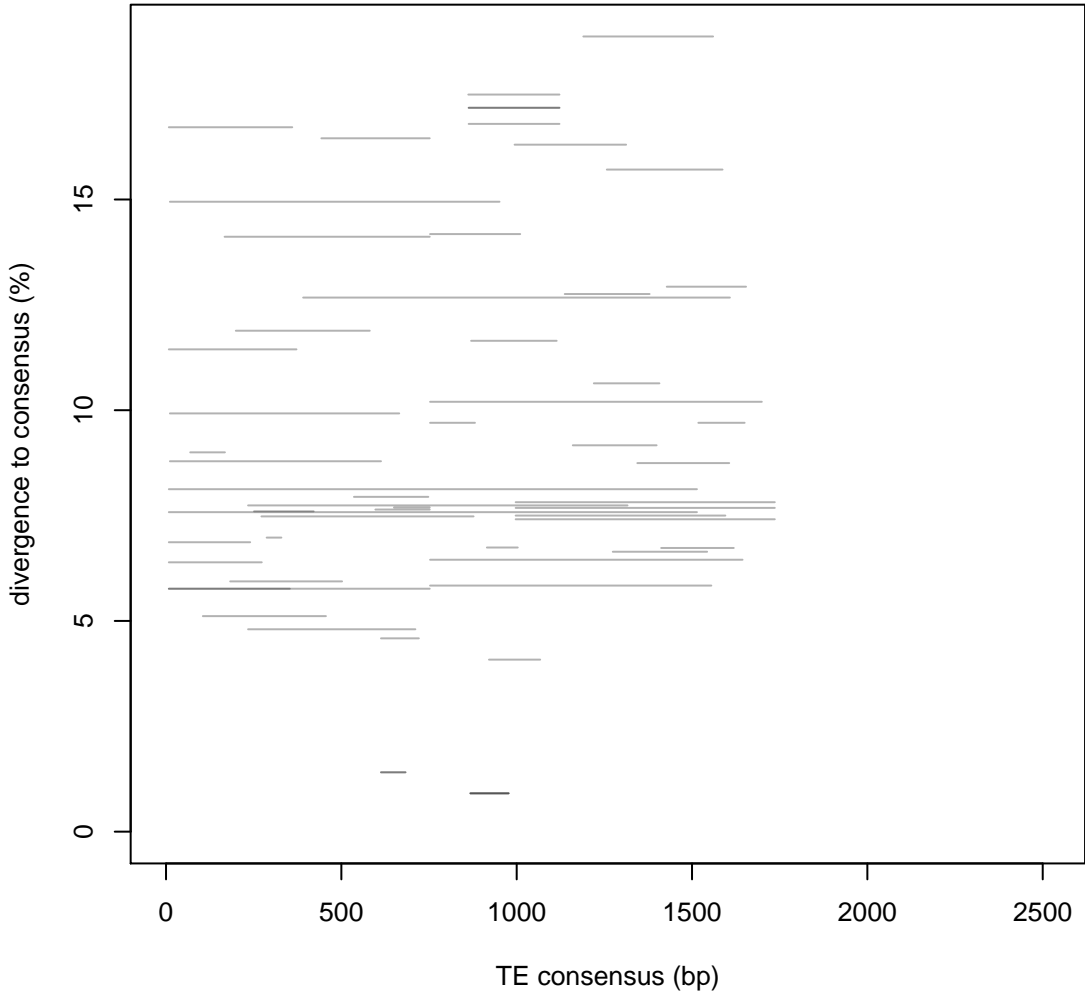
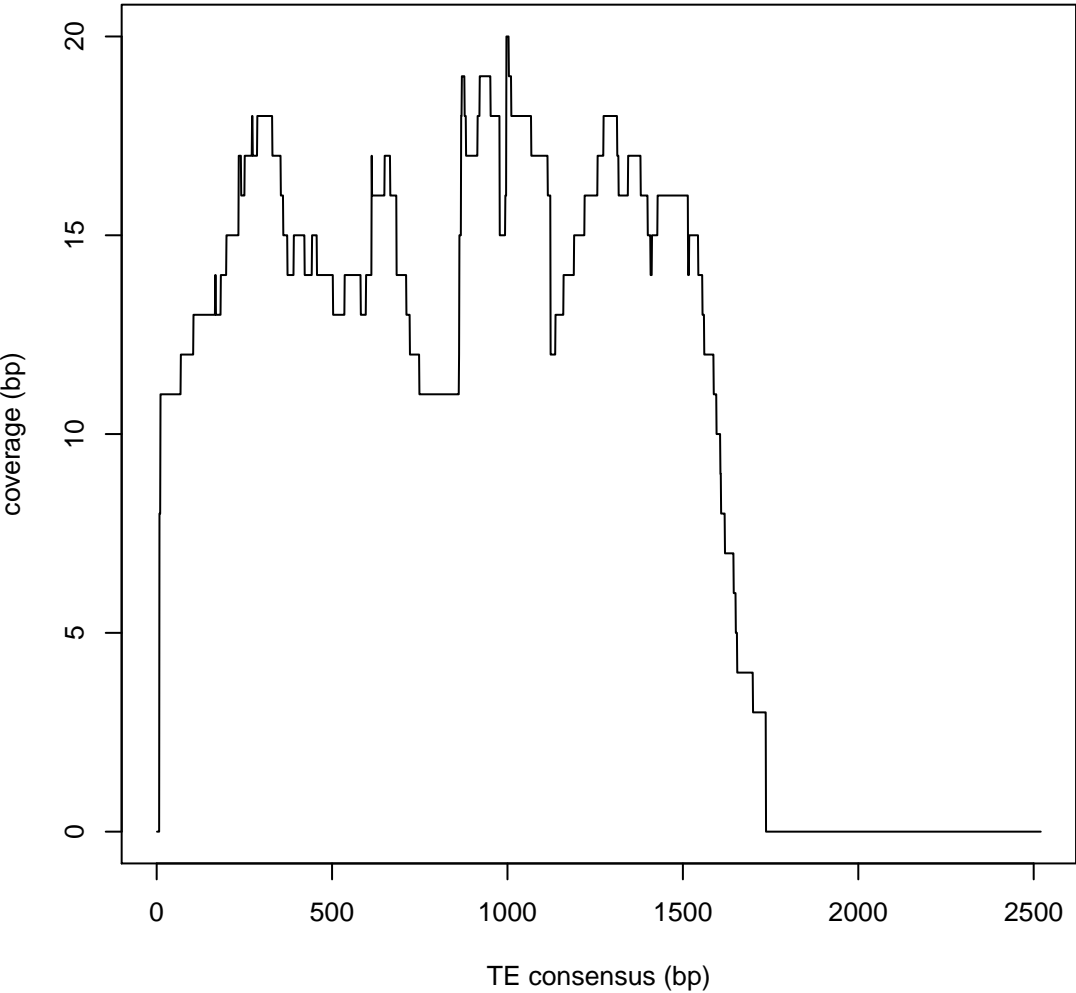


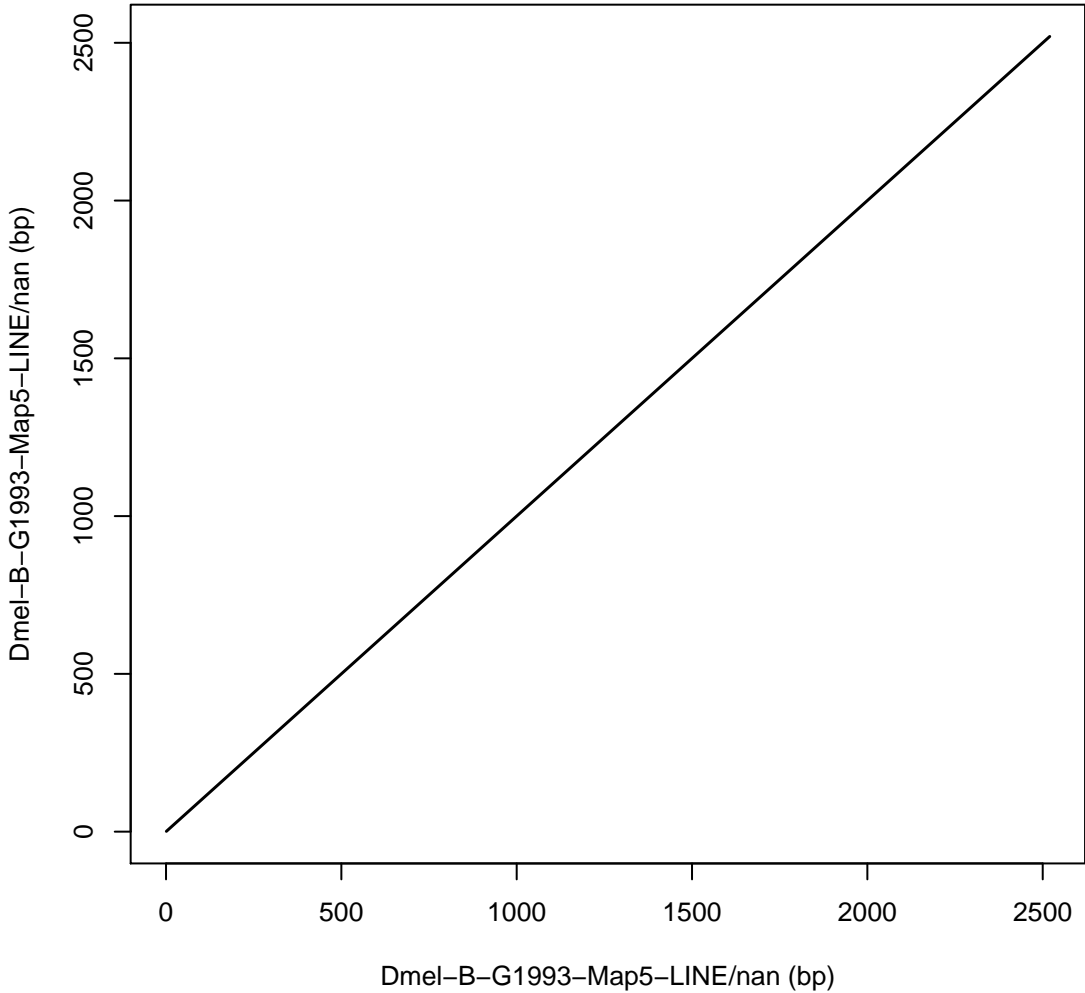
TE: Dmel-B-G1993-Map5-LINE/nan
consensus size: 2520bp; fragments: 59; full length: 0 (>=2268bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

