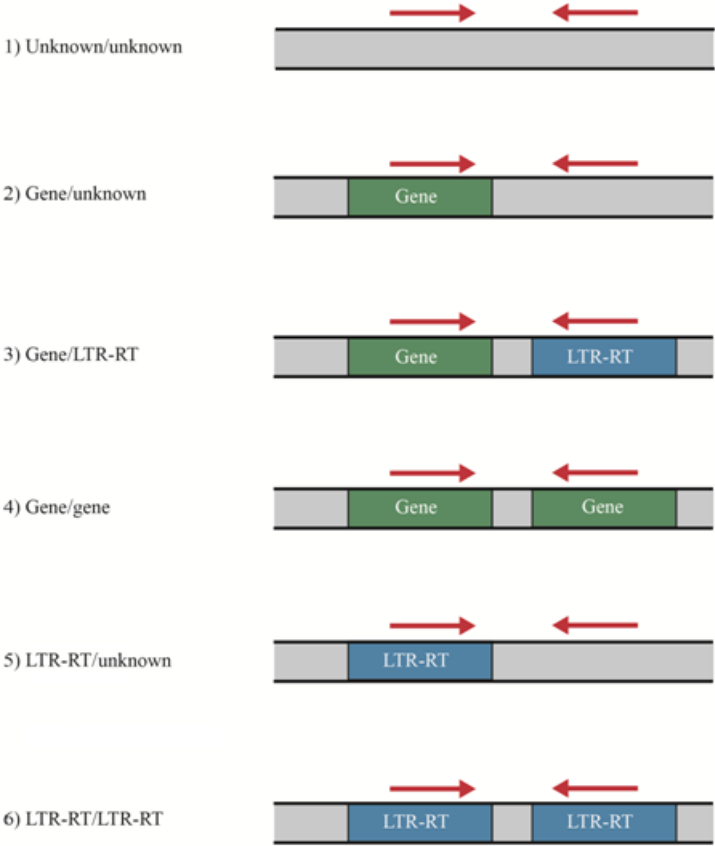


Supplementary material 4



Possible mapping patterns for paired-end reads onto genes, retrotransposons, or unknown (*i.e.*, not annotated) sequences.