

PO21: Precision Oncology Course

Exercise: Running PanDrugs

www.pandrugs.org



@PanDrugs_CNIO

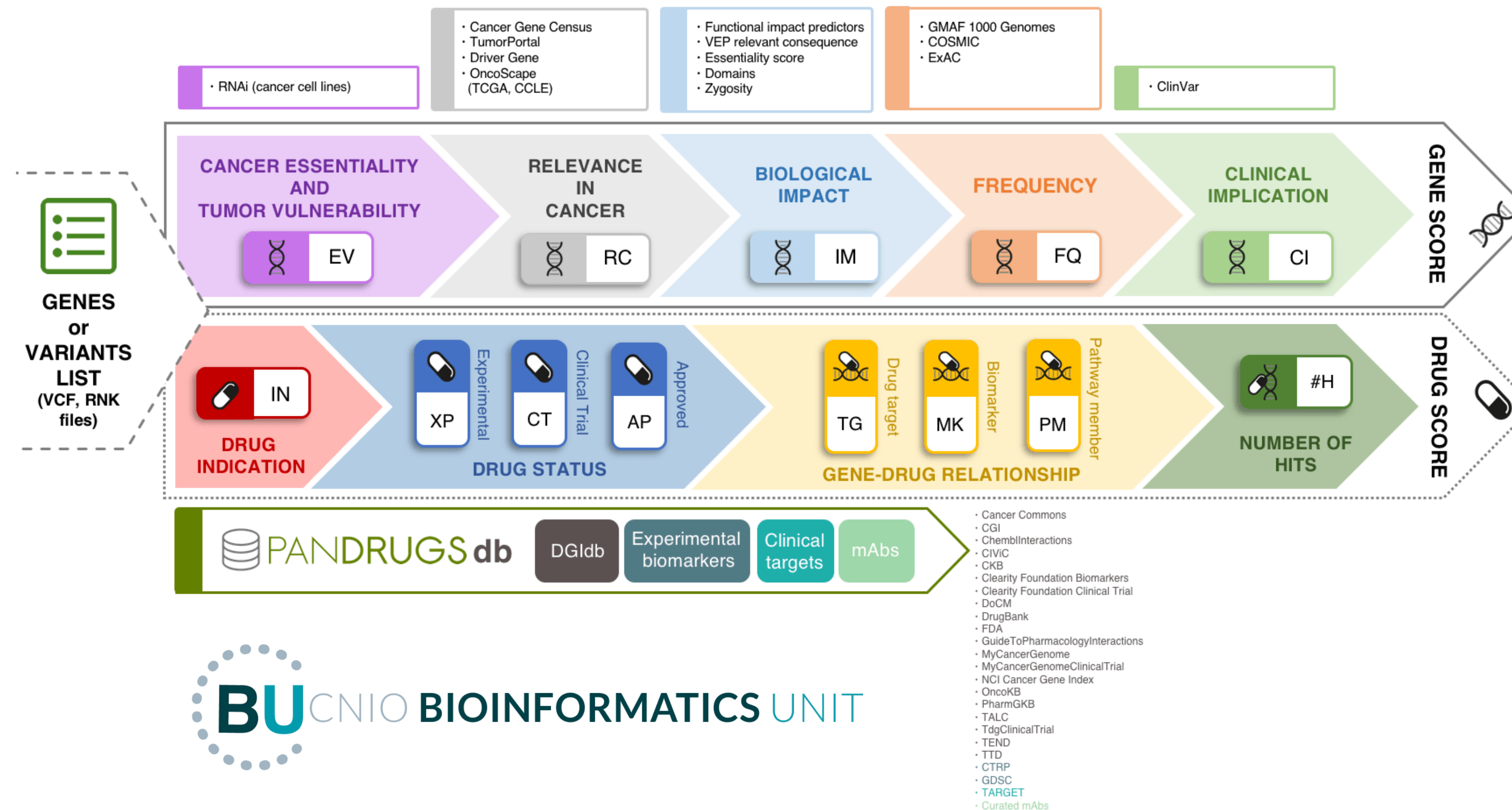
Welcome to PANDRUGS

A novel method for prioritizing therapies using individual
genomic data

Query! ✓



PanDrugs is a web tool for **prioritizing cancer therapies** based on **individual genomics data**.

