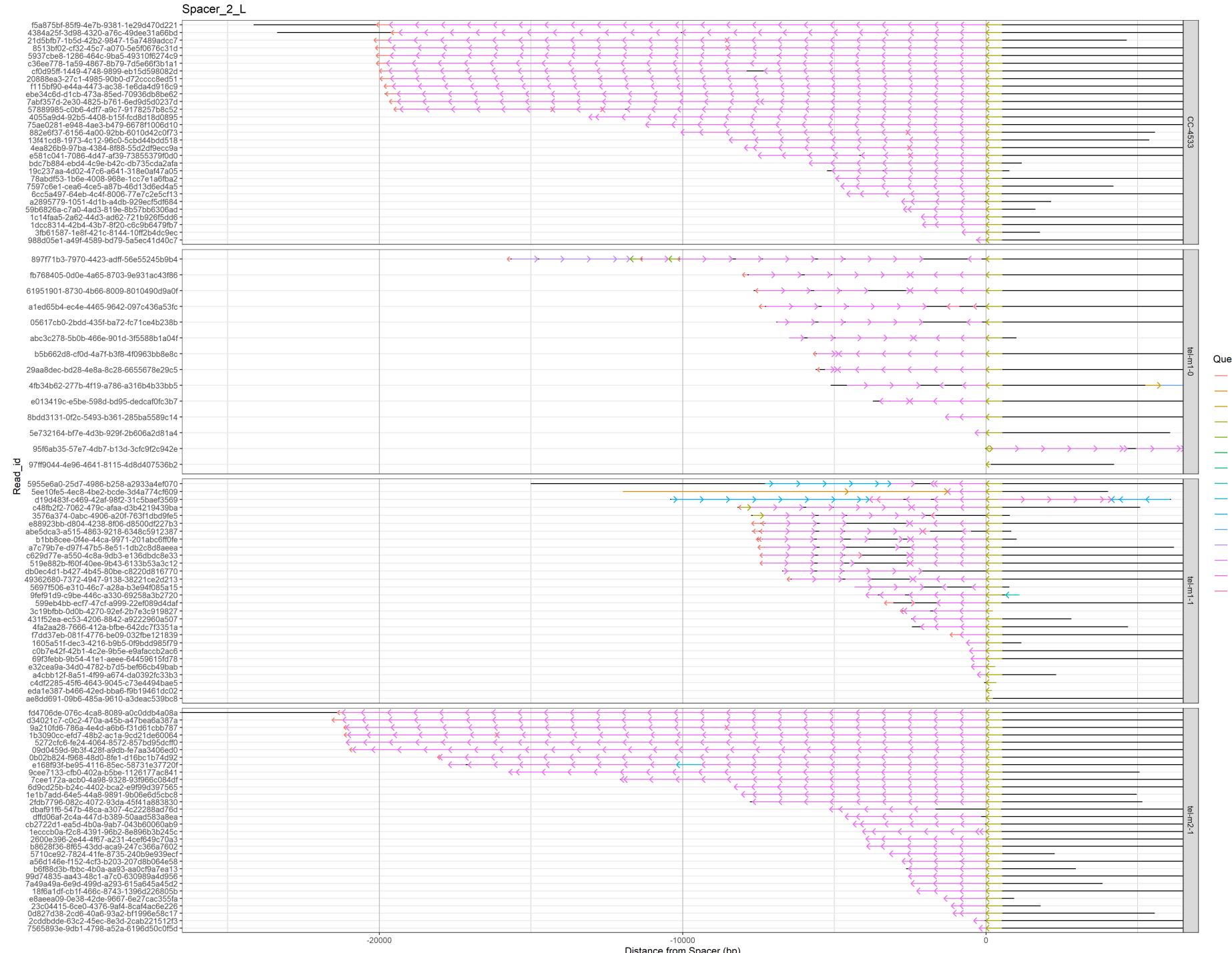
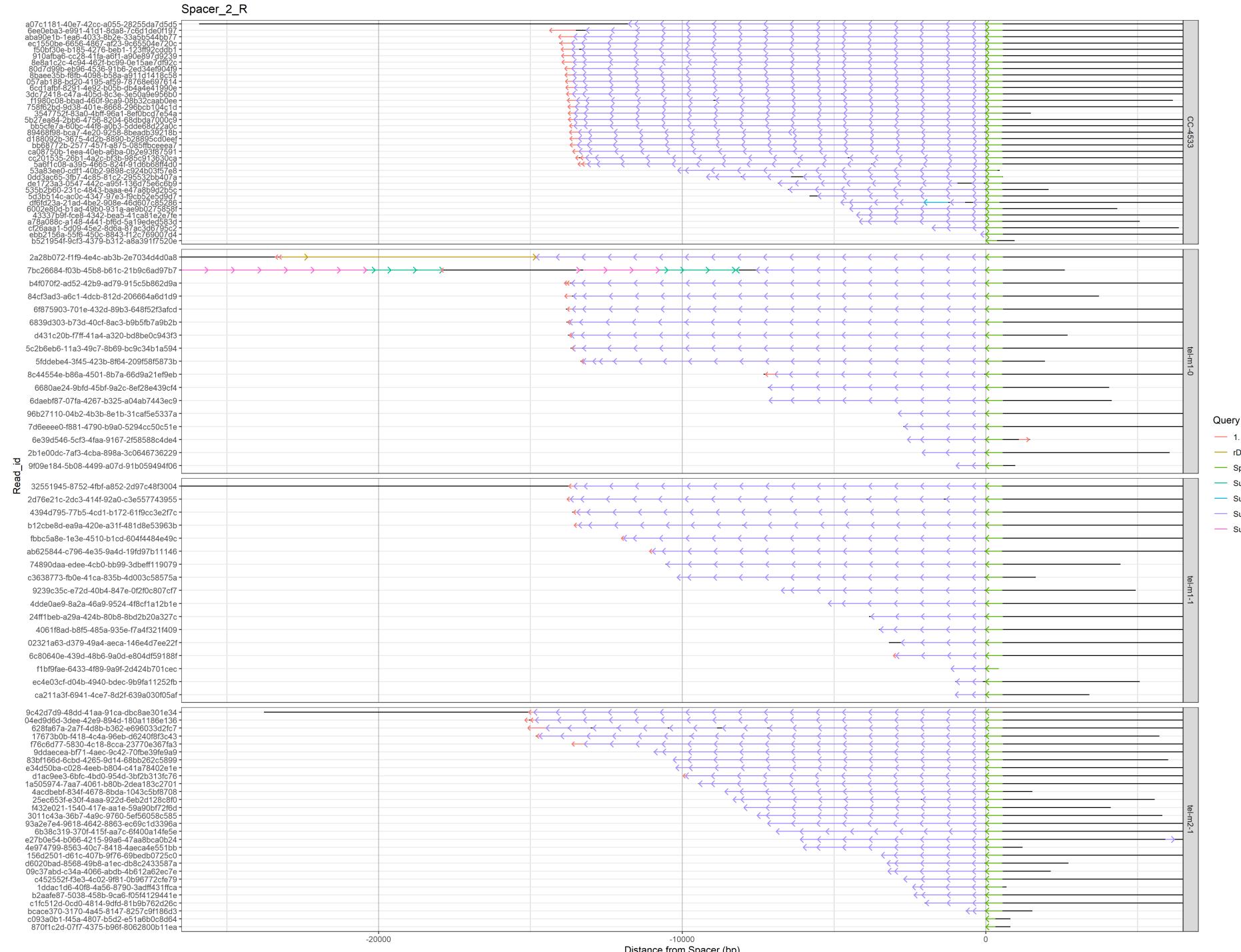
