

# ClustIRR

## Monthly meeting

June 2023

Recapitulation

Changes

Submission

Graph discussion

## Status after last meeting

- ▶ Pre-submission phase of ClustIRR
  - ▶ Integrate dataset into vignette
  - ▶ Submit package

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# Dataset integration

- ▶ Integrated reference dataset of Gliph2
- ▶ Demonstrated clustering with inserted ground truth
  - Reference dataset of  $10^4$  CDR3 $\beta$  sequences
  - Take random sample of  $n = 500$  CDR3s
  - Artificially enrich 20 sequences with motif *RQWW*
  - Simulate clonal expansion with two sequences :
    - ▷ *CATSRAAKPDGLAALETQYF* and *CATSRAAKPDGLAALSTQYF*
  - that get attached to the sample 15 times each
- ▶ Documented local and global clustering

# Trim flanks

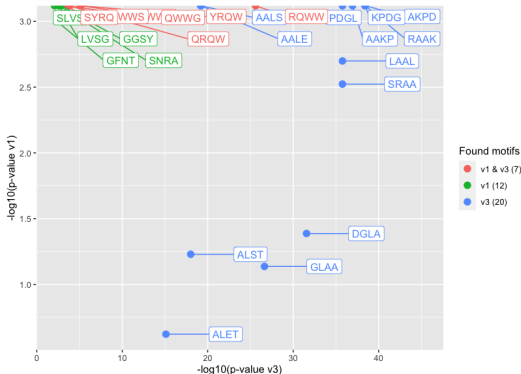
- ▶ Sequences could be "trimmed to death"
- ▶ E.g., *CASSCDRTQFV* (length 11) trimmed by  $2 * 6 \rightarrow NA$
- ▶ Solved by integrating warnings and error message
  - Warnings, if sequences are lost
  - Error message, if all sequences are trimmed

# Local clustering - v3 vs v2



- ▶ Motif *RQWW* gets found with high  $p$  by both versions
  - *QWWA*, *QWWG*, *QWWS*, *QRQW* and *YRQW* motif related
- ▶ Clonally expanded sequences also share motifs
  - Six motifs, counting only non-redundant sequences
  - 14 additional motifs, counting also redundant sequences

# Local clustering - v3 vs v1



- ▶ v1 also does not find clonally expanded sequence motifs
- ▶ But v1 finds 5 sequences not found by v2 or v3
  - *GFNT*, *GGSY*, *LVSG*, *SLVS*, and *SNRA*
- ▶ Bootstrapping related, high fdr-values both in v2 and v3
  - At least  $\text{fdr}=0.1524478$ , up to  $\text{fdr}=0.310575$



# Vignette

## Contents

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- 1 Introduction
  - 1.1 T cells
  - 1.2 B cells
  - 1.3 V(D)J genes
  - 1.4 IR sequencing
  - 1.5 Clustering
- 2 ClustIR algorithm
  - 2.1 Input
  - 2.2 Clustering
    - 2.2.1 Local clustering
    - 2.2.2 Global clustering
  - 2.3 Output
    - 2.3.1 Clustering output
    - 2.3.2 Graph output
- 3 Case study
  - 3.1 Input data
  - 3.2 Clustering
  - 3.3 Output
    - 3.3.1 Local motif clustering output
    - 3.3.2 Global CDR3 sequence clustering output
    - 3.3.3 Graph output
- 4 References
- 5 Session Info

- ▶ Added extended intro, detailed algorithm description
- ▶ Wrote additional vignette for detailed version comparison

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# Submission

- ▶ Integrate exemplary graph (optional)
- ▶ Submit package

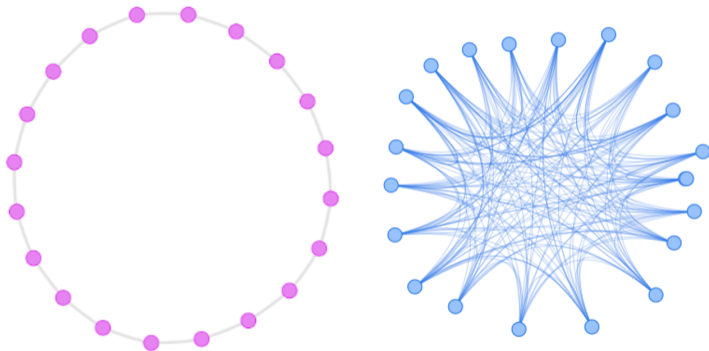
Recapitulation

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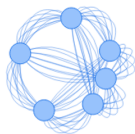
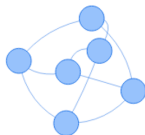
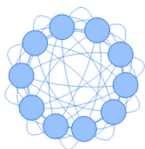
Graph discussion

# Motifs - ring vs fully connected



- ▶ Edges = shared motif, no information loss with ring (?)
- ▶ Could even be reduced to chain without information loss

## Motifs - Occluded motifs



- ▶ When two CDR3 share  $> 1$  motif, edges get reduced (left)
- ▶ Could be solved for example by using count weights

## Global similarity - long motif

- ▶ CDR3 of length 20 = essentially long motif
- ▶ Edges = similarity, enables hubs to exist
- ▶ Stop splitting into global and local, use "levels"?