



# Advances in Genomics of Monogenic Diseases: From Diagnostics to Therapeutics

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**NephKids 2025**

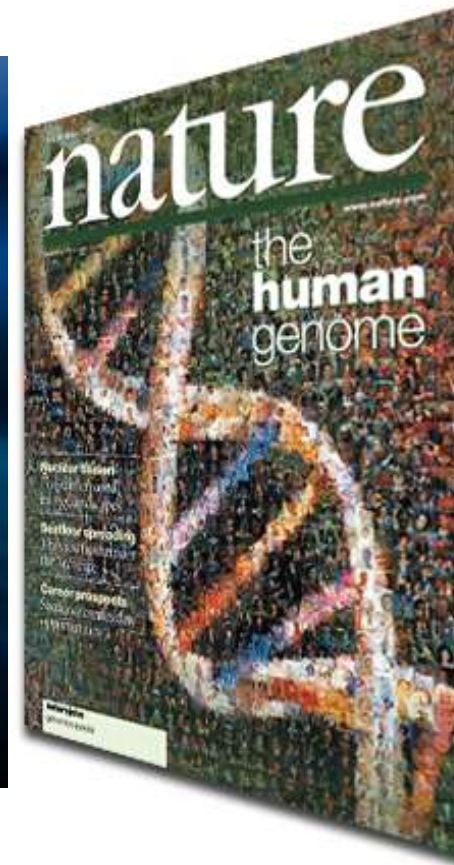
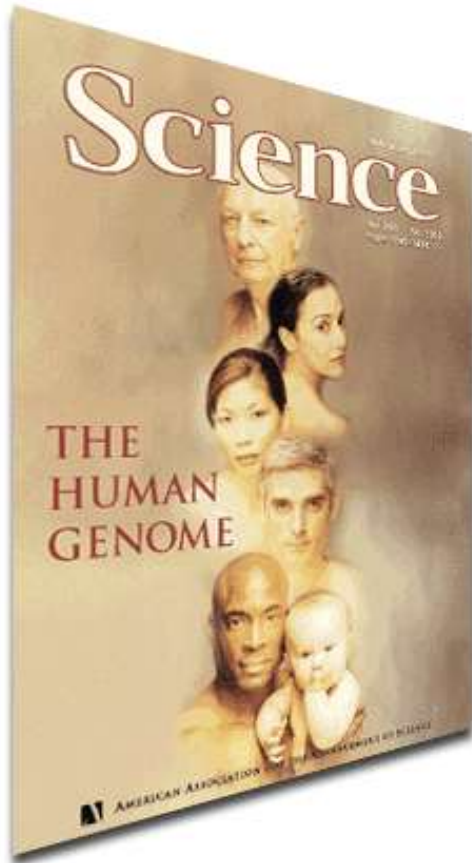
**Chennai**

**13<sup>th</sup> – 14<sup>th</sup> September 2025**

# OUTLINE

- Foundational events making genomic tests accessible and useful
- Genomics of kidney diseases and utility of genomic testing
- Leveraging genomics testing for therapeutics
- Challenges with integrating genomic testing in clinical workflow

# Mapping of Human Genome



3 billions base pairs (ATGC)

99.6% inter-individual identity (yet 4 millions differences)

***Genes [20,000-25000; 8000 mapped to Mendelian diseases]  
Sequences < 5 % - Exons - 1.5 % [Coding] Introns – 3.5 % [Non-coding]  
95 % disease causing variants identified in exonic regions***

INNOVATION > HEALTHCARE

# The First Child Saved By DNA Sequencing

By Matthew Herper, Former Staff. I cover science and medicine, and believe this is biology's century.  
Published Jan 05, 2011, 04:57pm EST, Updated Aug 11, 2011, 12:19pm EDT

GENETICS

## For the First Time, DNA Sequencing Technology Saves A Child's Life

The future of medicine has arrived.  
Posted January 10, 2011



Costed around 70,000 USD for exome sequencing

The hurdles to getting DNA sequencing from the laboratory to the hospital are huge.

