Precision medicine: NGS variant analysis and interpretation for translational research

Exercise 4: Running PanDrugs

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- Query PanDrugs using the Genes section
- Load Example 3 (Genes Involved in PI3K-AKT-mTOR pathway)
- Query the database with these genes

Which are the suggested therapies?

Which are the best candidates?

Which are the evidences for these best candidates?

 Is there any drug suggested by a gene that is discarded by another?

- Query PanDrugs using the Gene Ranking section
- Load Example file (Expression data for the top 500 up-regulated genes in a lung adenocarcinoma patient with EGFR alteration from TCGA)
- Query the database with this genes

- Search patient (TCGA-91-6847) in cBioPortal to visualize the alterations.
- Which are the suggested therapies?
- Which are the best candidates?
- Is there any drug approved for the same cancer type?
- Which are the evidences for the drugs that rank higher in the assignments?

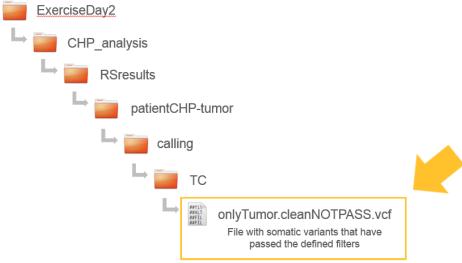
- Query PanDrugs using the Genomic Variant section
- Load Example file (vcf file with alterations in exome of melanoma patient with mutant BRAF from TCGA)
- Query the database with this file

- Search patient (TCGA-BF-A1PU) in cBioPortal to visualize the alterations.
- Which are the suggested therapies?
- Is there any drug suggested by a gene that is discarded by another?
- Is there any drug approved for the same cancer type?
- Is there any drug in cancer clinical trials for this cancer type?
- Could be used a drug repositioning approach?

Query PanDrugs using the Genomic Variant section

Load the vcf file without annotations from the exercise of CHP patient

Path to the input file



Query the database with this file

- Which are the suggested therapies?
- Is there any drug suggested by a gene that is discarded by another?
- Is there any drug approved for the same cancer type?
- Is there any drug in cancer clinical trials for this cancer type?
- Could be used a drug repositioning approach?