



Triple Diagnosis established by Whole Exome Sequencing

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Whole exome sequencing (WES) was requested to determine the possible genetic cause of symptoms (table 1, figure 1) for a 37-year-old Lebanese patient. WES performed on Illumina Platform only for the index (Nextera Rapid Capture Exome Kit, ~95% of target bases were covered at ≥10x). All disease-causing variants reported in HGMD®, in ClinVar or in CentoMD® (class 1), as well as all variants with minor allele frequency (MAF) of less than 1% in ExAC/gnomAD were considered. Using WES as a diagnostic tool, three different molecular diagnoses (table 2) have been discovered for this patient with a complex phenotype (table 3).

| | FEATURES |
|---------------------|--|
| DYSMORPHIC FEATURES | Hypertelorism |
| | Microphthalmia |
| | Blepharphimosis |
| | Epicanthus |
| | Short palpebral fissures |
| | Broad nasal root |
| | Long nose |
| | Hypoplastic alae nasi |
| | Thick vermilion of upper and lower lips |
| | Mild mandibular prognathism |
| DENTAL | Caries |
| | Hypodontia |
| | Microdontia |
| NEUROLOGIC | Developmental delay |
| | Intellectual Disability |
| | Hydrocephaly |
| | Seizures (clonic) |
| SKIN | Smooth, thin, dry skin |
| | Hypotrichosis (scalp) |
| SKELETAL | Hypoplastic ribs first ribs from both side |
| | Brachydactyly |
| | Radio cubital synostosis |
| | Delayed bone age |
| GENITAL | Micropenis |
| | Ectopic testis |

Table 1: Clinical Findings

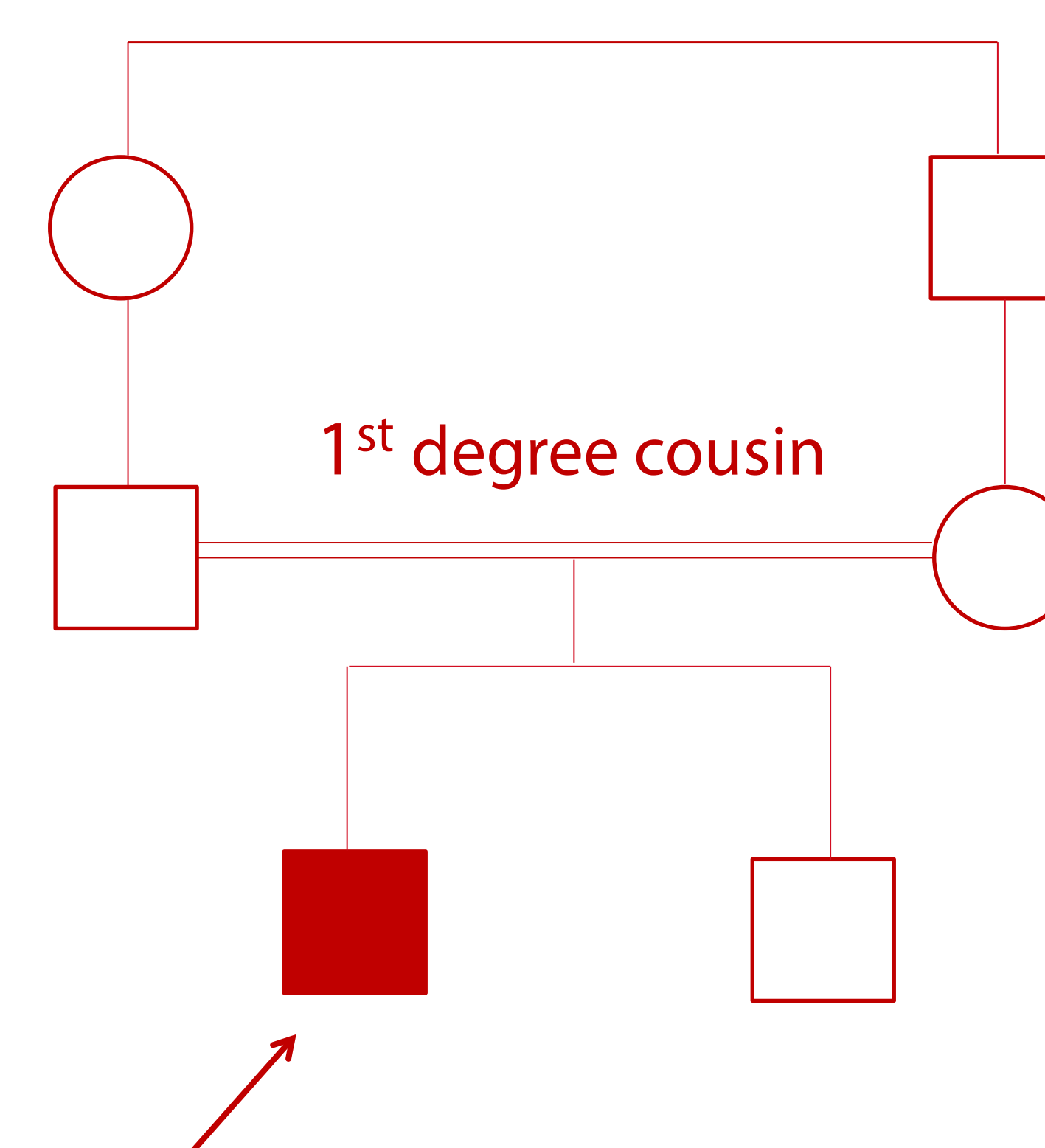


Figure 1: Pedigree

| SYSTEMS | FEATURES | THOC6 | EDAR | PTCH2 |
|-------------|--|-------|------|-------|
| Dysmorphism | Short palpebral fissures | + | | |
| | Broad nasal root | | | + |
| | Long nose | + | | |
| | Thick vermilion of upper and lower lips | + | | |
| | Mild mandibular prognathism | | | + |
| Dental | Caries | + | | |
| | Hypodontia | | + | |
| | Microdontia | | + | |
| Neurologic | Developmental delay | + | | |
| | Intellectual Disability | + | | + |
| Skin | Smooth, thin, dry skin | | + | |
| | Hypotrichosis (scalp) | | + | |
| Skeletal | Hypoplastic ribs first ribs from both side | | | + |
| | Brachydactyly | | | + |

Table 3: Features explained by WES Results

| GENE (TRANSCRIPT) | NUCLEOTIDE [PROTEIN] | ZYGOSITY | CLASSIFICATION ¹ | DISORDER (OMIM ² , INHERITANCE) |
|-------------------------------|----------------------------------|----------|-----------------------------|---|
| THOC6 (NM_024339.3) | c.893del [p.(Pro298Glnfs*20)] | Hom. | Likely pathogenic | Beaulieu-Boycott-Innes Syndrome (613680, AR) |
| PTCH2 (NM_003738.4) | c.528del [p.(Met176Ilefs*9)] | Het. | Likely pathogenic | Basal cell nevus syndrome (109400, AD) |
| EDAR (NM_022336.3) | c.486del [p.(Ser163Argfs*26)] | Het. | Pathogenic | Ectodermal dysplasia type 10A (129490, AD/AR) |

Table 2: WES Results

References

- Richards S. *et al.* 2015
- Online Mendelian Inheritance in Man, OMIM®. McKusick-Nathans Institute of Genetic Medicine, Johns Hopkins University (Baltimore, MD), [20.01.2017]. World Wide Web URL: <https://omim.org/>

Disclosures

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