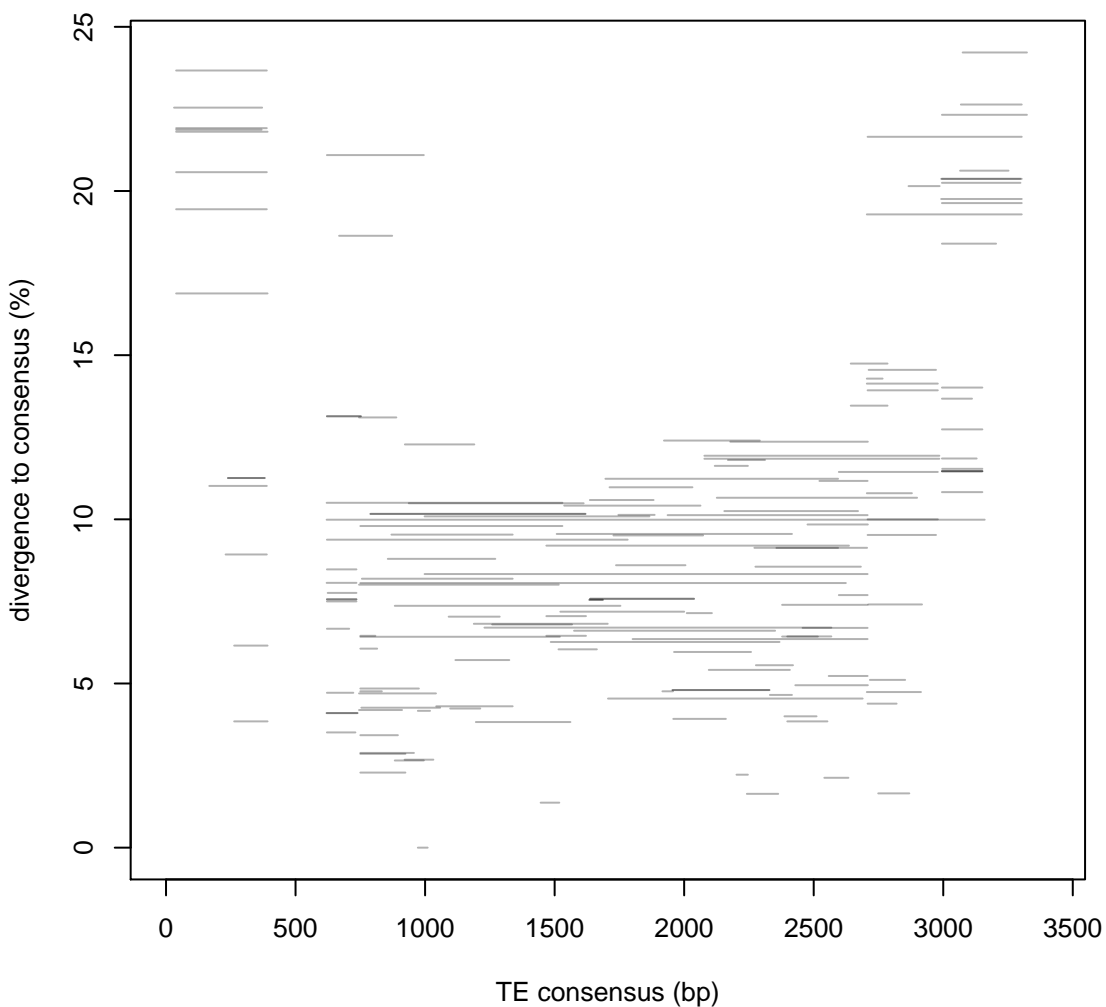
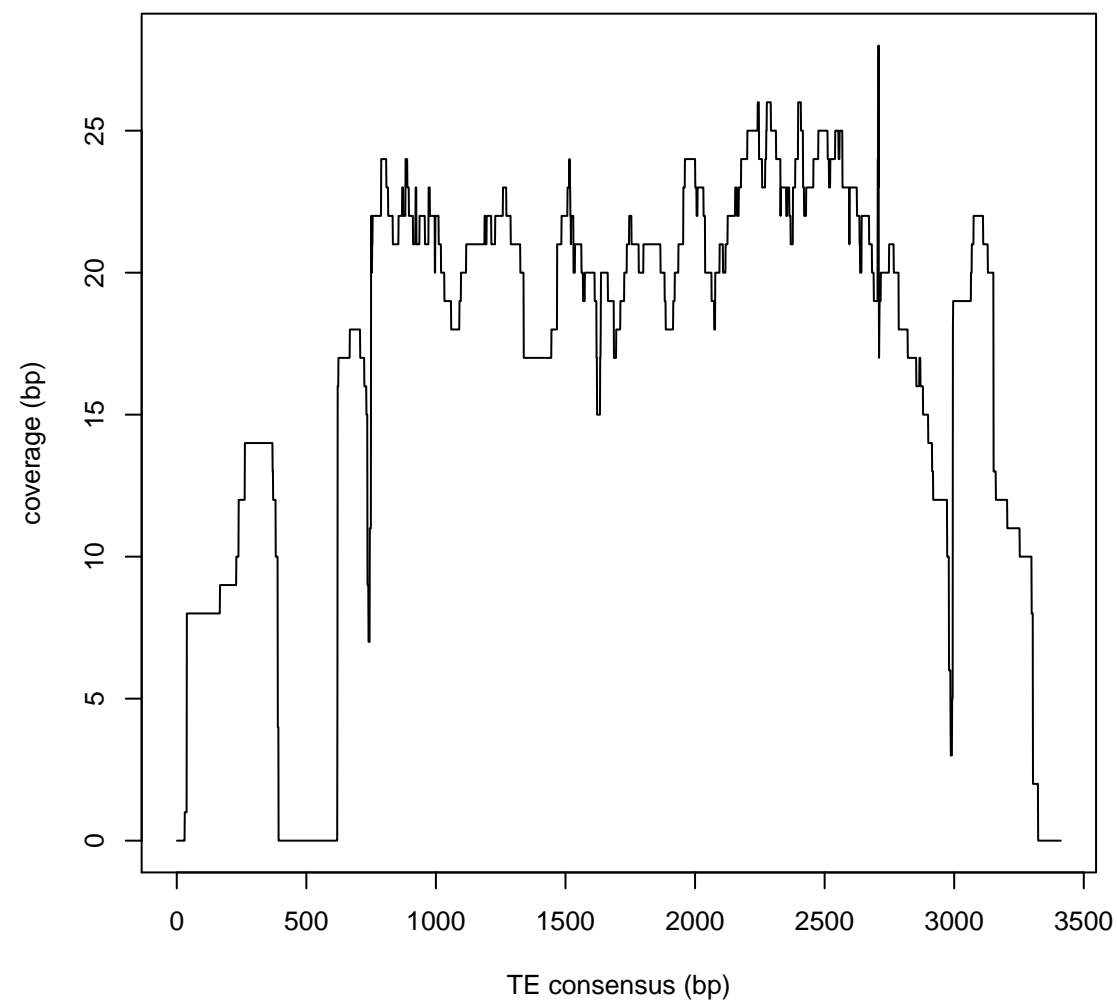


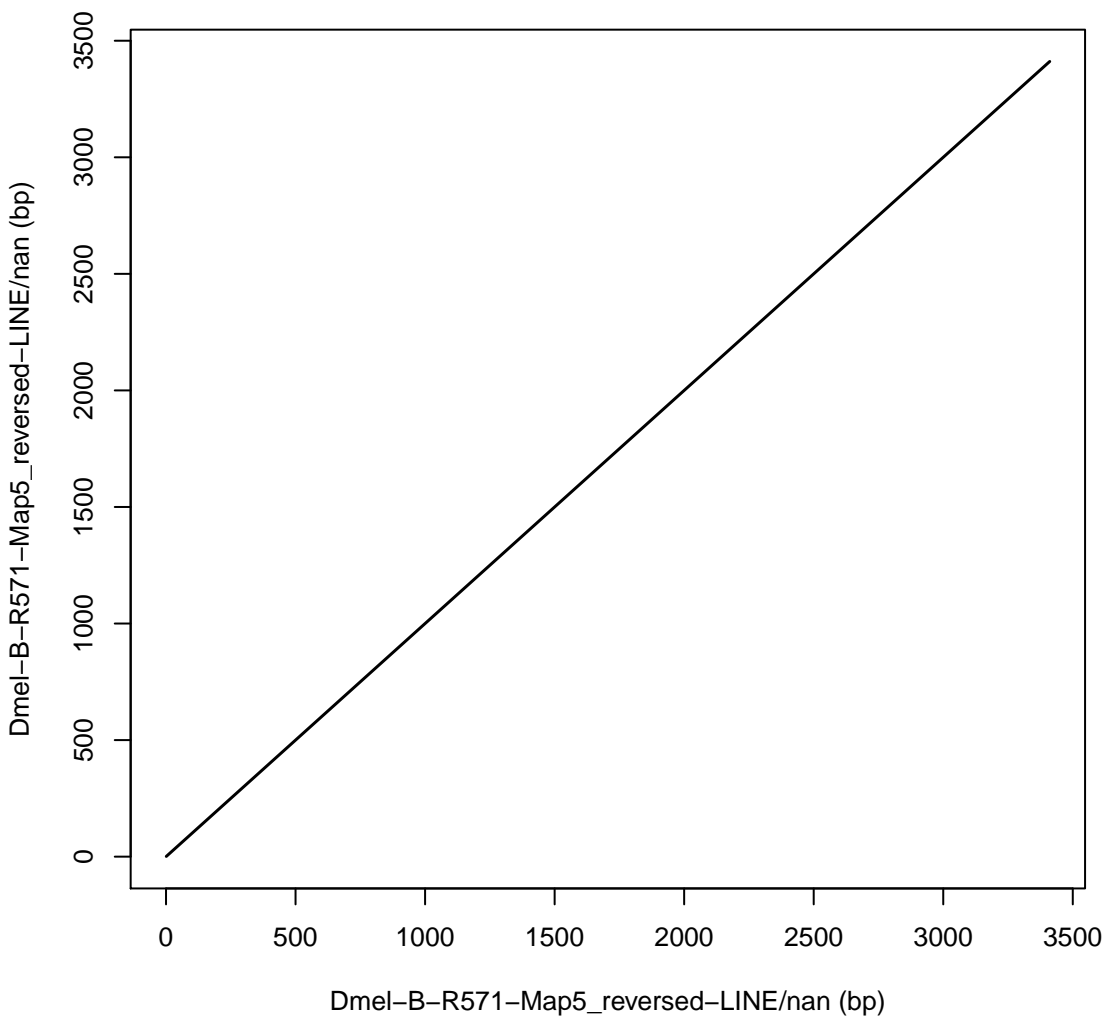
TE: Dmel-B-R571-Map5_reversed-LINE/nan
consensus size: 3411bp; fragments: 164; full length: 0 (>=3069.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

