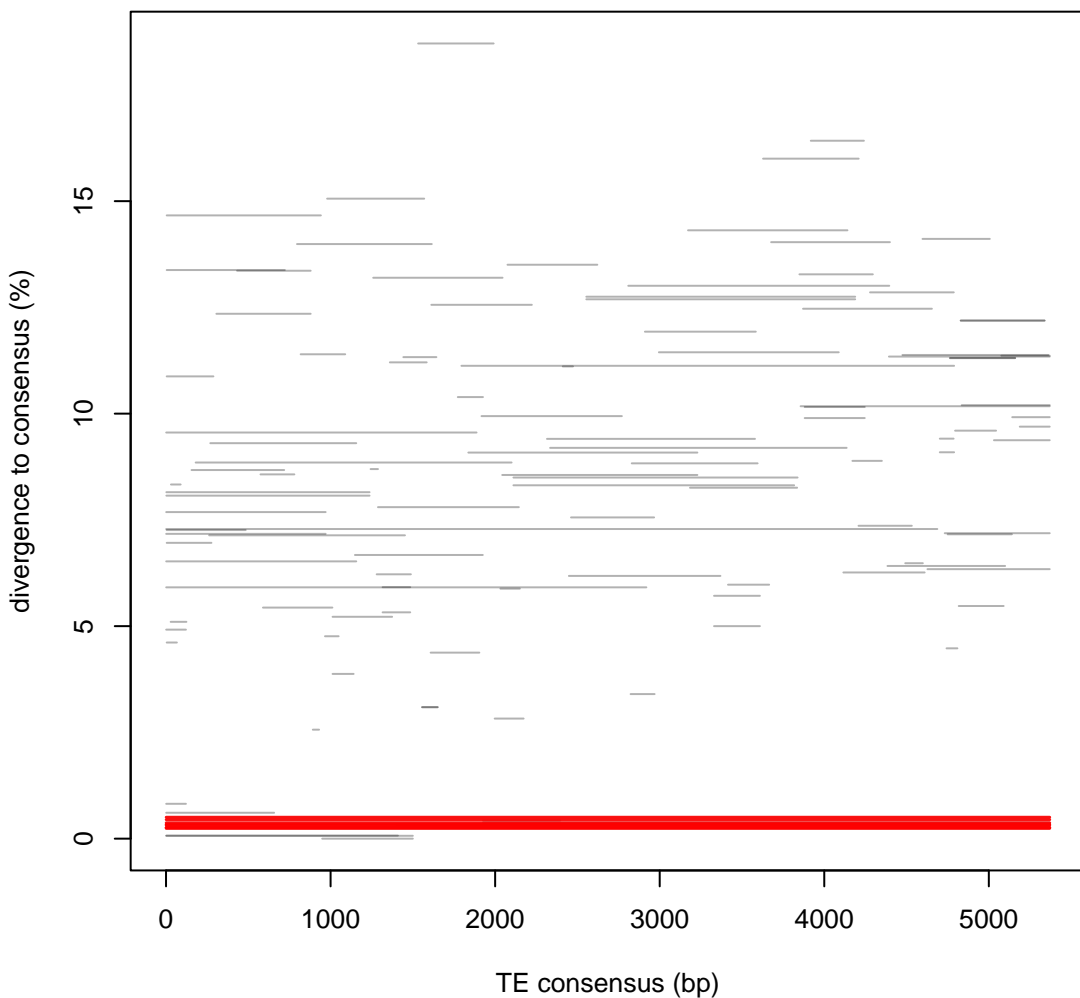
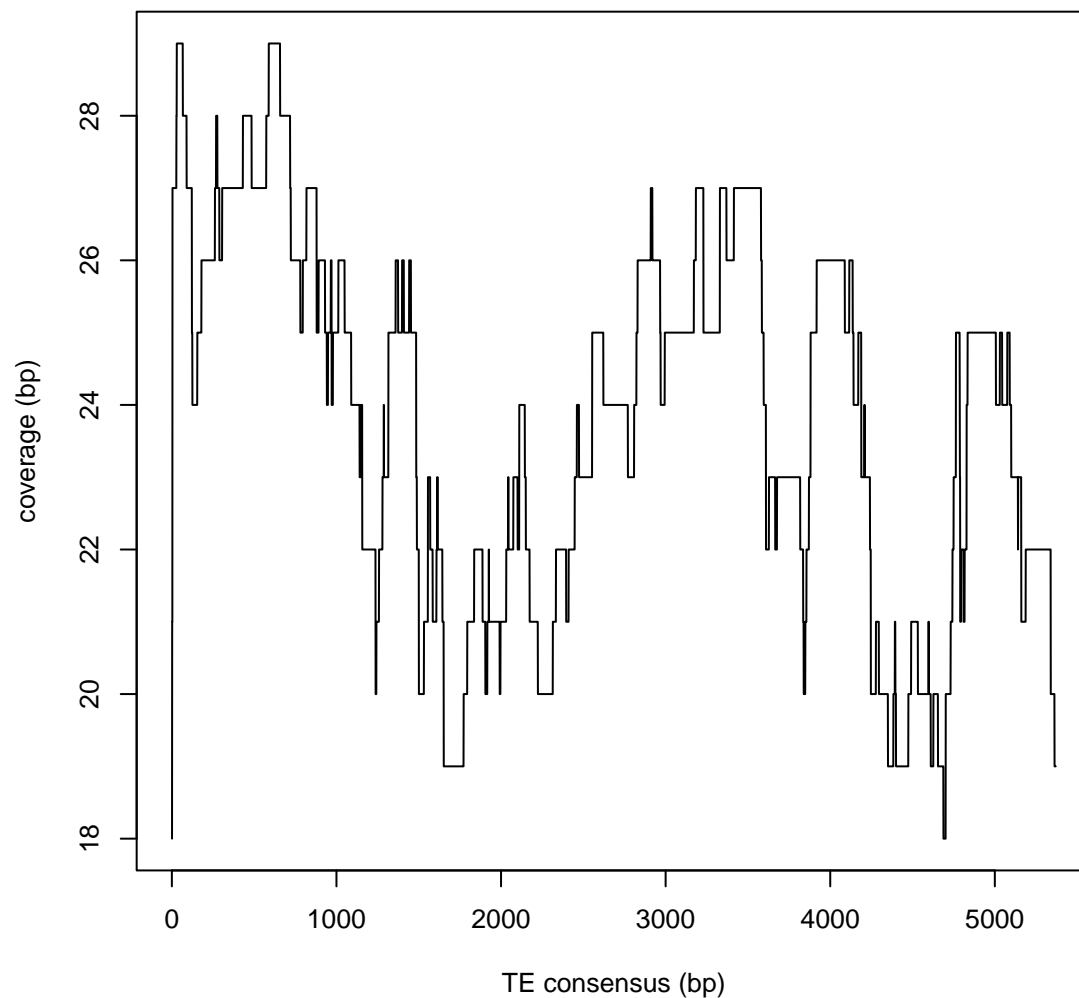


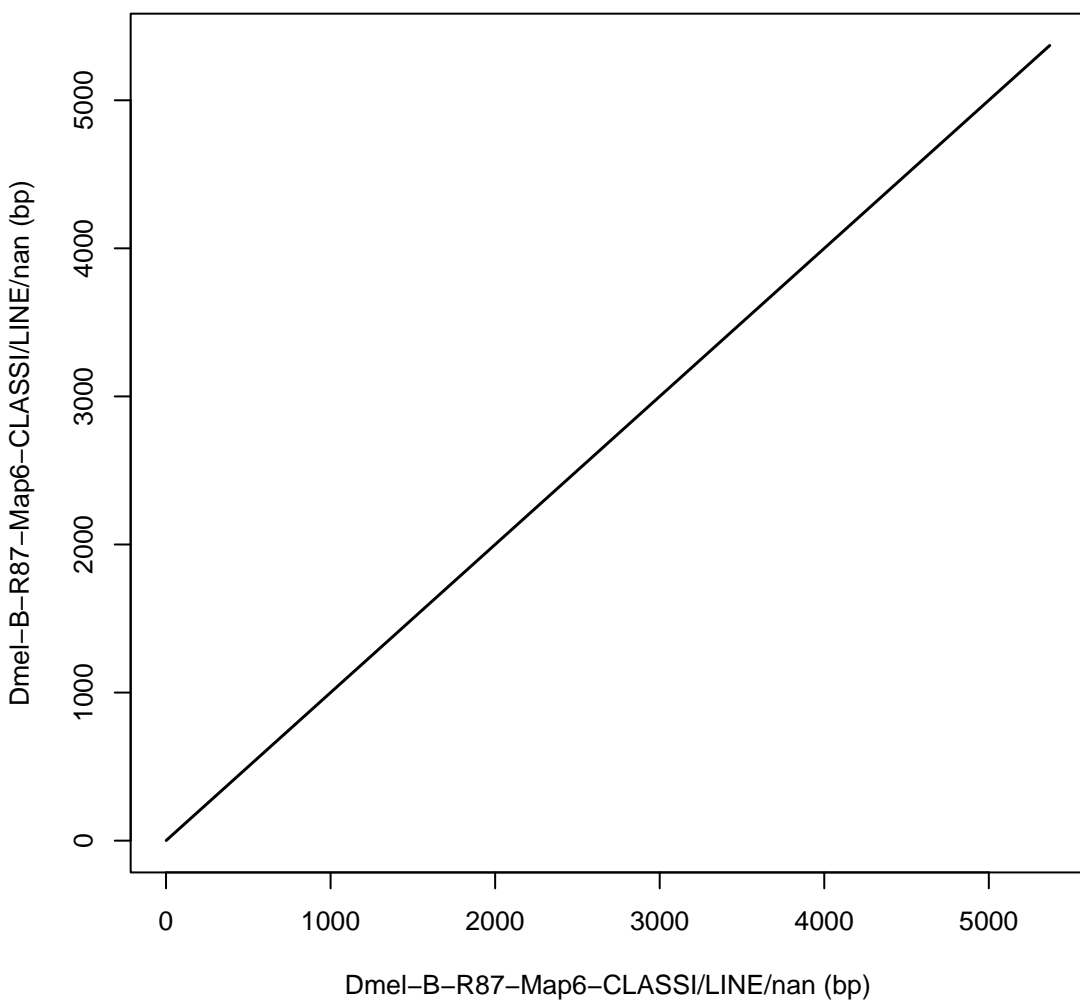
TE: Dmel-B-R87-Map6-CLASSI/LINE/nan
consensus size: 5370bp; fragments: 123; full length: 8 (>=4833bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