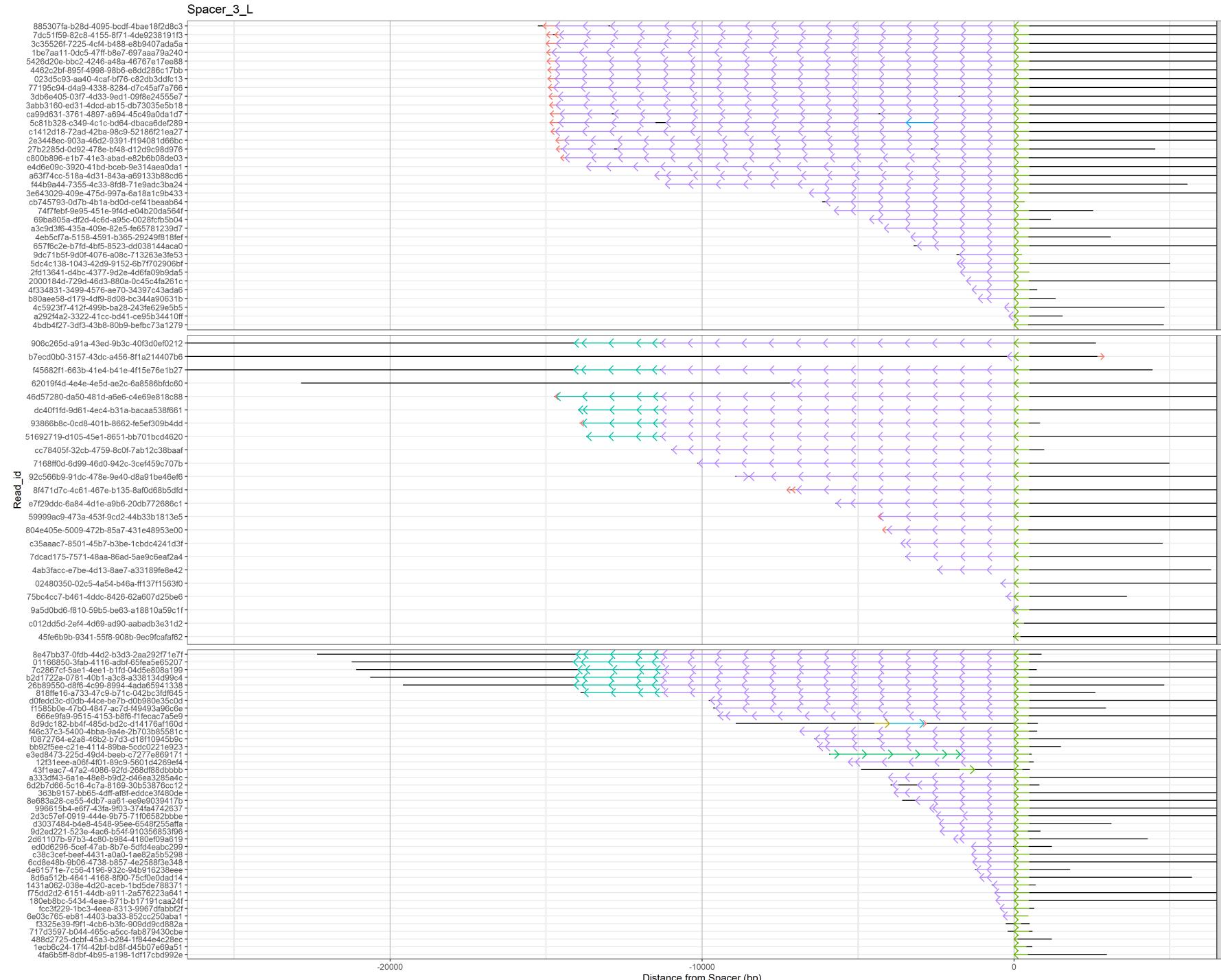
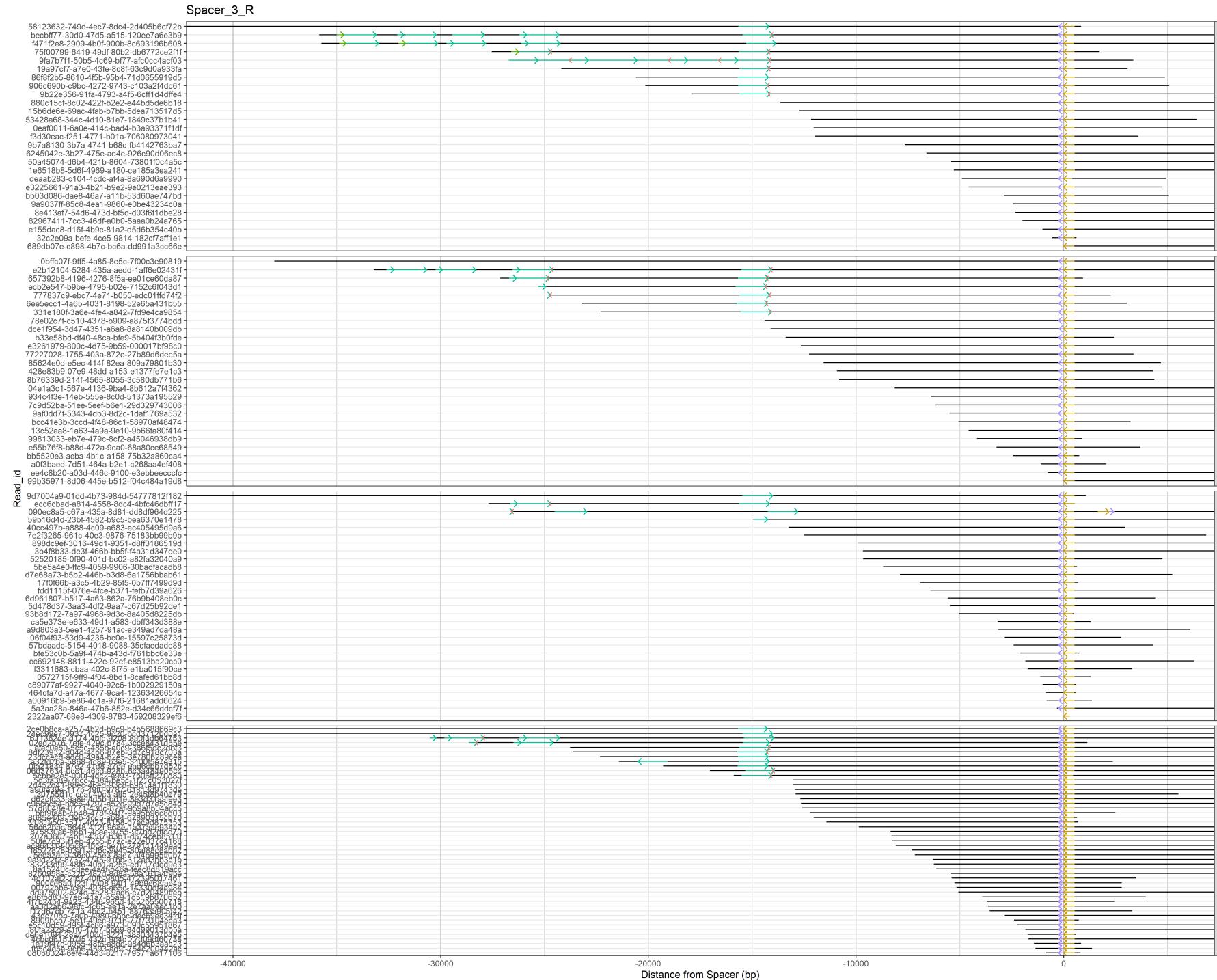


