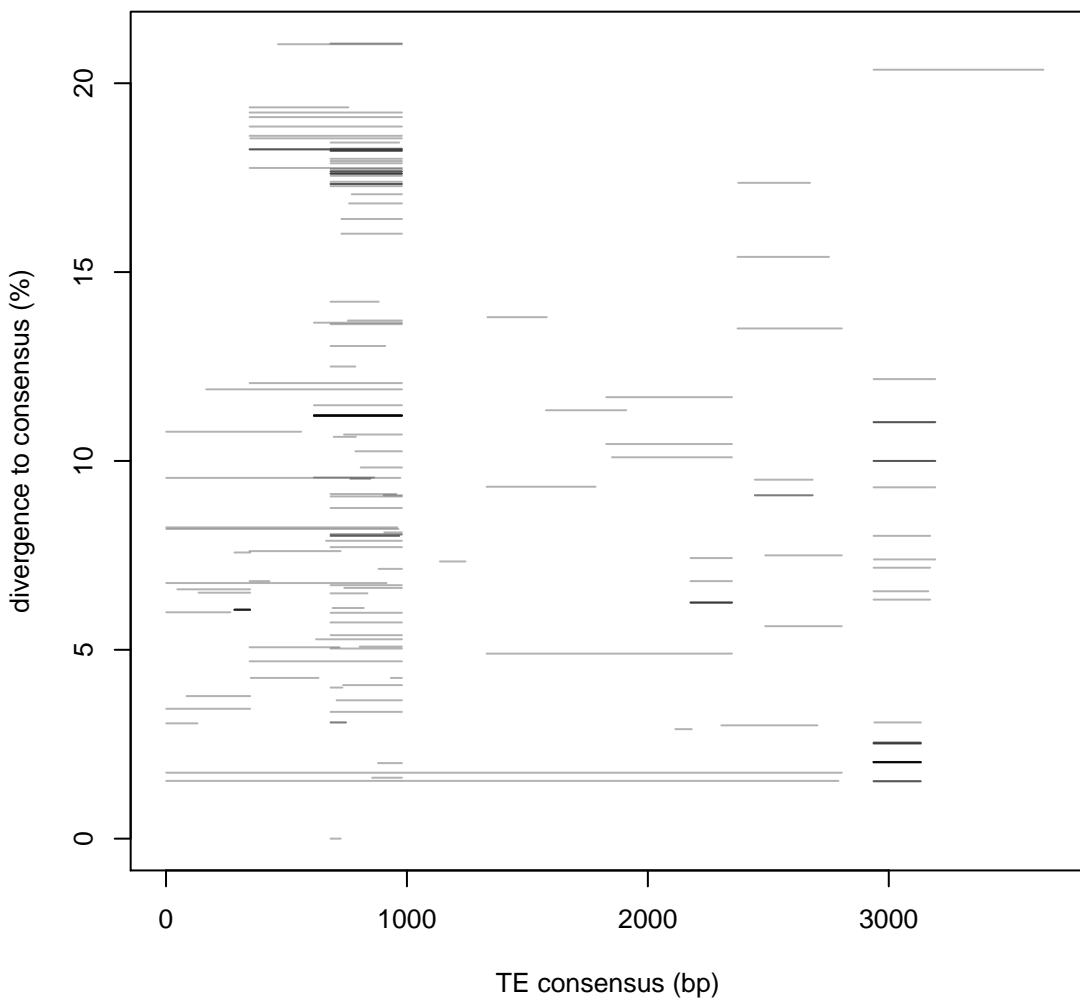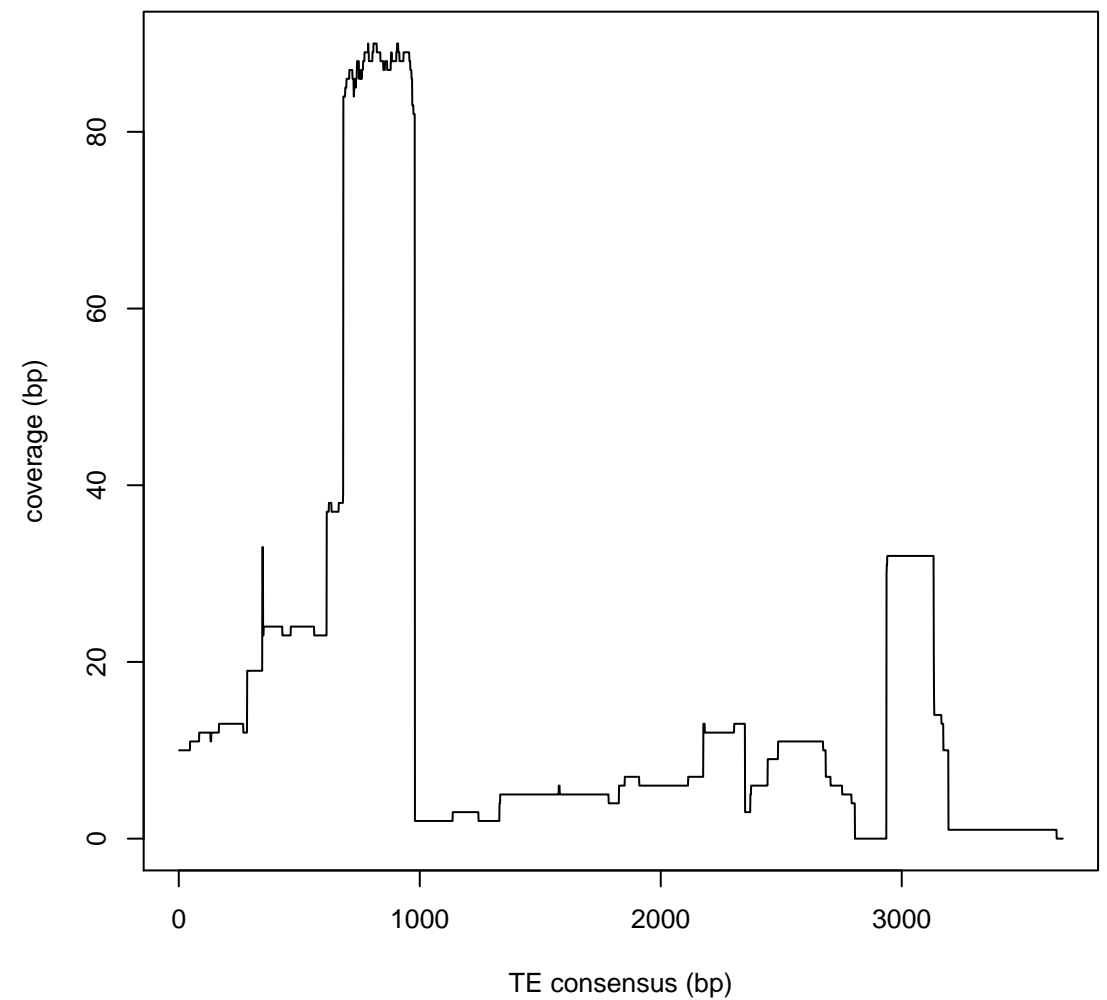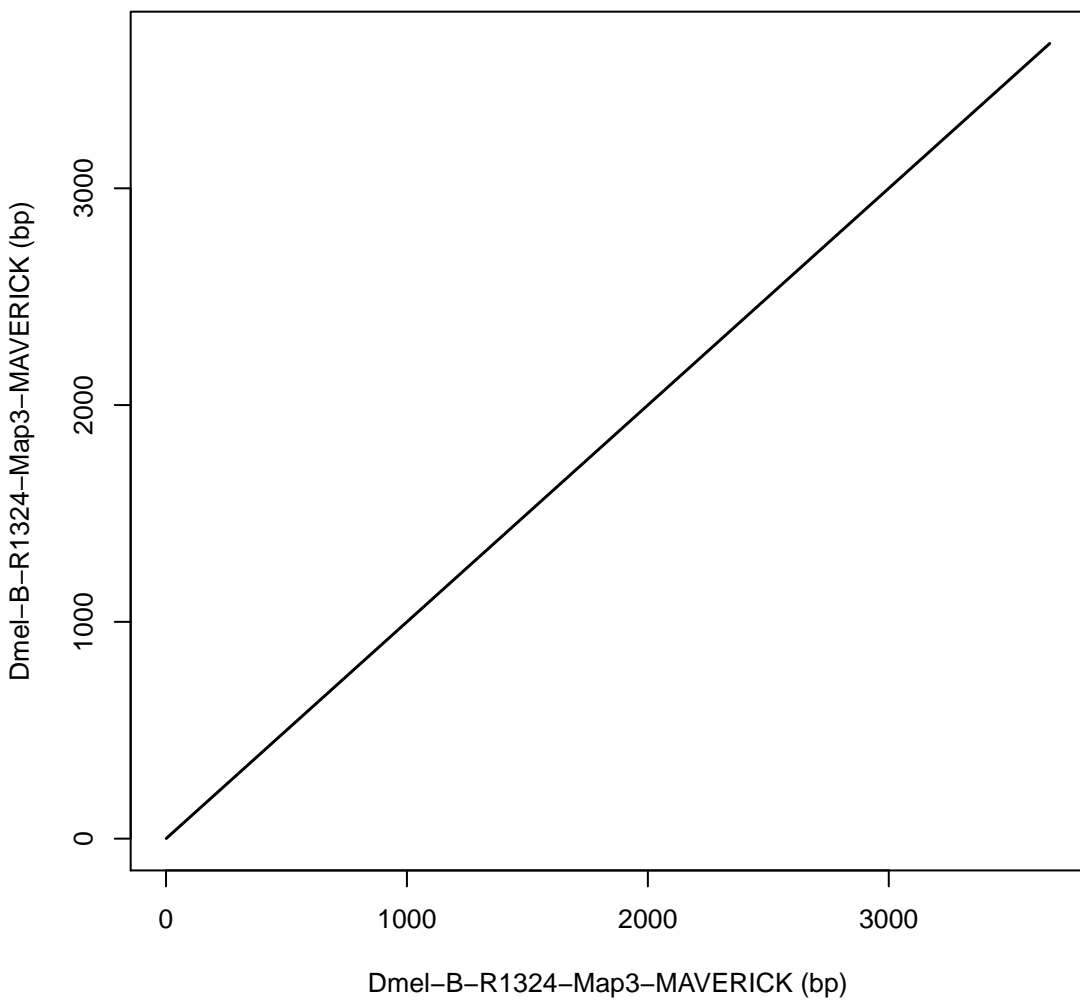**TE: Dmel–B–R1324–Map3–MAVERICK**
consensus size: 3668bp; fragments: 177; full length: 0 (>=3301.2bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

hAT–19_HM_tp TIR/hAT–Charlie