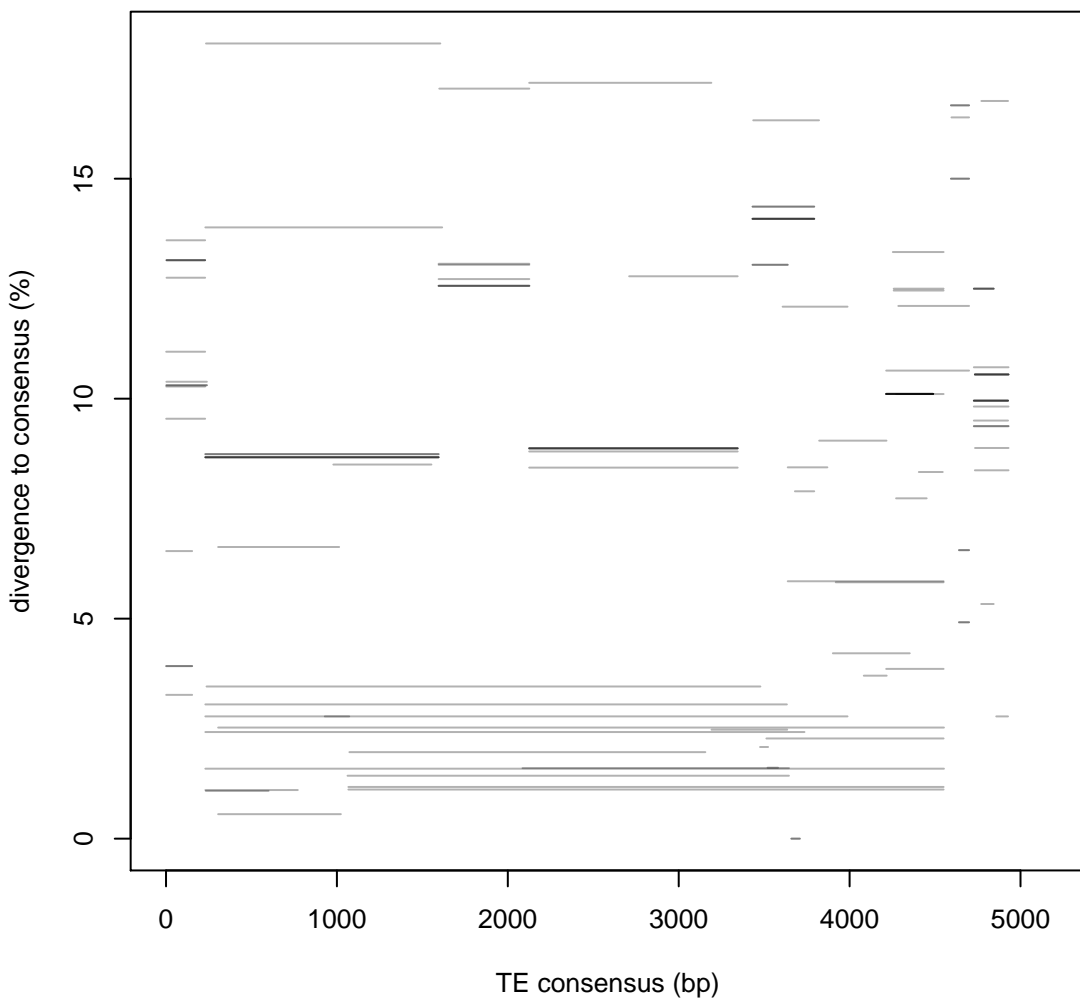
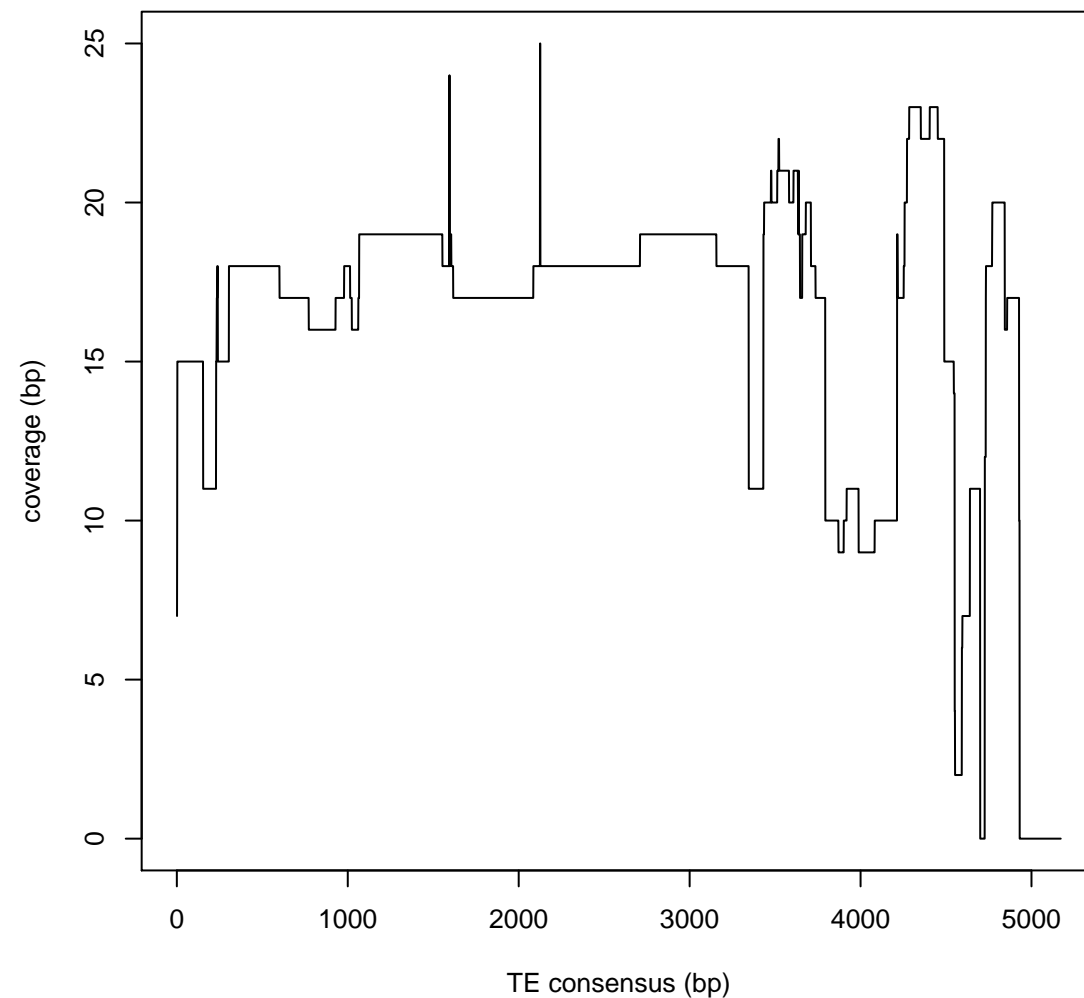


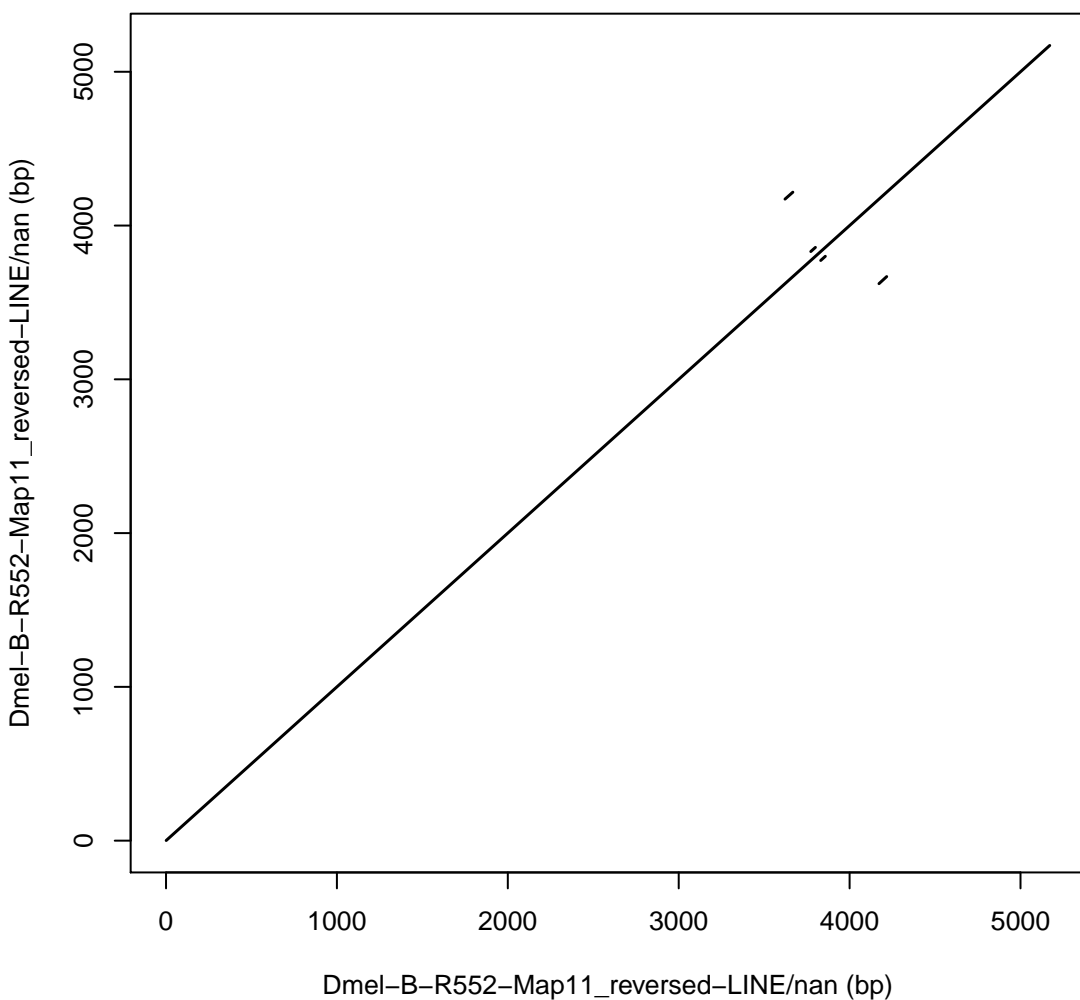
TE: Dmel-B-R552-Map11_reversed-LINE/nan
consensus size: 5171bp; fragments: 124; full length: 0 (>=4653.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