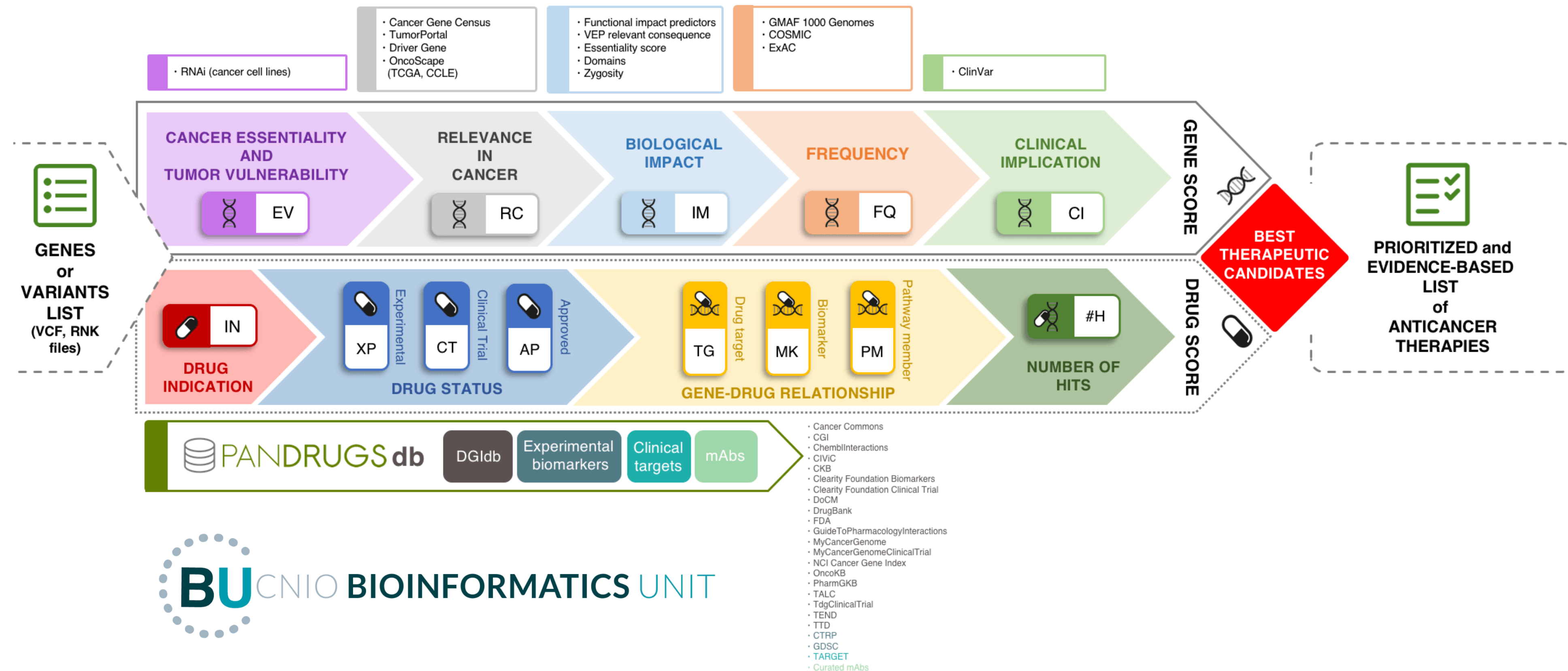


- **GScore (0-1):** Measures the **relevance of genetic variants in cancer**.
- **DScore (0-1):** Measures the **drug efficacy** against different targets.

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Based on these two scores, PanDrugs outputs a ranking of the **best therapeutic candidates** for a particular patient.

Input 1: Gene list

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

Input 1: Gene list

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- Are all the genes in PanDrugs? How many drugs are in each approval status?
- Which is the most abundant mechanism of action?
- Which are the best therapeutic candidates?
- What are the evidences for these best therapeutic candidates?
- Is there any drug suggested by a gene that is discarded by another?