Supplemental Dataset 1. *Spacer*-based analysis of subtelomeres. All *Spacer*-containing reads anchored at the *Spacer* element are depicted for each subtelomere for CC-4533, *tel-m1-0*, *tel-m1-1* and *tel-m2-1*. The queried elements are displayed in color. The identifier for each read (“Read_id”) is also shown, to facilitate retrieval from the sequencing data.

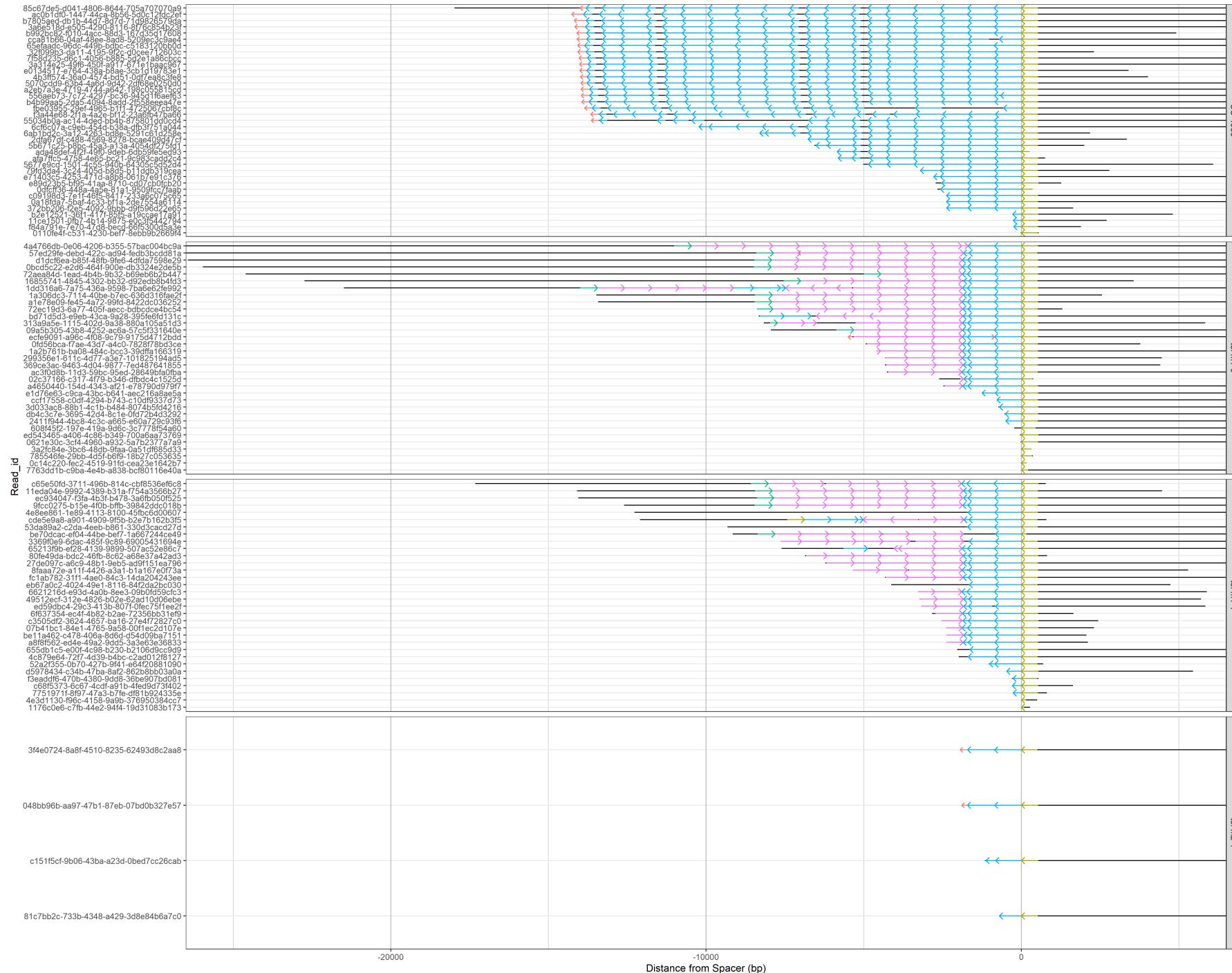




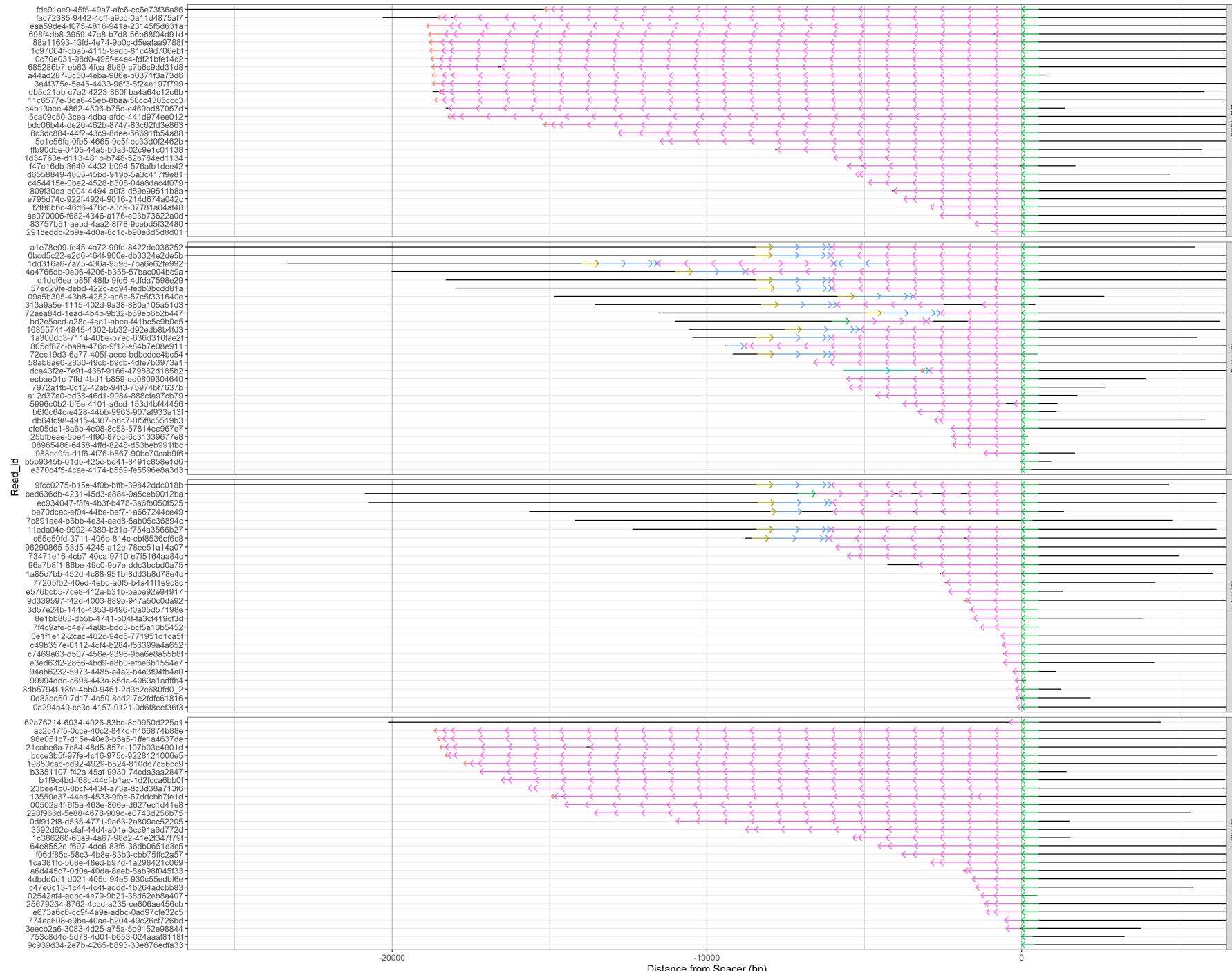


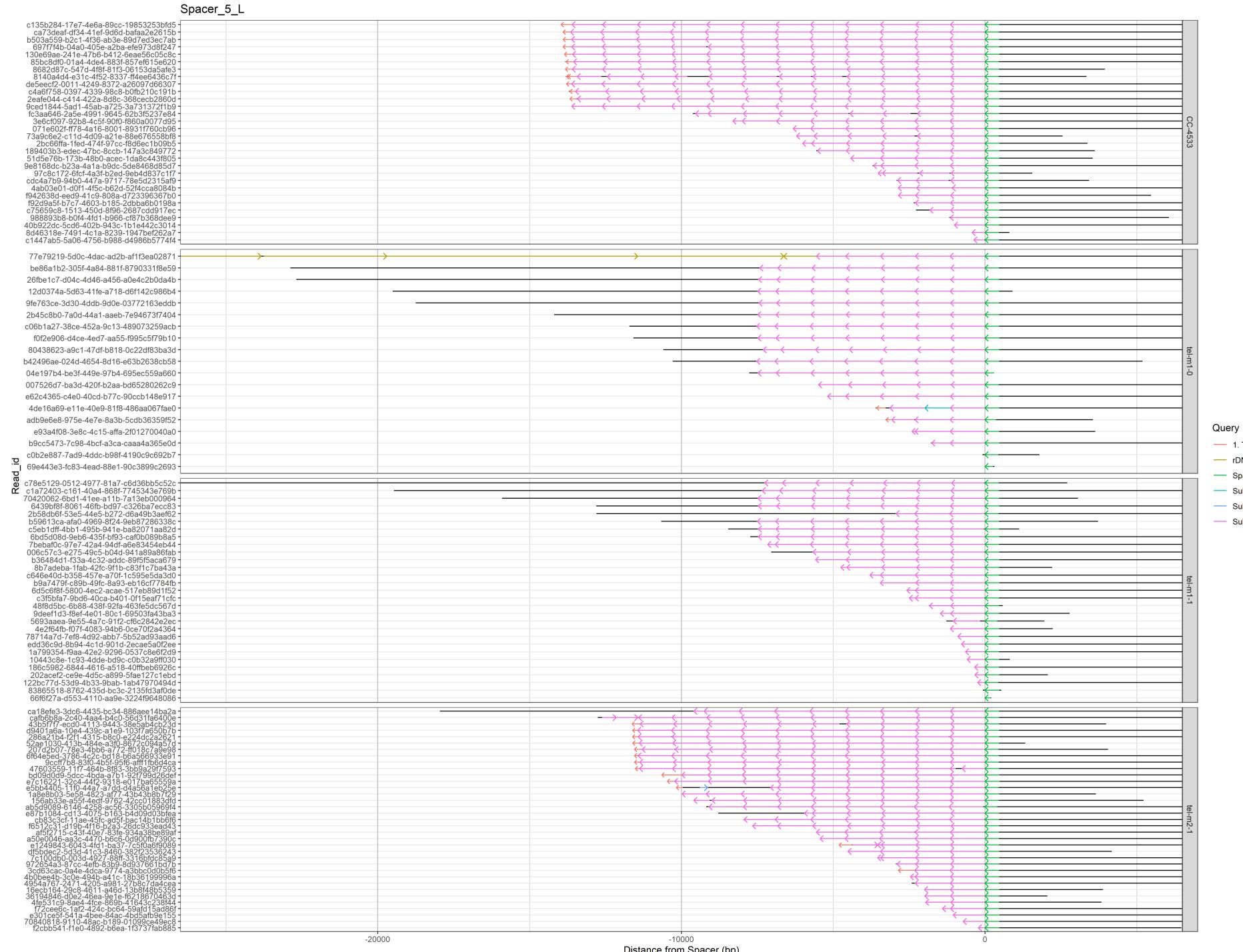


Spacer_4_L

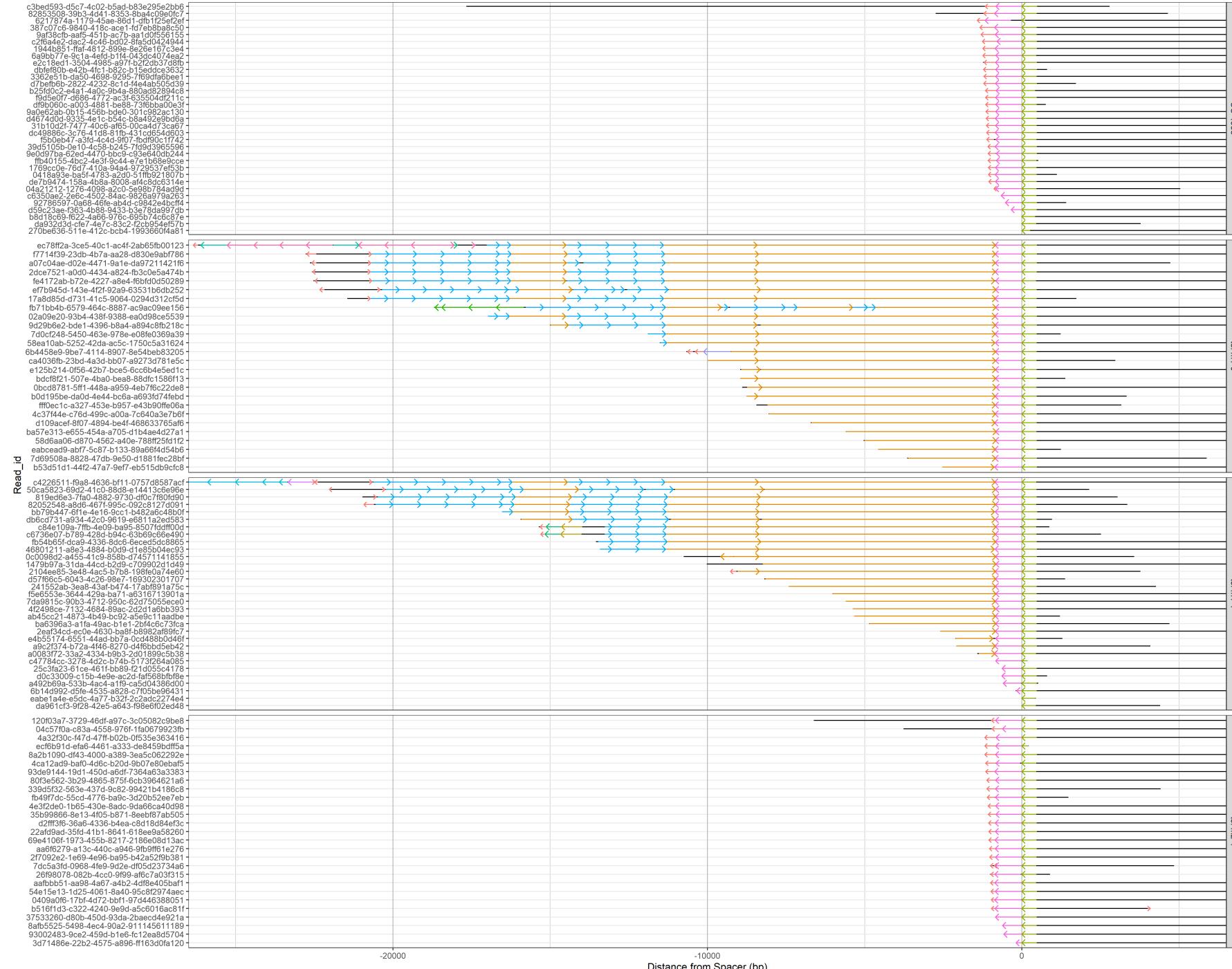


