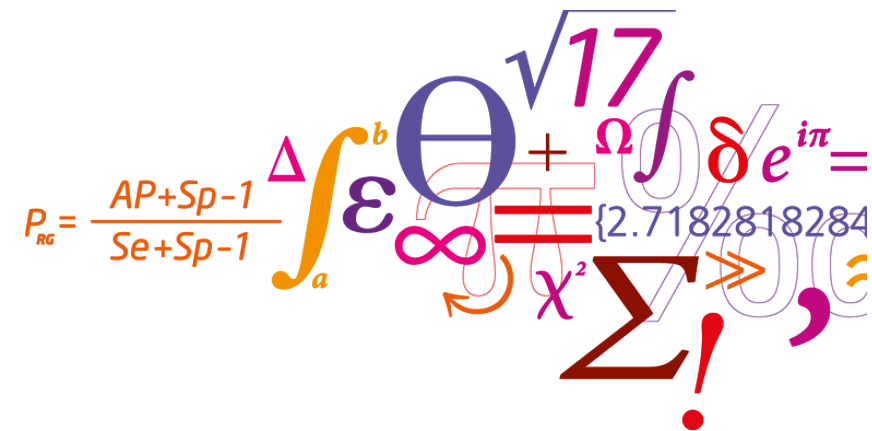


36685 Immunological Bioinformatics

January 2018

Introduction to Project Work



Introduction to Project Work

- Project time frame
 - Thursday January 11th – Thursday January 18th
- Groups
 - The predefined groups of 3-4 students will work together on the project
- Outcome
 - A project presentation (PowerPoint) presented at the exam January 19th
 - Note, you should *not* hand in a project report

Groups

- G01 - Line, Solveig and Monica
- G02 - Mikael, Jacob and Kathrine
- G03 - Sandra, Esteban, Narasimha and Taner
- G04 - Signe, Julie, Ina and Aimilia
- G05 - Christian, Mikkel and Frank
- G06 - Keith, Nadia and Jeppe
- G07 - Sofie, Natasja and Ulla

Group folders on DTU inside

DTU **INSIDE**
Leon Eyrich Jessen
DTU Bioinformatics

MY COURSES AND GROUPS
TOOLBOX
STRUCTURE AND RULES
ACADEMIC OFFERS AND GUIDANCE
SOCIAL AND POLITICAL
EDUCATIONAL ADMINISTRATION

Front page > DTU Bioinformatics > 36685 Immunological Bioinformatics Jan 18 > [File sharing](#) ★

36685 IMMUNOLOGICAL BIOI...

- List of participants
- Messages
- Calendar
- File sharing
- Study activity req.
- Links
- Assignments
- Home page
- Polls
- Course Base
- Wiki setup
- Adobe Connect
- Reporting

File sharing: Student folder

Tree structure
Upload file
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Your browser supports drag and drop upload. Drag files onto the page to upload them to this folder.

Path: [Filesharing](#) / [Student folder](#) /

<input type="checkbox"/> NAME	AUTHOR	LAST MODIFIED	
<input type="checkbox"/> Parent folder (Top)			
<input type="checkbox"/> G01 - Line, Solveig and Monica	Leon Eyrich Jessen	03/01/18 12:50	...
<input type="checkbox"/> G02 - Mikael, Jacob and Kathrine	Leon Eyrich Jessen	03/01/18 12:52	...
<input type="checkbox"/> G03 - Sandra, Esteban, Narasimha and Taner	Leon Eyrich Jessen	03/01/18 18:48	...
<input type="checkbox"/> G04 - Signe, Julie, Ina and Aimilia	Leon Eyrich Jessen	03/01/18 12:52	...
<input type="checkbox"/> G05 - Christian, Mikkel and Frank	Leon Eyrich Jessen	03/01/18 12:53	...
<input type="checkbox"/> G06 - Keith, Nadia and Jeppe	Leon Eyrich Jessen	03/01/18 12:53	...
<input type="checkbox"/> G07 - Sofie, Natasja and Ulla	Leon Eyrich Jessen	03/01/18 12:53	...

Delete selected
Move selected
Download selected as zip

Exam

- Exam is on Friday January 19th 9 – 17 in building 208, room 062 (Where we are now)
- Examinators will be Morten and Leon
- The group will give the presentation ~10 minutes followed by 5 min of project questioning (Note, everyone is responsible for every aspect of the group project)
- Following the presentation, there will be an individual oral exam in the project and course curriculum, also ~15 minutes (Total time per group will be ~1 hour)
- In case you would like a specific exam time during the day (for a good reason), please contact Leon (jessen@bioinformatics.dtu.dk)

Project Examples

- **Project I – Pathogen vaccine**

- a) Choose a single protein from a human pathogen, which would be the best suited for a vaccine considering B-cell epitopes
 - b) Make a peptide based t-cell vaccine with both broad HLA* and pathogen coverage. E.g. a polytope consisting of class-I and class-II epitopes. Select from all proteins in the organism. Consider the processing of the final polytope to avoid the presence of neo-epitopes and if relevant check to similarity to self
- **You are free to limit your focus to a special population and genotypes, but should justify your choices*

Project Examples

- **Project II – Cancer vaccine**

- As project I, but aimed at cancer specific proteins, e.g. in testis cancer

Project Examples

- **Project III – Cancer Immunotherapy**

- Humanization project: CAR T-cell humanisation and de-immunisation
- Similar to the “Anti-CD19 CAR T-cells” exercise you did
- E.g. using the tabhu server in the context of eliminating t-cell epitopes (cd4+ class II binders)
- (Don’t just follow the questions from the exercise – Make the project your own)

Project Examples

- **Project X**

- Any other great ideas that will cover several of the methods introduced are also welcome!
- You should however include a part about HLA immune activation

Project Examples

- Examples of pathogens with fully sequenced genomes:
 - HIV
 - HCV
 - HPV
 - Ebola/Marburg
 - Smallpox
 - Mycobacterium tuberculosis
 - Influenza
 - Chlamydomonas reinhardtii
 - Measles virus
 - Salmonella typhi

Guide to Presentation

- The IMRAD approach
 - Introduction
 - Materials and methods
 - Results And Discussion
- Brief and clear (limit amount of text on each slide)
- Be sure to think about the information you wish to convey with each slide
- Note, everyone is responsible for every aspect of the project
- Remember to include references, e.g.
 - **NetMHCpan-4.0: Improved Peptide–MHC Class I Interaction Predictions Integrating Eluted Ligand and Peptide Binding Affinity Data.** Jurtz, V., Paul, S., Andreatta, M., Marcatili, P., Peters, B. and Nielsen, M. J. Immunol. (2017)

Project supervision

- There is a very limited time frame for the project, so you need to start the project today! (Thursday)
- If you get stuck in the project, you should as a point of reference try to solve the challenge in the group
- If this is not possible, then contact:
 - Protein structure and antibodies
 - Kamilla (kamjen@bioinformatics.dtu.dk)
 - Leon (jessen@bioinformatics.dtu.dk)
 - T-cell epitopes and HLAs
 - Leon (jessen@bioinformatics.dtu.dk)
 - Morten (mniel@bioinformatics.dtu.dk)

Course Evaluation

- I will soon open for course evaluation
- Please spend the 5-10 minutes it takes to evaluate
- Your feedback is extremely valuable!

Enjoy the project work! 😊

