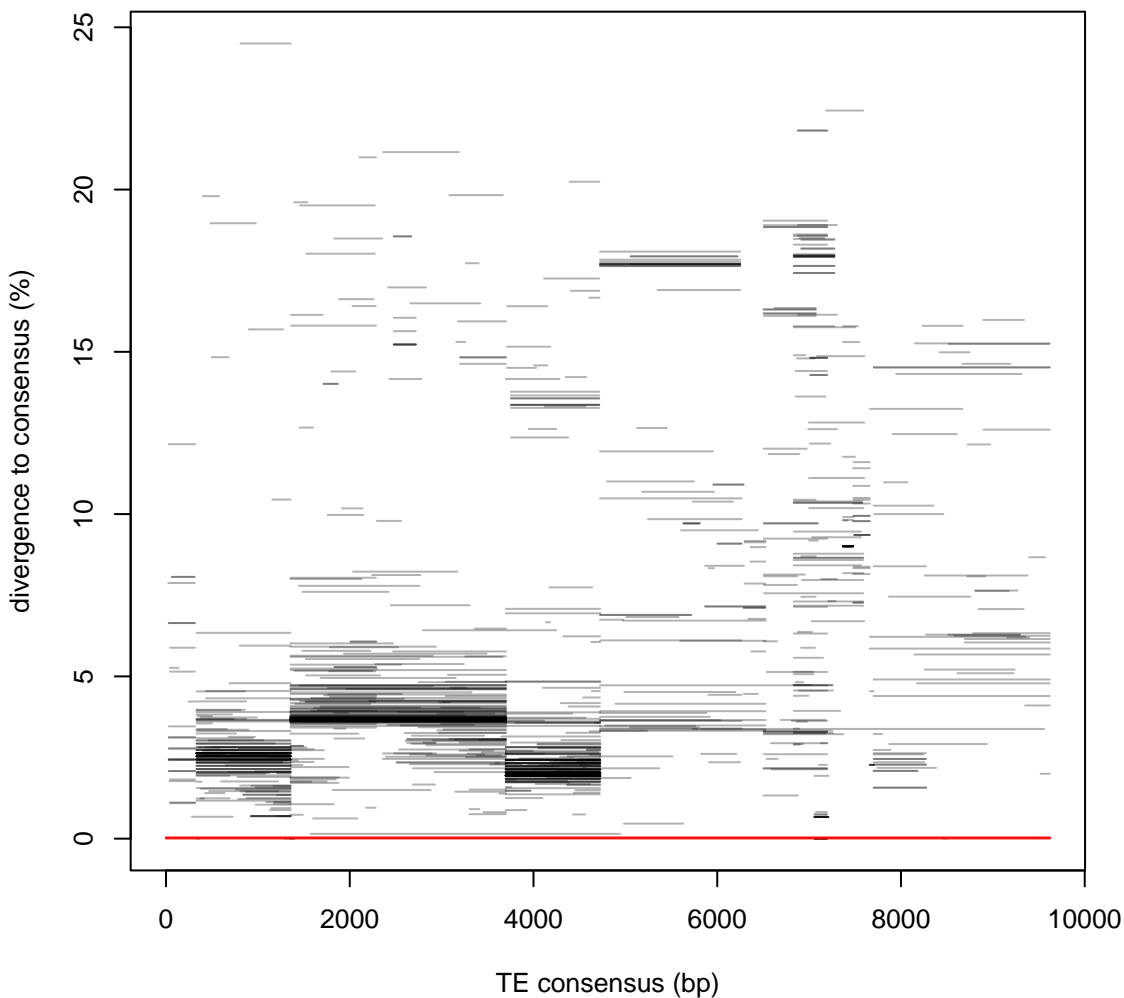
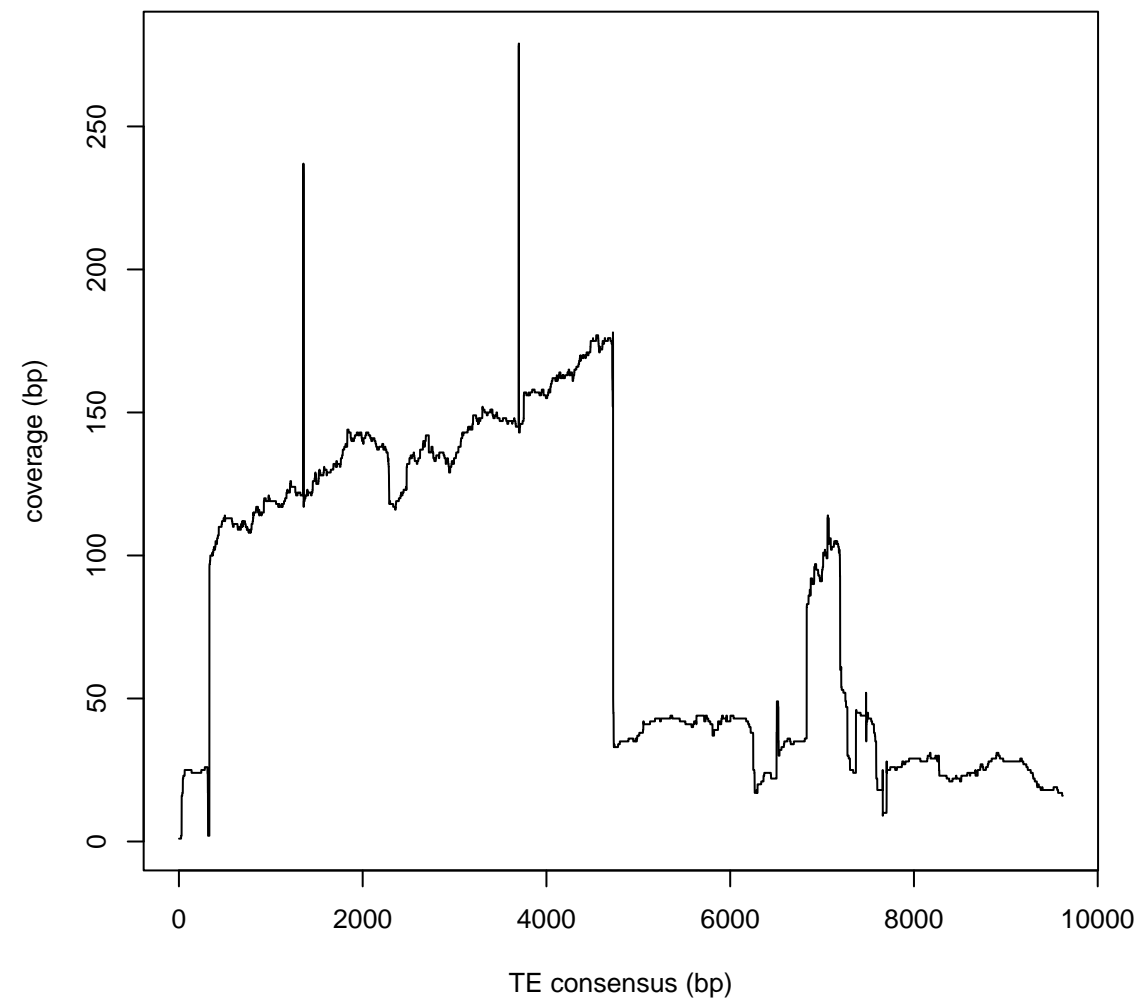


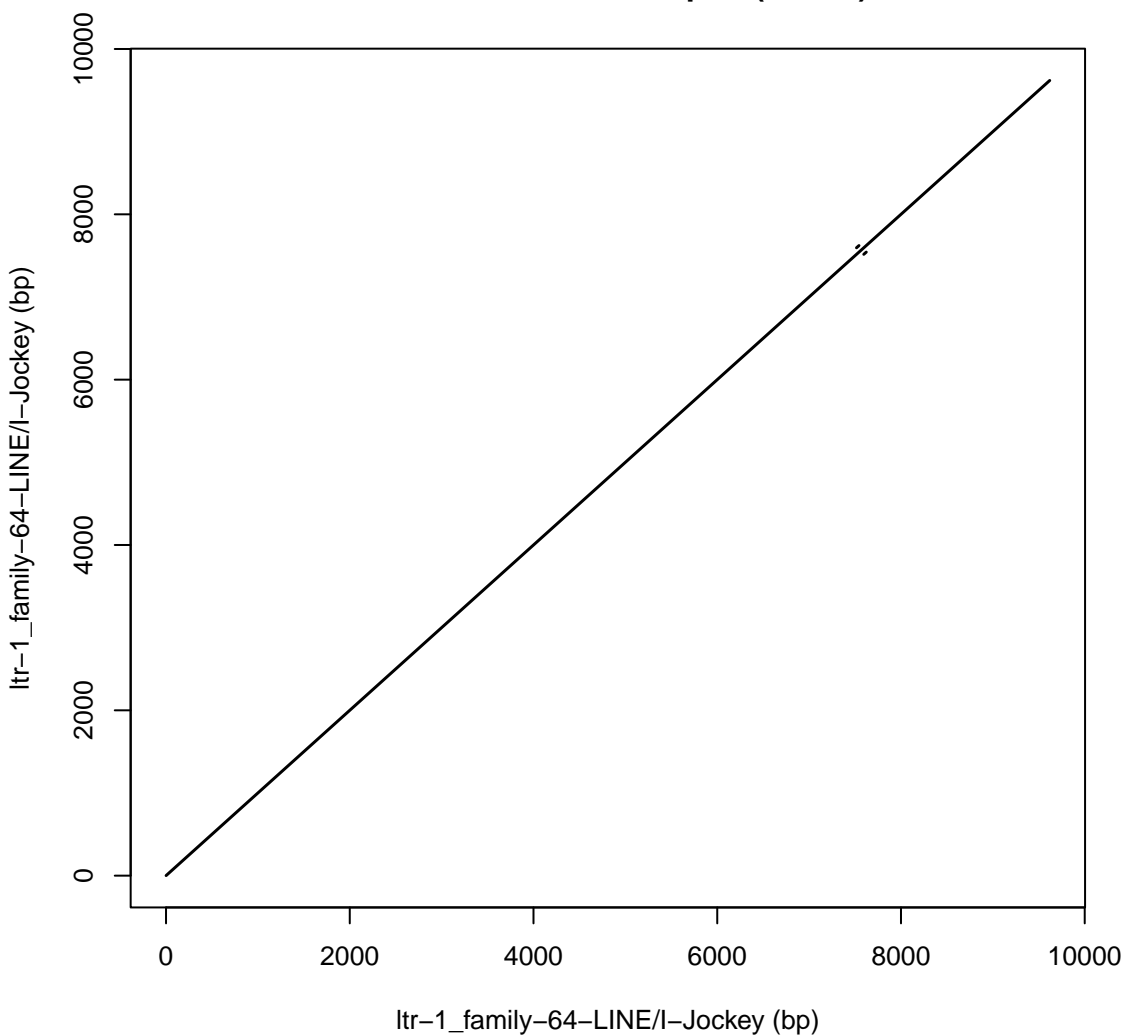
TE: ltr-1_family-64-LINE/l-Jockey
consensus size: 9619bp; fragments: 970; full length: 1 (>=8657.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

