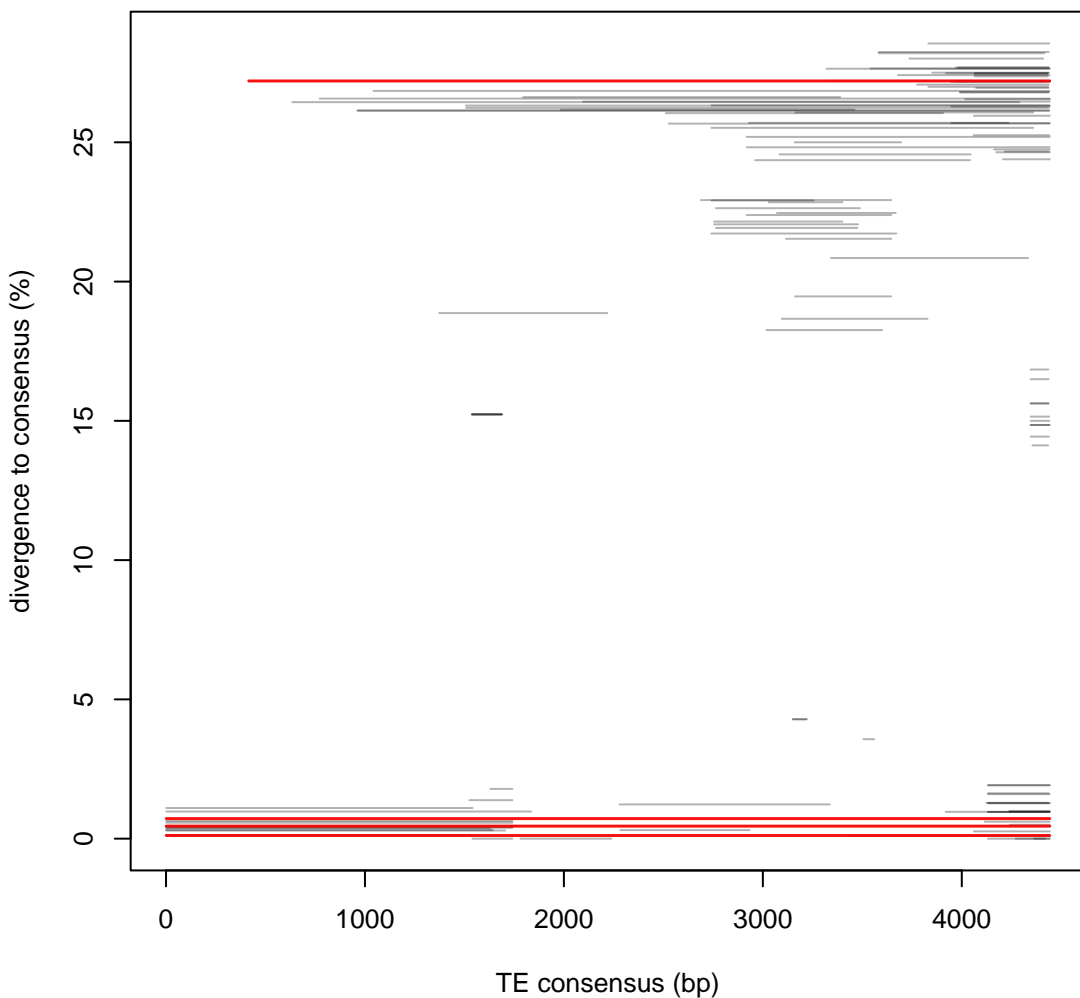
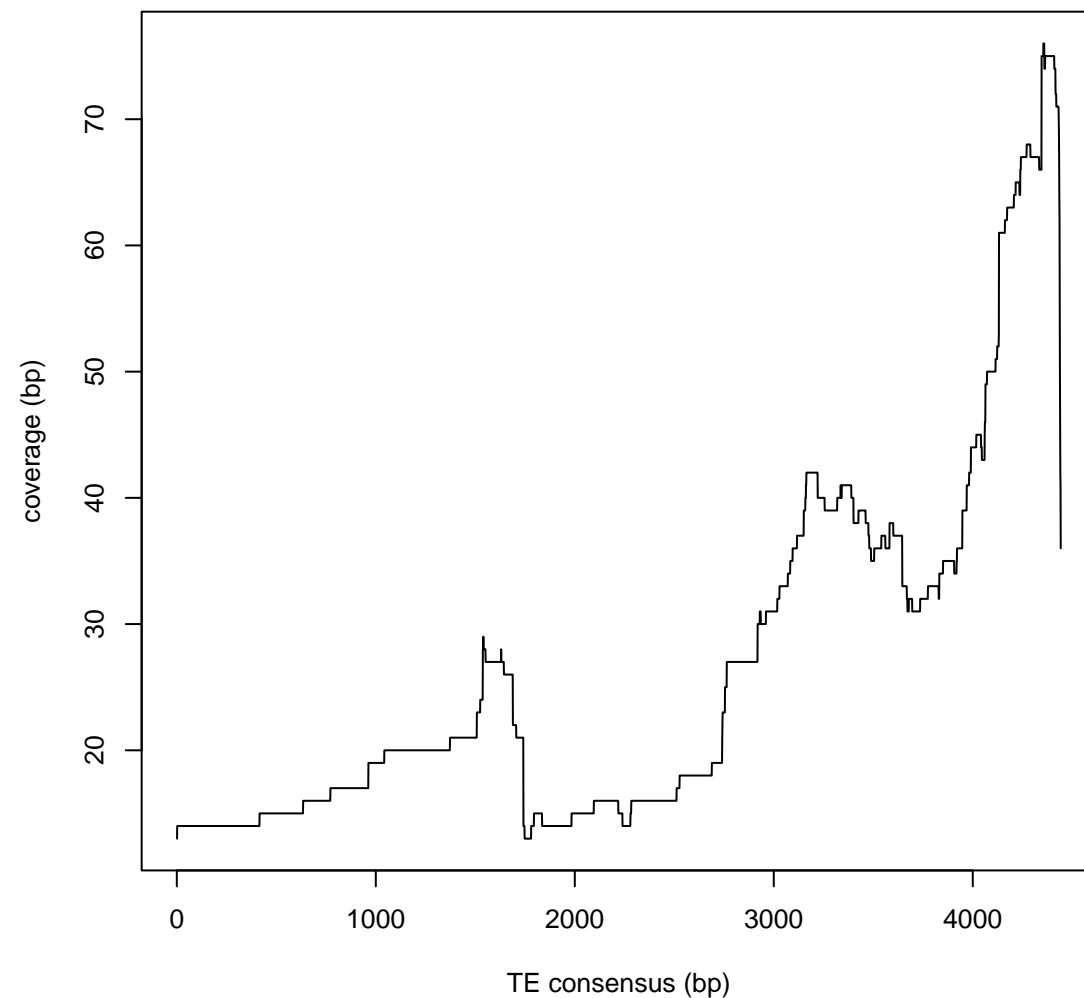


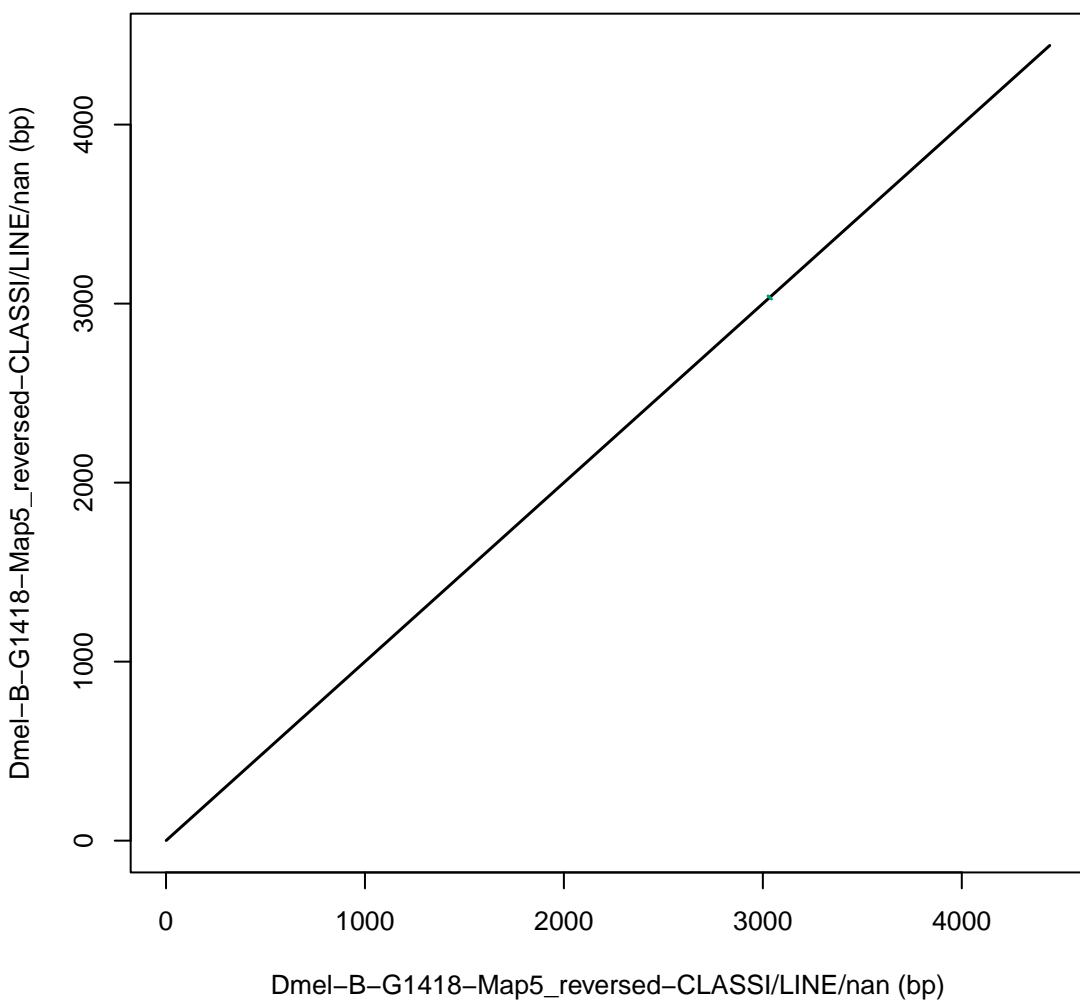
TE: Dmel-B-G1418-Map5_reversed-CLASSI/LINE/nan
consensus size: 4442bp; fragments: 125; full length: 4 (>=3997.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

