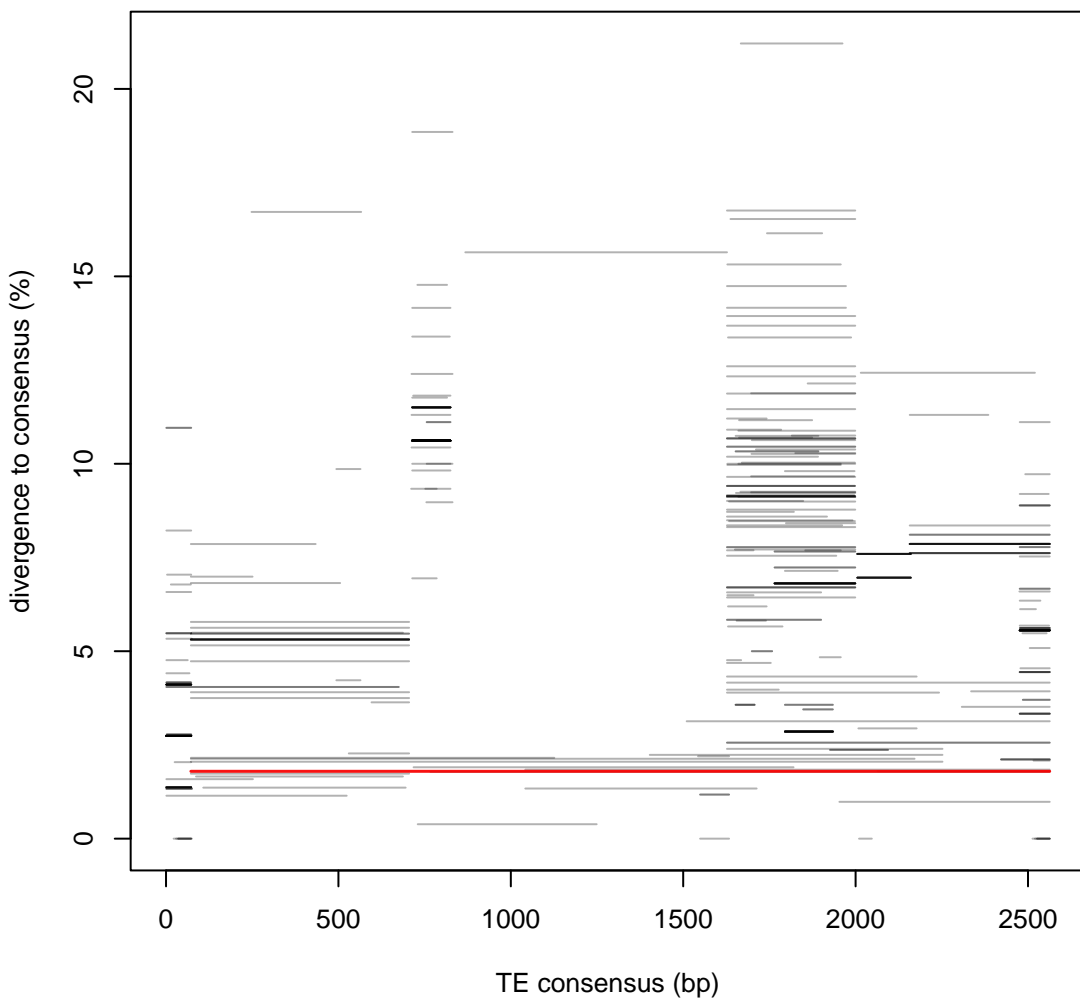
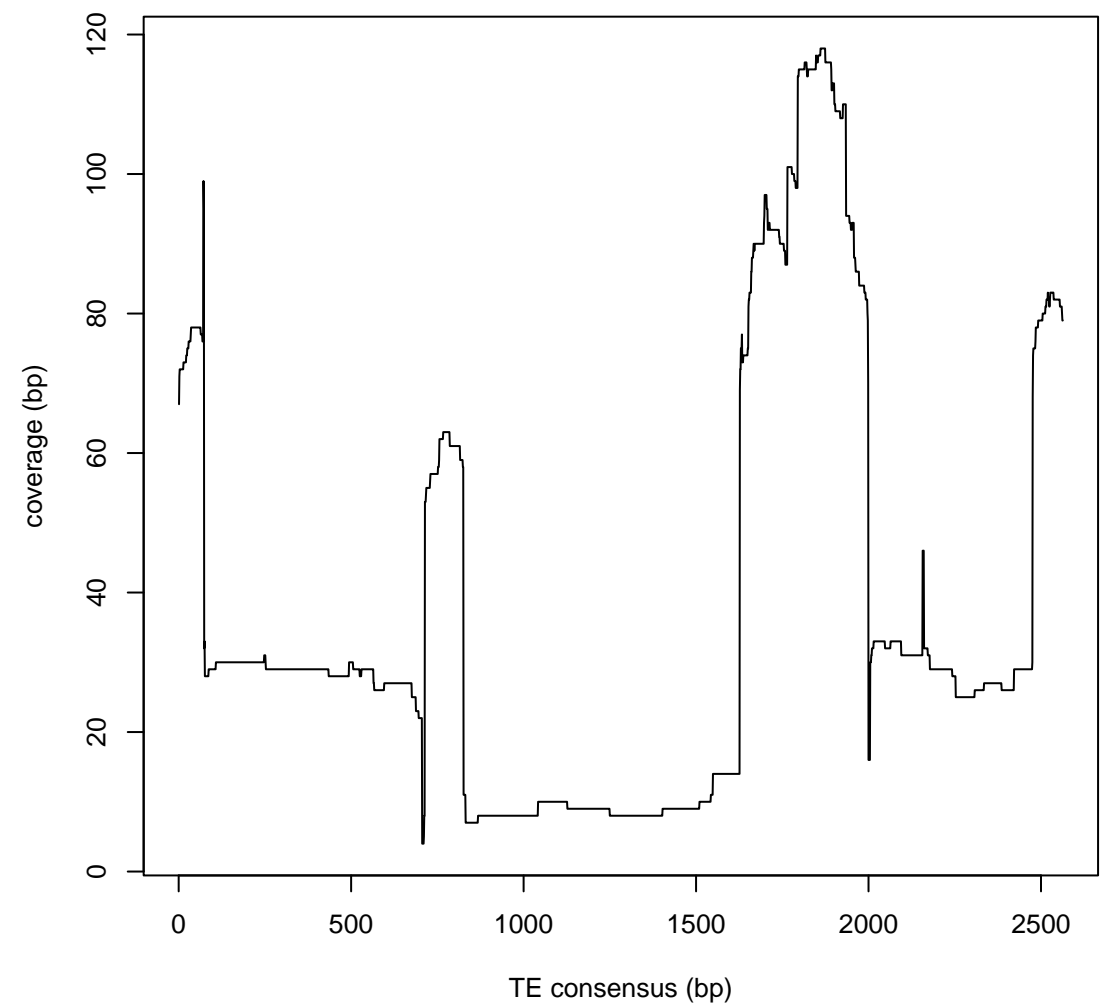


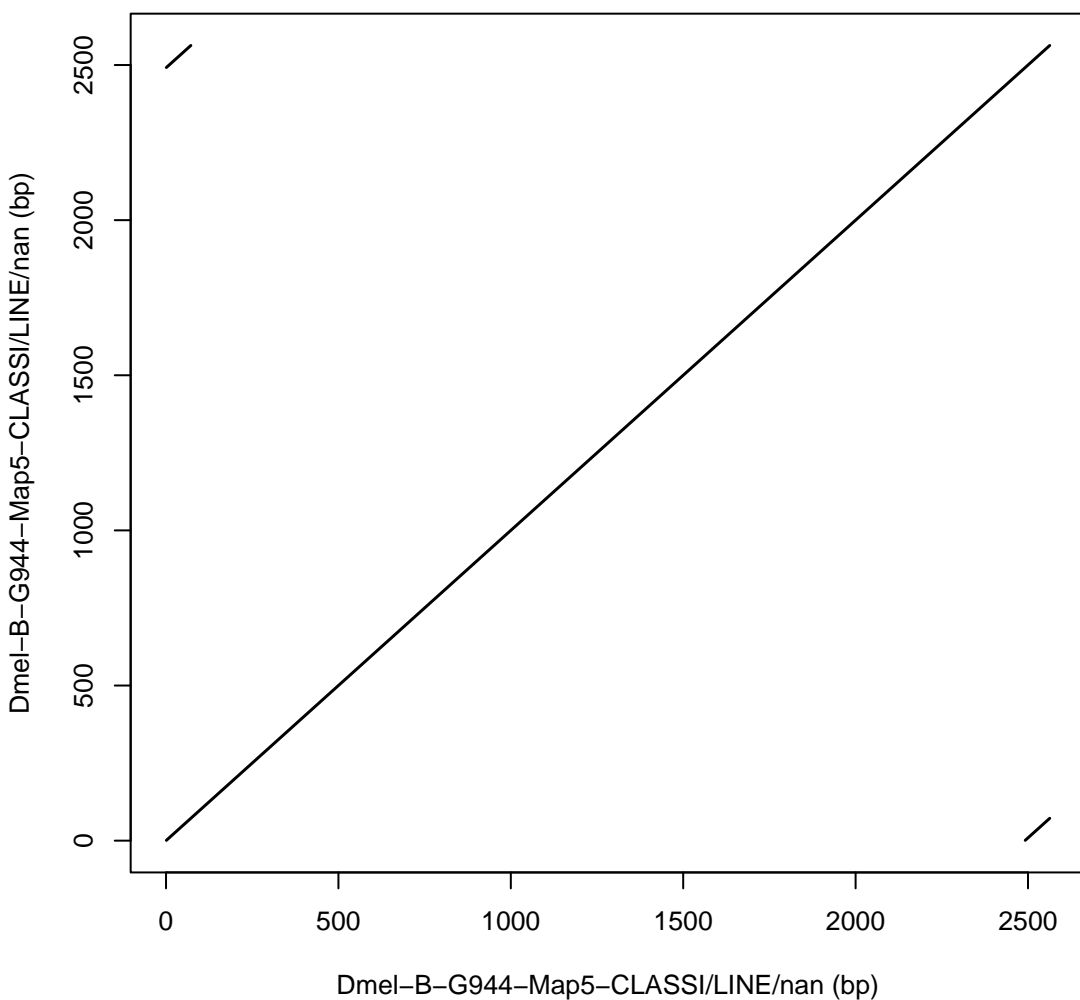
TE: Dmel-B-G944-Map5-CLASSI/LINE/nan
consensus size: 2563bp; fragments: 403; full length: 1 (≥ 2306.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