Spacer_4_R

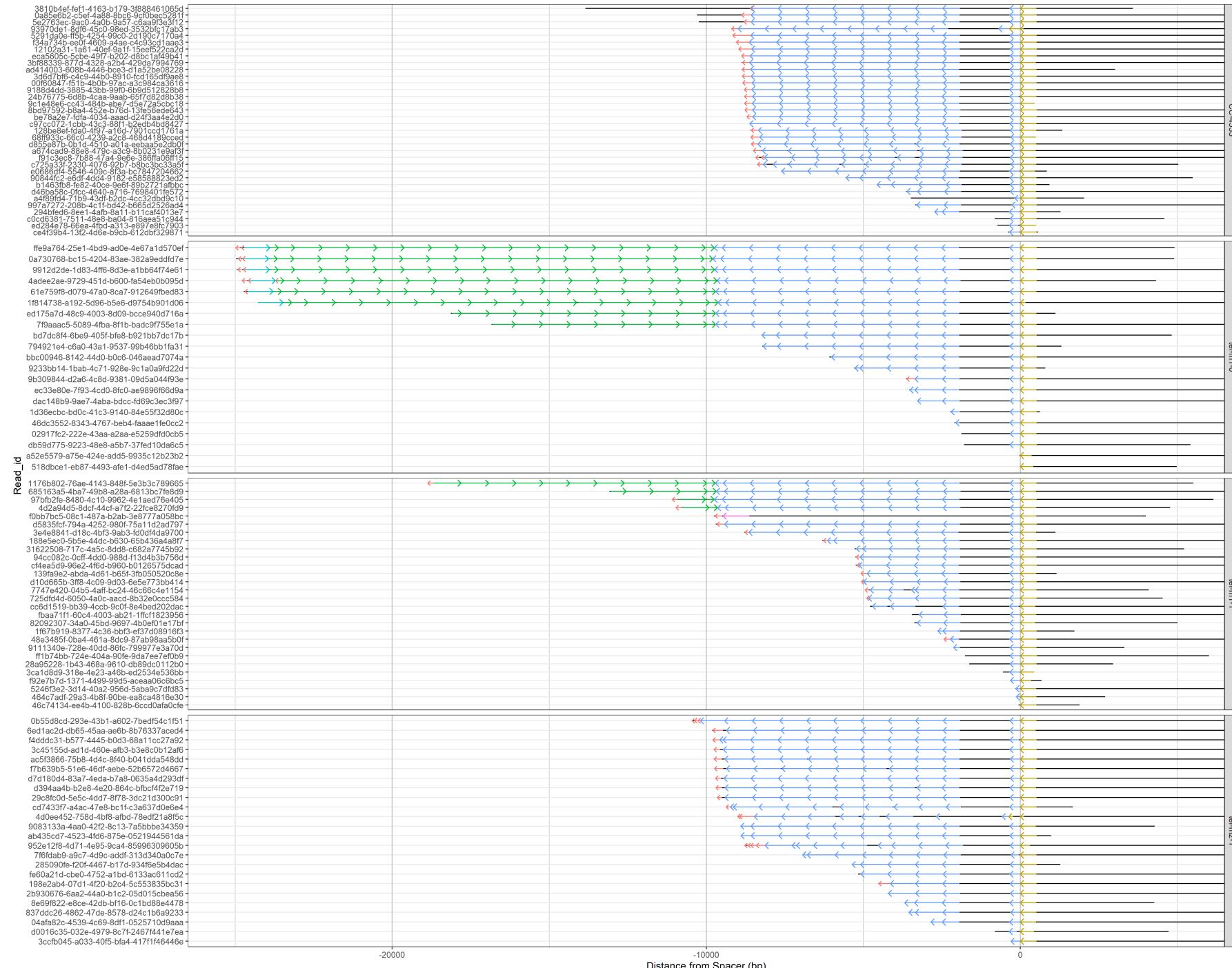


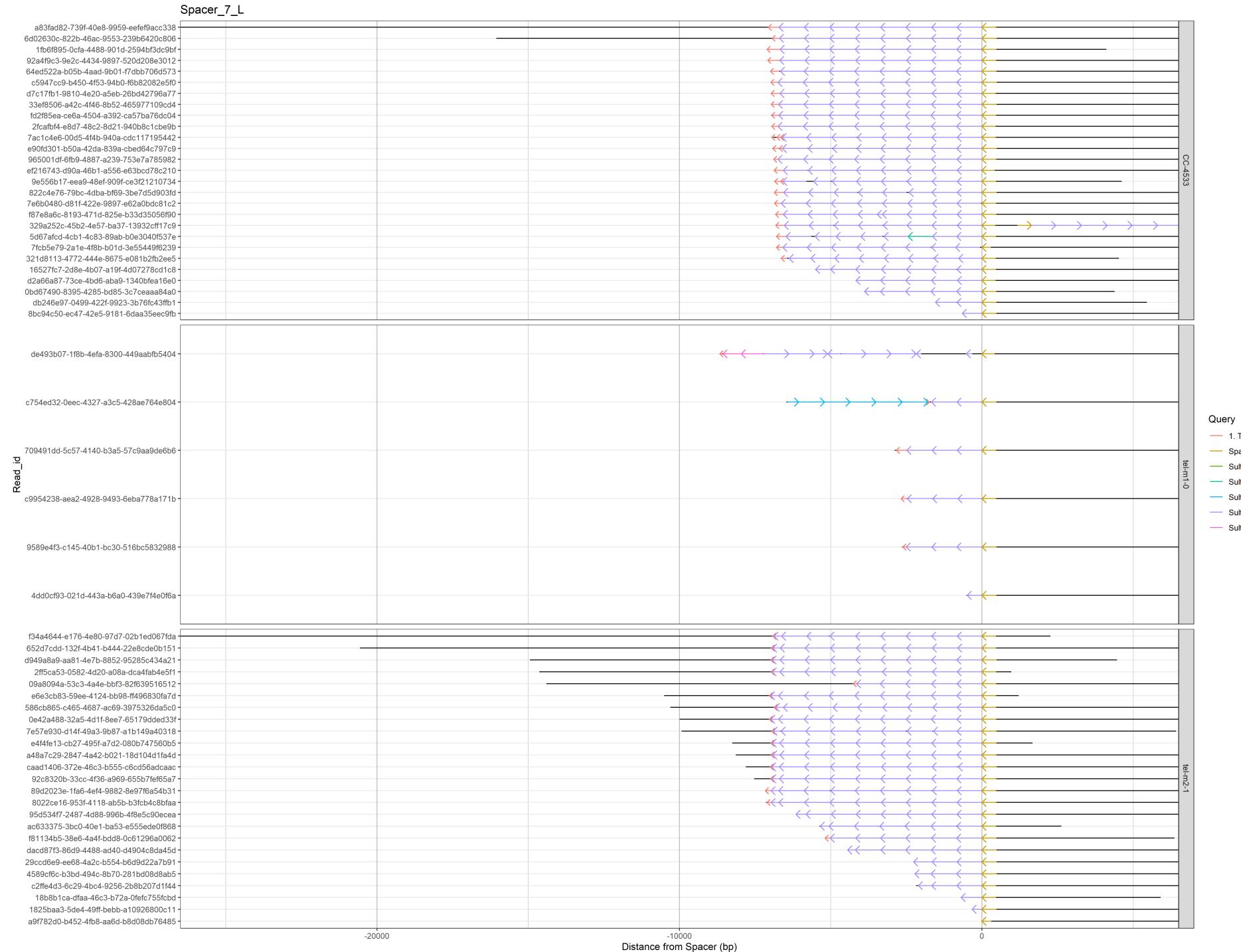


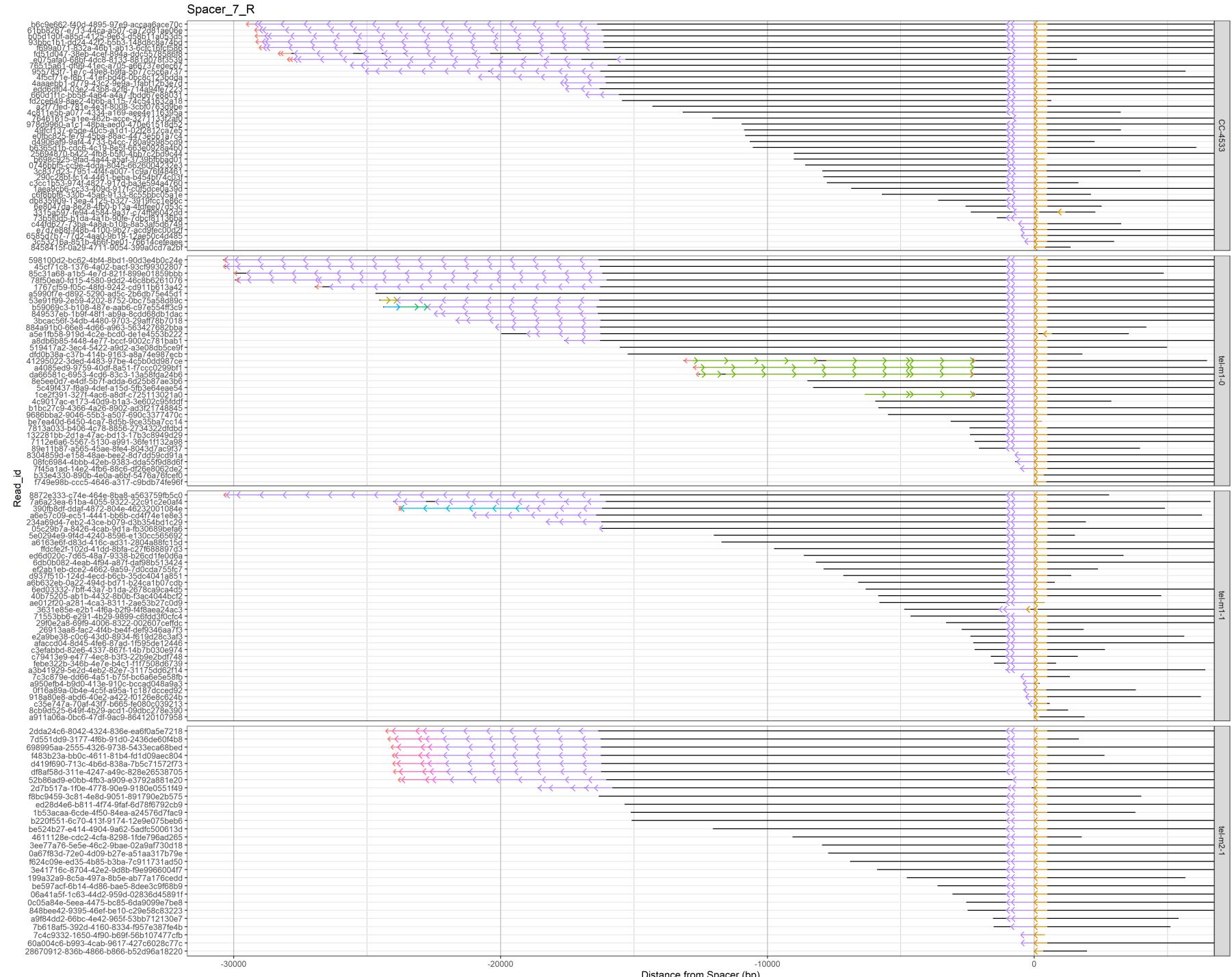
Spacer_6_L

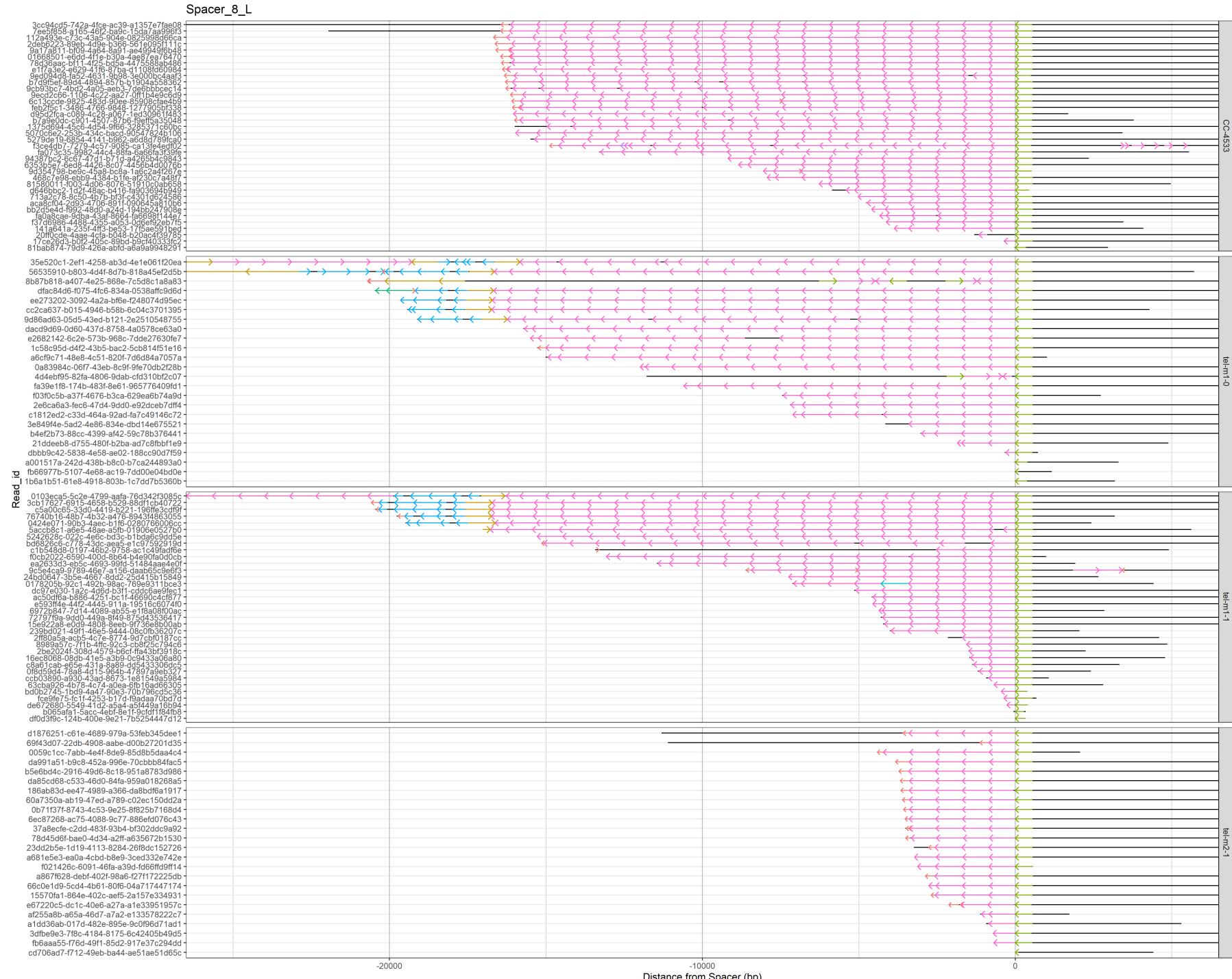


Spacer_6_R

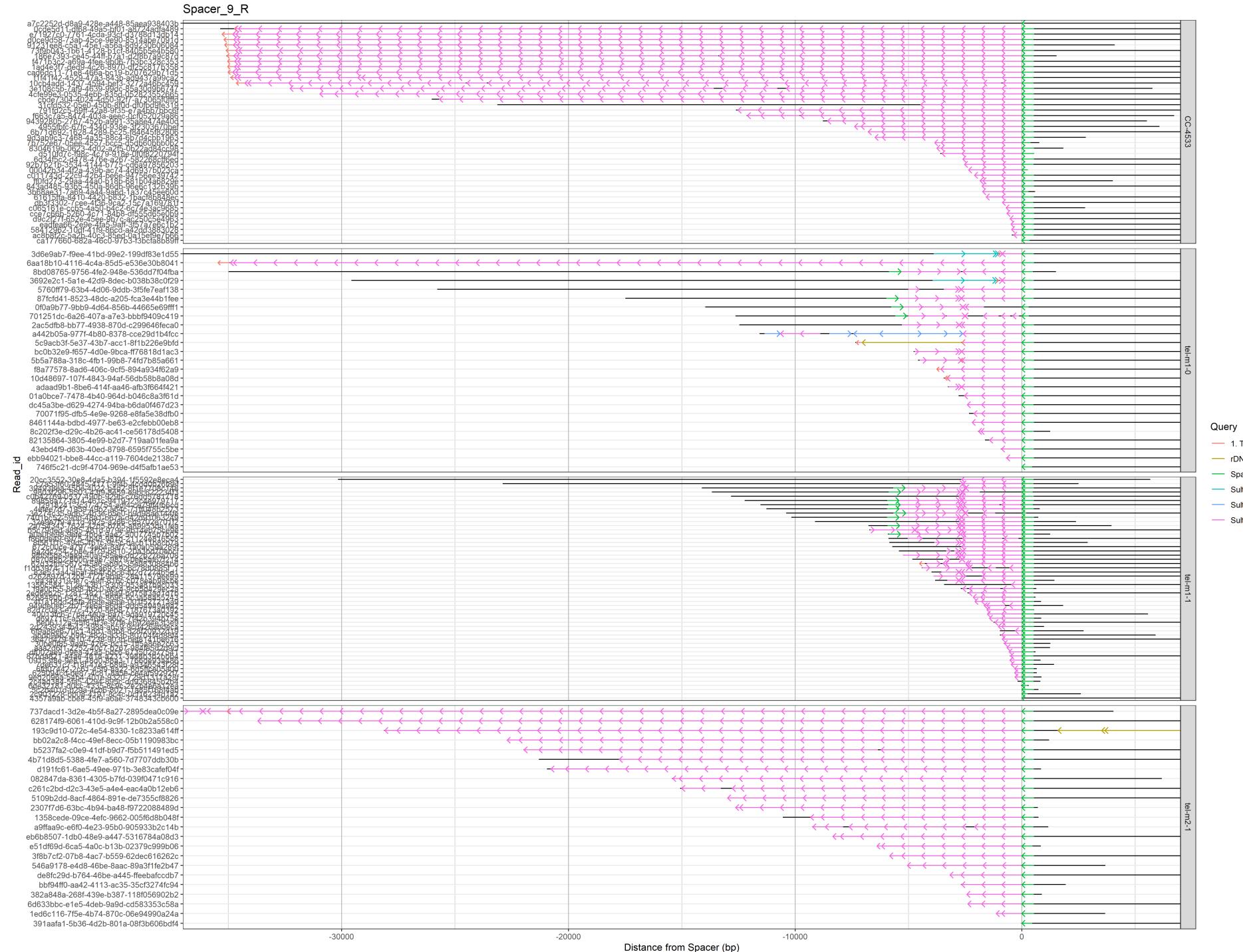




