



# COGEN

Preconception, Preimplantation and Prenatal Genetic Diagnosis (CoGEN)

# Prenatal exome sequencing (WES). A retrospective cohort study in fetuses with congenital defects detected by antenatal ultrasound.

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## INTRODUCTION

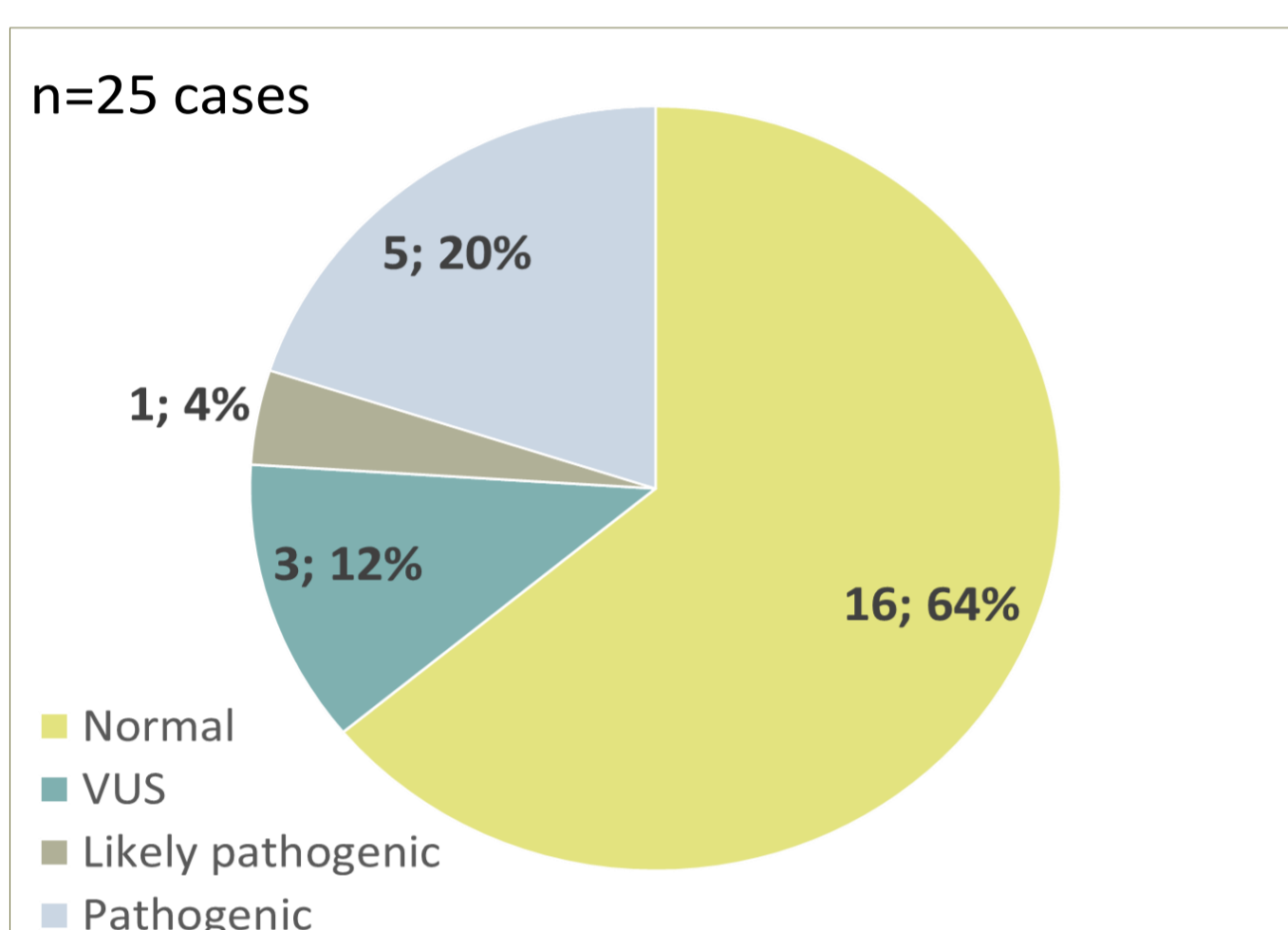
Whole exome sequencing (WES) is a diagnostic tool in postnatal settings for individuals with a suspected genetic condition. In prenatal settings, WES diagnostic yield ranges from 15% to 35%. The aim is to evaluate the usefulness of WES in identifying the genetic etiology of congenital defects.

