

Case reports

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Clinical features. Phenotype. GeneReviews®

Test requested. Check main genes. See if you can filter other genes out

Phenotype vs Gene vs Variant

ClinGen expert panel

Other parameters, variant type, MAF, ClinVar,
In-silico protein function predictors

Classify Variant using ACMG-
AMP and Report

NGS test

- NGS method = Targeted panel / WES
- Phenotype
- Only proband
- Sanger only when necessary
 - Parental DNA
 - Poor coverage
- Clin Gen expert panel
- ACMG-AMP
- Analysis = Phenotype

Example: Case 1

- Six month old female patient presenting with failure to thrive, Webbed neck, widely spaced nipples, low set ears and dysmorphic features.
- Cardiac assessment showed pulmonary valve stenosis and atrial septal defect
- DNA testing for Noonan syndrome was requested.

<https://www.ncbi.nlm.nih.gov/books/NBK1124/>

On GeneReviews®, check genes on the Molecular genetic test table

Go back to the excel file and find genes associated with the disorder and filter accordingly.

Excel file: Example case 1

- Annotated data
- NCBI/GeneReviews[®] link for the disorder
 - Phenotype details
 - Gene details
- Find gene associated with the phenotype
- Go back to the Excel file
- Prioritise potential variant
- Apply ACMG-AMP rules where you can

Example: Case 2

- Suspected to have neurodevelopmental genetic disorder of unknown cause and presented with the following clinical
- features: atrial septal defect and ventricular septal defect, macrocephaly,
- developmental delay, cryptorchidism, hypoplastic nails and dysmorphic features
- including a broad chest, short palpebral fissures, hooded eyelids and cup shaped ears.

<https://www.ncbi.nlm.nih.gov/medgen/934663>

Excel file: Example case 2

- Annotated data
- NCBI/GeneReviews® link for the disorder
 - Phenotype details
 - Gene details
- Find gene associated with the phenotype
- Go back to the Excel file
- Prioritise potential variant
- Apply ACMG-AMP rules where you can