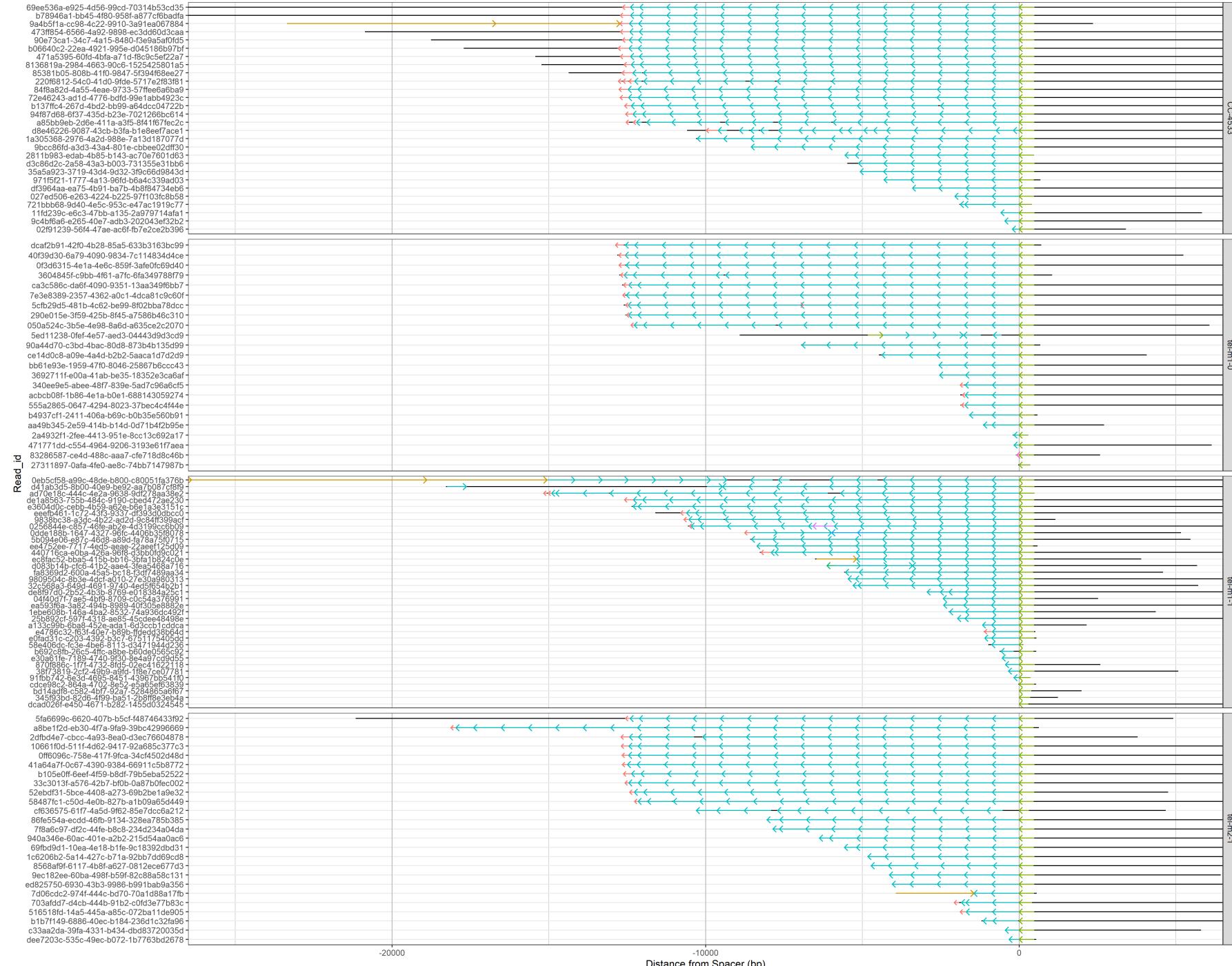


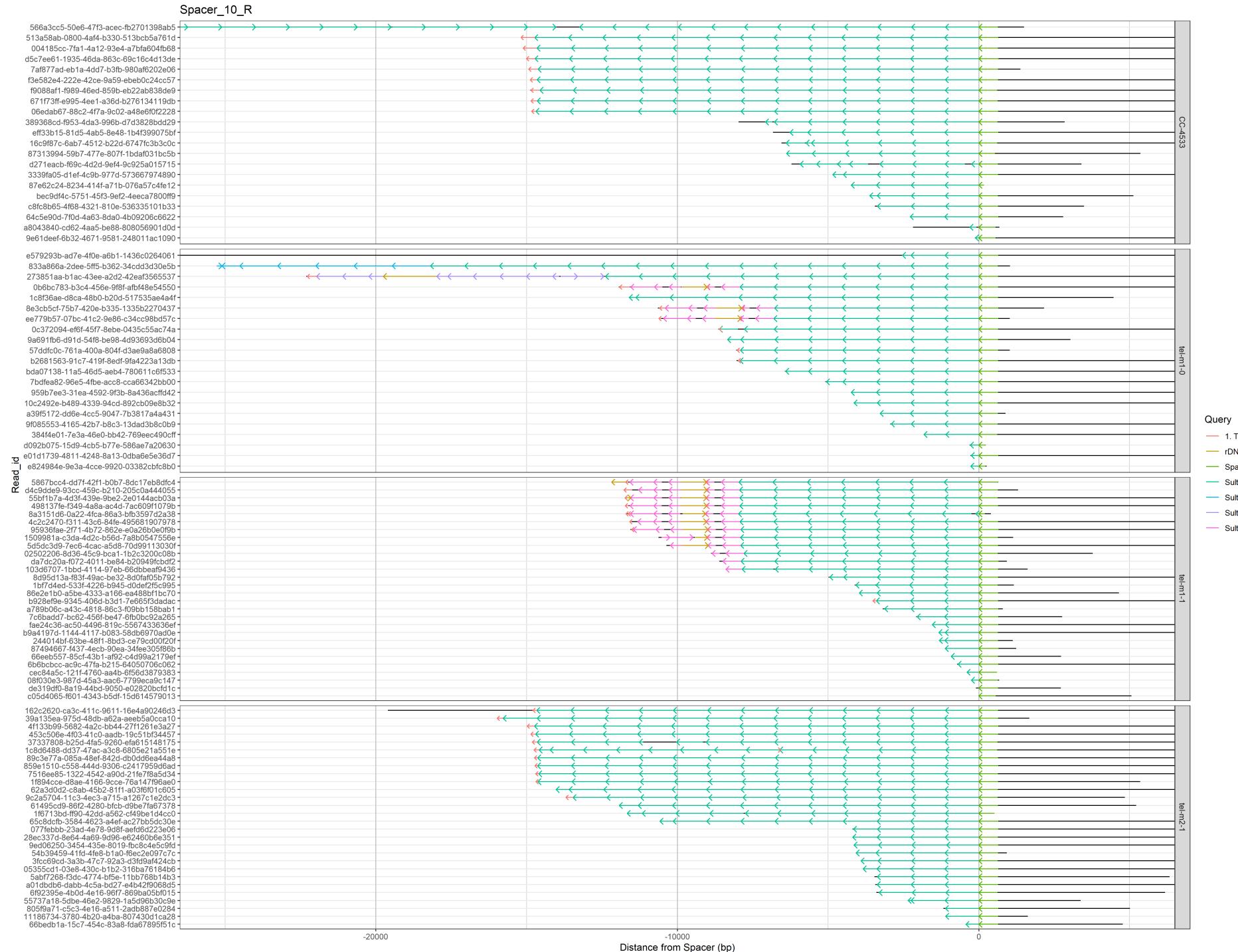




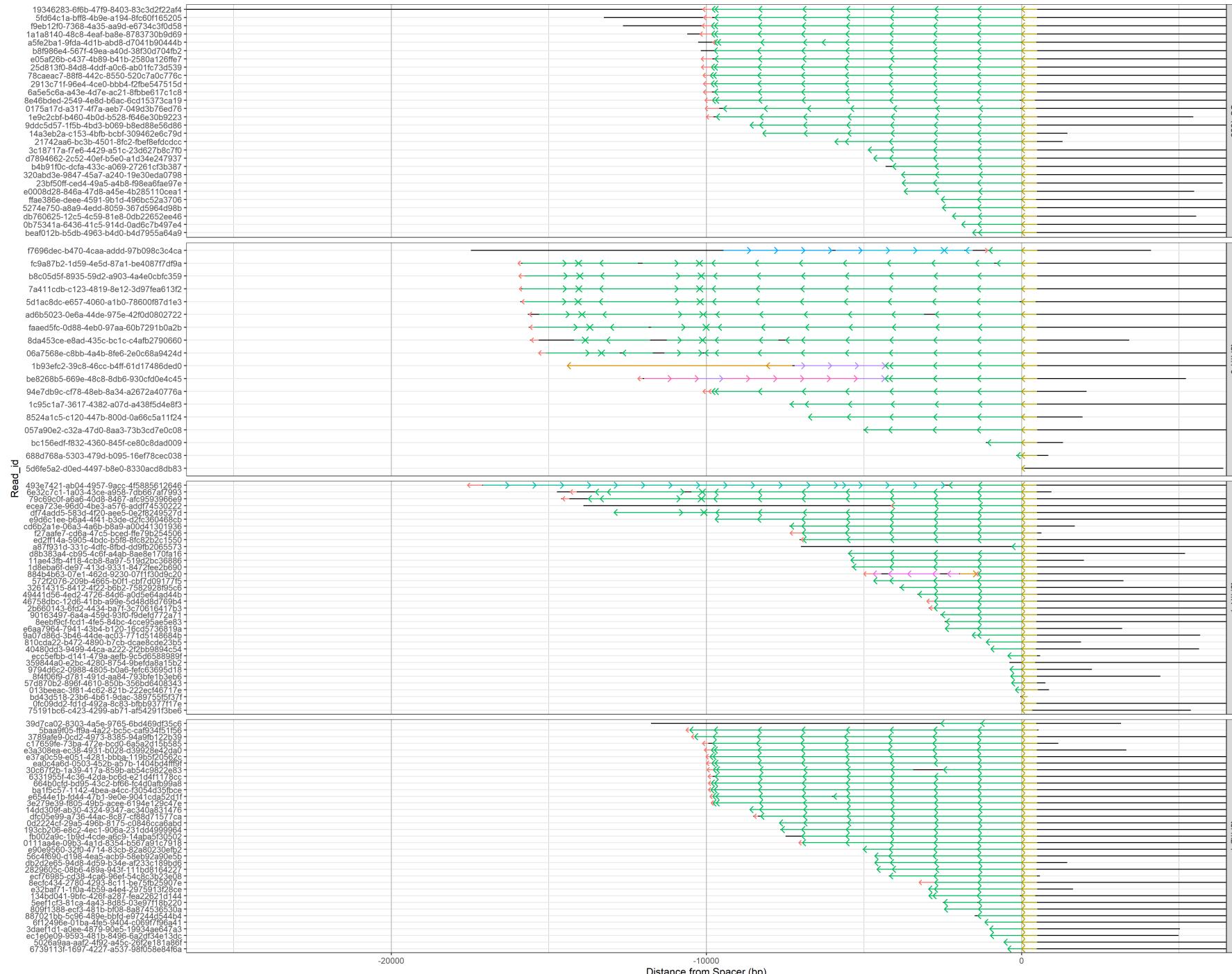


Spacer_10_L

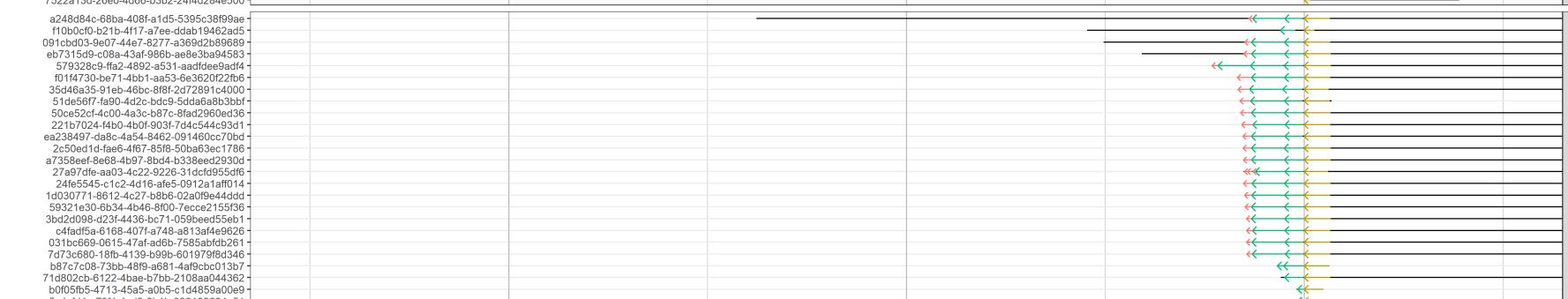
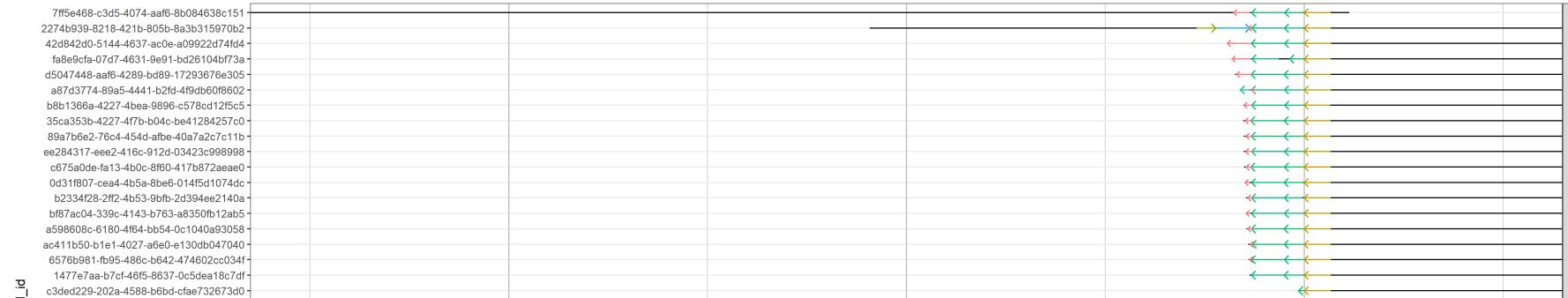
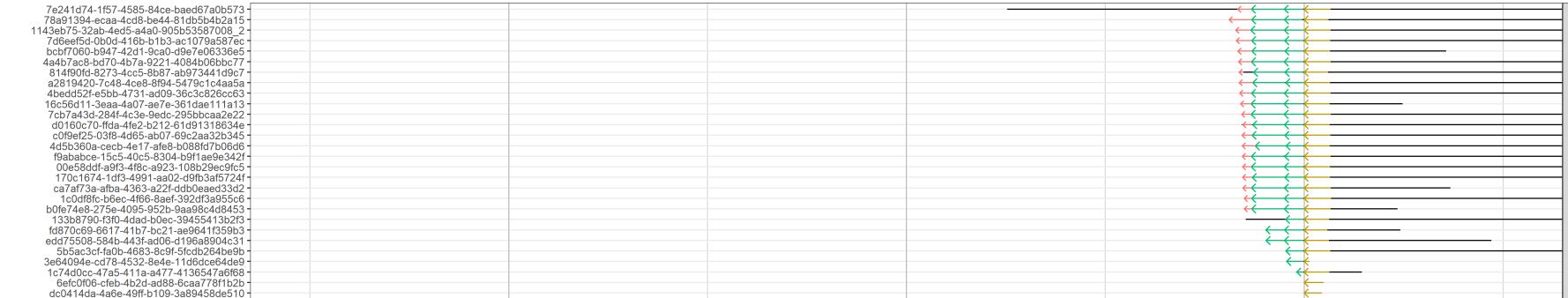




Spacer_11_R



Spacer_12_L

