



Alignment for typel-B2 self-targeting spacer

```

gcaaaatccgtgacggttgaatagtctag
|||||.|||||.|||||.|||||.
gcaaagtcactgccagatcattattctcg
agctaaaagtgttactgttgaatgattag
||.|||||.|||||.|||||.|||||.
agataaaatttatagtagtgcttgcttag
  
```

Information I want to deliver:

- 1, tniQ1 (tniQ between tnsC and cas protein) cluster together.
- 2, tniQ2(tniQ behind CRISPR array) cluster together.
- 3, tniQ 1 are close to tniQ from CRISPR tn7 systems (Type I-F tniQ).
- 4, tniQ 2 are close to tniQ from normal tn7 systems (Tn7 tniQ)

