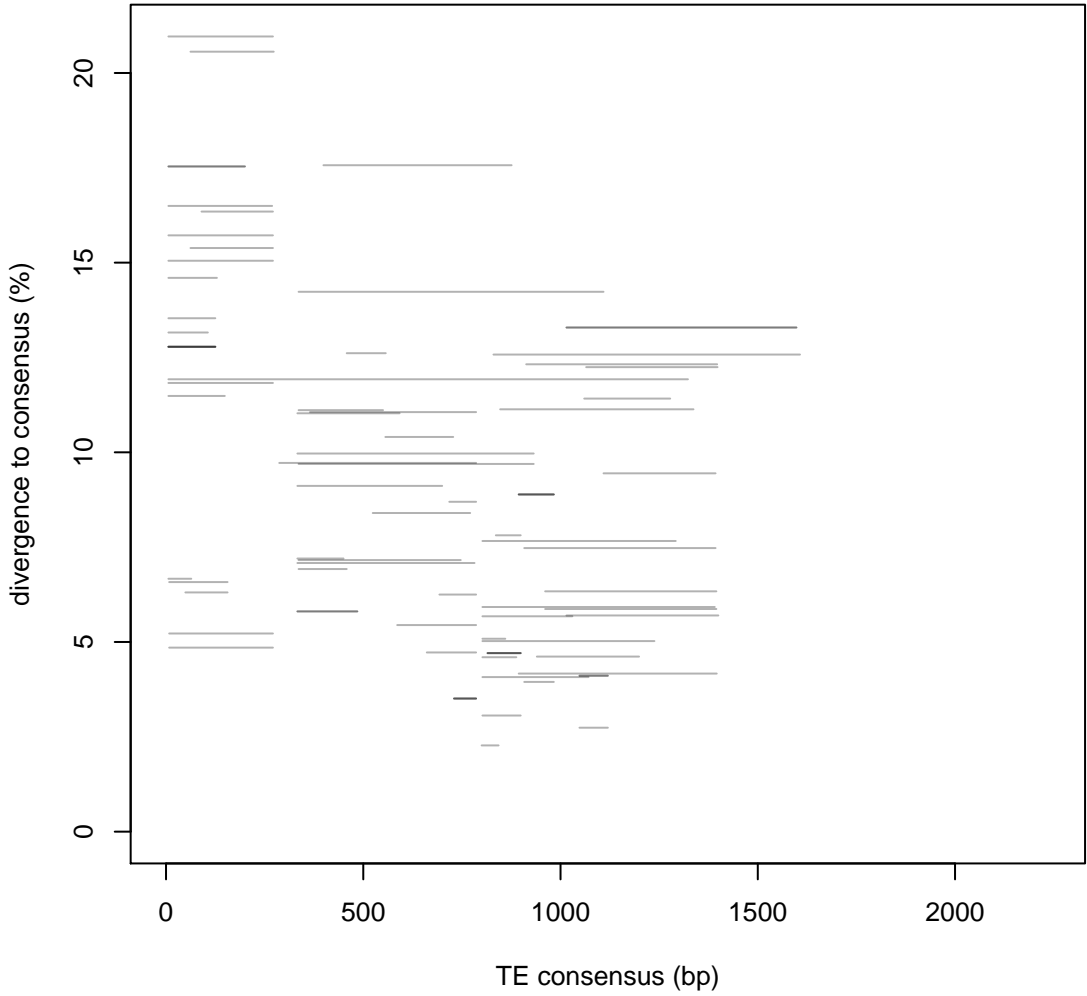
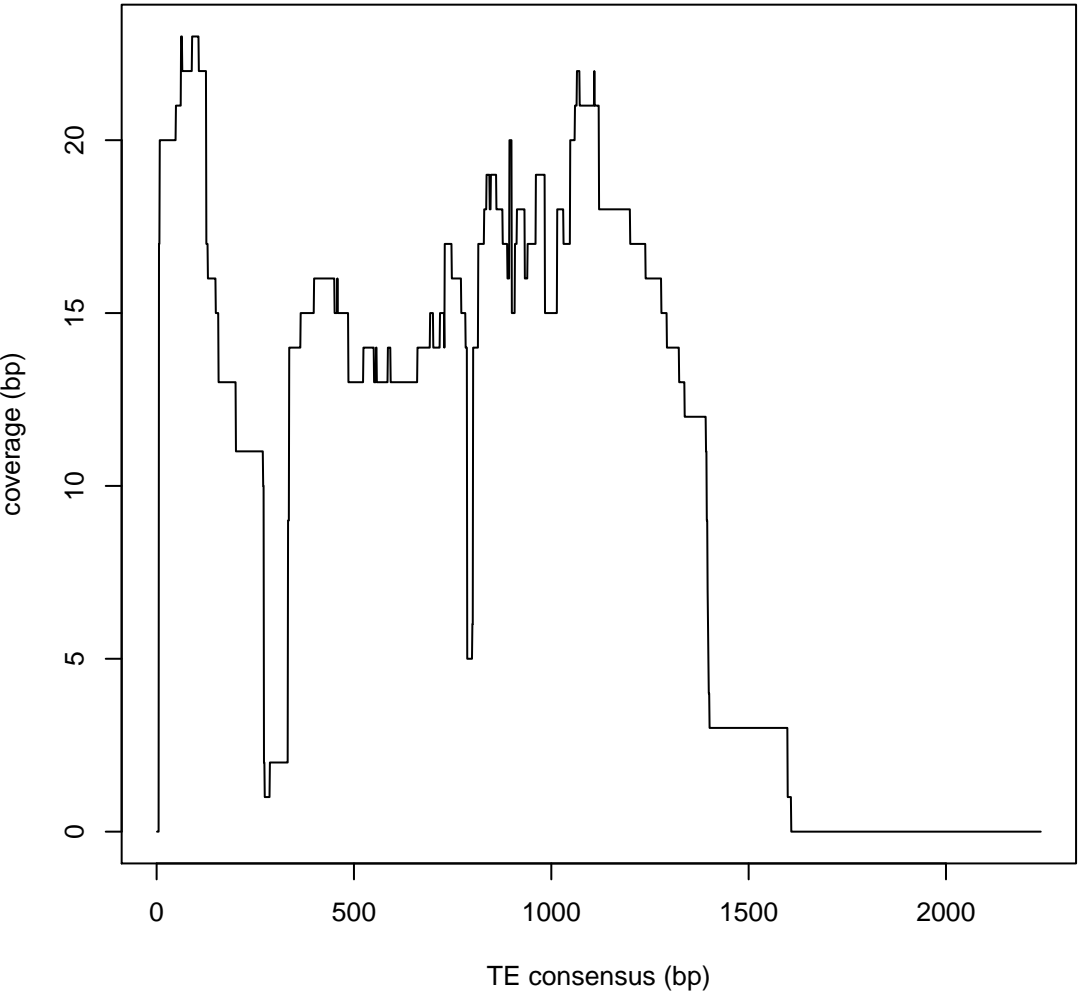


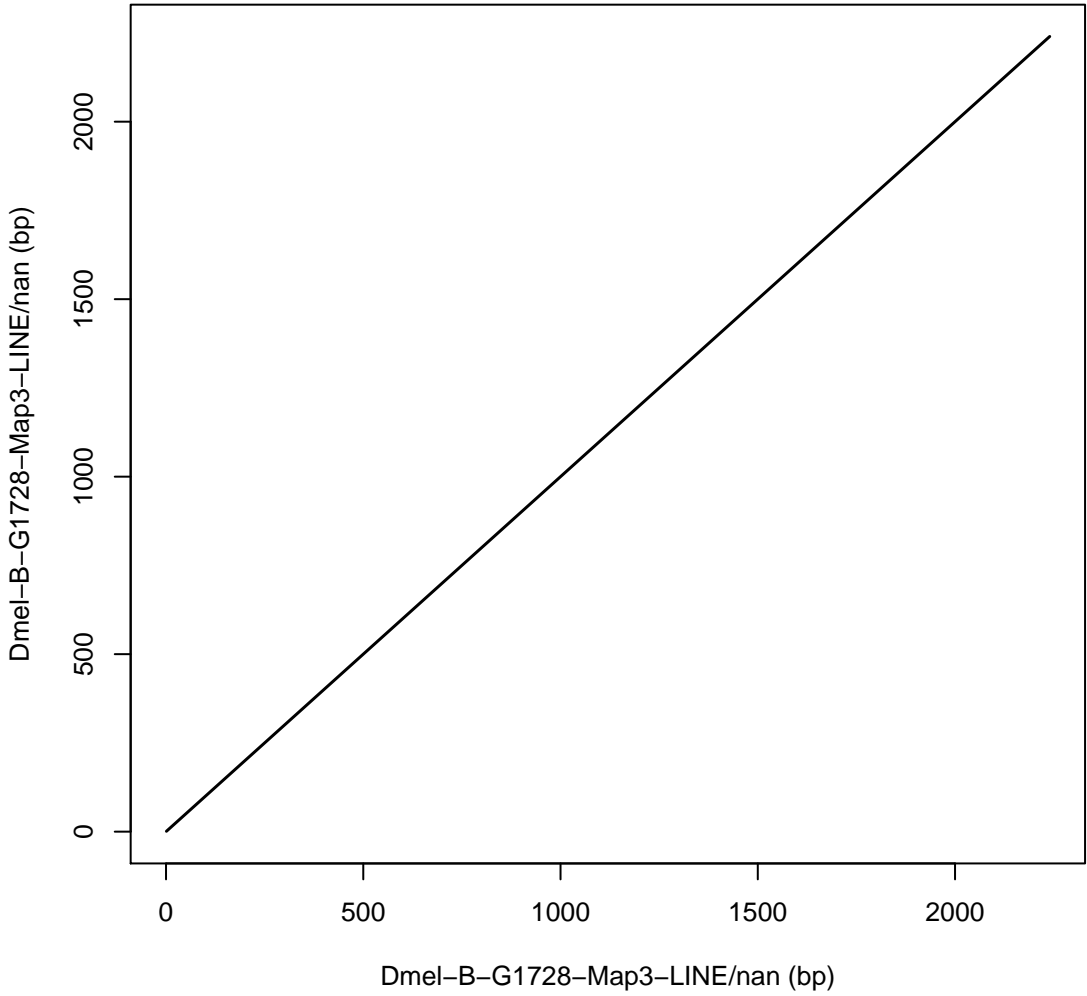
TE: Dmel-B-G1728-Map3-LINE/nan  
consensus size: 2240bp; fragments: 83; full length: 0 (>=2016bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