## METHOD

WES was performed in 25 fetal samples with ultrasound anomalies and previous prenatal CGH-array with a normal result. Kit Nextera Flex for enrichment, oligos Illumina Exome using NextSeq (Illumina). Variant and segregation studies were confirmed by Sanger sequencing.

## RESULTS

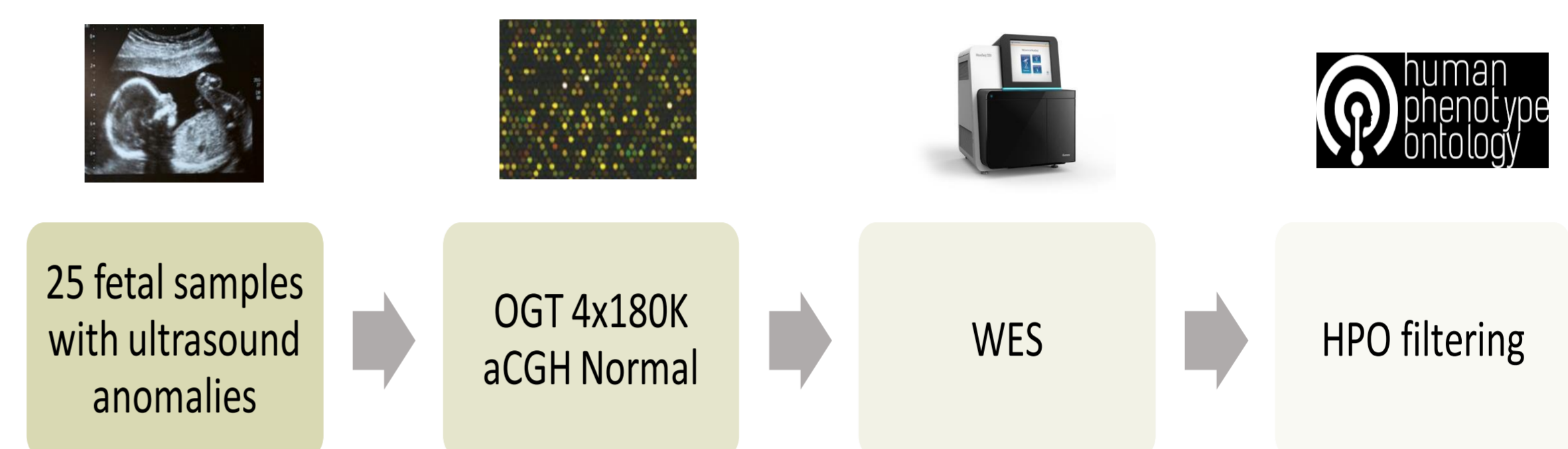
### 1. Global results:



### 2. Pathogenic results:

Fetal phenotype	Variant	Gene	Pathogenicity	Inheritance
Hidrocephaly, retrognathia, lordosis	c.2287_2321del	<i>EFTUD2</i>	Pathogenic	<i>de novo</i> dominant
Focomelia	c.663+1G>A	<i>TBX5</i>	Pathogenic	paternal dominant
Severe ventriculomegaly	c.2509C>T	<i>OFD1</i>	Pathogenic	paternal dominant
Hydrops	c.716A>G hom	<i>SEC23B</i>	Pathogenic	carrier parents recessive
IUGR, VSD, ectopic kidneys	c.2303T>C c.3788_3790del	<i>FANCA</i>	Pathogenic	carrier parents recessive
NT>p99, hydrops	c.226G>C	<i>PTPN11</i>	Likely Pathogenic	<i>de novo</i> dominant