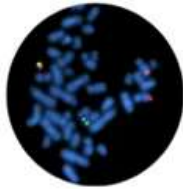


# Types of genetic (genomic) testing methods

Chromosomal anomalies

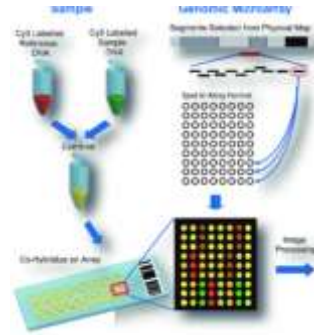
Copy Number Variants

Single nucleotide variants/Small InDels



Karyotyping

FISH



Microarray (CGH)

- ❑ Multiplex ligation dependent probe amplification (MLPA)



Sequencing



Sanger Sequencing

Massively Parallel Sequencing (Formerly called Next Generation Sequencing)

- ❑ Exome Sequencing
  - Targeted Panel
  - Clinical exome (CES)
  - Whole exome analysis (WES)
- ❑ Whole genome analysis (WGS)

Nanopore technology, in situ nucleic acid sequencing, and microscopy-based sequencing



Sanger Seq

1975

Southern Blot

1977

PCR

1983

454

2005

Solexa

2006

2007

Solid

2010

Ion Torrent

2012

Ion Proton

2014

Oxford Nanopores

2015

HiSeq X

2017

NovaSeq



BGISeq-500

GeneReader

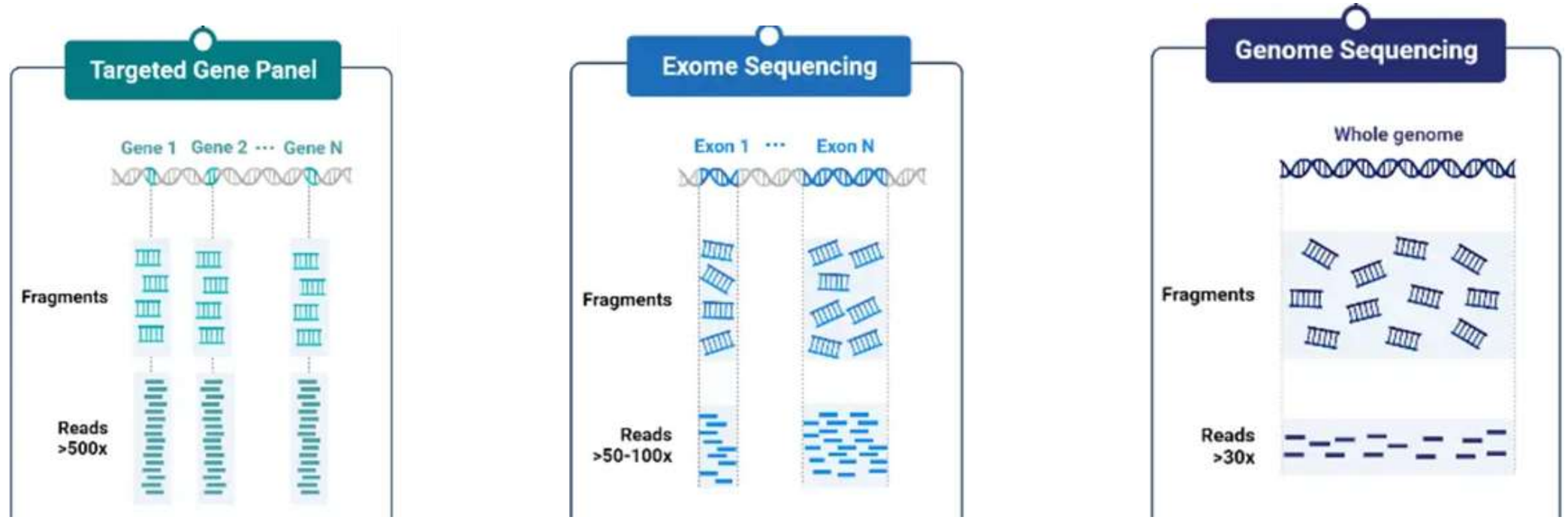


Oxford Nanopore

(MinION)

