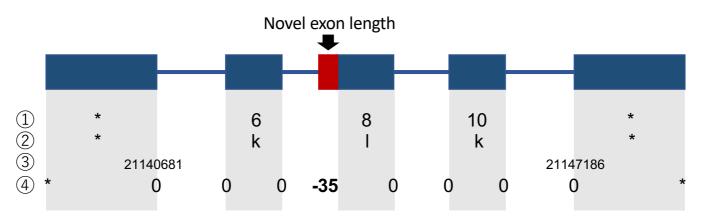


- 1 Exon numbers
- ② Exon length (k: known, s: short, l: long, sl: short&long)
- ③ Positions of insides of both ends of the exons
- 4 Changes of exon length



^{*:} We did not annotate both end exons (UTRs) because of the low quality at the ends of the reads. If you want to determine the position of the both ends, refer to the position of the reference exon.