## Analysis of high-throughput sequencing data



IT4Innovations national0\$%@&@ supercomputing center\$@0@&1@%



When November 27 - 28, 2014

Venue VŠB – Technical University of Ostrava, IT4Innovations building, room 450

TutorPaolo Ribeca<br/>(The Pirbright Institute, UK and Centro Nacional de Análisis Genómico, Spain )

| Thursday    | November 27, 2014                             |
|-------------|---|
| 09:30–10:00 | registration                                  |
| 10:00–11:30 | Basics of high-<br>throughput sequencing      |
| 11:30–13:00 | time for lunch                                |
| 13:00–14:30 | Alignment<br>(basics; mapping)                |
| 14:30–15:00 | coffee break                                  |
| 15:00–16:30 | DNA-seq mapping<br>hands-on                   |
| 16:30–17:00 | coffee break                                  |
| 17:00–18:00 | Mapping, revisited<br>(algorithms; protocols) |

| Friday      | November 28, 2014                                      |
|-------------|--|
| 09:00–10:30 | RNA-seq mapping<br>hands-on                            |
| 10:30–11:00 | coffee break   |
| 11:00–12:45 | Downstream<br>analysis. Differential<br>quantification |
| 12:45–14:00 | time for lunch   |
| 14:00–15:30 | Basics of assembly                                     |

More information and registration at: http://prace.it4i.cz/BIO-11-2014

Deadline for registration: November 20, or exhausted capacity

Number of attendees: 20

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