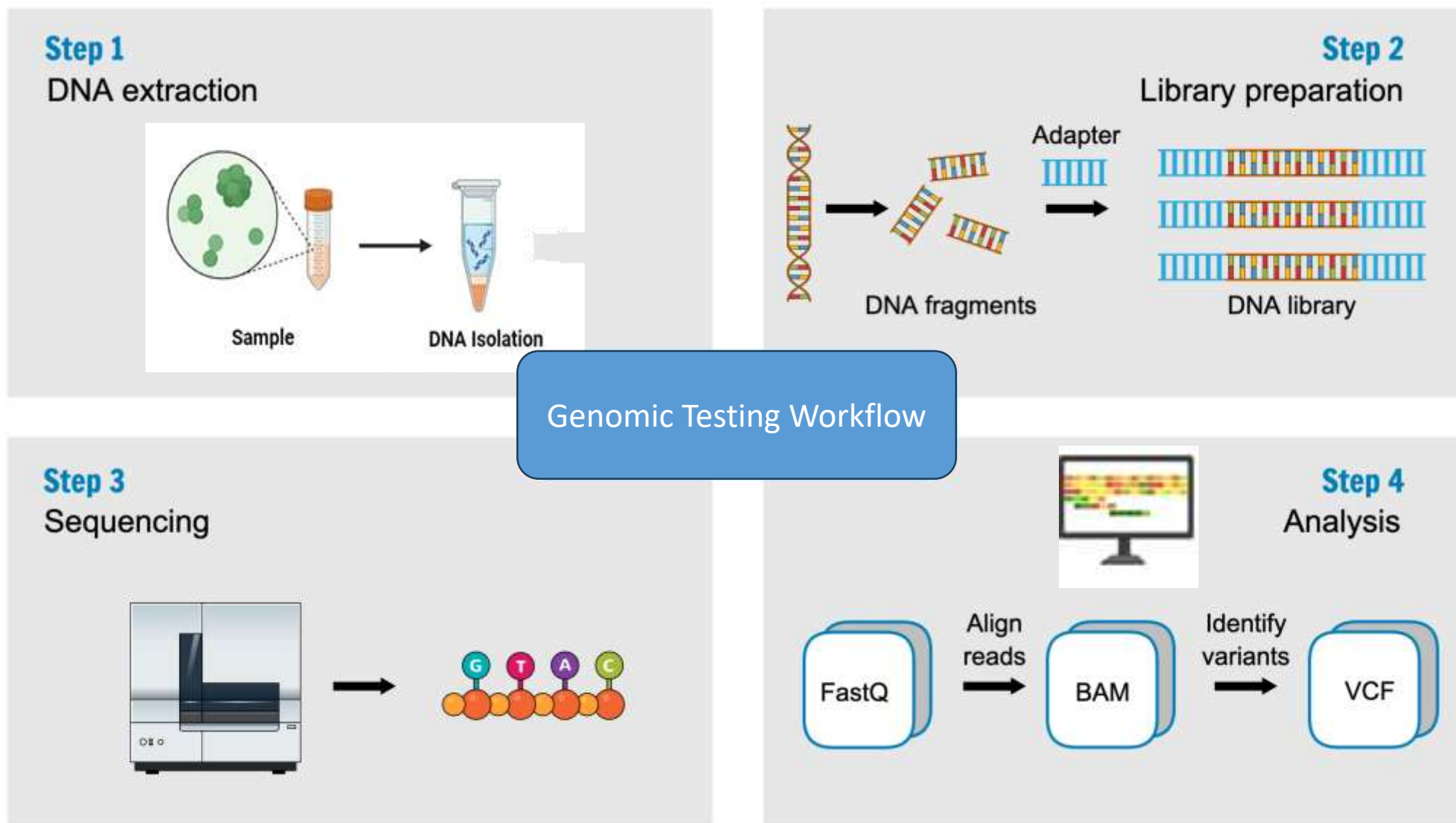
in situ nucleic acid sequencing

# Choosing a genomic testing method



Coverage	All exons of selected genes (10-500)	Exons of all genes in OMIM (~8000) Exons of all genes known (~25000)	All genes and noncoding DNA
Accuracy	High	Good	Low to Good
Time (TAT)	Quick (2-4 weeks)	Long (2-6 weeks)	Longest
Cost	Most cost effective	Cost effective	Most expensive
Utility	Disease phenotype/clinical diagnosis certain	Phenotype overlap/heterogeneity; Unknown clinical diagnosis; CNVs	Phenotype suggestive of genetic cause and exome sequencing is negative

# General workflow of clinical genomic sequencing tests



## Evidence Types

### Human observational data

Population Observations

Unaffected Observations

Affected Observations

Specific Phenotype

Segregation

### Predictive/Functional data

Predicted Effect

Inferred Effect

Functional Studies

## ACMG Classification

Pathogenic

>0.99

Variants that are known to cause disease.

