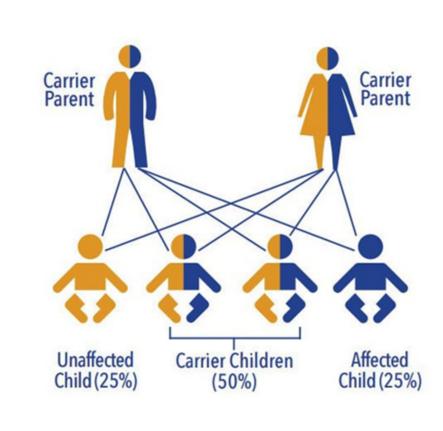
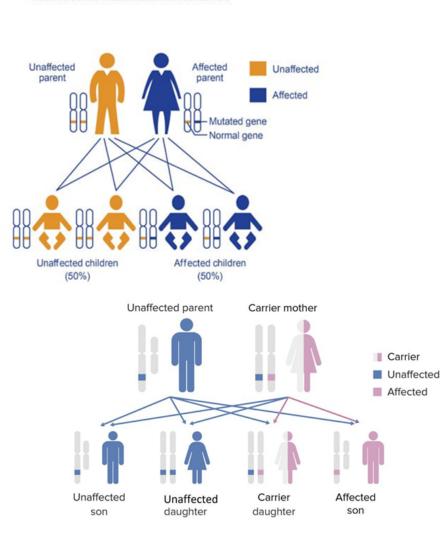
#### COUPLE CARRIER SCREENING

Carrier screening aims to determine the carrier status of healthy parents for recessive disorders, including X-linked recessive diseases. Even the most complex scenarios are well covered in a single carrier screening filter in ADVAT Pipeline so that cases can be effortlessly solved in a matter of time.

#### 1.AUTOMATED SCREENING FOR RECESSIVE DISORDERS

A single-click workflow considers factors such as the affection status and genotype so that only variants that fit the segregation pattern for carrier screening are displayed. Therefore, no manual work in ruling out variants based on segregation.





Autosomal dominant inheritance

#### 2.SCREEN CASES WITH A FAMILY HISTORY EFFORTLESSLY

Screen cases with a family history swiftly by recording the clinical information. In cases where family history and phenotypes are known, our proprietary HPO similarity score helps a great deal in identification of causative variants. Prioritize the list of variants with respect to the score and focus on the most relevant variants first.

#### 3.VIRTUAL PANELS TO APPLY ALL TIERS

ADVAT have cutomized gene panel that has been designed by the laboratory or one that has been recommended by organizations such as ACMG and ACOG.

2271-genes cover our carrier screening panel targets the full coding region of all genes, enabling the analysis of >36,000 nonbenign ClinVar variants for single-nucleotide variants (SNVs), insertions and deletions (indels).



## clence Summary. The stop gained NR\_000137 AlfAH): c709ChTip R237\* is observed in 1/15274 (0.0065N) alleles from individuals of poom AD Genomes v3 Latino background in the genomAD Genomes v3 distant (Genome Appropriation Distalbase et al., 2000, but was not seen in the homorapposis state. The variant was added to double as x 1269503015 in version 144. This variant was found in Cliniviry (variant exast double and observed in 1/15274 (0.0065N) alleles from individuals of promise production of the production of th

# ACMG Scoring Scored Criteria: PM2 PVS1 PP5 Probability of Pathogenic: 84.6% - Predicted Classification: Pathogenic Classification: Pathogenic ©

ADVAT Pipeline have facilitates clinical variant interpretation in concordance with the ACMG standards and guidelines. The evidence for classification becomes part of our ACMG knowledge base.

**4.ACMG CALCULATOR IN ADVAT** 

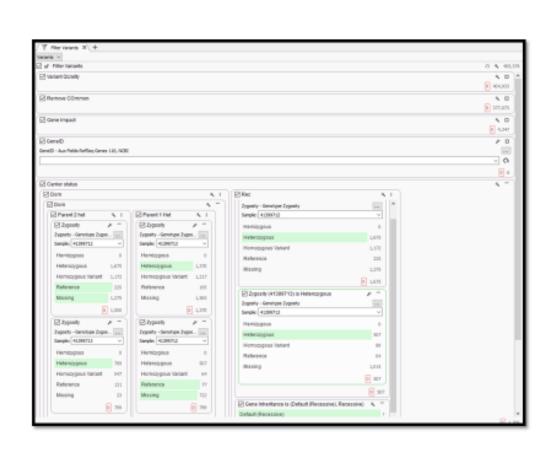


### EXPANDED CARRIER SCREENING GENES

- Carrier panel genes (2271)
- X- linked genes (16)
- Pan ethnic gene (12)
- Male infertility genes (126)
- Female Infertility genes (144)
- RPL genes (38)
- PCOS genes (15)
- Primary Ovarian insufficiency genes(25)



VarSeq Couple carrier Workflow-Major filter Steps



- Step1: Variant Quality
- Step 2: Remove Common Variants
- Step3: Gene Impact
- Step4: Gene panels -Phenotype
- Step5:Carrier Status -Inheritance

