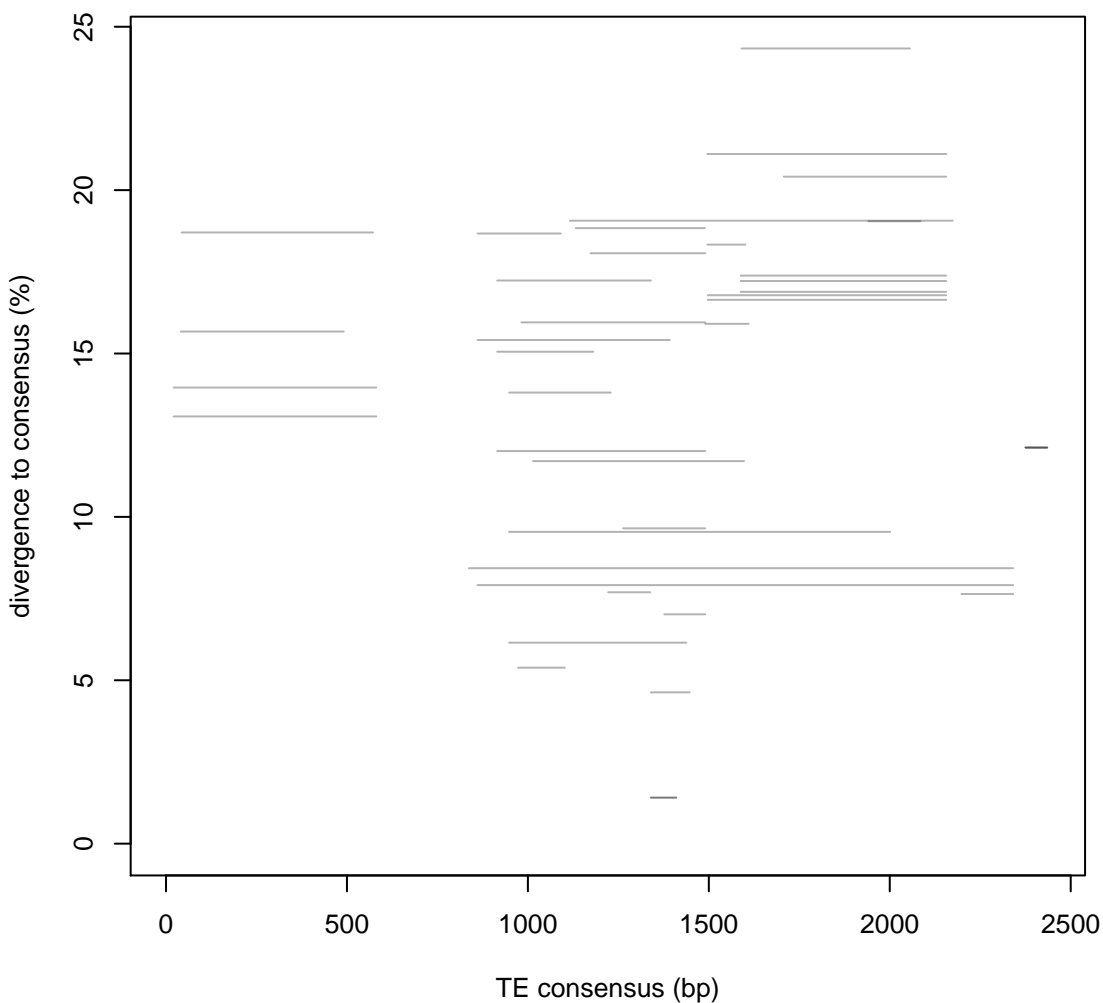
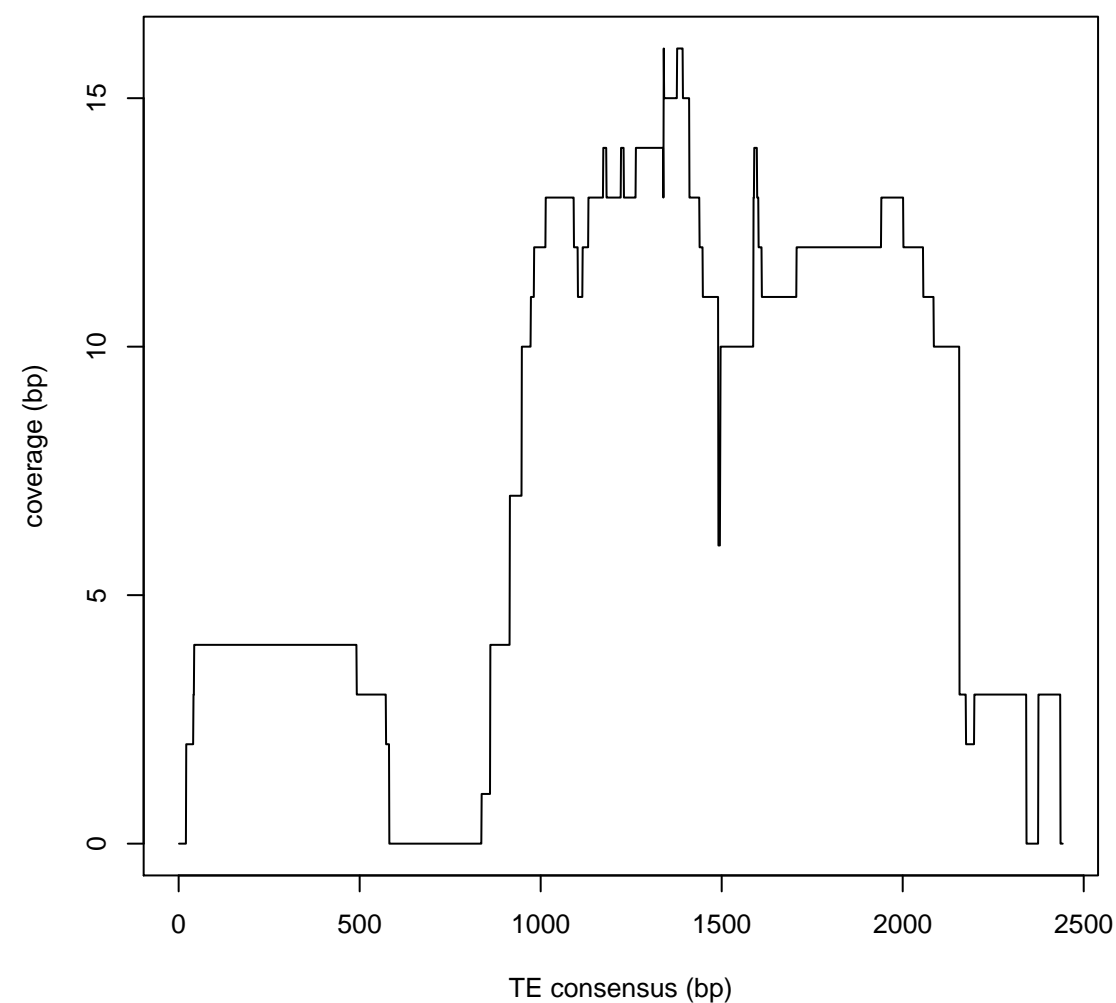


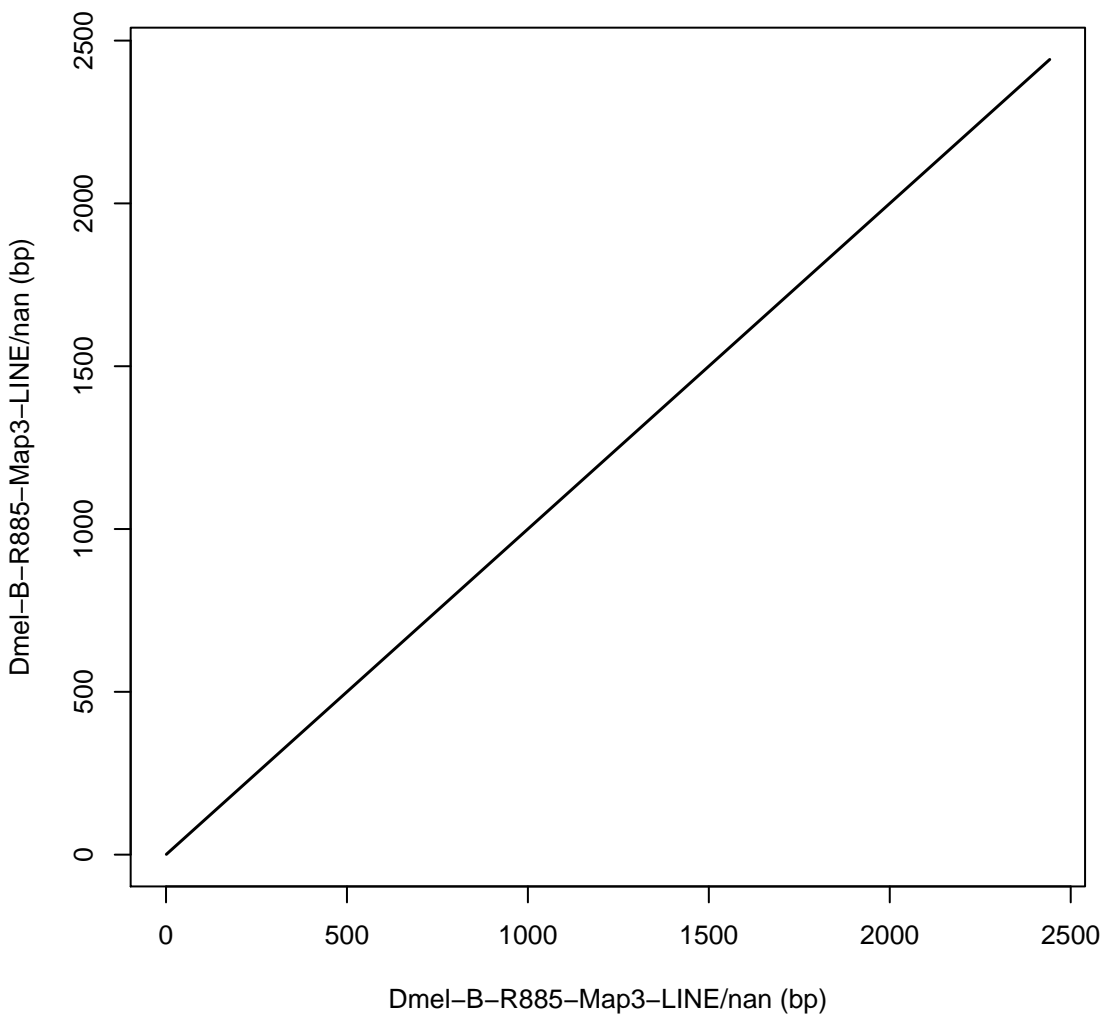
TE: Dmel-B-R885-Map3-LINE/nan  
consensus size: 2442bp; fragments: 41; full length: 0 (>=2197.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

