### DRESDEN, IMMREP22

TCR specificity workshop July 28th- 30th

Organisers: Anne Eugster and Virag Sharma

## **Program**

## Thursday, July28th:

#### 9.30am

Anne Eugster, Short introduction Presentation round by all on site participants

Short presentations on models from participants – 15 min

# Chair: Anne Eugster

Alessandro Montemurro (Morten Nielsen), Similarity-based model (TCR-baseline) and NetTCR, (virtual)

Yoram Louzou, ERGO, (virtual)

Anna Weber (and Maria Rodriguez Martinez), TITAN and DECODE (virtual)

Barbara Bravi, Model to be announced (virtual)

Pieter Meysman, TCRex Random Forest, TCRmatch

#### <u>11am</u>

coffee break

#### 11.30am

Thierry Mora, soNNIA
Liel Cohen, SETE and TCRGP
Vadim, pMTnet
Rose, ICON
Paul, TcellMatch
Eli, DeepTCR

## ??, iSMART

??, Triplet Network and Modal Alignment Network subunits. tcrdist3

#### <u>1pm</u>

Lunch at the PKS

### 2pm

Chair: Pieter Meysman

Virag Sharma:

- -Short presentation on training- and test data preparation
- -Pieter Meysman, short presentation of the evaluation strategy
- -Introduction of discussion points regarding goals we try to address with the working group:
  - 1 What is the impact on the performance of models when including the alpha chain?

- 2 Do CDRs 1 and 2 impact predictive performance of models?
- 3 Should we do peptide or pan-specific models?
- 4 How do be best compare predictions between peptides (most relevant when defining the best peptide target for a given TCR)?
- 5 How to best construct negative data (swapped, negative controls, healthy controls, combinations)?
- 6 What baseline models to apply to demonstrate the value of "deep" models vs baseline models?
- -Release of test data and start individual work on models and short intro to evaluation

Evening:

Dinner in town

### Friday, July 29<sup>th</sup>:

Deadline to send modelling outputs: 7am

9.00am

Chair: Thierry Mora

Short presentation: Vadim Kaurnakov own model

- -Short feedback from each participant regarding the model used, technical issues?
- -Evaluation of points regarding the goals we try to address -after the results
  - 1 What is the impact on the performance of models when including the alpha chain?
  - 2 Do CDRs 1 and 2 impact predictive performance of models?
  - 3 How do we best build models: clustering or classification/epitope specific models?
  - 6 How to best construct negative data (swapped, negative controls, healthy controls, combinations)?
  - 7 What baseline models to apply to demonstrate the value of "deep" models vs baseline models?

#### 11am

coffee break

Pieter Meysman, Presentation of evaluation of results obtained by participants on each model

#### 11.30am

Continuation of the discussion

#### 1pm

Lunch at the PKS

# 2pm -4.30pm

Chair: Virag Sharma

Discussions around the used models:

- -Can we group the top models, are there similarities?
- -Can we crystallise out important parameters?
- -Preparing the ideal next workshop" Dataset required?

Discussion and wrap up

### **Evening**

Dinner individually

# Saturday, July 30th

Continuation of discussion of required, else individual meetings or free time