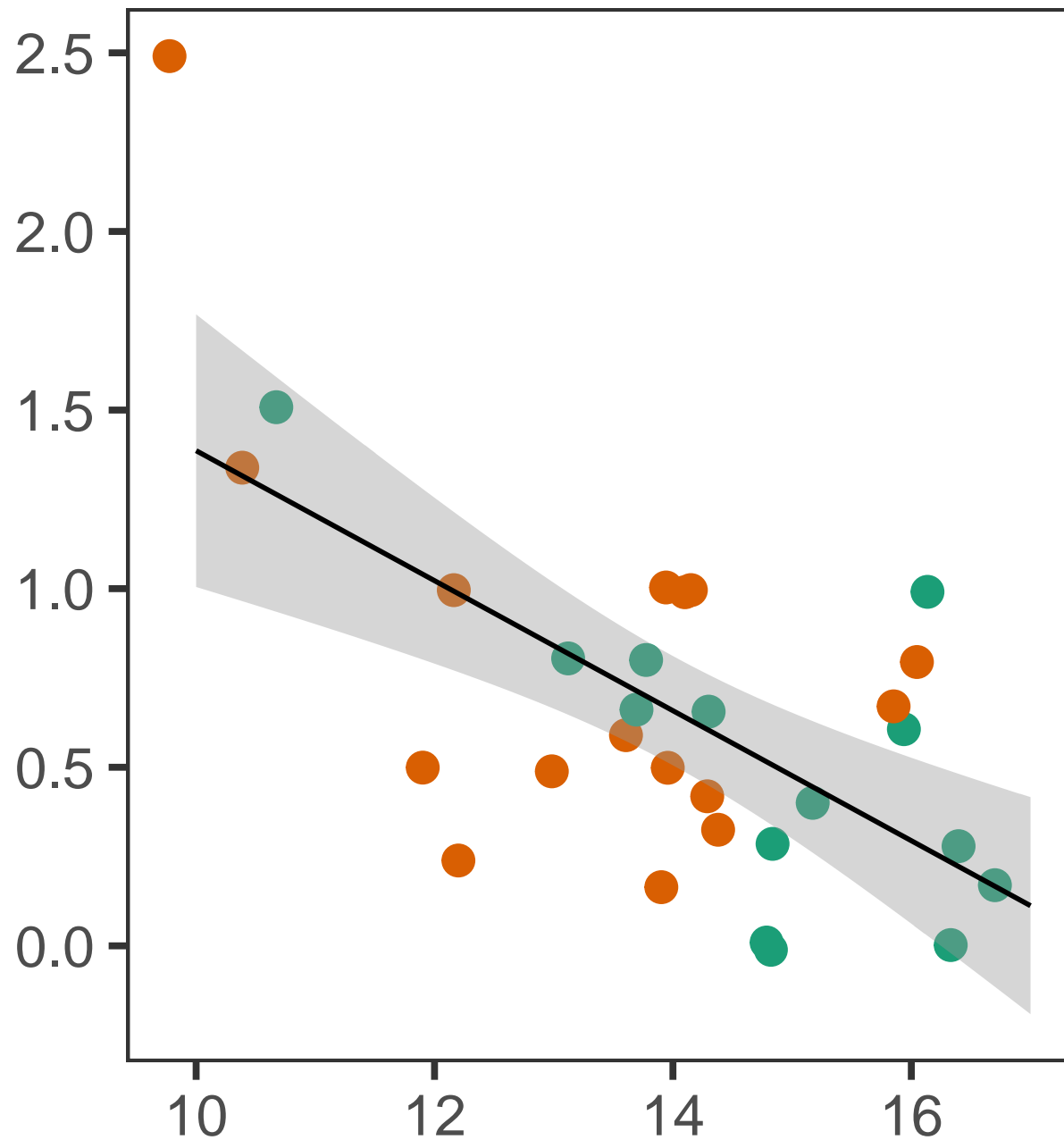


Peptide contacts,
CDR1/CDR3 ratio



PITmatch

- non-PIT-matched
- PIT-matched

TRA junction length (AA)