Likely Pathogenic

0.95-0.99

Variants that are probably disease-causing but not confirmed.

Uncertain Significance

0.05-0.949

Variants of uncertain significance with unclear implications.

Likely Benign

<0.001-0.049

Variants that are probably not disease-causing.

Benign

<0.001

Variants that are confirmed to not cause disease.

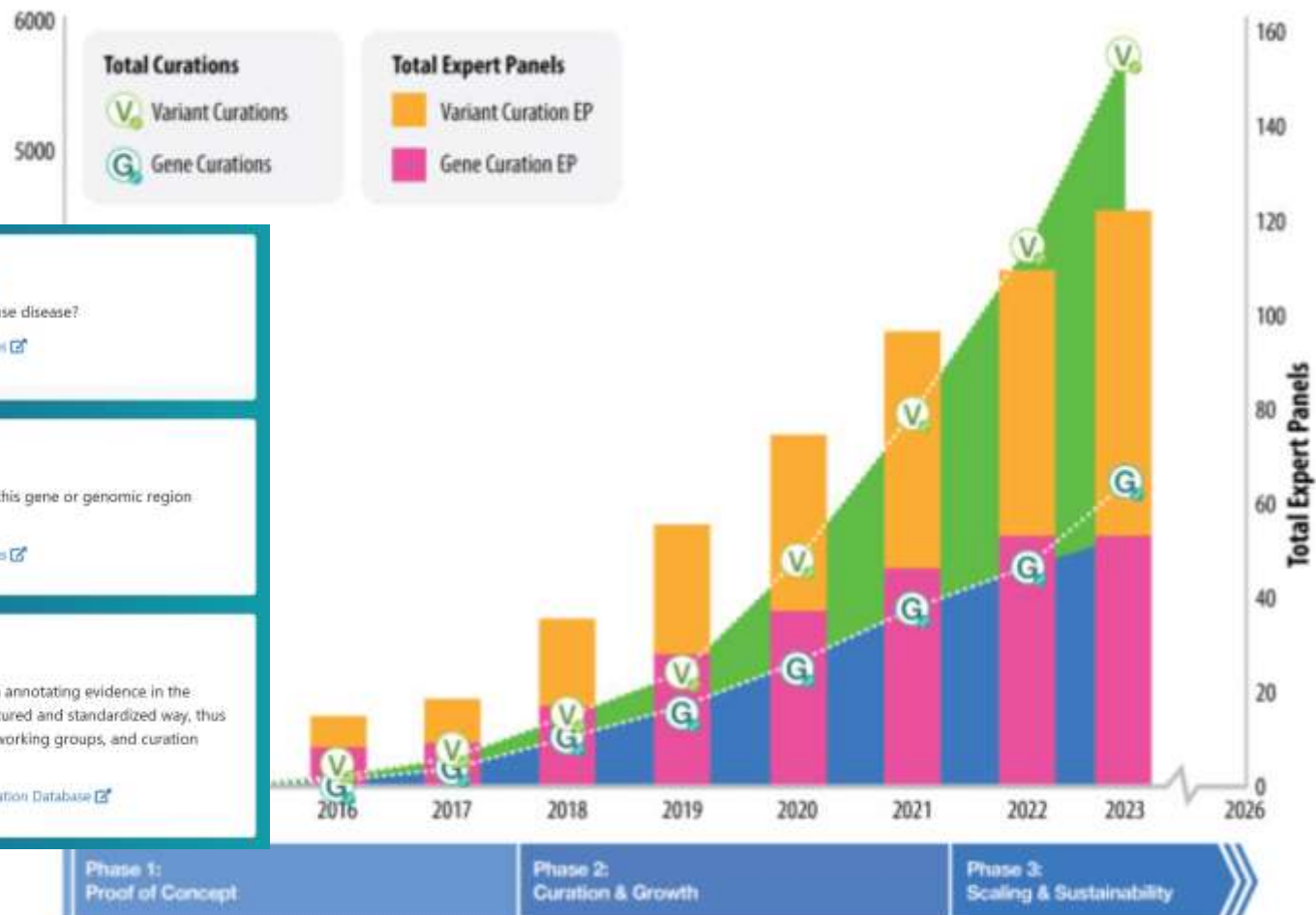


# Explore the clinical relevance of genes & variants

ClinGen is a National Institutes of Health (NIH)-funded resource dedicated to building a central resource that defines the clinical relevance of genes and variants for use in precision medicine and research.

