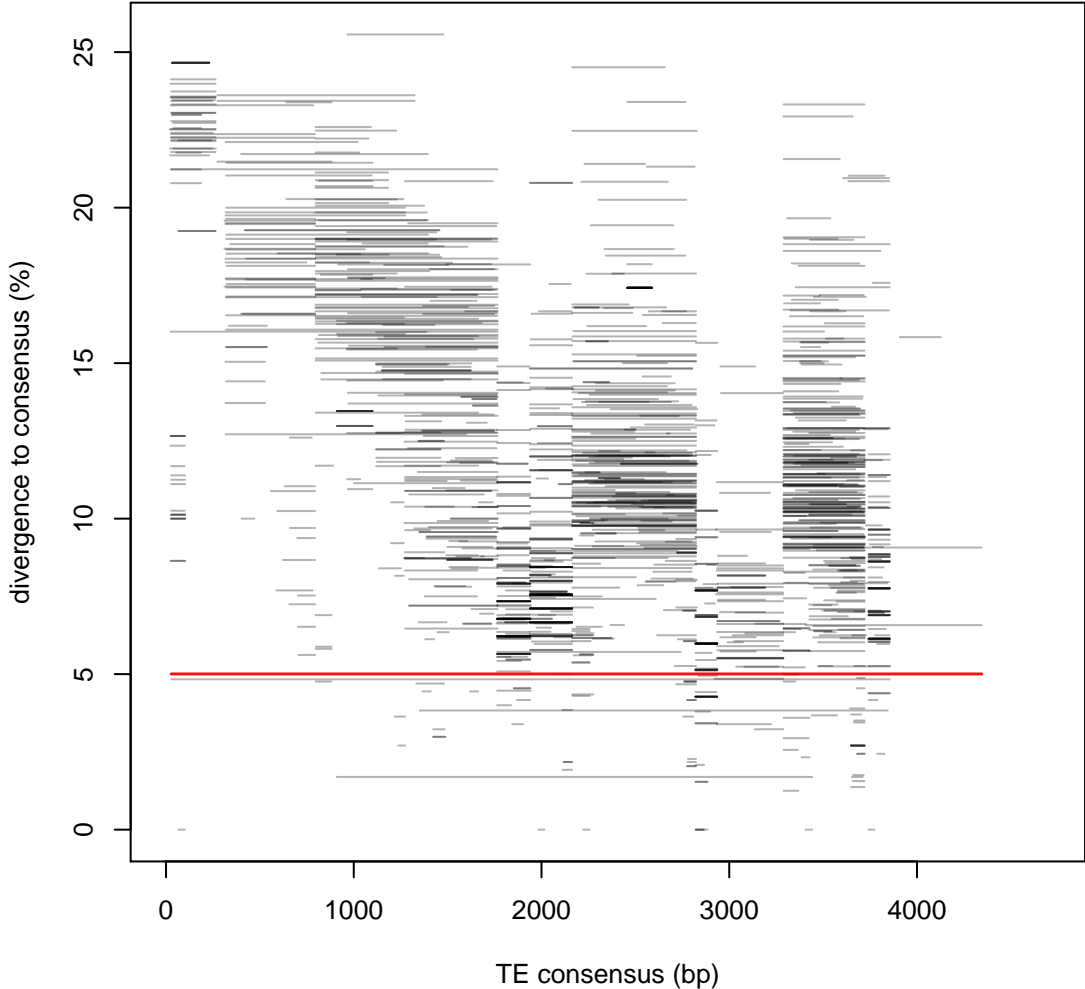
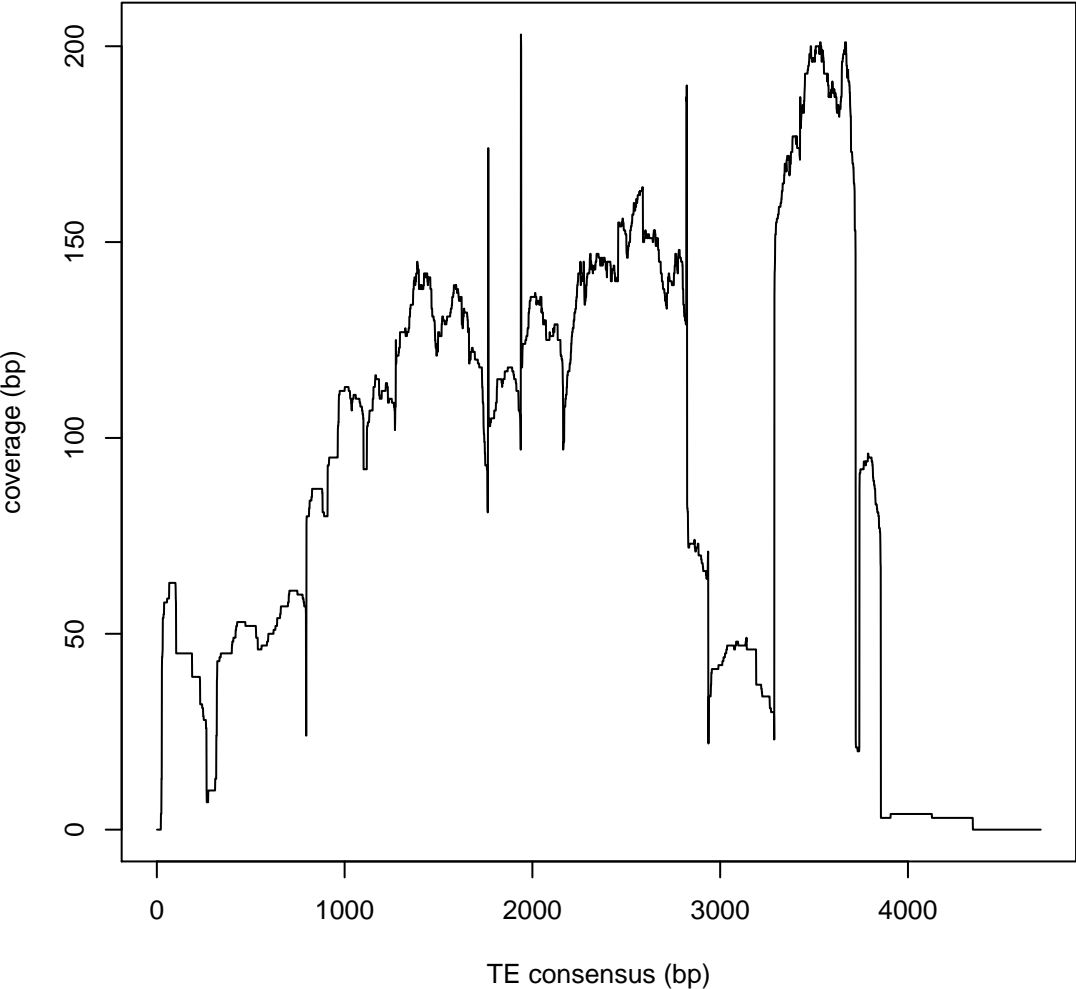


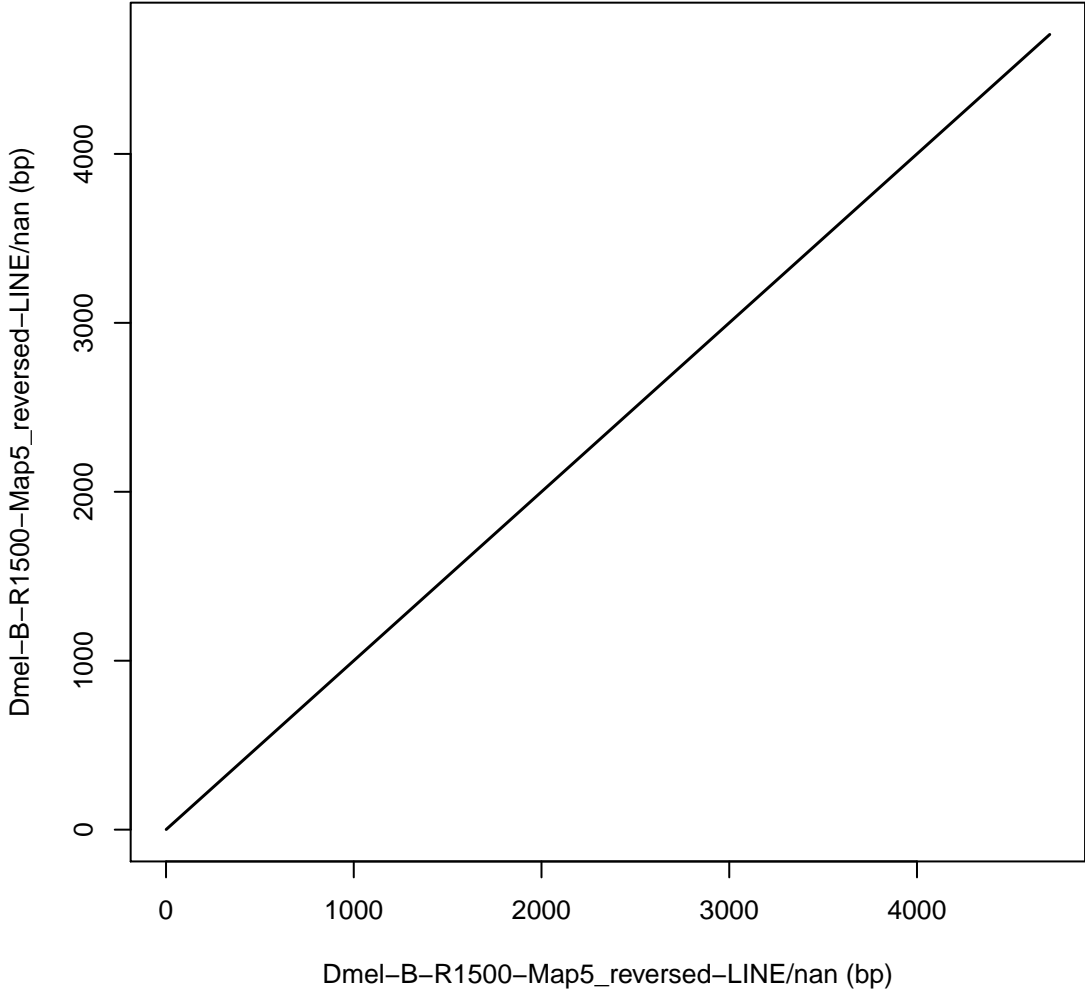
TE: Dmel-B-R1500-Map5_reversed-LINE/nan
consensus size: 4707bp; fragments: 1363; full length: 1 (>=4236.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

