Practical - Fellay Lab
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<table>
<thead>
<tr>
<th>Cursus</th>
<th>Language</th>
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<tbody>
<tr>
<td>Approches moléculaires du vivant</td>
<td>English</td>
<td>Obl.</td>
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**Sem.**

**Type**

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<tr>
<th>Language</th>
<th>Credits</th>
<th>Session</th>
<th>Exam</th>
<th>Workload</th>
<th>Hours</th>
<th>Lecture</th>
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<tbody>
<tr>
<td>English</td>
<td>1</td>
<td>Project report</td>
<td>30h</td>
<td>24</td>
<td>6</td>
<td>18</td>
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**Frequency**

Every year

**Remarque**

3-day Block course, every year in January. To register, contact EDMS Administration

**Summary**

Next time in January 2019 Exploration of human genetic variation in large-scale sequencing data.

**Content**

Students will be asked to access large-scale data from sequenced human exomes, to focus on one gene (or pathway) of interest, and to catalog and describe all variants present in the study population. The potential associations between the identified variants and clinically relevant phenotypes will also be explored.

**Keywords**

Human genomics, bioinformatics.

**Learning Prerequisites**

Recommended courses
Knowledge in genetics and bioinformatics.

**Assessment methods**

Project report.

**Resources**

Websites
- http://fellay-lab.epfl.ch/