Gene Symbol:  Search

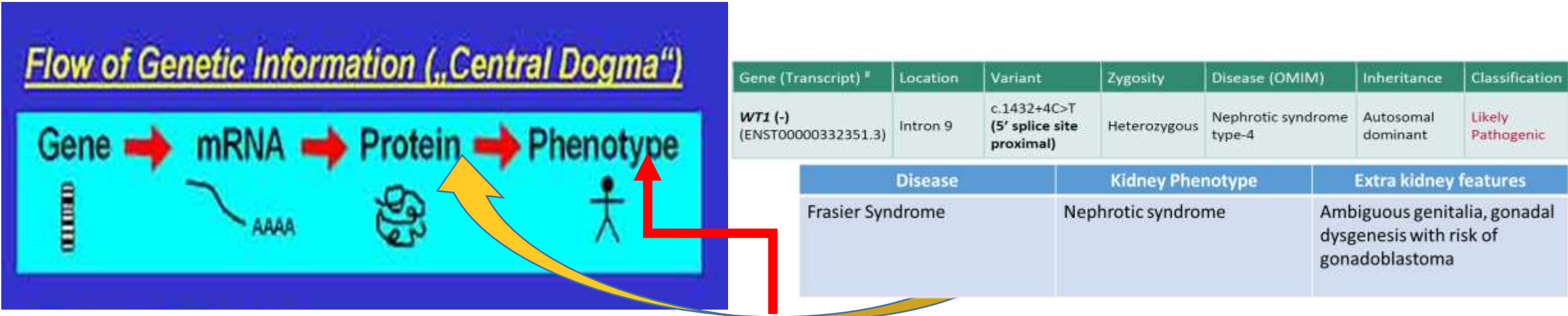
Curated Genes - Gene-Disease Validity - Dosage Sensitivity - Clinical Actionability - Curated Variants - Statistics - More - ?



Bayesian Points Classification  
for variant classification

# Interpretation of Genomic Test Report

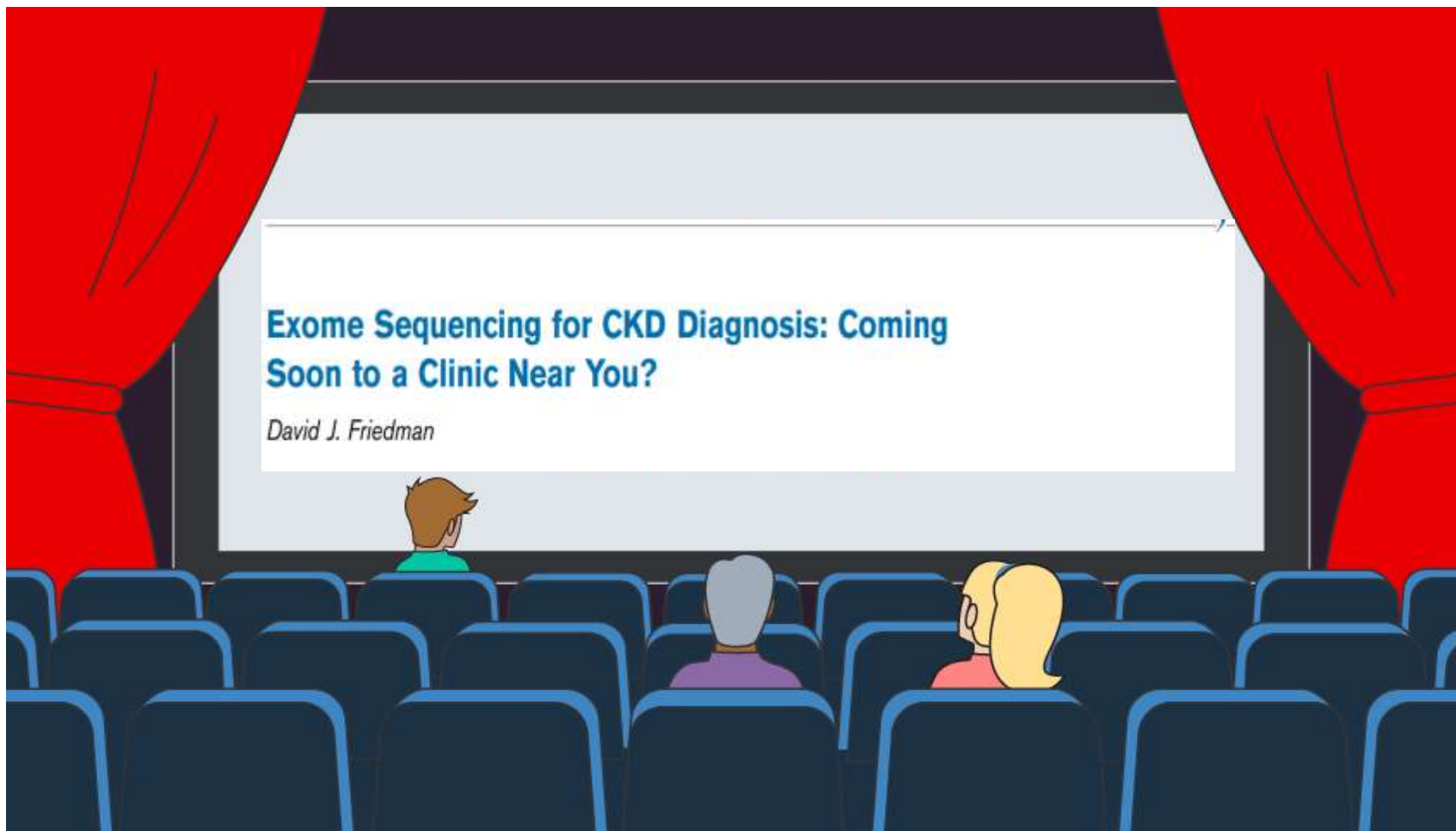
Variants identified must be evaluated for causation



