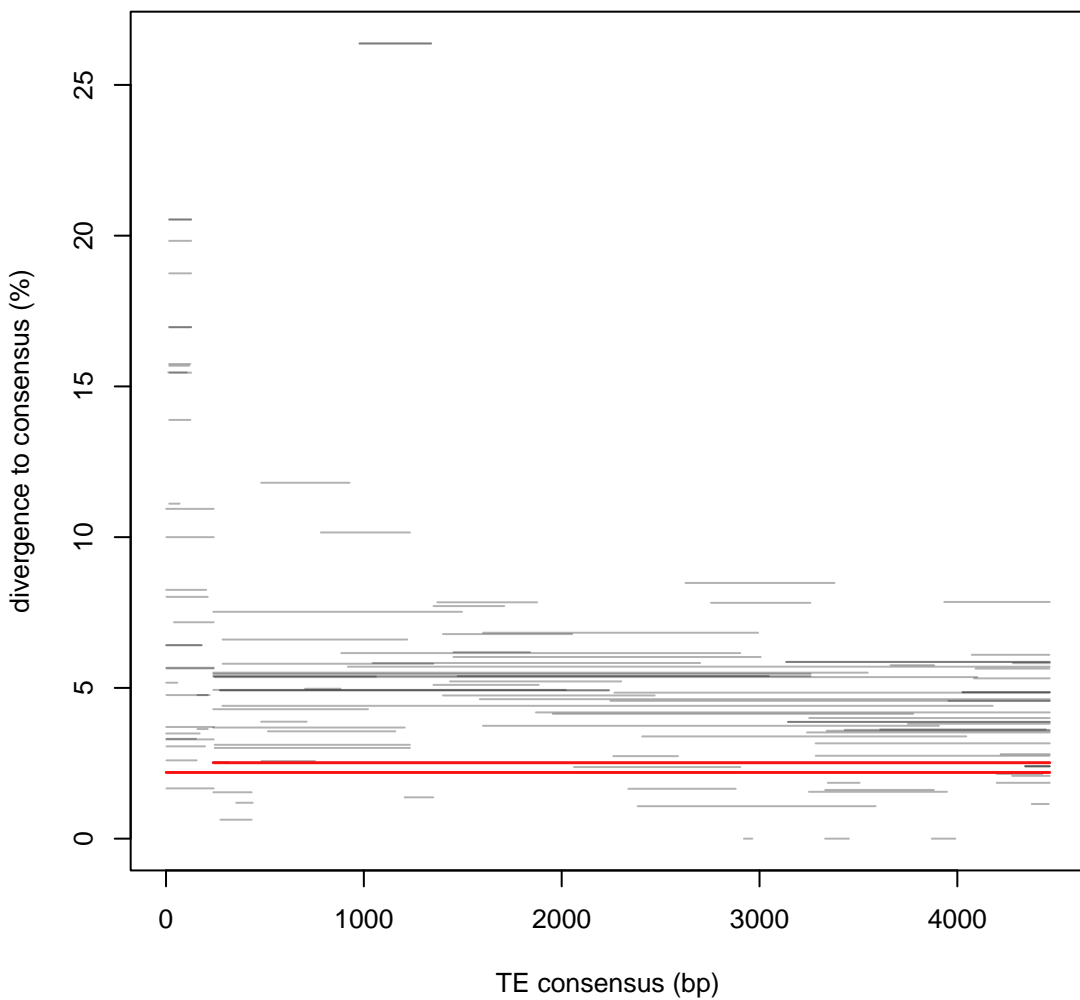
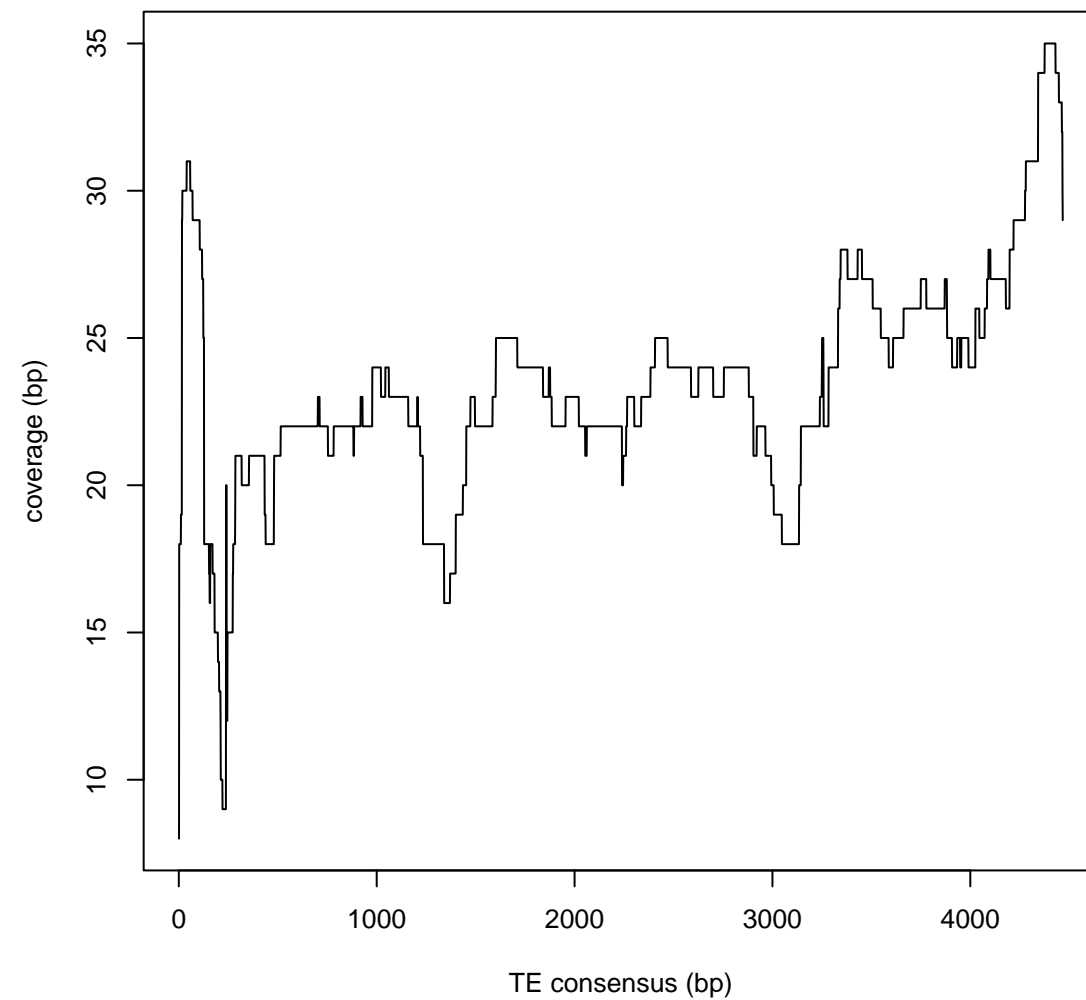


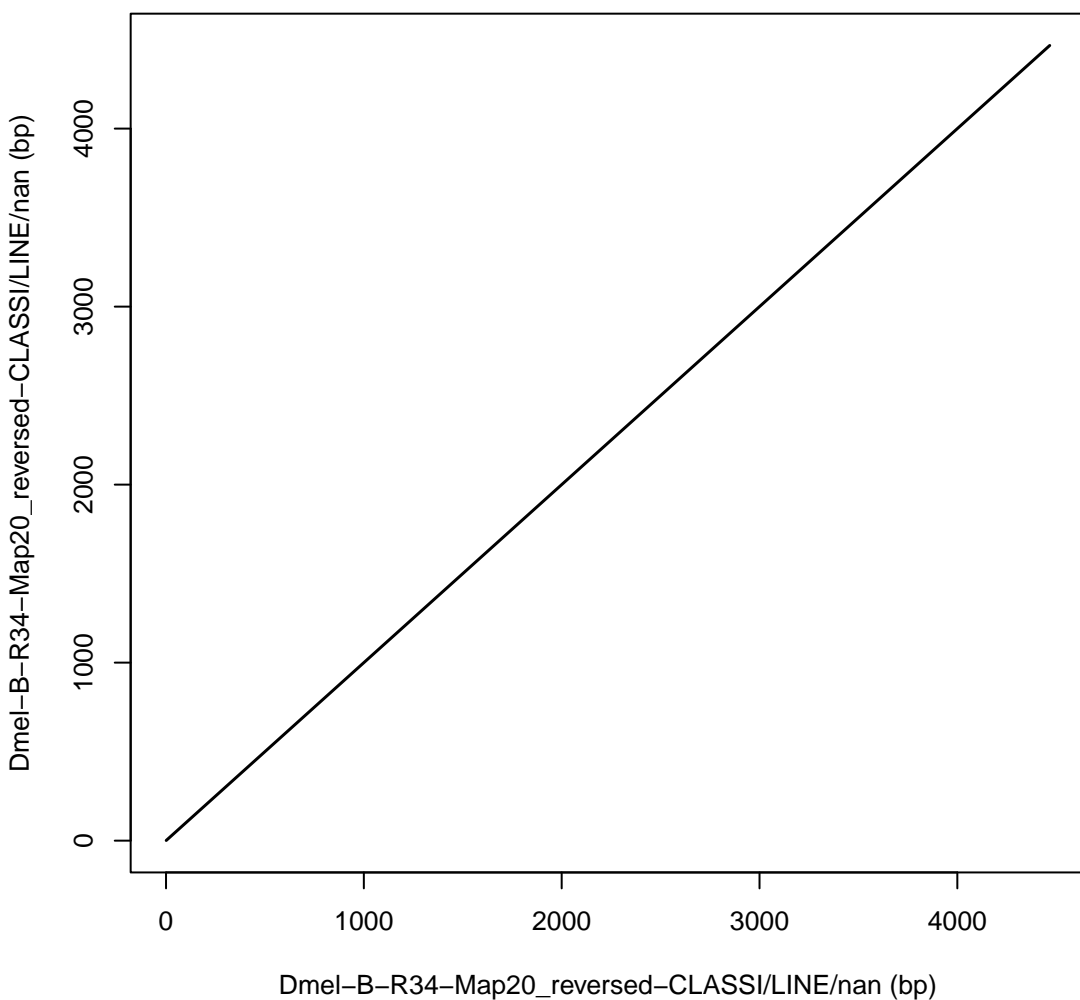
TE: Dmel-B-R34-Map20_reversed-CLASSI/LINE/nan
consensus size: 4467bp; fragments: 124; full length: 2 (>=4020.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

