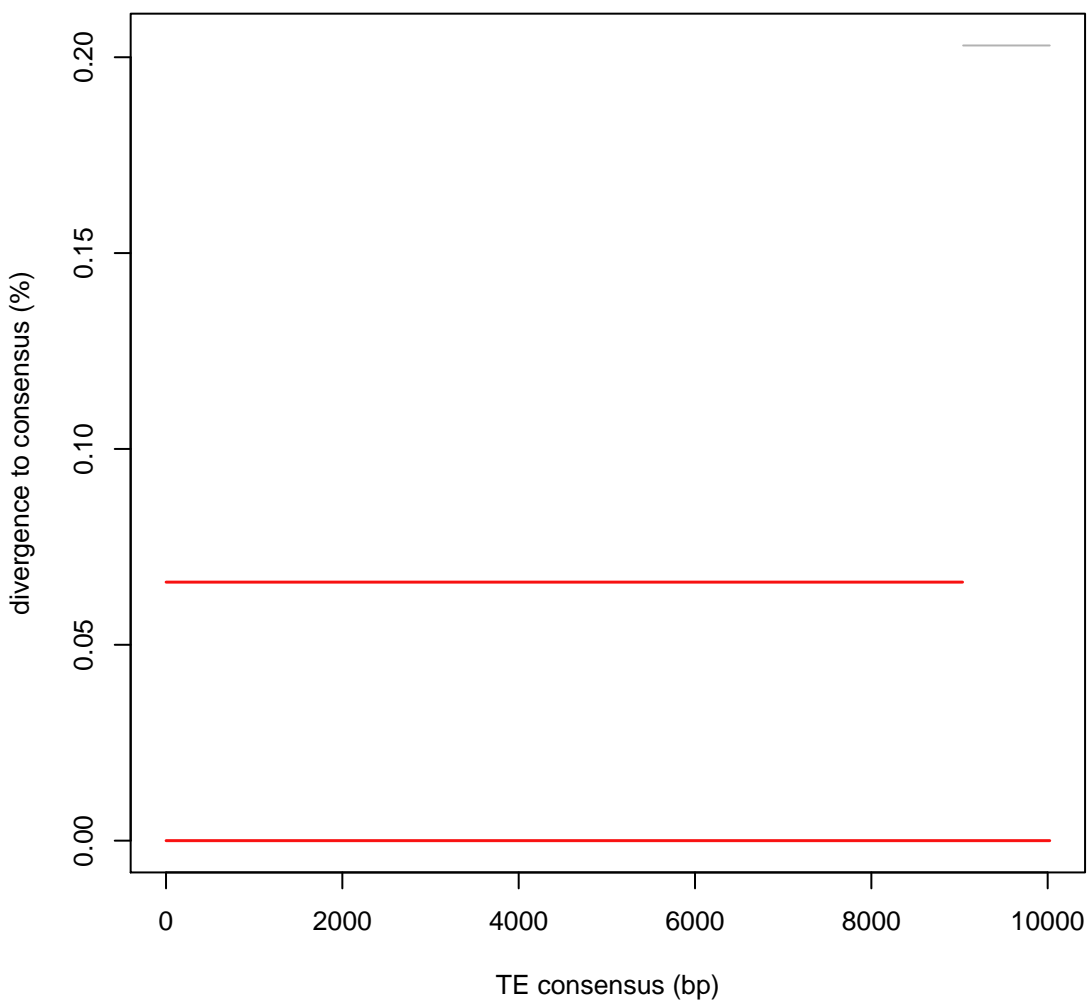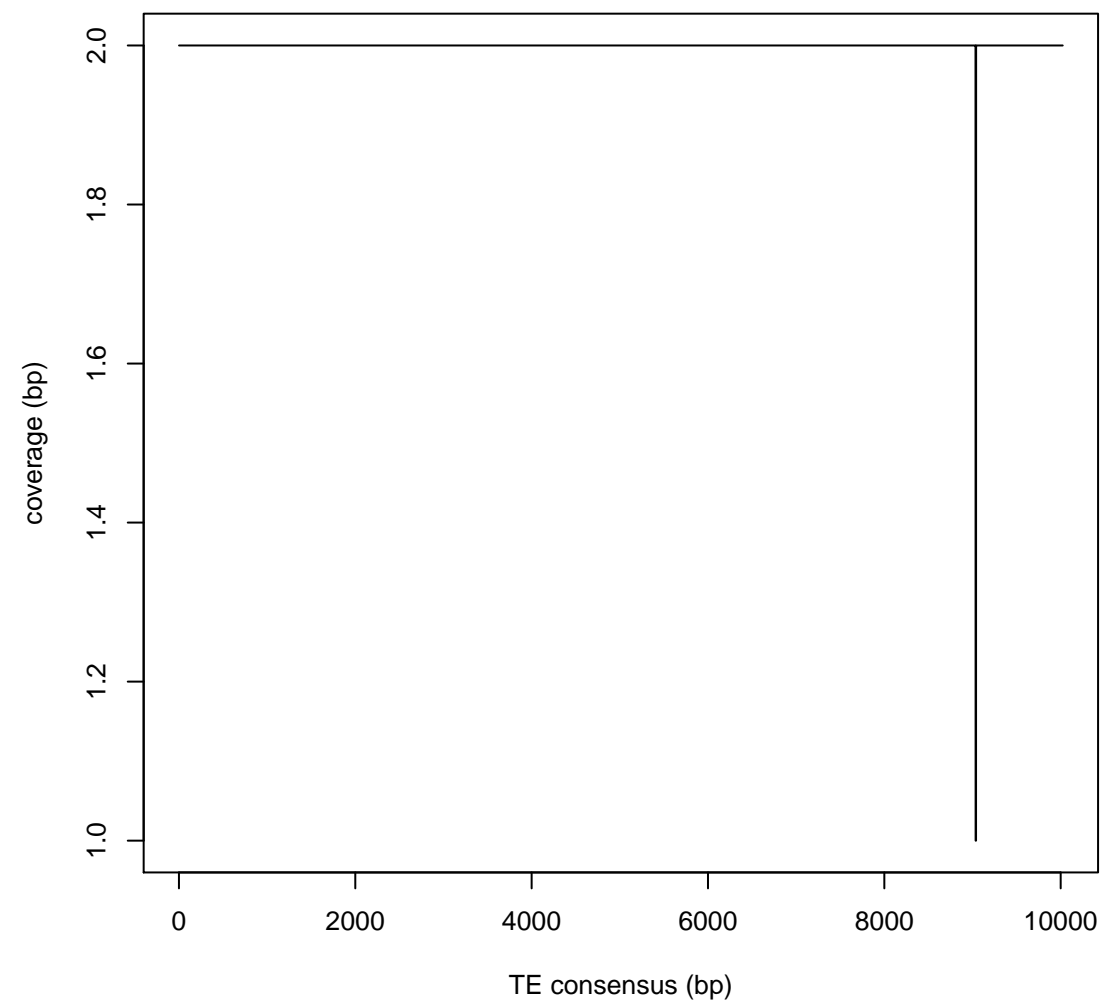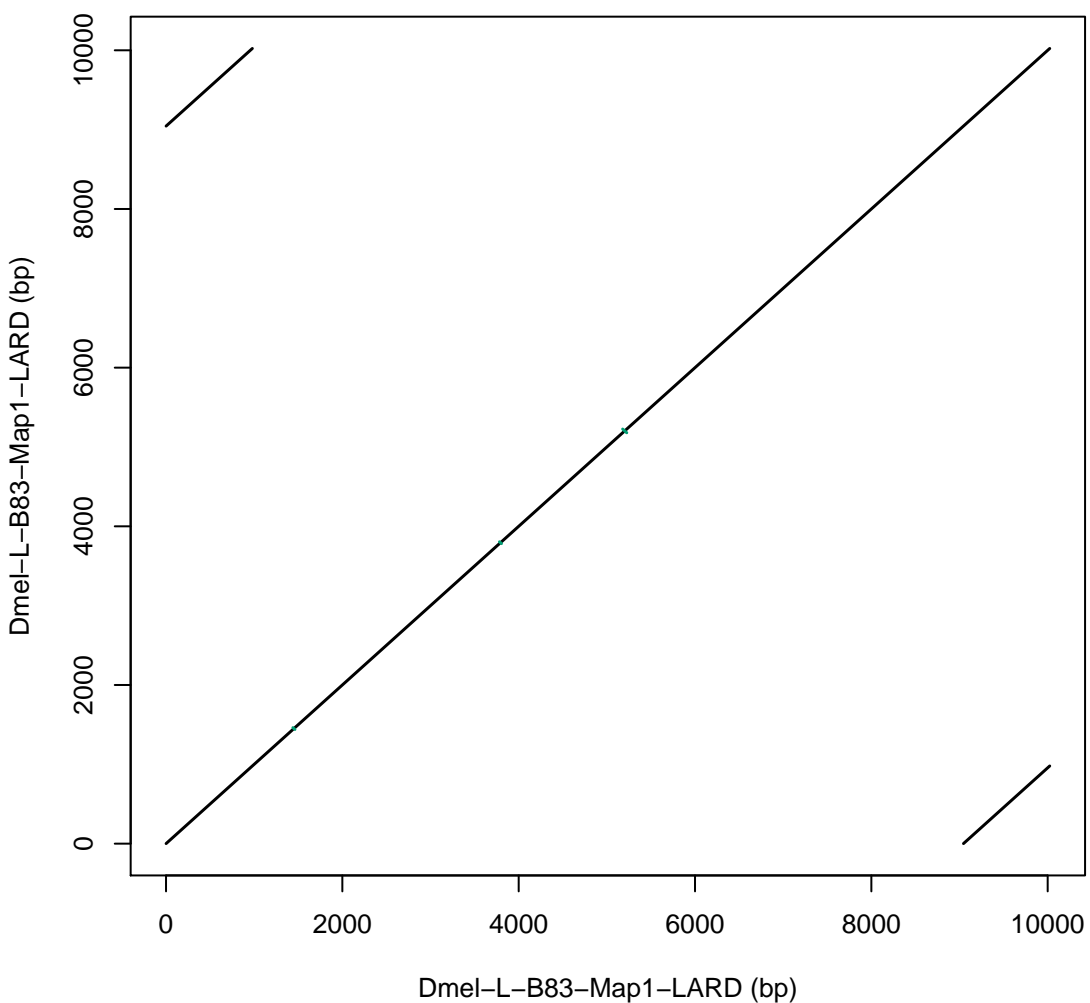**TE: Dmel−L−B83−Map1−LARD**
consensus size: 10023bp; fragments: 3; full length: 2 (>=9020.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

Crack−1_CP_pol LINE/L2

Gypsy−8_ECa_1p LTR/Gypsy

UN−4538901_Ath_pol LINE/L1

L1−39_XT_pol LINE/L1