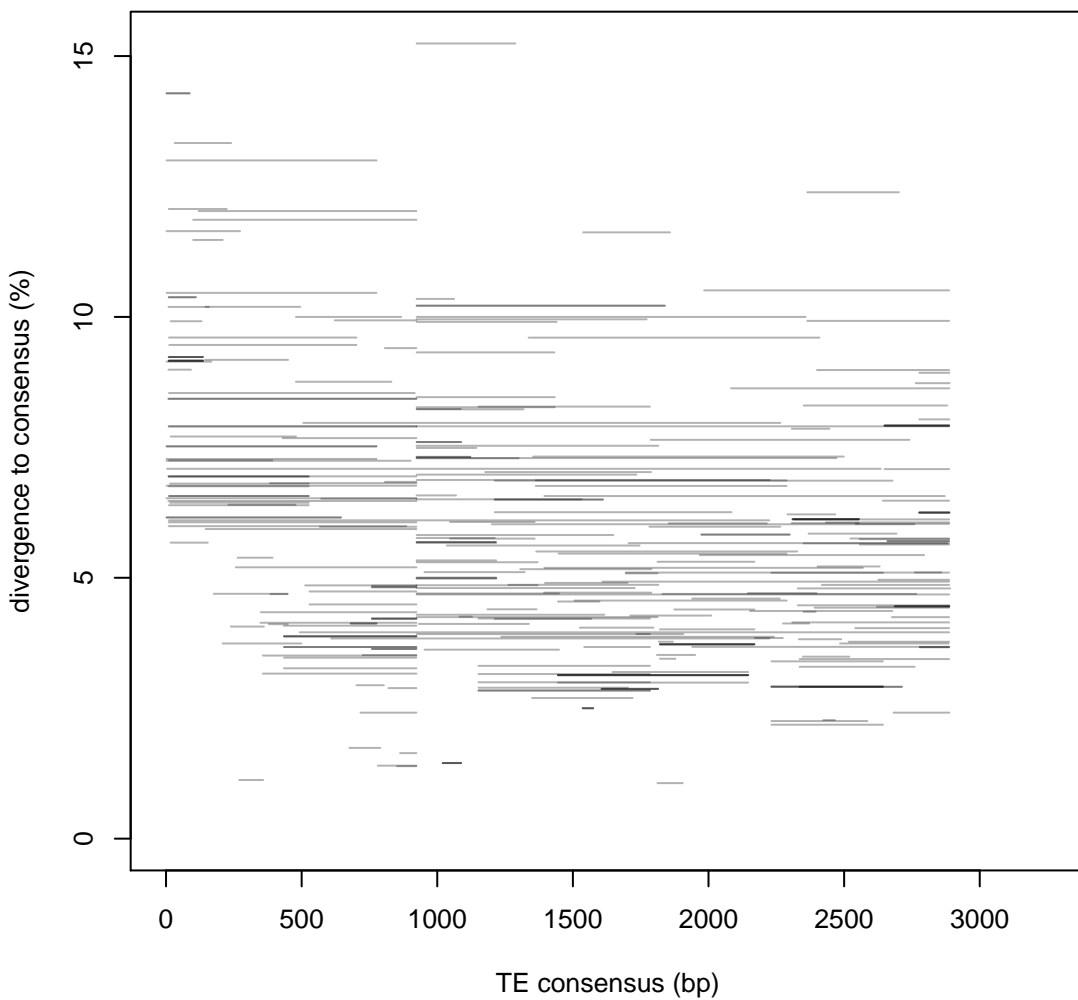
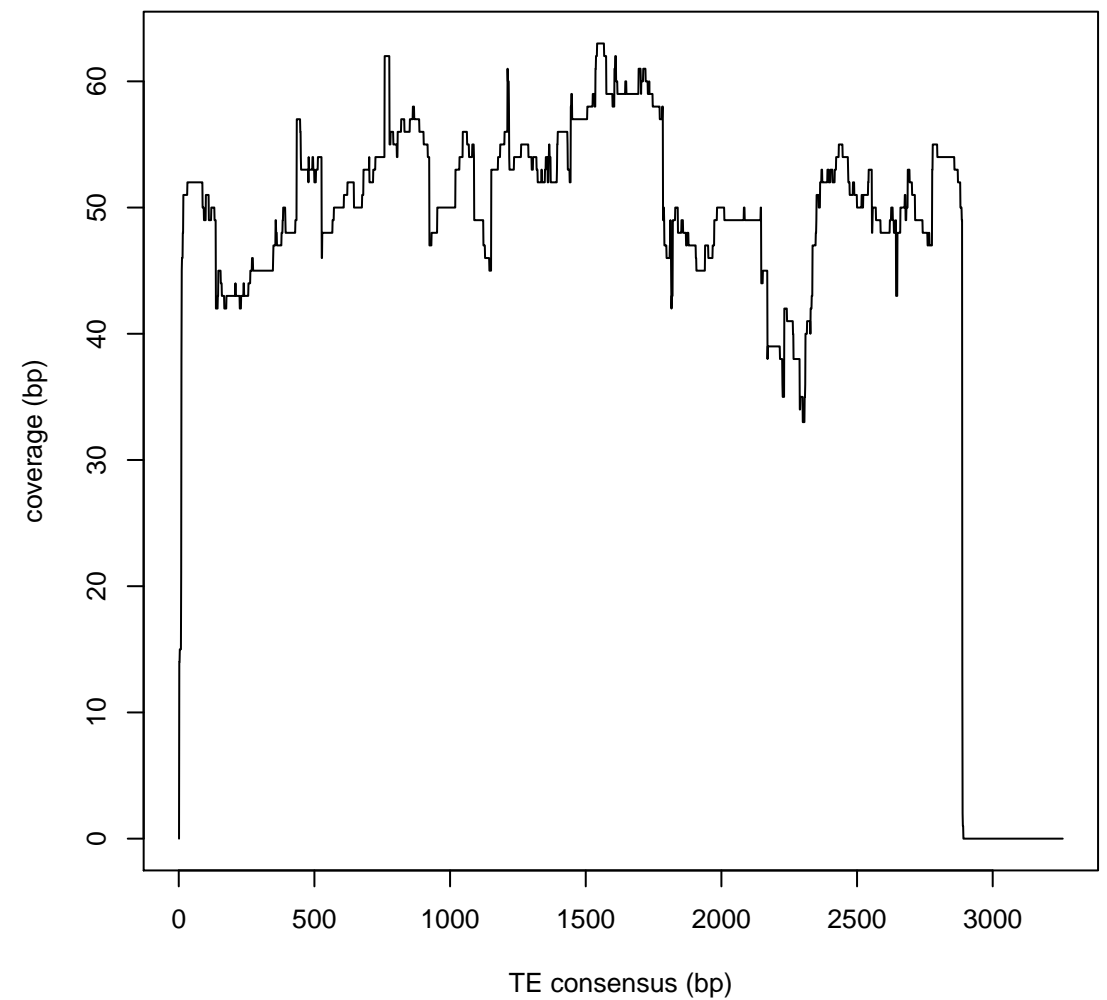


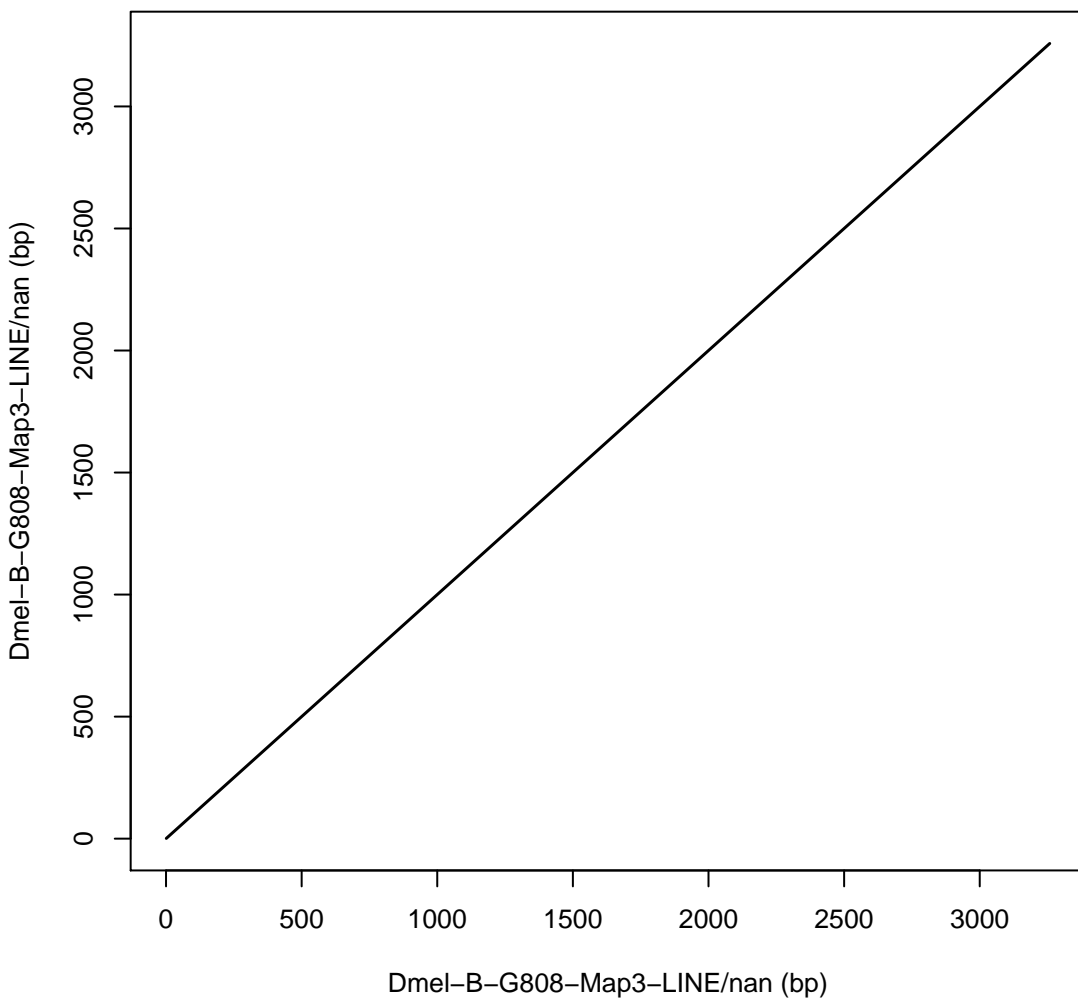
TE: Dmel-B-G808-Map3-LINE/nan
consensus size: 3258bp; fragments: 311; full length: 0 (>=2932.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