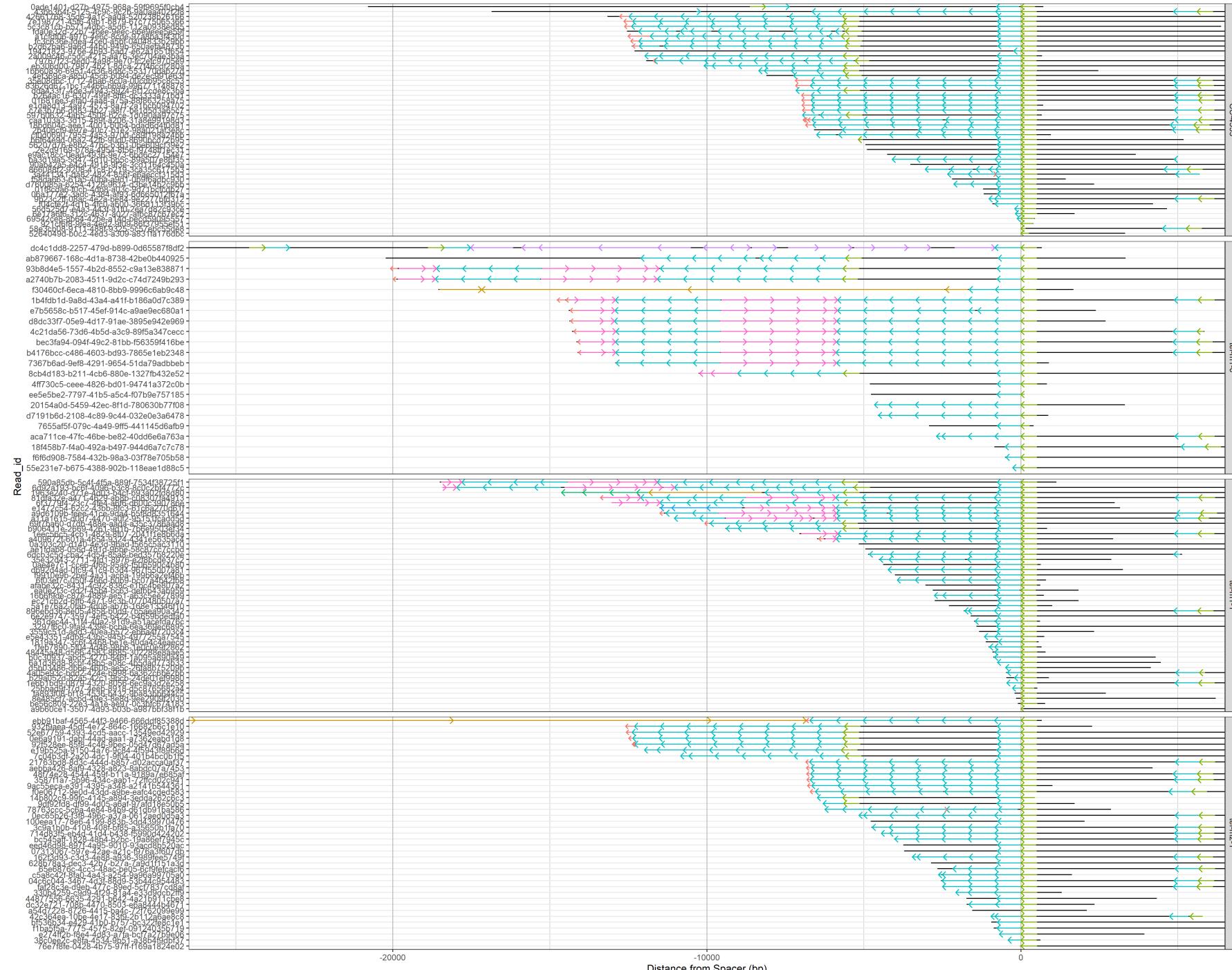


Query

- 1. Telomere
- 2. rRNA
- 3. Spacer_12_L
- 4. Spacer_15_L
- 5. Spacer_6_L
- 6. Sultan_12_L
- 7. Sultan_14_L
- 8. Sultan_15_L
- 9. Sultan_15_R
- 10. Sultan_5_R
- 11. Sultan_6_L
- 12. Sultan_7_L

Distance from Spacer (bp)

