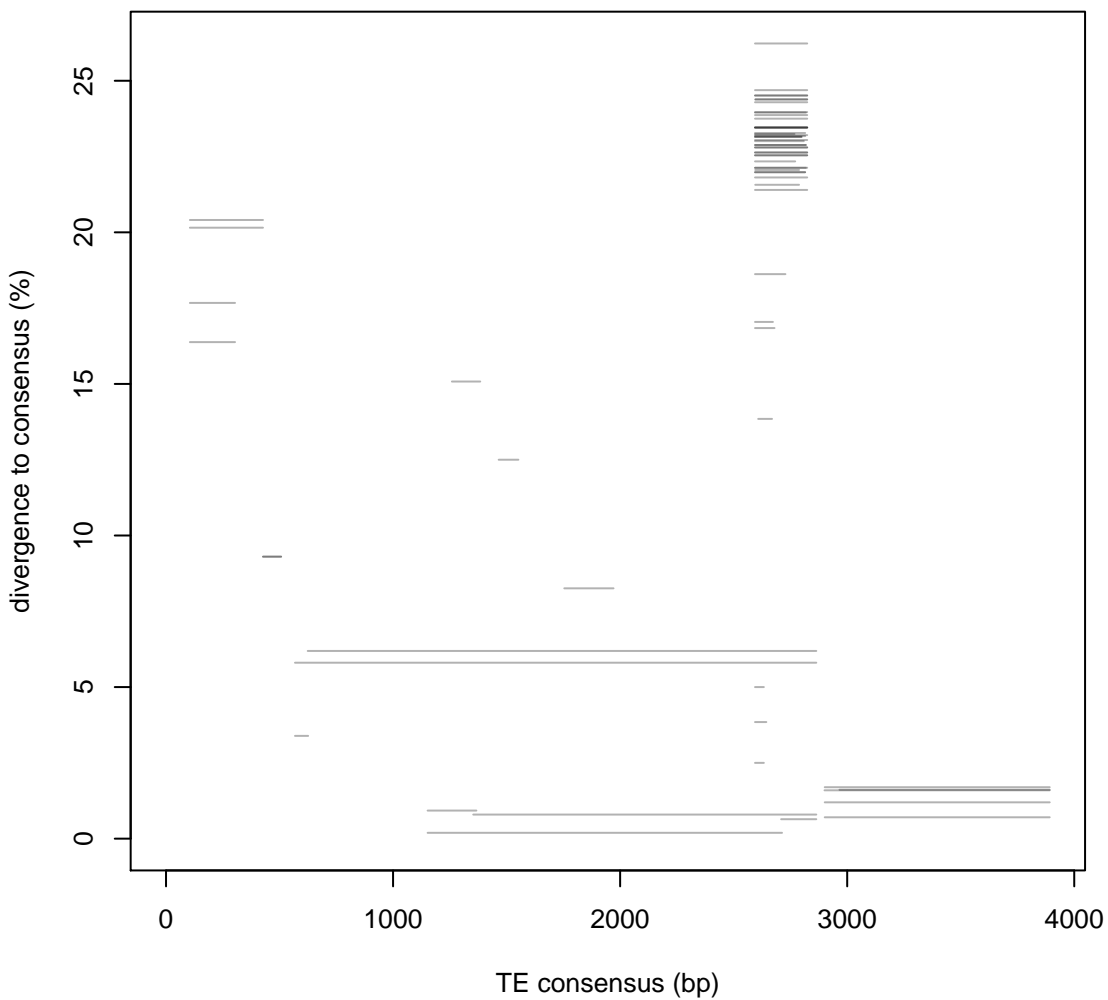
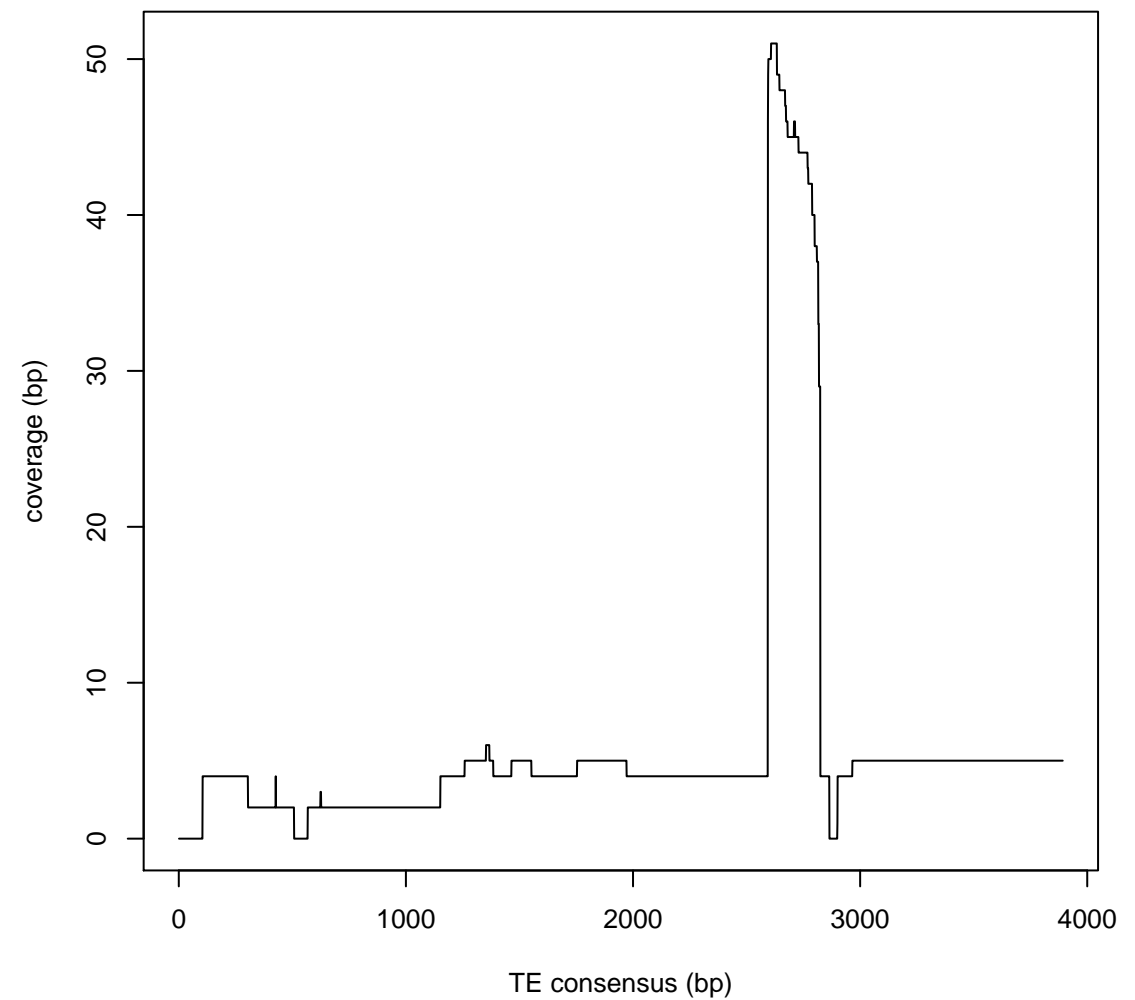


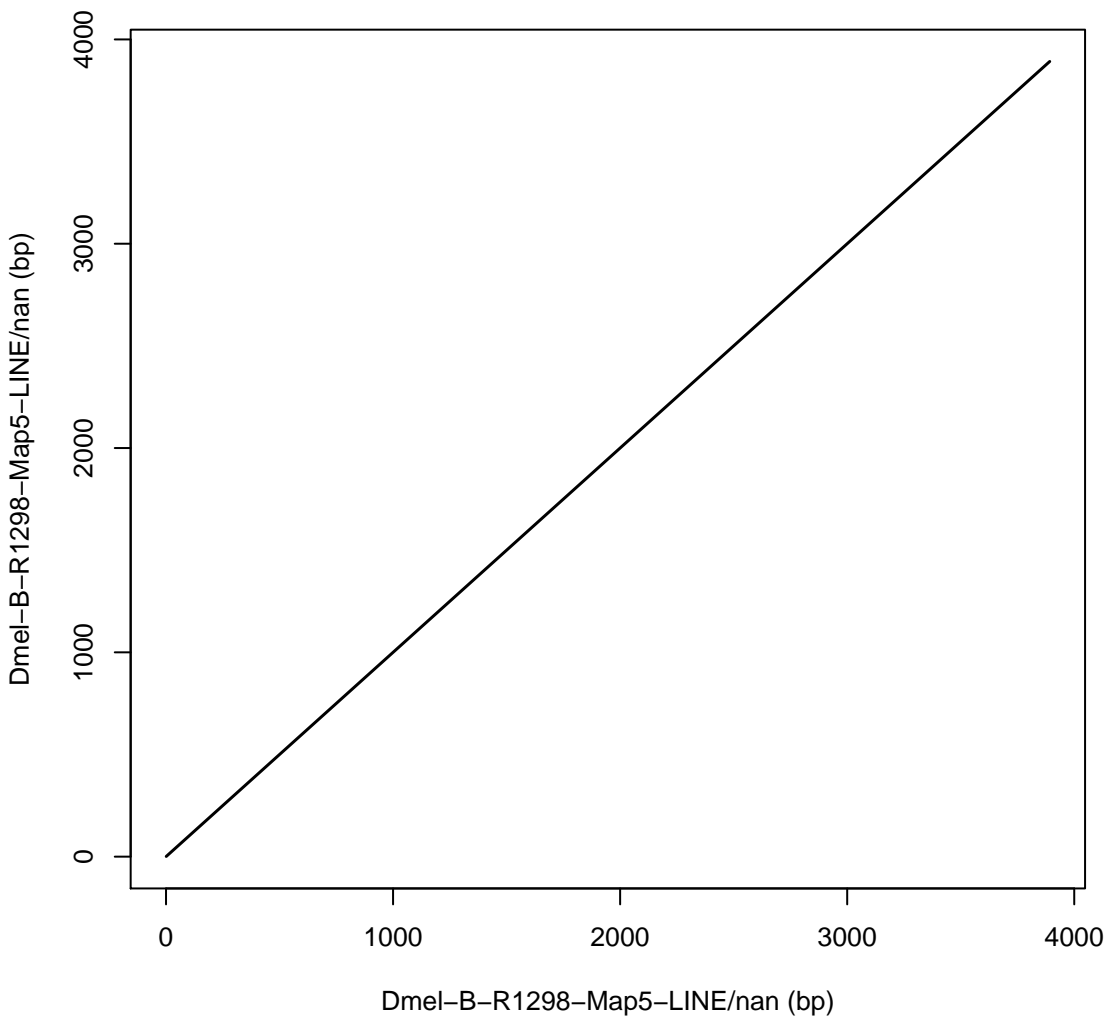
TE: Dmel-B-R1298-Map5-LINE/nan
consensus size: 3892bp; fragments: 68; full length: 0 (>=3502.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

