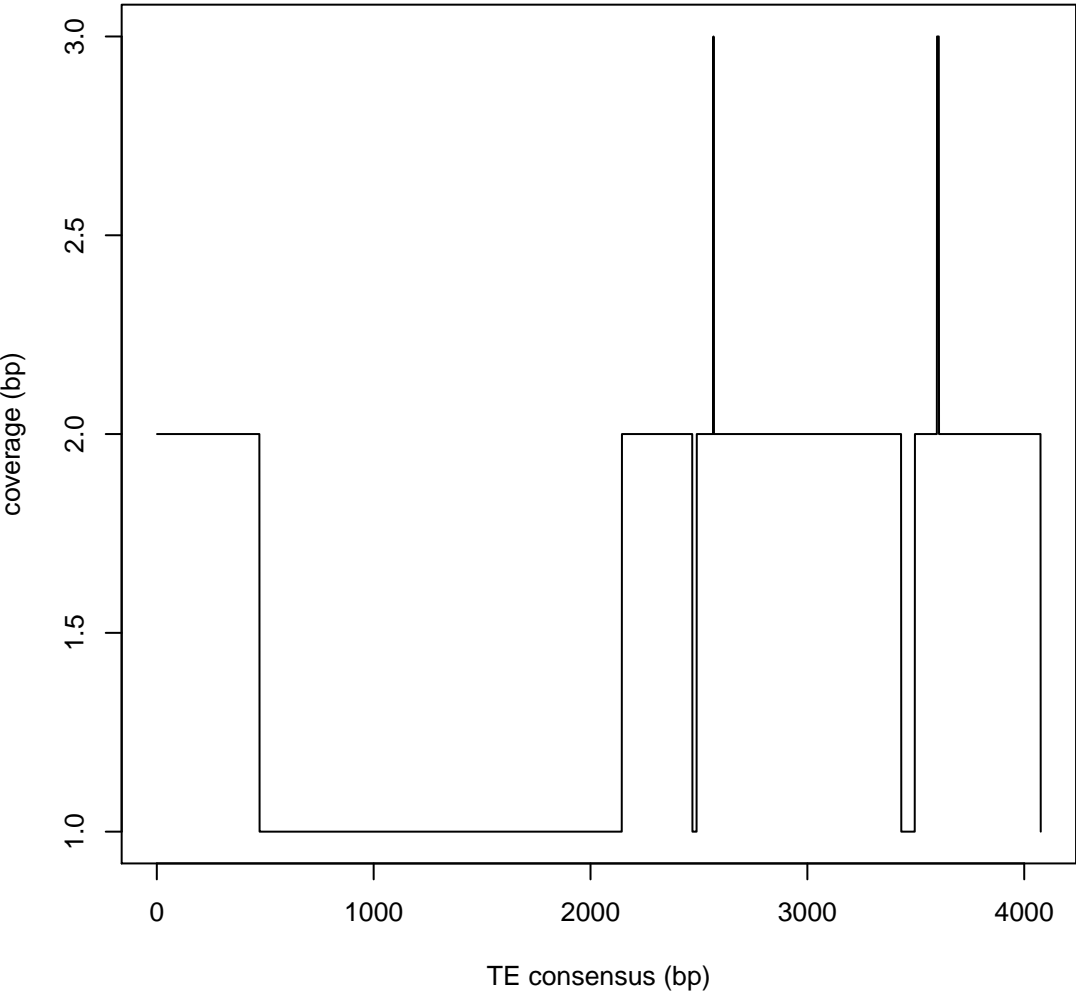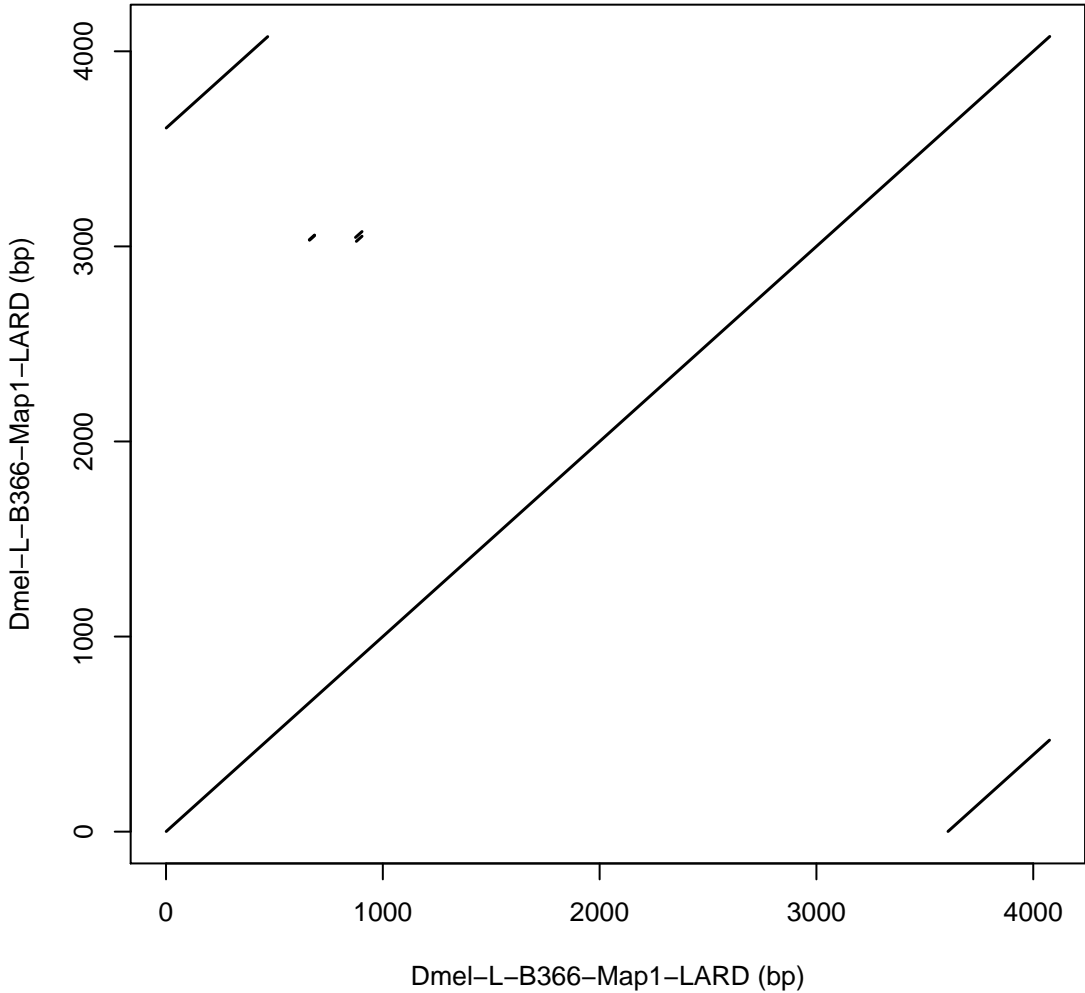**TE: Dmel−L−B366−Map1−LARD**
consensus size: 4076bp; fragments: 7; full length: 1 (>=3668.4bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

L1−5_Alp_pol LINE/L1
L1−33_XT_pol LINE/L1
SC−5 LTR/Copia
RTE−7_AMi_pol LINE/RTE−BovB
L1−96_Cis_pol LINE/L1−Tx1
CATCH2_DR_pol LTR/Pao
DIRS−1_PPo_yr DIRS/DIRS
L1−33_XT_pol LINE