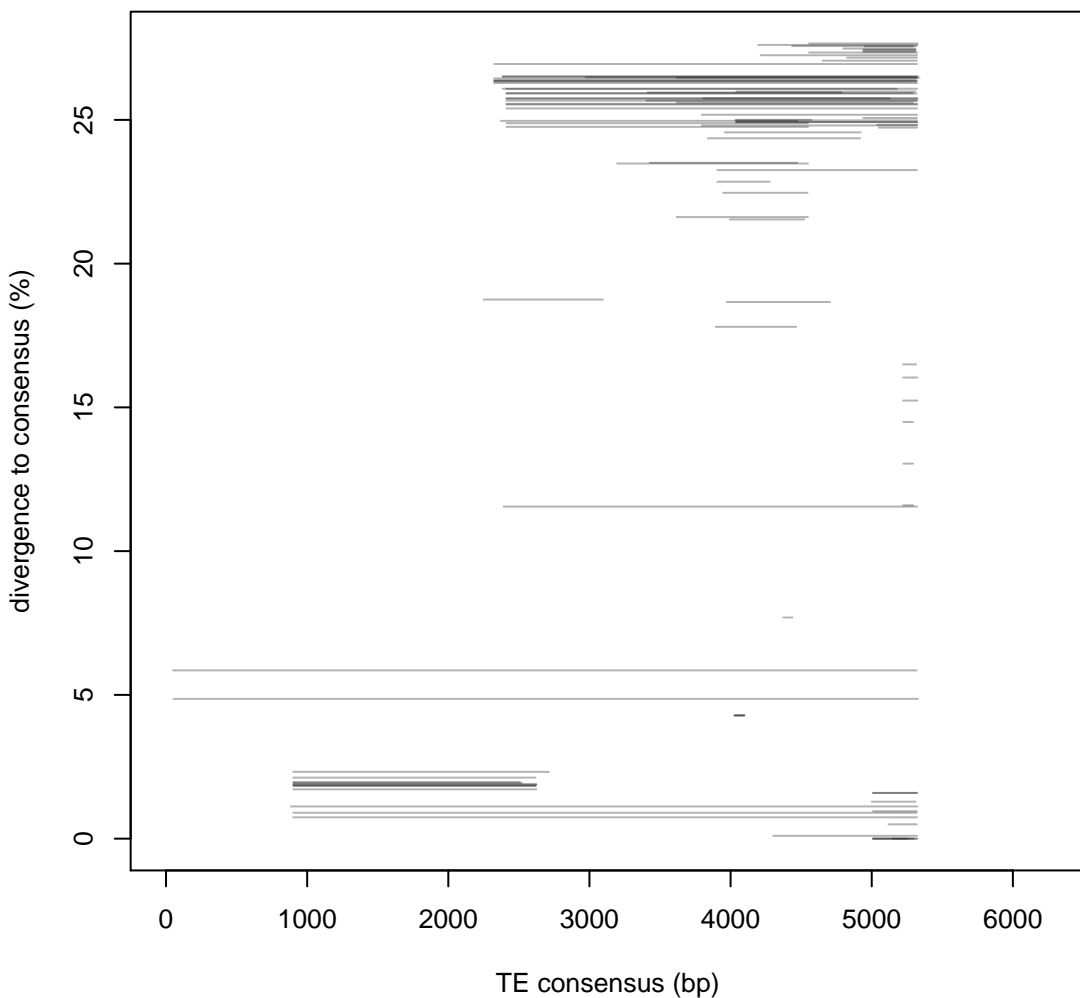
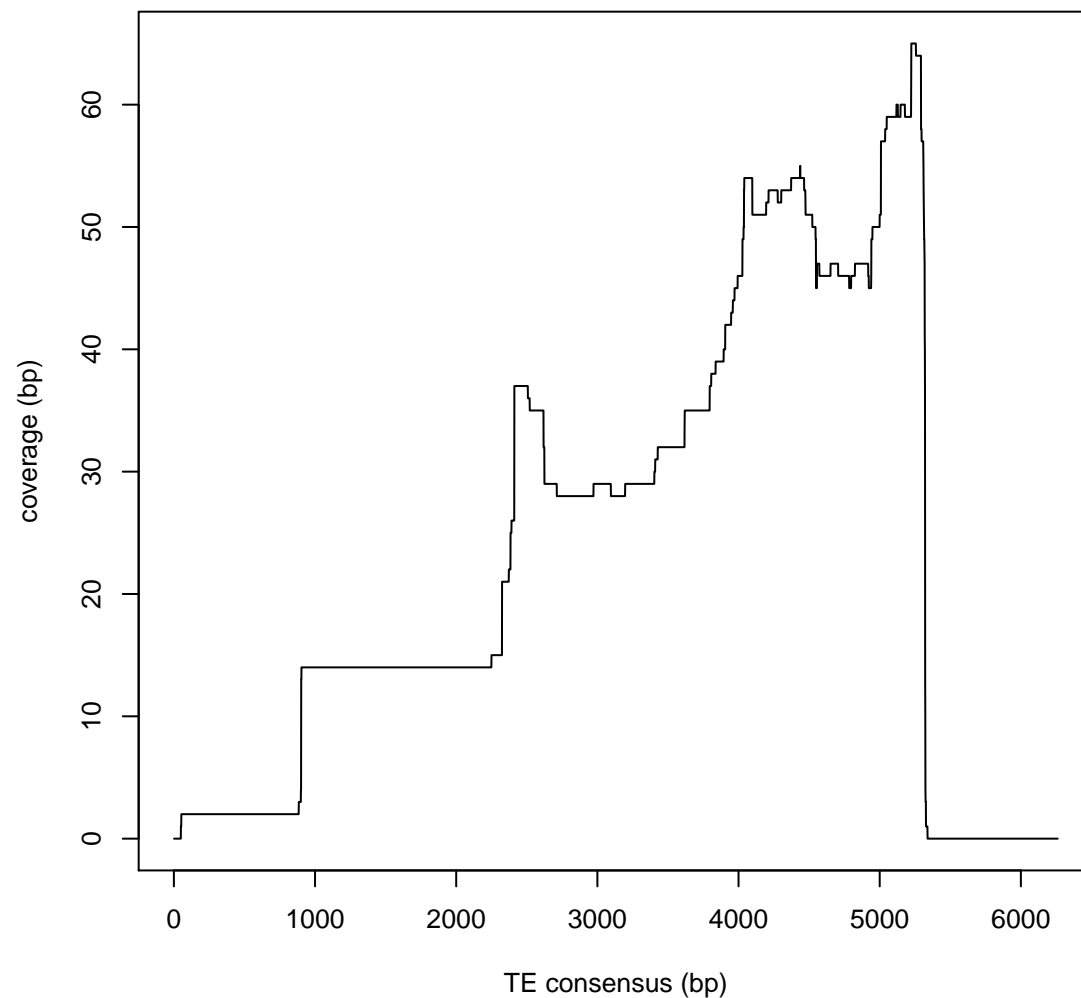


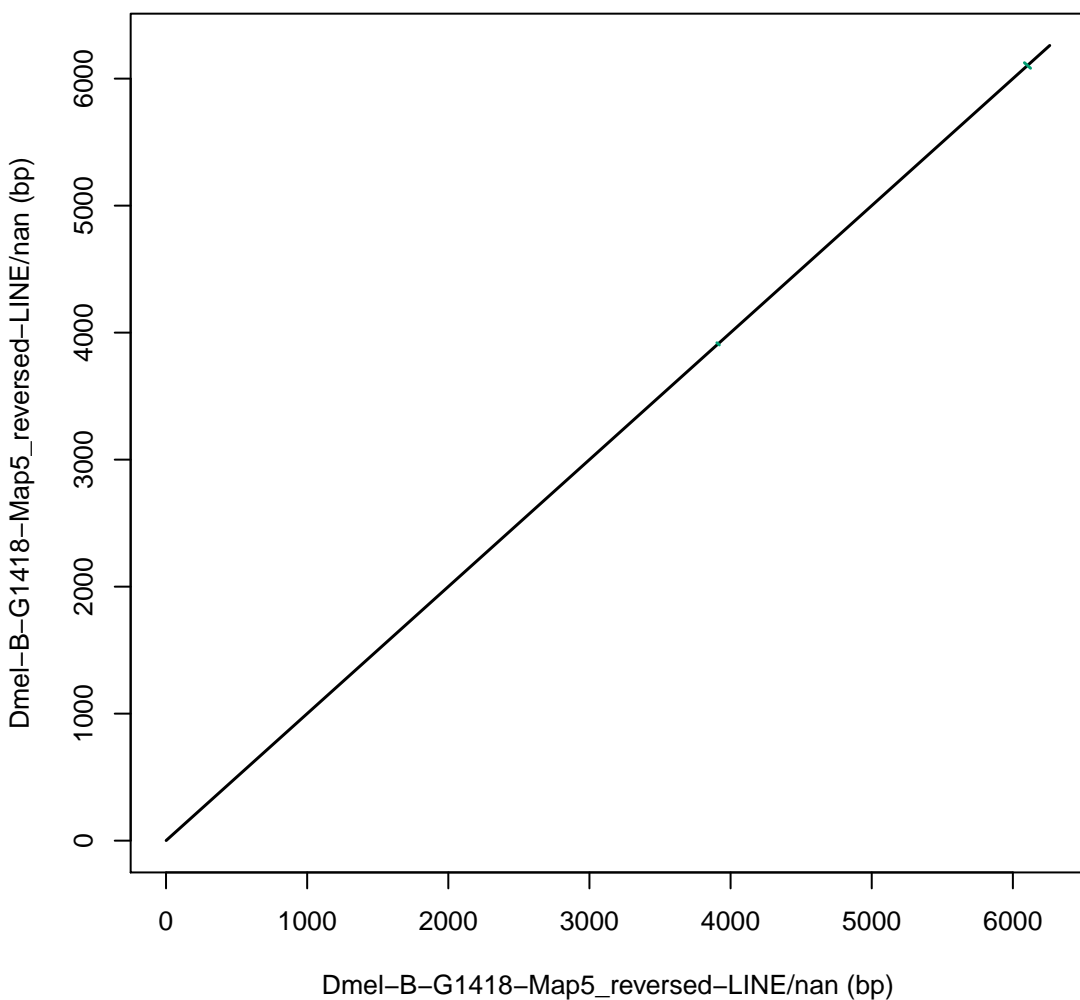
TE: Dmel-B-G1418-Map5\_reversed-LINE/nan  
consensus size: 6261bp; fragments: 96; full length: 0 (>=5634.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