Variant level assessment and interpretation

Case level interpretation

*Case level interpretation incorporates both pathogenicity of variants and how well the variant matches the phenotype*

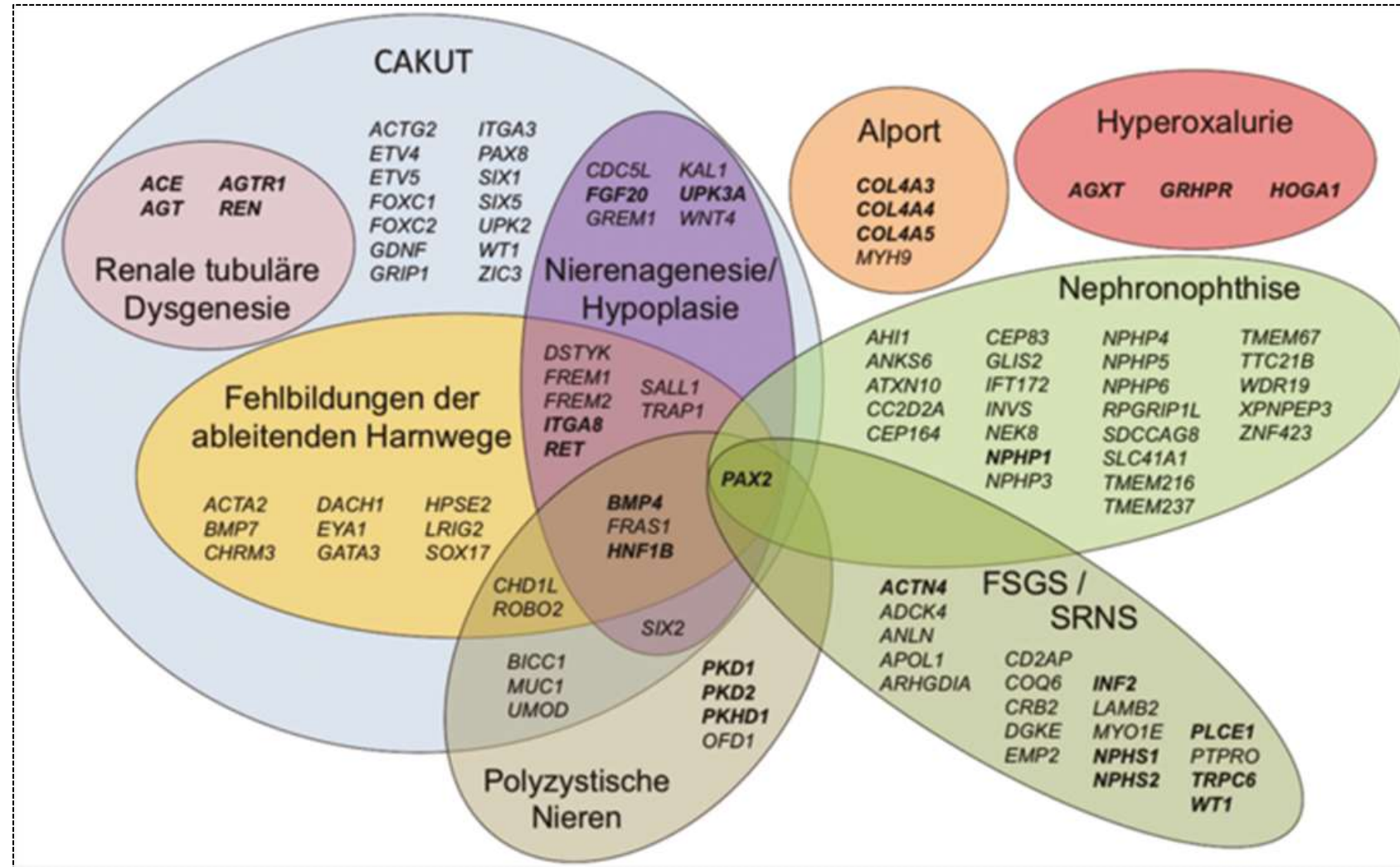


# Monogenic Kidney Diseases

- 625 Mendelian disorders associated with kidney and urological traits

Over 70% of CKD cases under the age of 25 years can be attributed to a genetic etiology

> 500 genes associated with kidney diseases





# Patient characteristics that increase the diagnostic yield of genetic testing

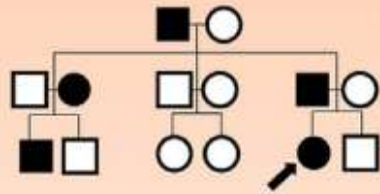
Young age of onset



**Examples:**

- Nephrotic syndrome
- Microscopic hematuria
- CKD
- HTN/electrolyte abnormalities

Strong family history



**Examples:**

- ADPKD
- Alport syndrome
