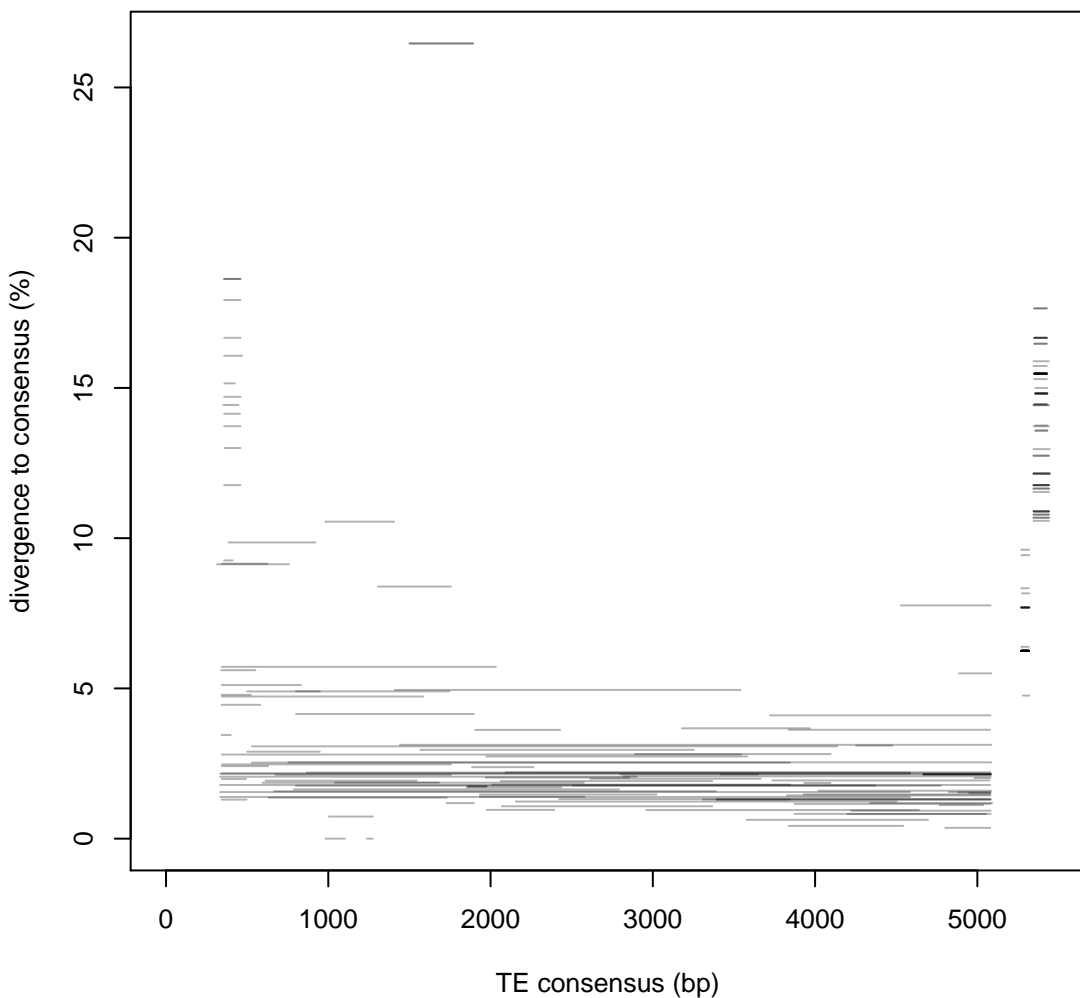
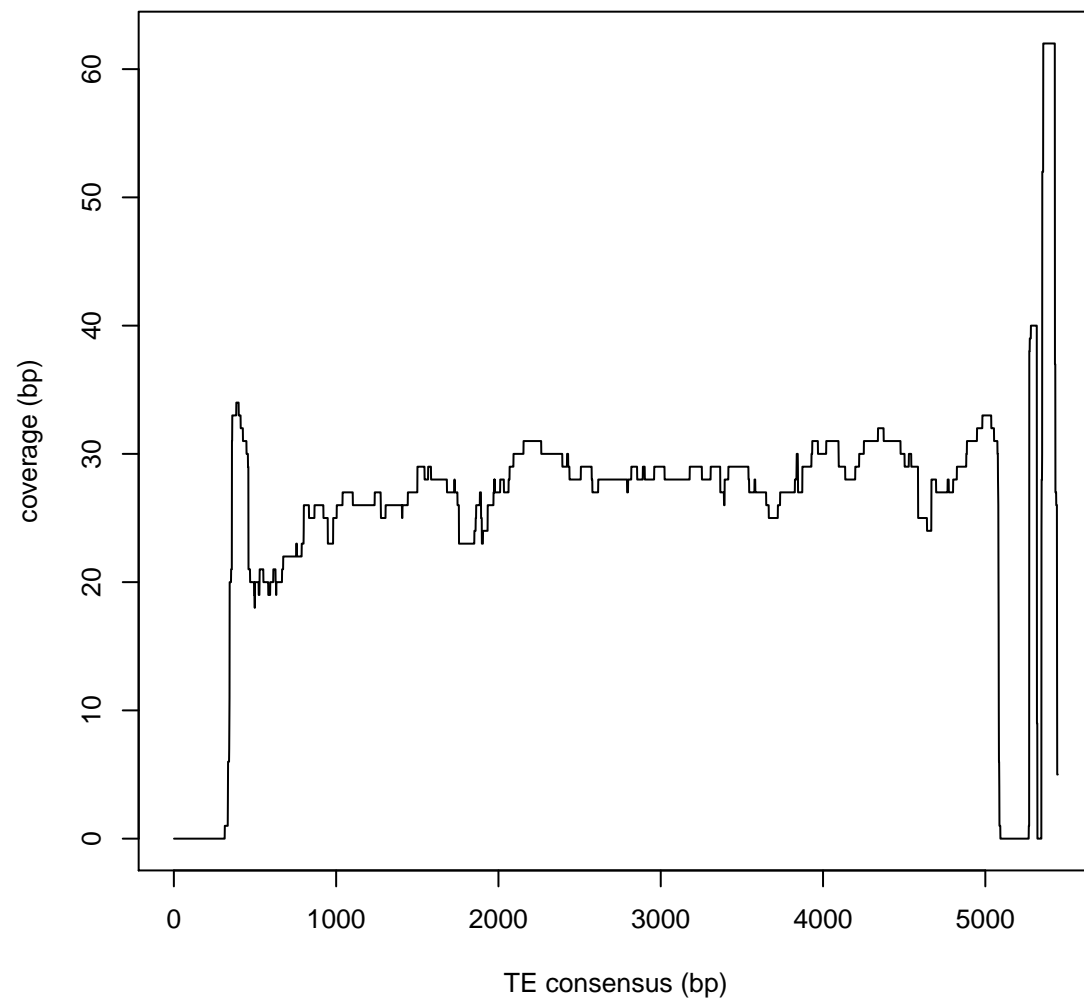


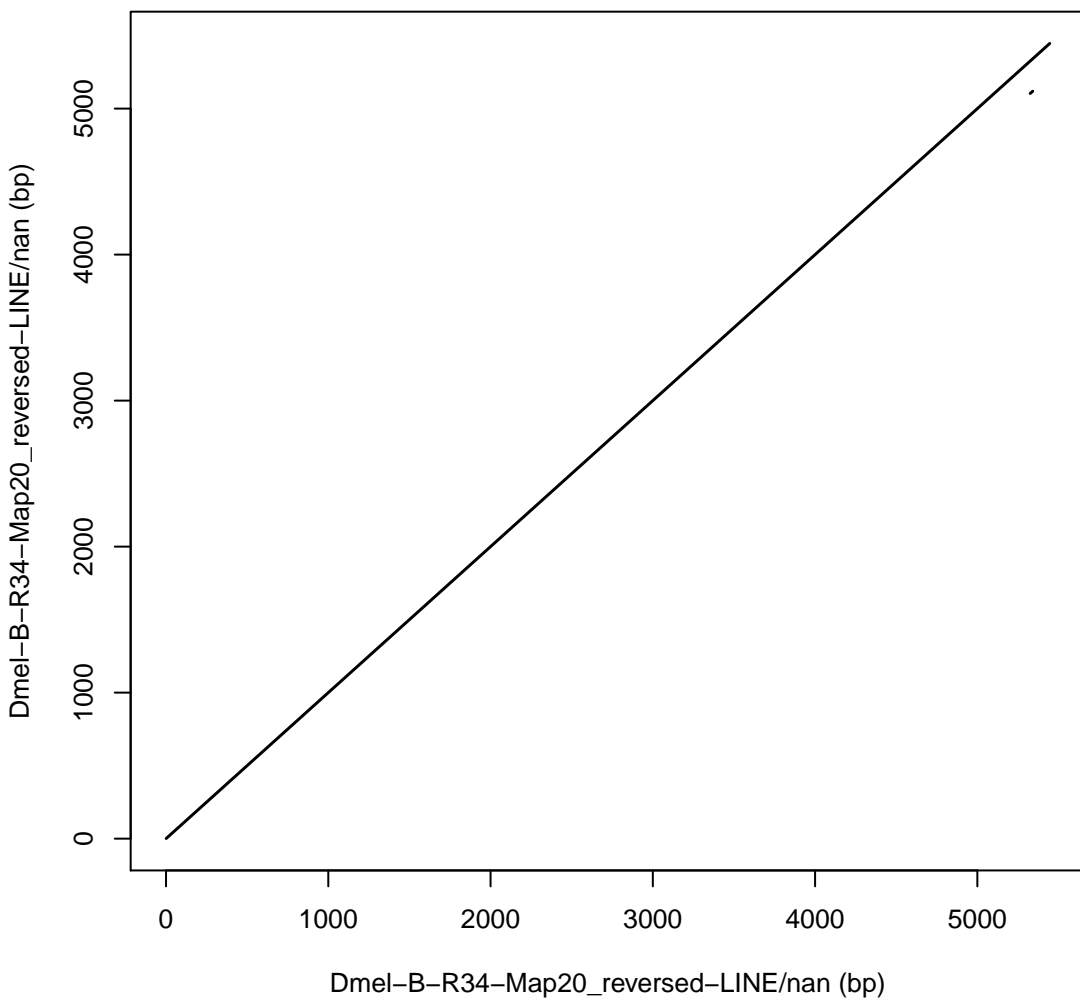
TE: Dmel-B-R34-Map20_reversed-LINE/nan
consensus size: 5446bp; fragments: 217; full length: 0 (>=4901.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

