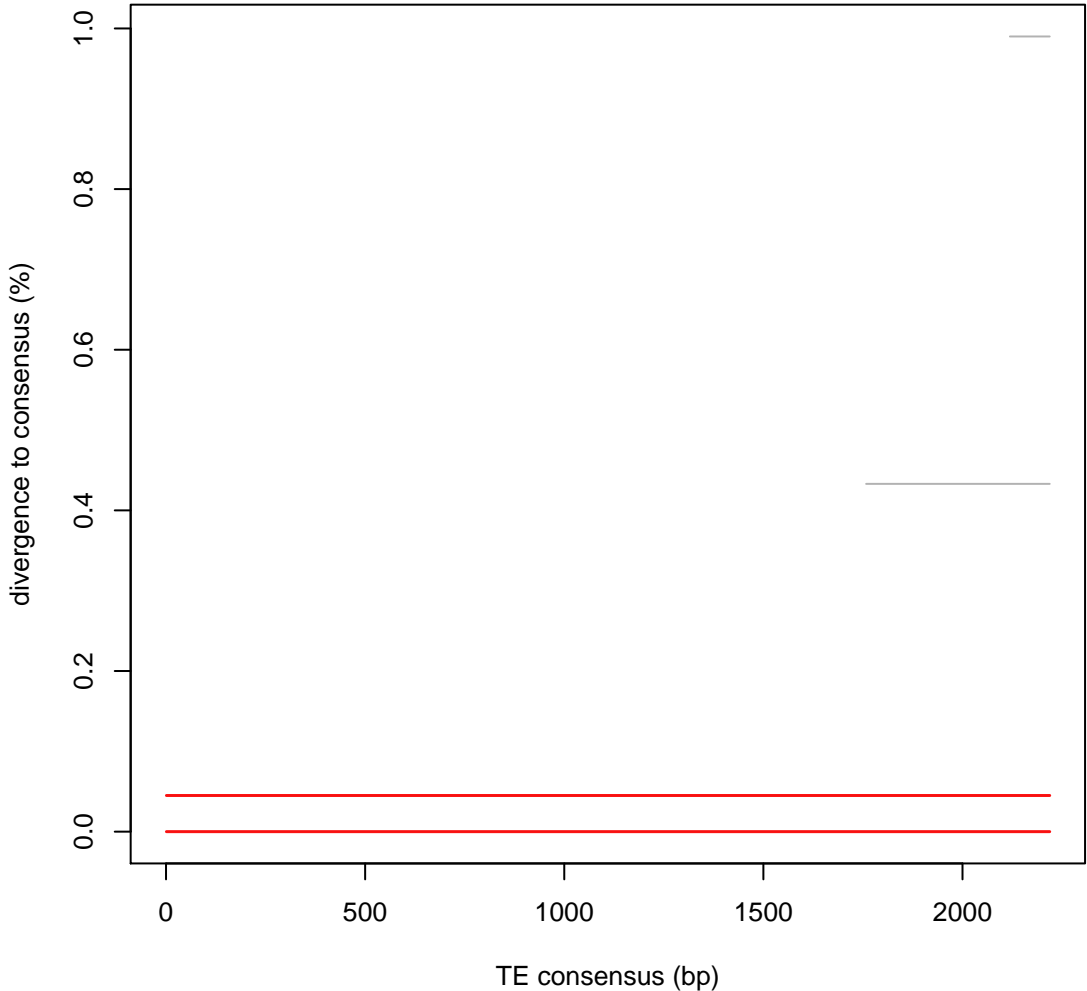
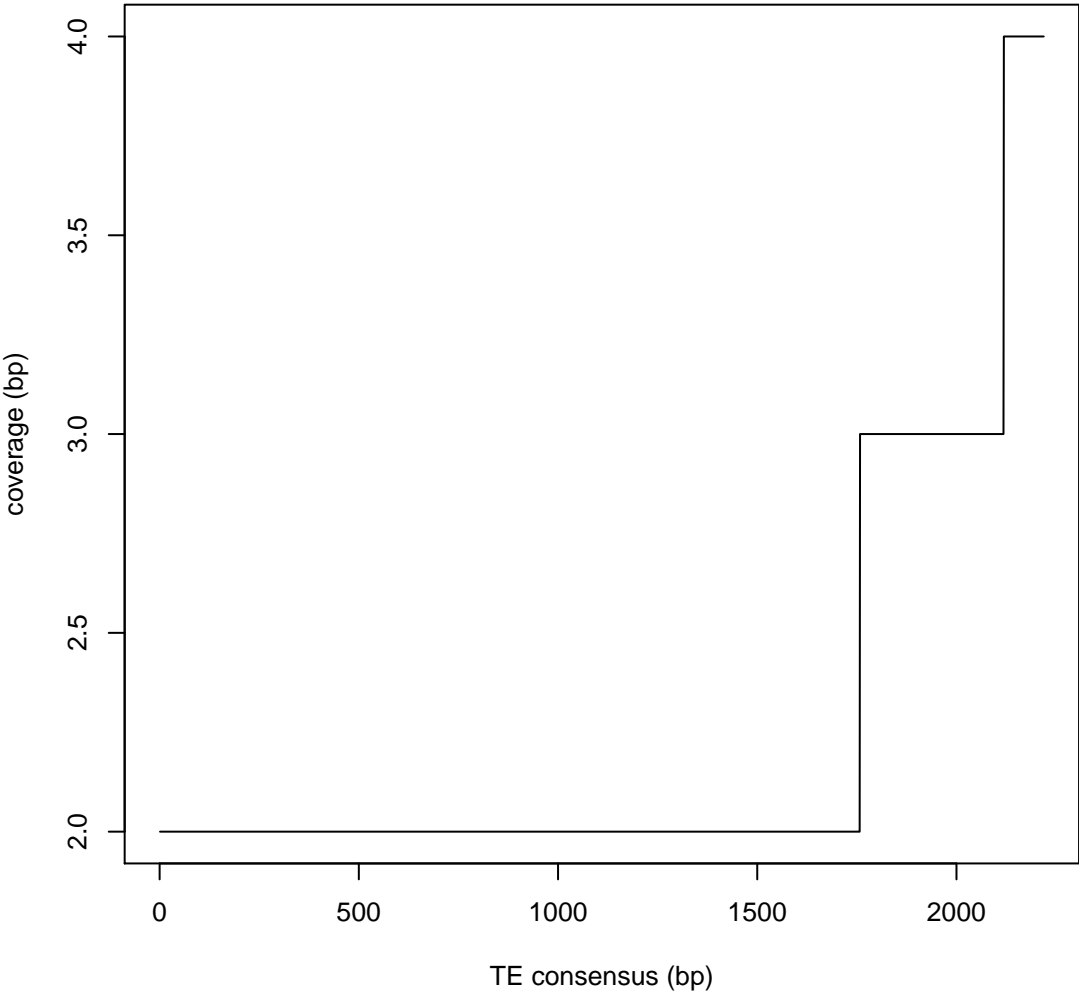


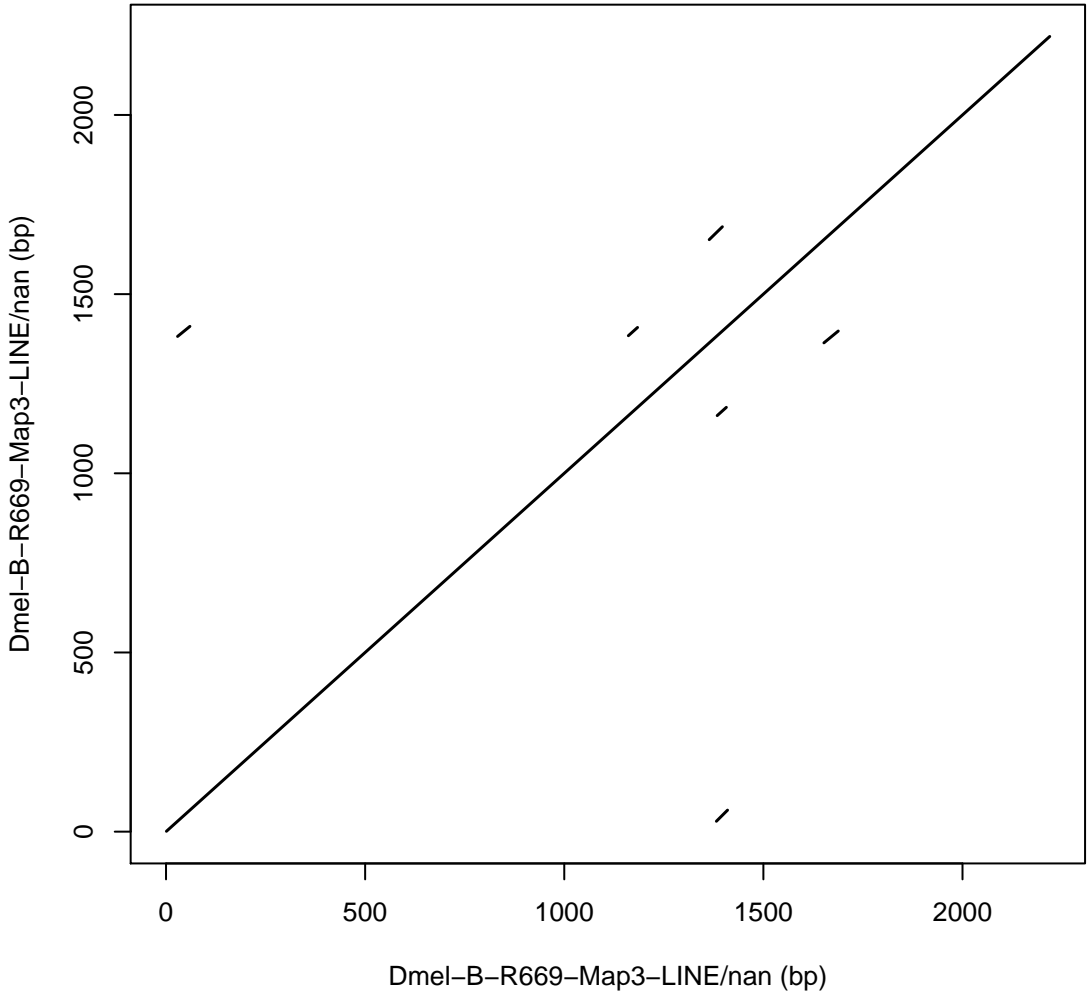
TE: Dmel-B-R669-Map3-LINE/nan
consensus size: 2219bp; fragments: 4; full length: 2 (>=1997.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

