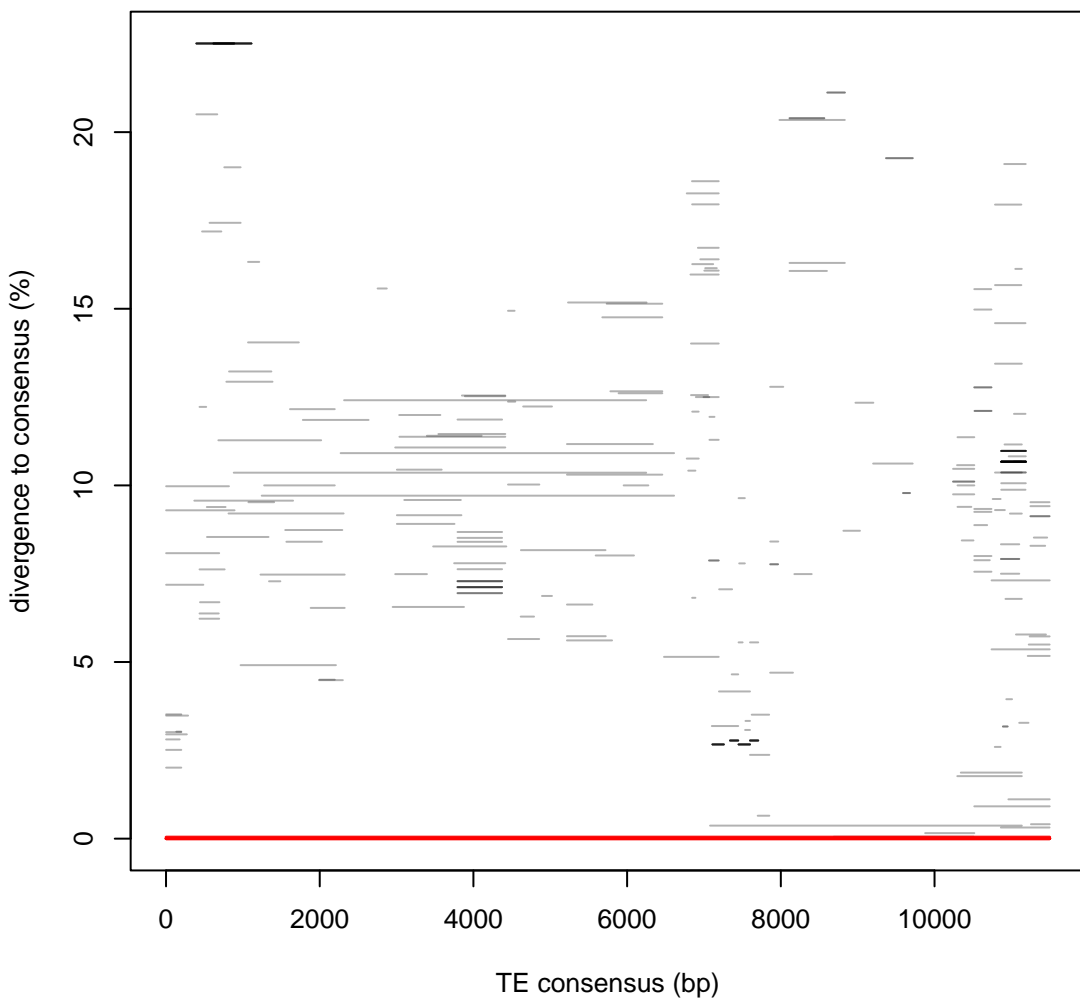
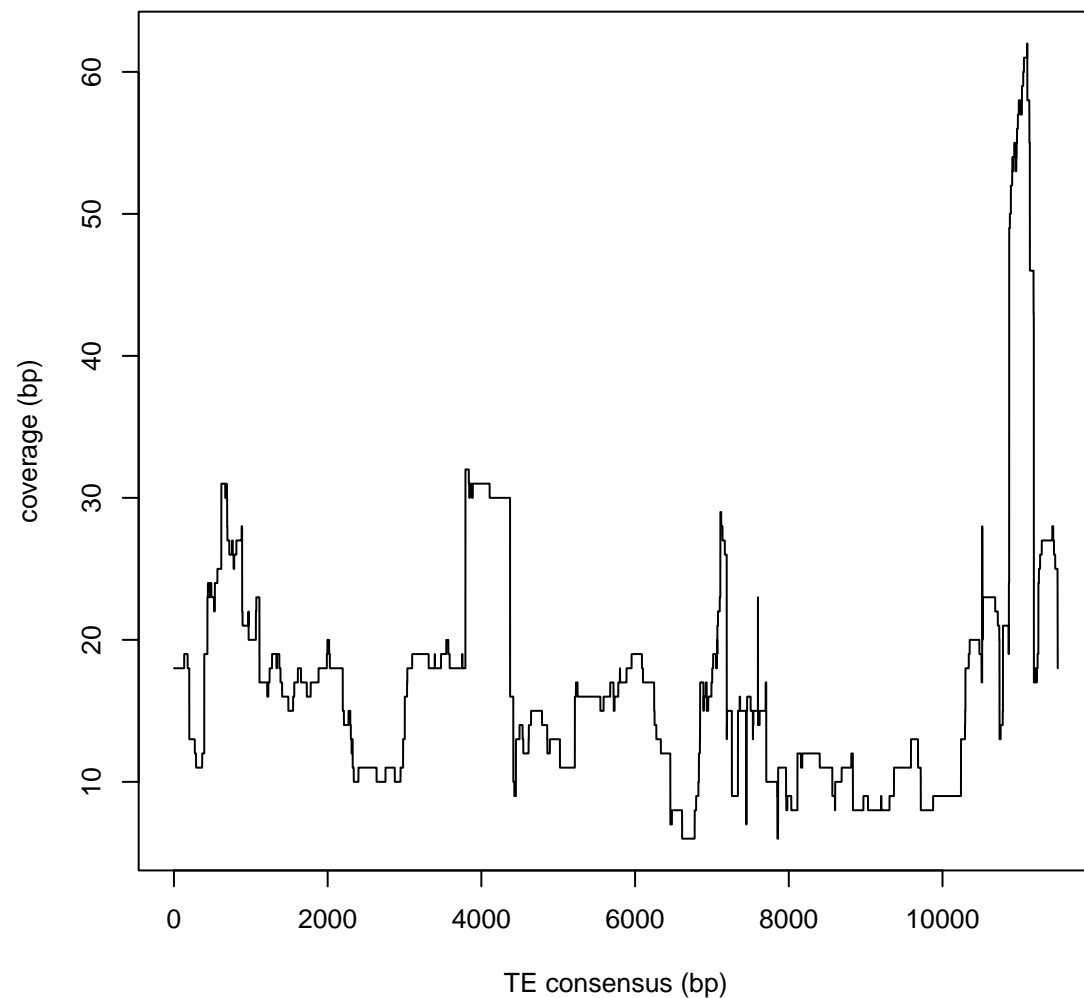


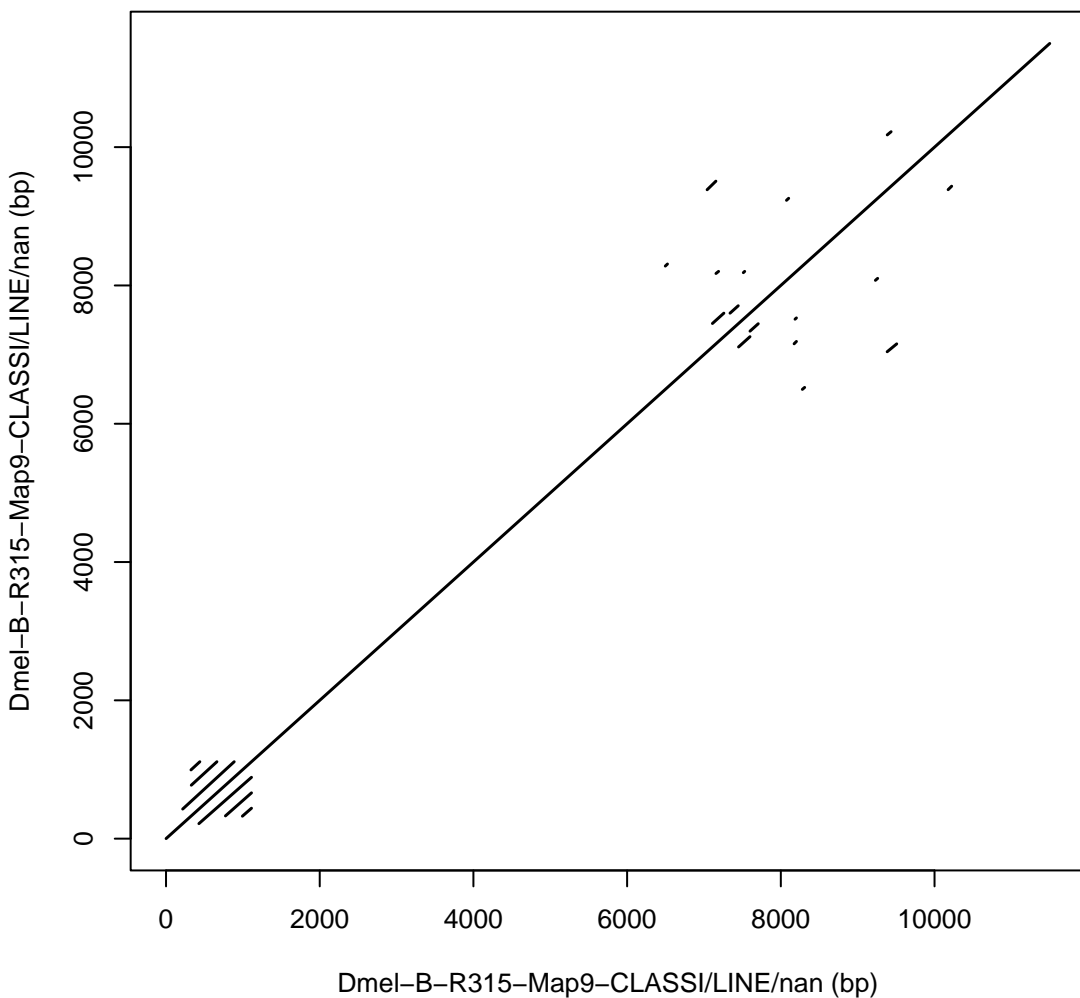
TE: Dmel-B-R315-Map9-CLASSI/LINE/nan
consensus size: 11499bp; fragments: 296; full length: 5 (>=10349.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

