

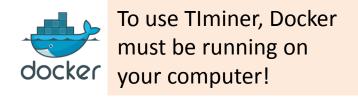
IO17 | Large Scale Bioinformatics for Immuno-Oncology

Neoantigens: exercise 1

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We have access to single-end RNA-seq data from a melanoma patient:

Patient 1 RNAseq.fastq

Perform HLA typing with Tlminer (function TlminerAPI.executeOptitype) using "Patient_1" as subject ID)

Hint: you can perform parallel computation using up to 8 CPUs.

Once you get the results, answer the questions related to this exercise at:

https://b.socrative.com/login/student/

How to access the questions on Socrative (1)

- Access Socrative with student access at: https://b.socrative.com/login/student/
- 2. Join the IO17 room



Student Login			
Room Name			
1017			
	JOIN		
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How to access the questions on Socrative (2)

3. Enter your name and click "done" to start the quiz

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