TE: Dmel−L−B379−Map1−LARD
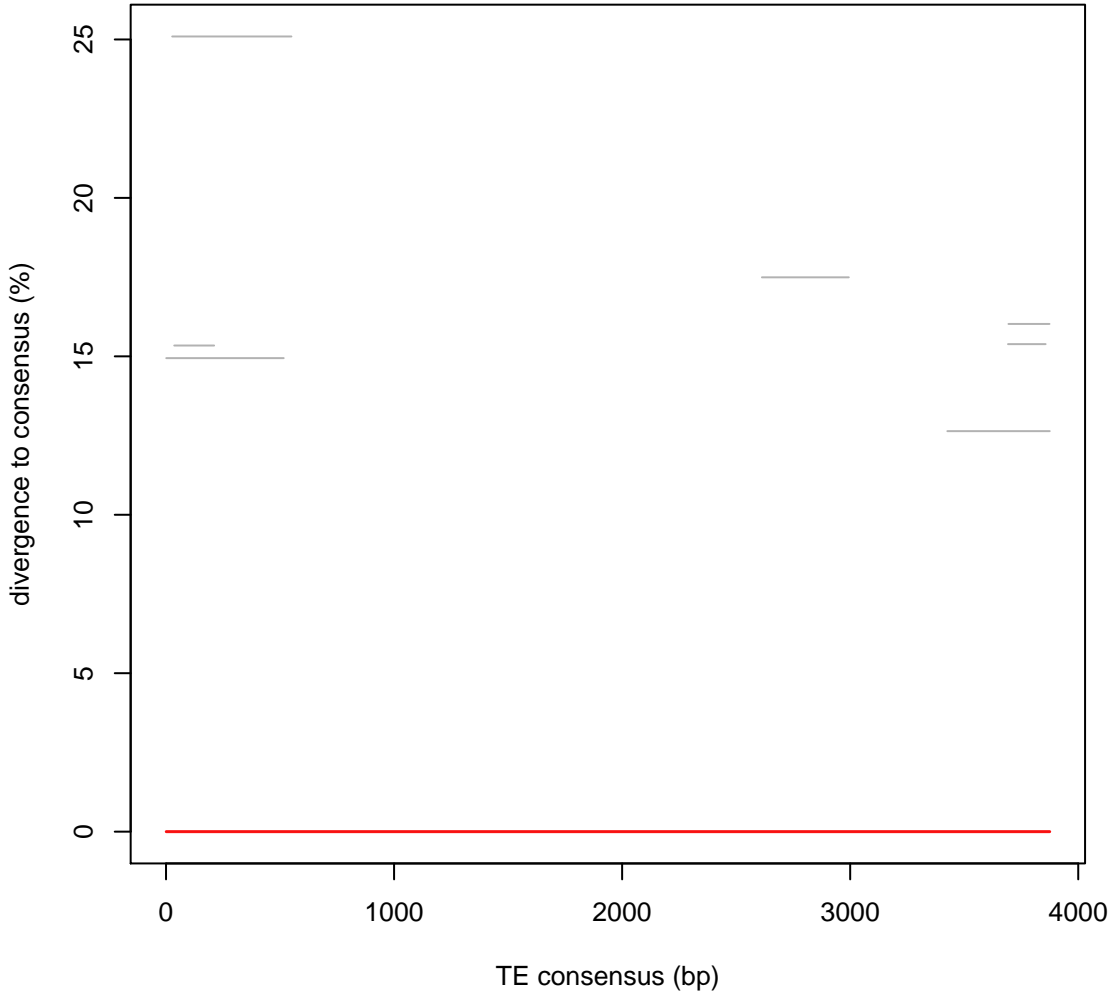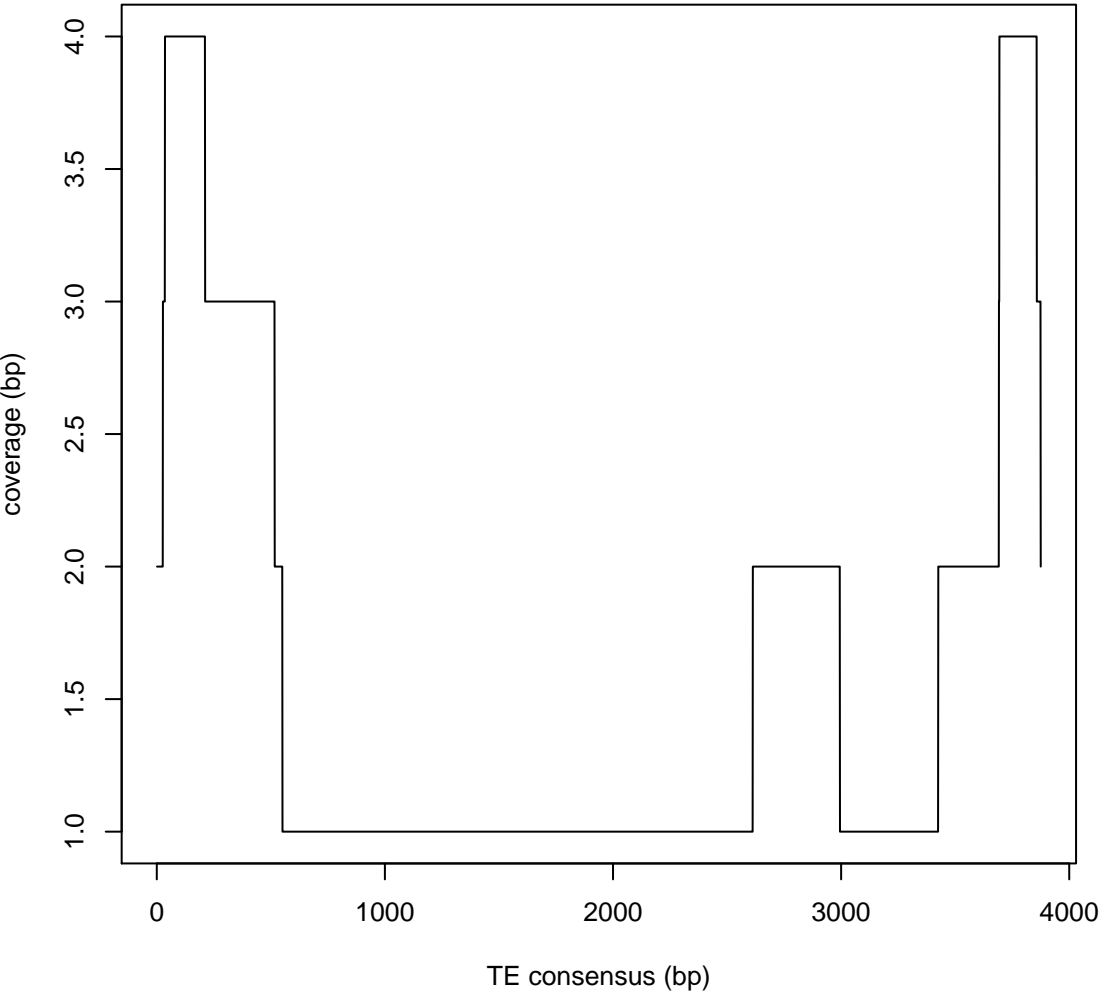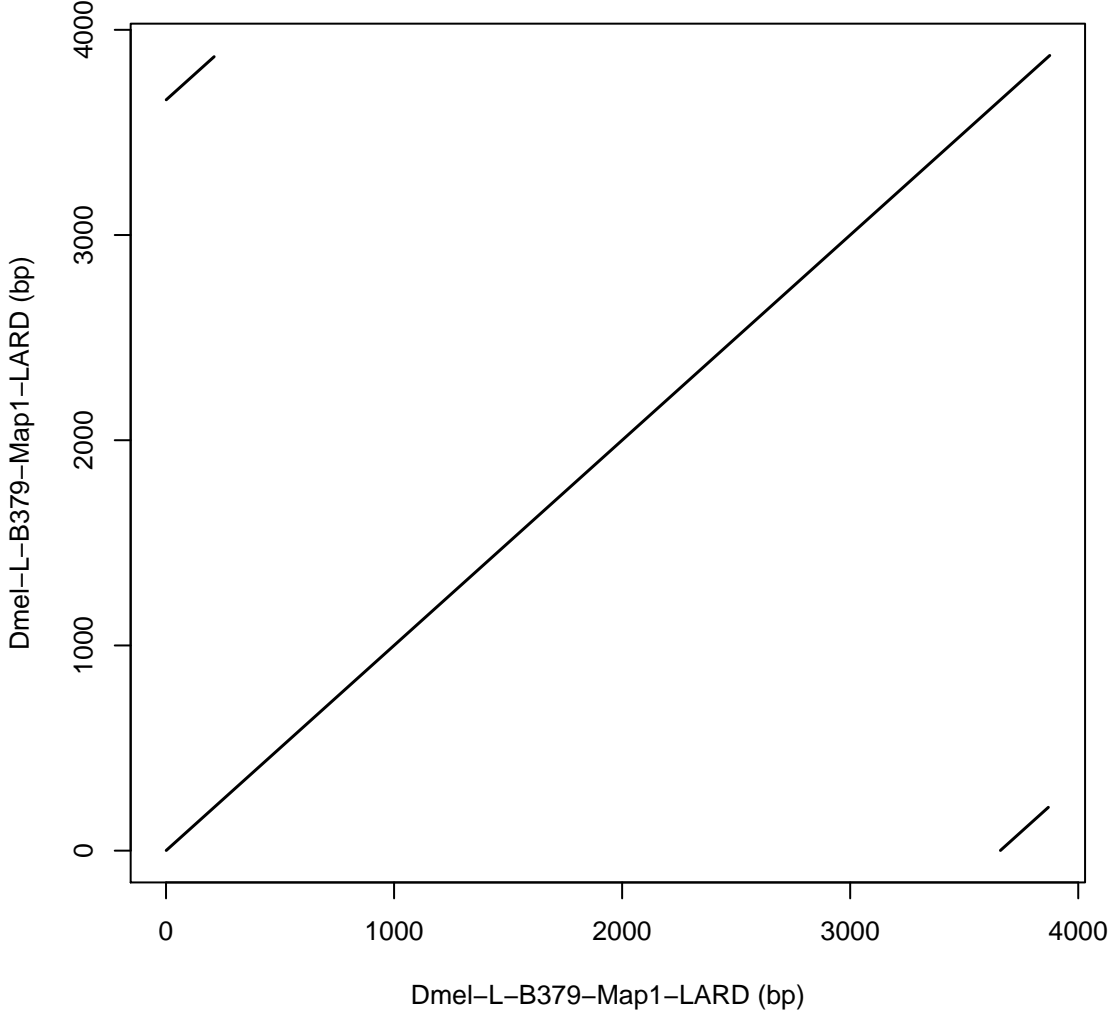**consensus size: 3875bp; fragments: 8; full length: 1 (>=3487.5bp)**

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

UN−294896690_tp TIR/hAT−hAT19