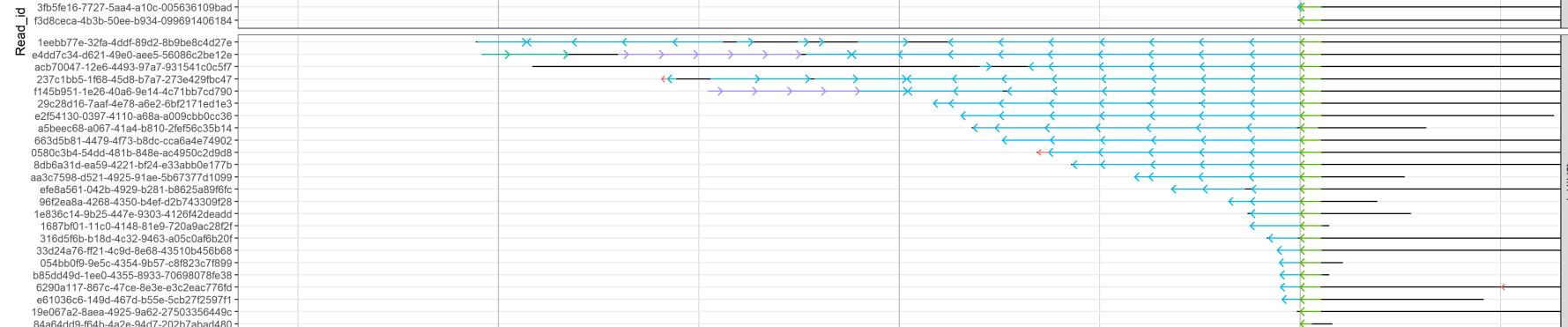
Spacer_12_R



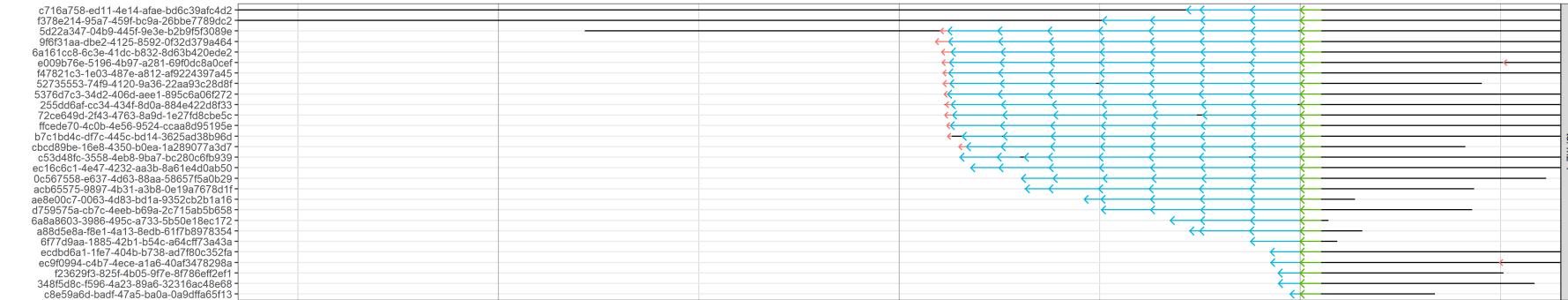
Spacer_13_L



CC-4533



tel-m1-0



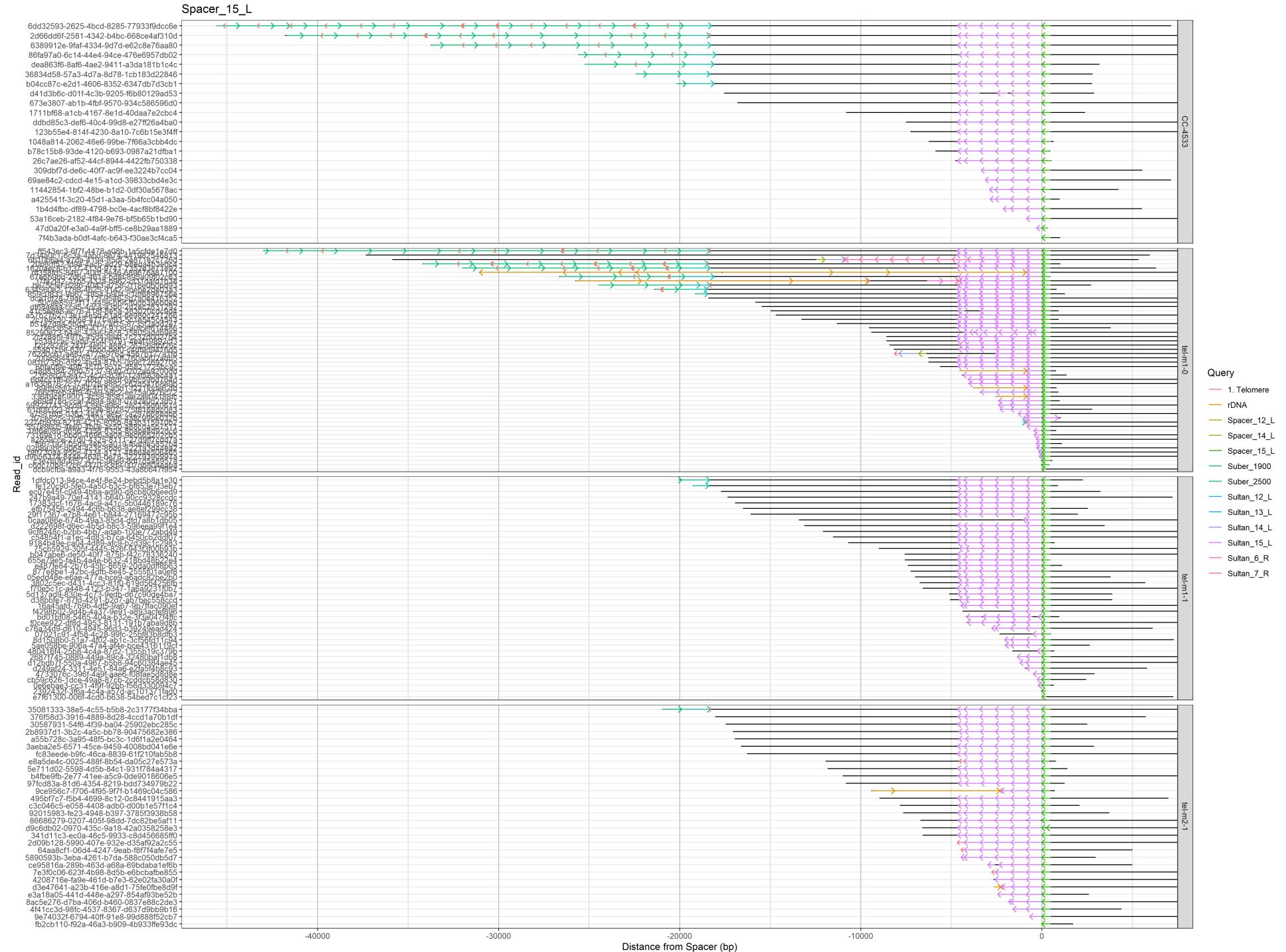
tel-m1-1

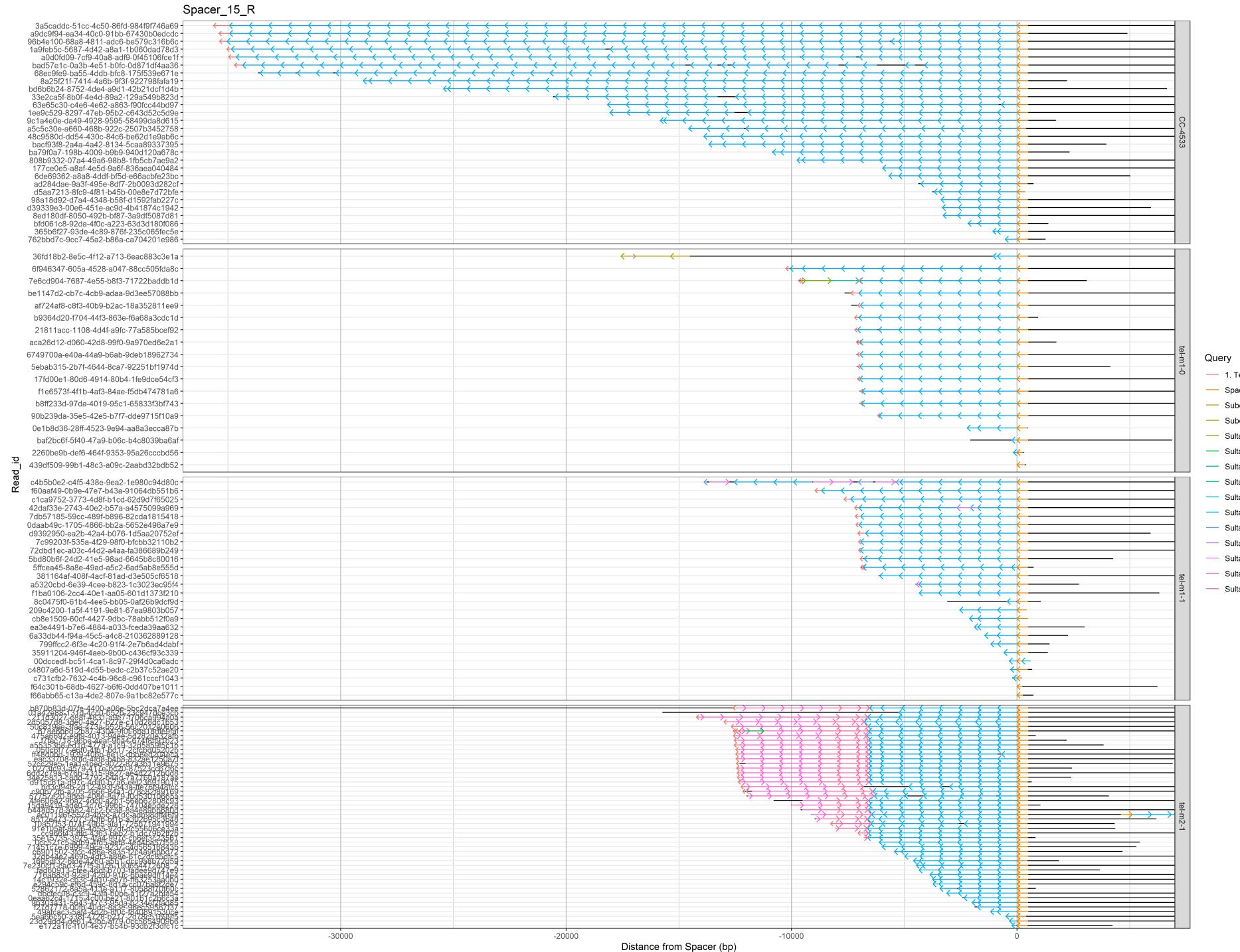


tel-m2-1

Distance from Spacer (bp)







Spacer_16_L

