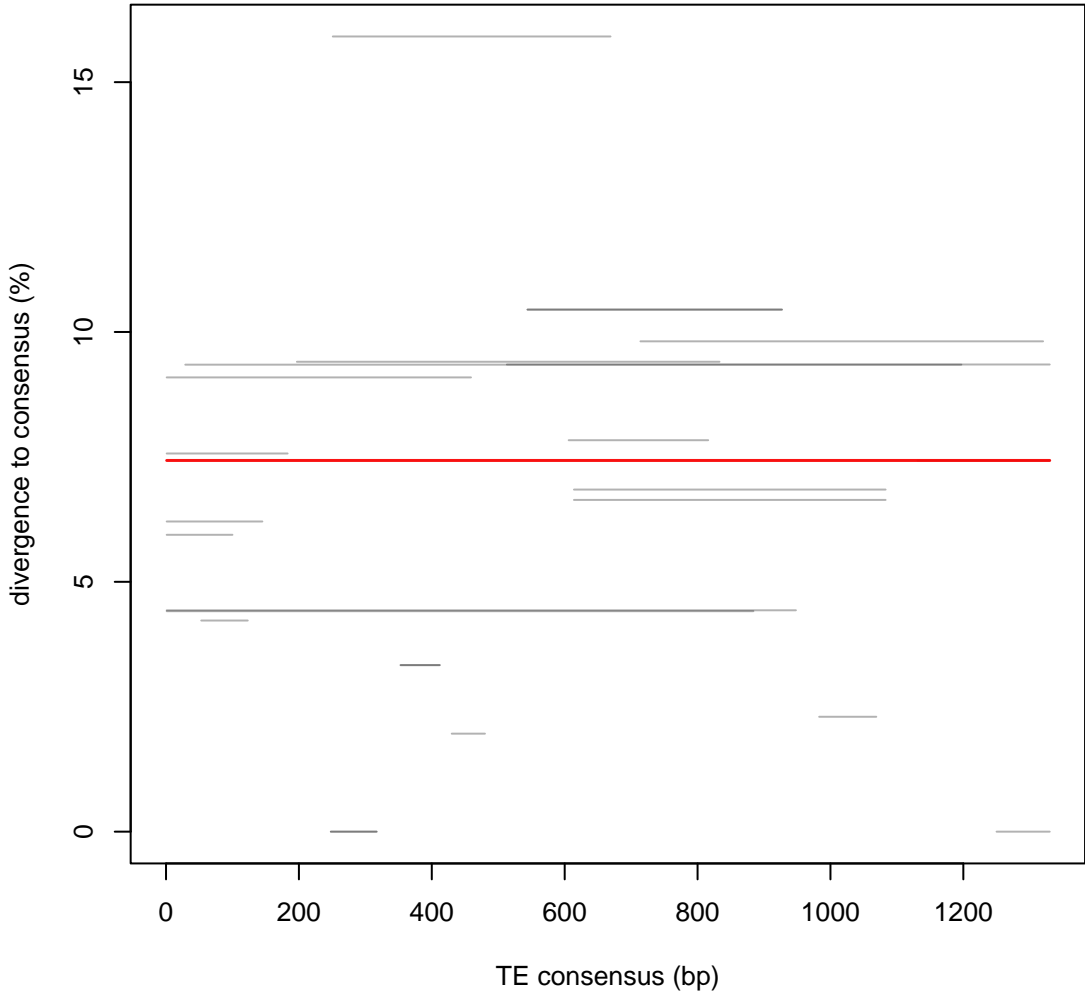
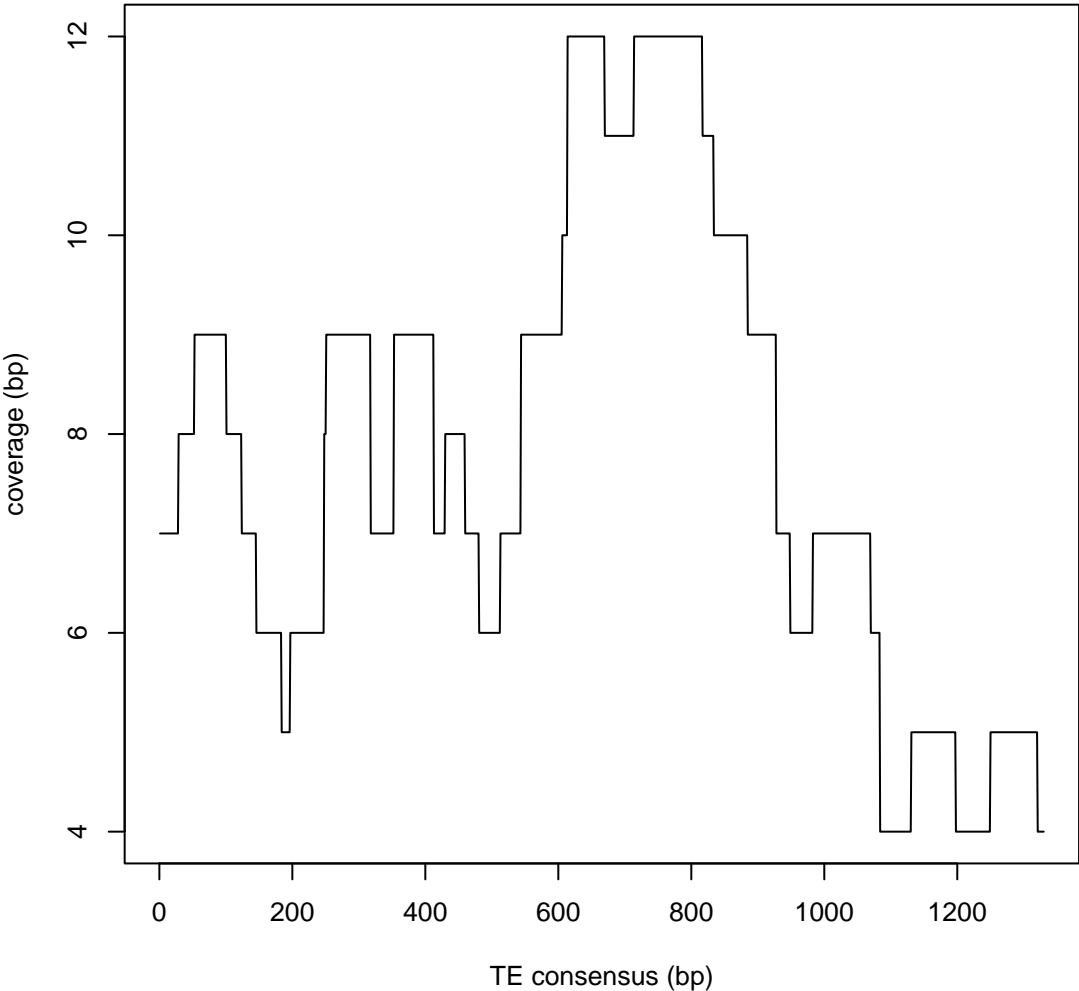


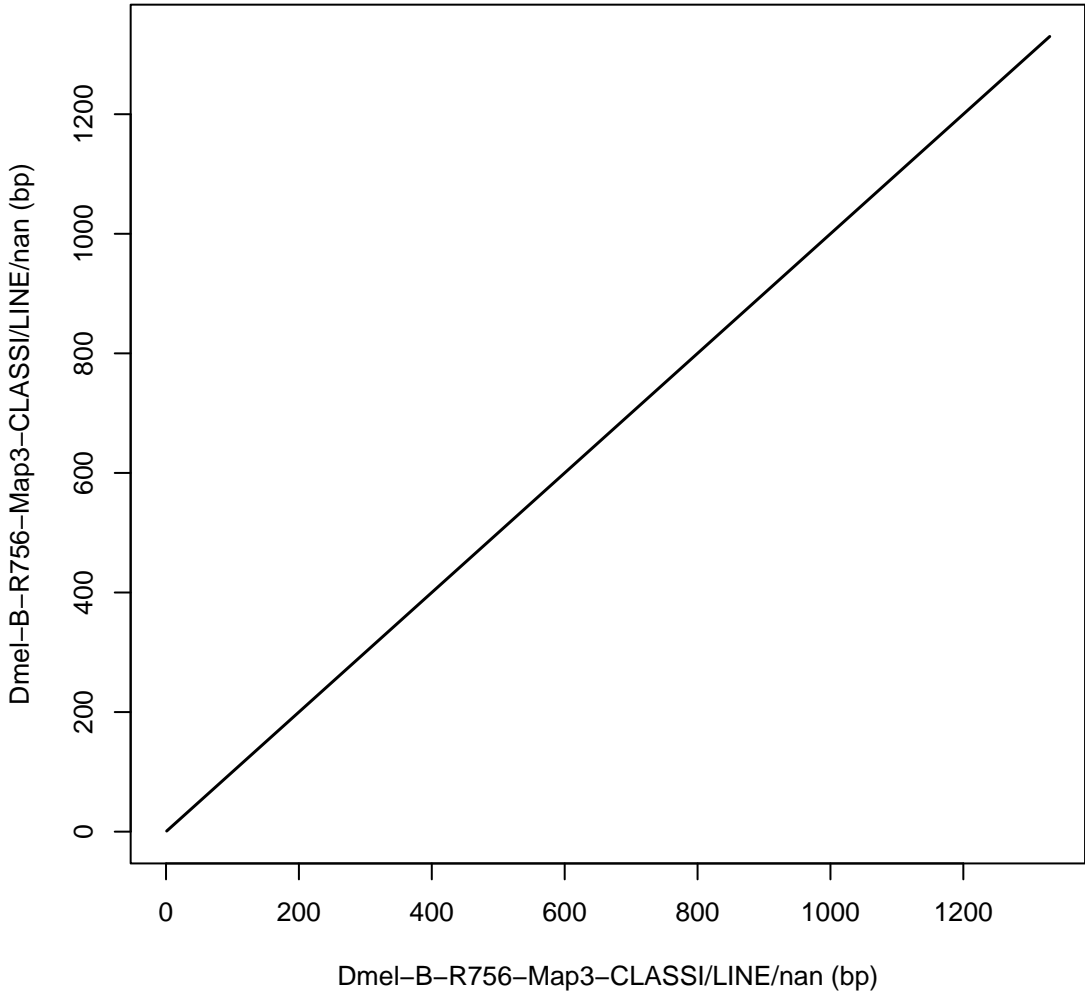
TE: Dmel-B-R756-Map3-CLASSI/LINE/nan
consensus size: 1330bp; fragments: 26; full length: 1 (>=1197bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

