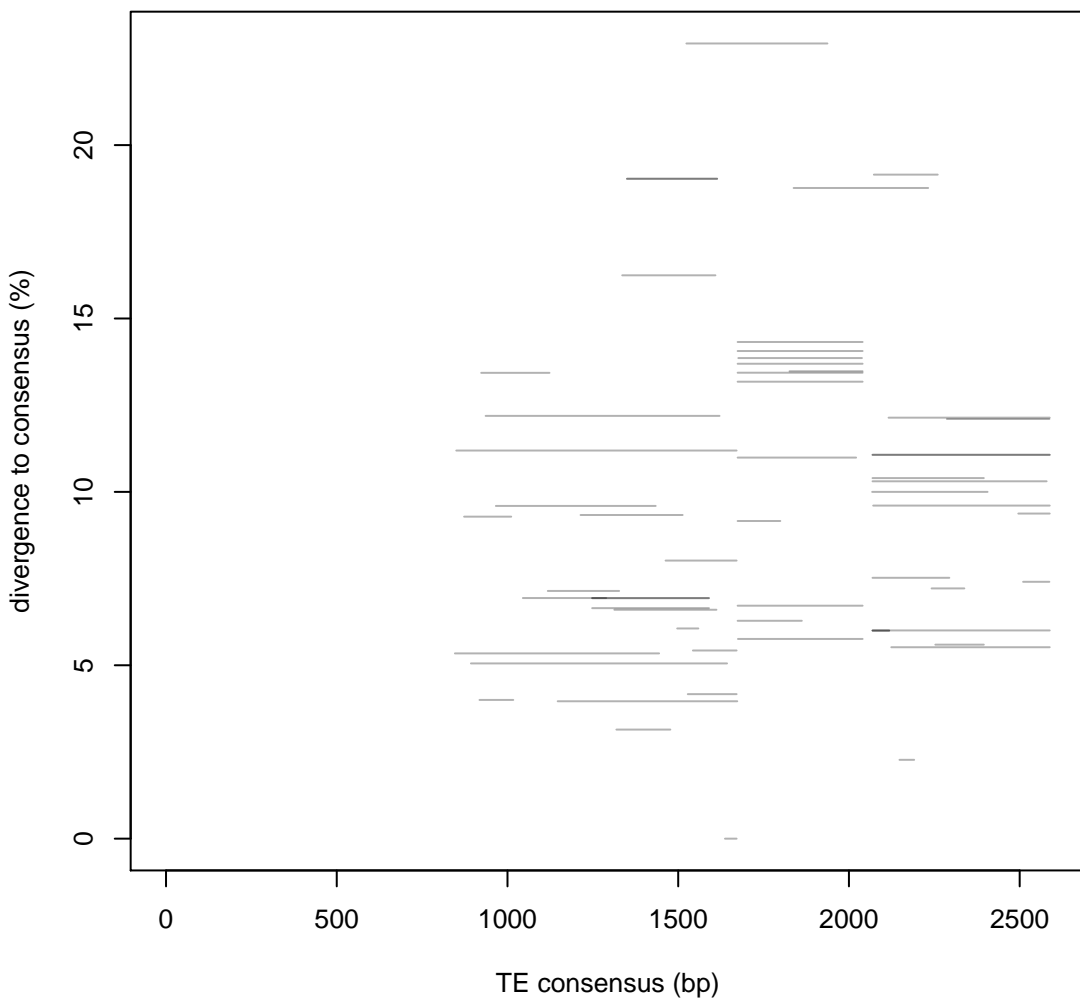
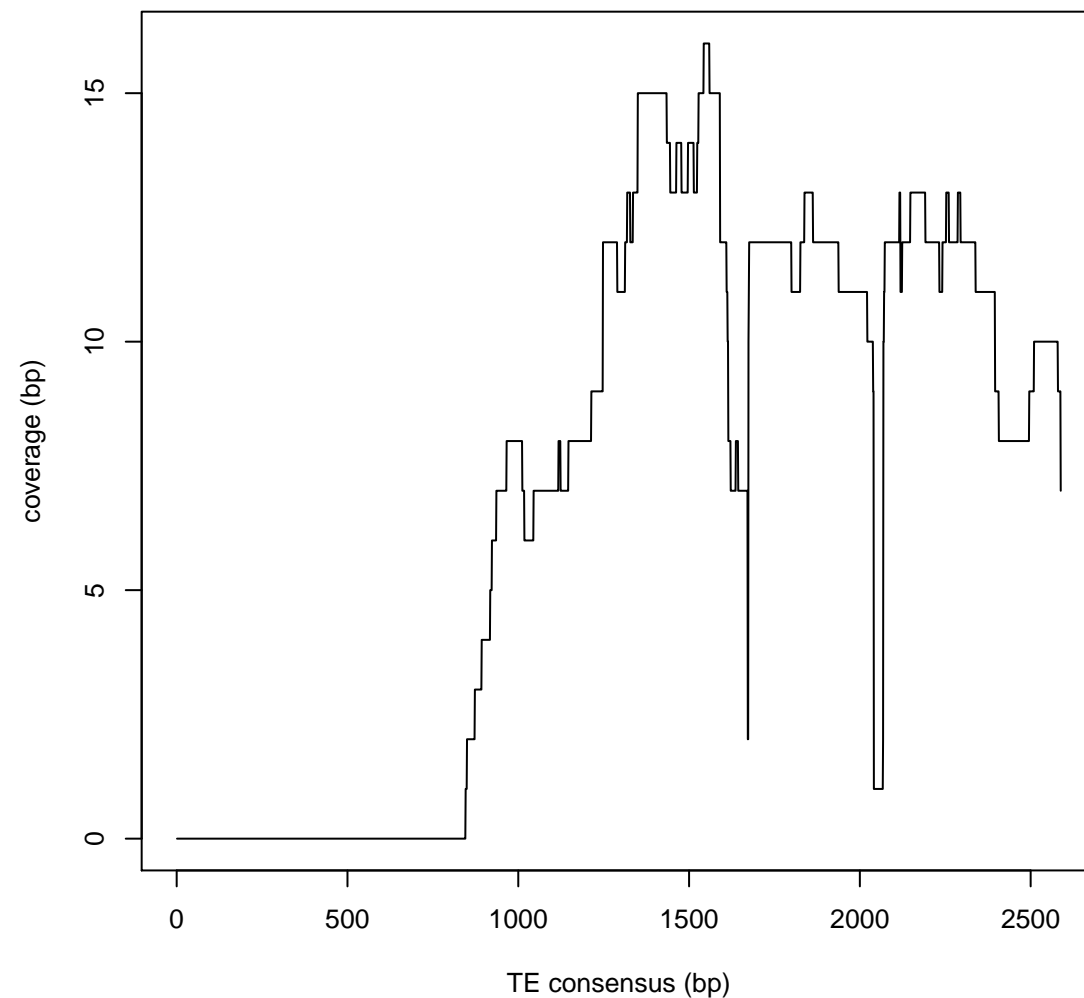


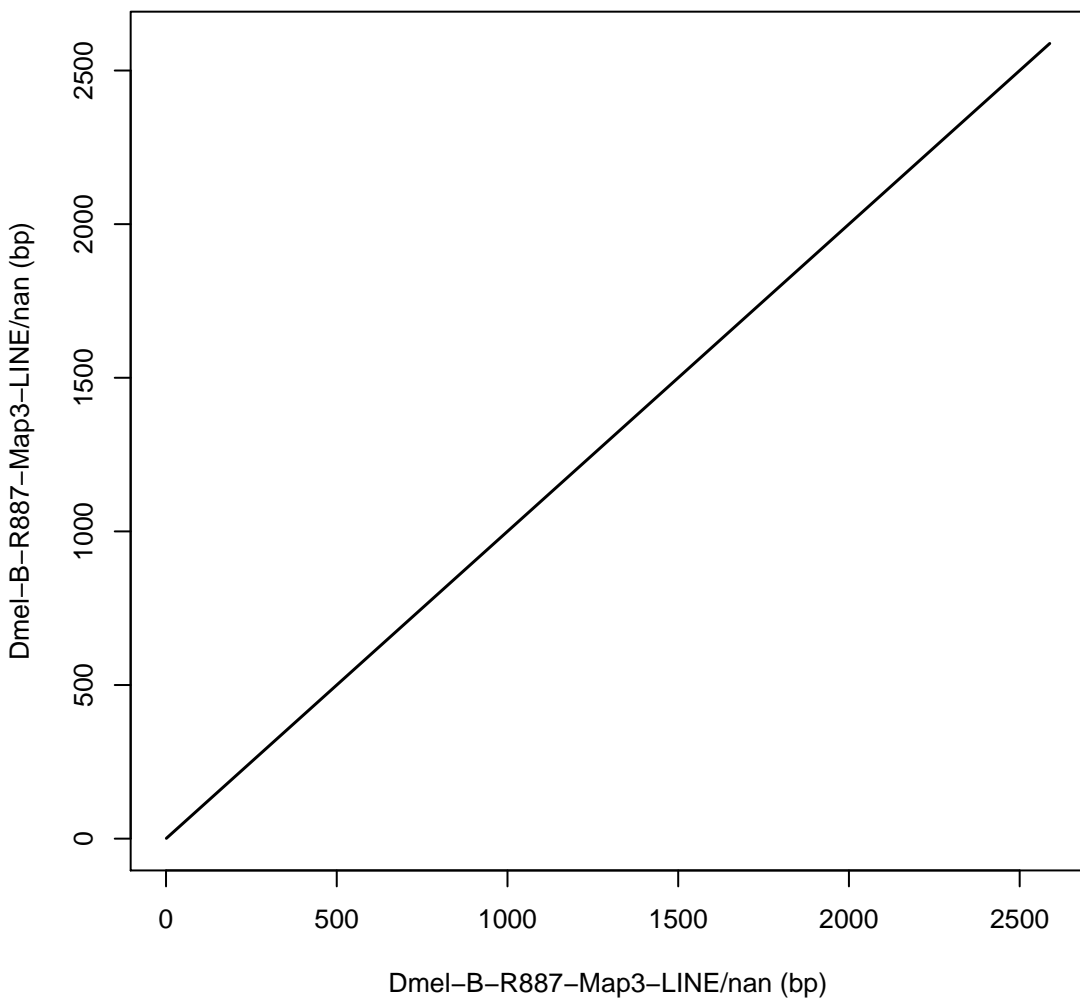
TE: Dmel-B-R887-Map3-LINE/nan
consensus size: 2588bp; fragments: 58; full length: 0 (>=2329.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

