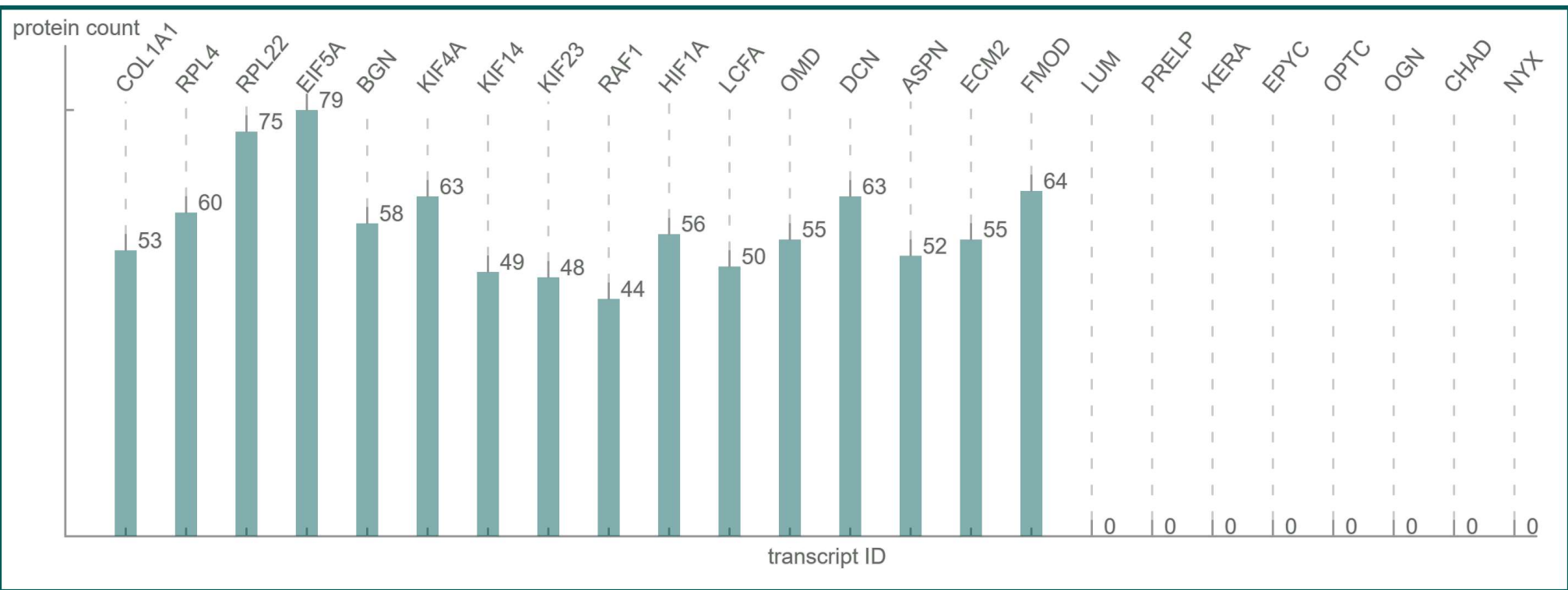


- [Home](#)
- [Settings](#)
- [Ribosomer](#)
- [Ribo-Seq output](#)
- [Protein levels](#)

Protein levels: cumulated protein counts and translation efficiencies (TE)



Translation efficiency (TE = # protein count / # transcript copy number)