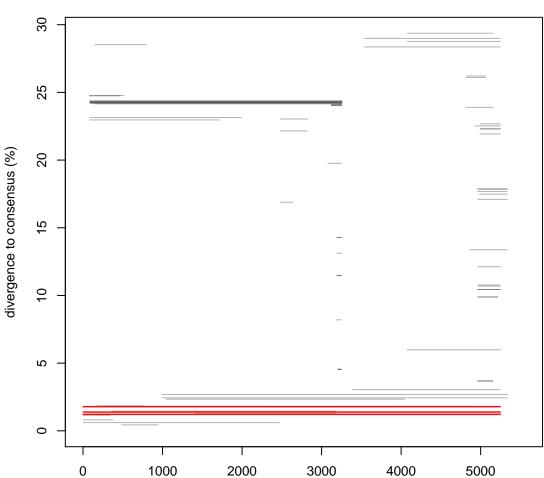
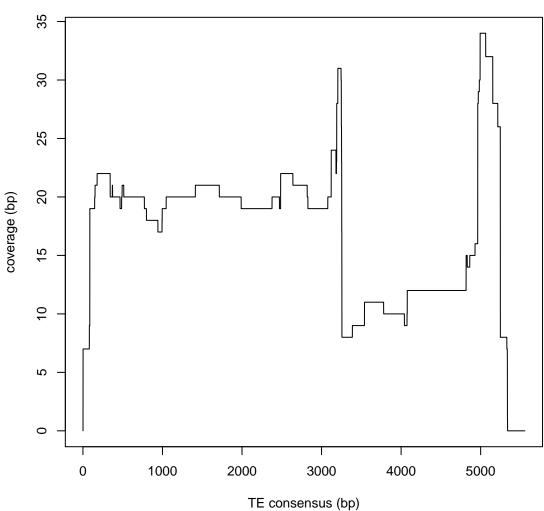
TE: rnd-6_family-1433-LTR/Gypsy consensus size: 5556bp; fragments: 76; full length: 3 (>=5000.4bp)

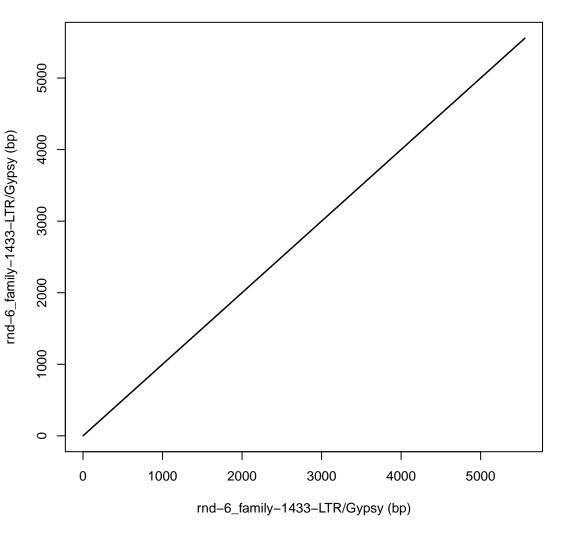


TE consensus genomic coverage



TE consensus self dotplot (blastn)

TE consensus (bp)



TE consensus structure and protein hits