Input 2: Gene ranking

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

Input 2: Gene ranking

PANDRUGS

- **How many genes are in PanDrugs?**
- **Which are the best candidates according to PanDrugs?**
- **What are the evidences for the drugs that rank higher in the assignments?**
- **Is there any drug approved for the same cancer type?**

Input 3: VCF

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1. Query PanDrugs using the Genomic Variant section
2. Load Example file (VCF of melanoma patient with **mutant** **BRAF** from TCGA)
3. Query the database with this file

Input 3: VCF

PANDRUGS

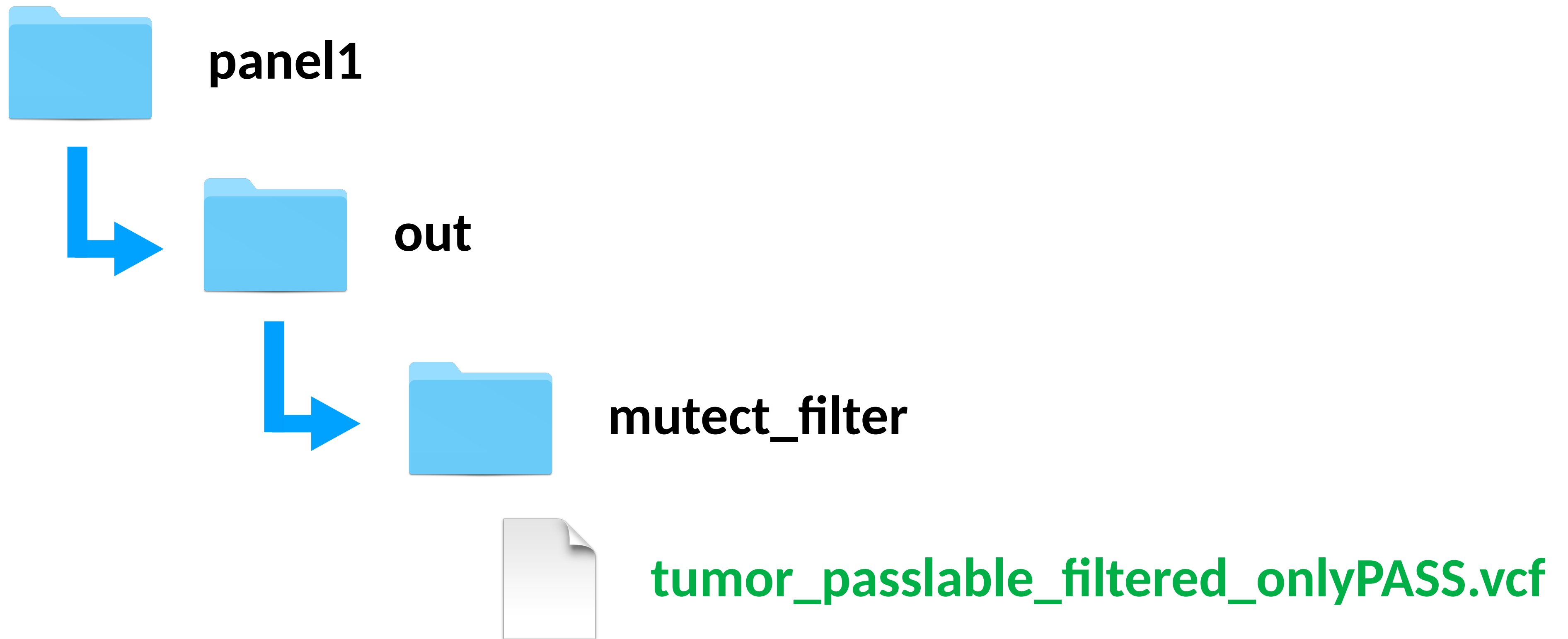
- Which are the suggested therapies? Do they match the specific alterations?
- What are the evidences for the gene with the highest GScore?
- Is there any drug suggested by a gene that is discarded by another?
- Is there any drug approved for the same cancer type? Is there any drug in cancer clinical trials for this cancer type?
- Could a drug repositioning approach be used?

Input 4: VCF

PANDRUGS

1. Query PanDrugs using the Genomic Variant section
2. Load the VCF without annotations from the exercise of the **CHP patient**
3. Query the database with this file

Input 4: VCF



Input 4: VCF

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- 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 a drug repositioning approach be used?