IUGR: Intrauterine growth retardation, VSD: ventricular septal defect, NT: Nuchal translucency



**Fetus with Fanconi anemia (c.2303T>C *FANCA*)**

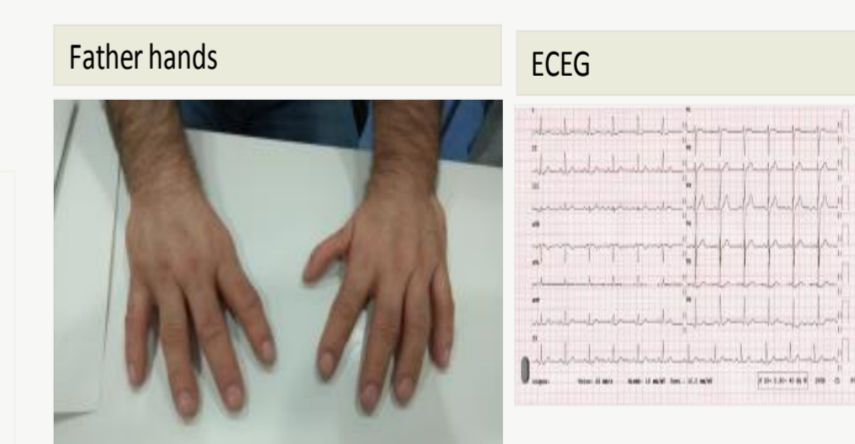
Ultrasound: Pelvic fused ectopic kidney  
Ultrasound: muscular VSD  
Magnetic resonance: Pelvic fused ectopic kidney (red arrow right kidney, green arrow left kidney)

**Fetus with Holt Oram syndrome (c.663+1G>A *TBX5*)**  
Focomelia, oligodactyly and congenital heart defect (ASD type OS and VSD).

### Holt Oram syndrome

- Limb defects (100%)
- Congenital defects (75%)
- Cardiac conduction abnormalities (sporadic)
- Complete penetrance
- Intrafamilial variability

**Father**  
Bilateral hypoplasia of thumbs  
Normal EEG



Fetal autopsy  
WES  
TBX5:NM\_181486:c.663+1G>A  
Mother  
GGC TTA CCT TGT G  
Foetus  
GGC TTA CCT TGT G  
Father  
GGC TTA CCT TGT G

ASD: atrial septal defect, OS: ostium secundum

### 3. VUS results:

Fetal phenotype	Variant	Gene	Inheritance
Aortic stenosis vs aorta coartation	c.13817A>G	<i>KMT2D</i>	maternal dominant
Multicystic kidney	c.214A>T	<i>GLI3</i>	paternal dominant
Lobar holoprosencephaly	c.5050T>C	<i>CNOT1</i>	No maternal dominant

## CONCLUSION

- Despite the small sample size, the results show the clinical utility of prenatal WES for detecting pathogenic/likely pathogenic variants in fetuses with congenital defects.
- As in postnatal WES, detailed phenotyping is necessary to filter and prioritize variants to obtain a molecular diagnosis.

## REFERENCES

- References: Lord J et al. Prenatal exome sequencing analysis in fetal structural anomalies detected by ultrasonography (PAGE): a cohort study. *Lancet* 2019;393(10173):747-757. doi: 10.1016/S0140-6736(18)31940-8
- Petrovski S et al. Whole-exome sequencing in the evaluation of fetal structural anomalies: a prospective cohort study *Lancet* 2019;393(10173):758-767. doi: 10.1016/S0140-6736(18)32042-7.
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