- Young onset of ESKD

Cystic/anatomic abnormalities



**Examples:**

- Multiple renal/hepatic cysts
- CAKUT

CKD of unclear etiology



**Examples:**

- No diagnosis despite thorough work up
- Tubulointerstitial disease of unclear cause (ADTKD)

Extrarenal manifestations



**Examples:**

- Liver cysts
- Developmental delay
- Skeletal abnormalities
- Vision/hearing loss



# Varying yield using molecular genetic analysis and genetic heterogeneity

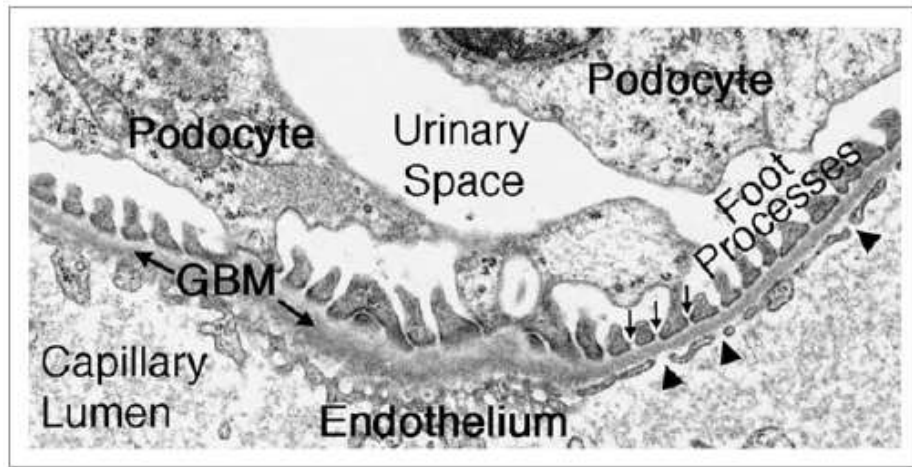
Glomerular	25 - 80 %
Tubular Diseases	30 - 90 %
Nephrolithiasis	10 - 60 %
<i>Cystic Diseases</i>	20 - 80 %
Ciliopathies	
CAKUT	5 - 22 %



- ❖ Genotypic heterogeneity
- ❖ Phenocopy
- ❖ Pleiotropy

# CKD is characterized by significant phenotypic heterogeneity and expanded phenotype

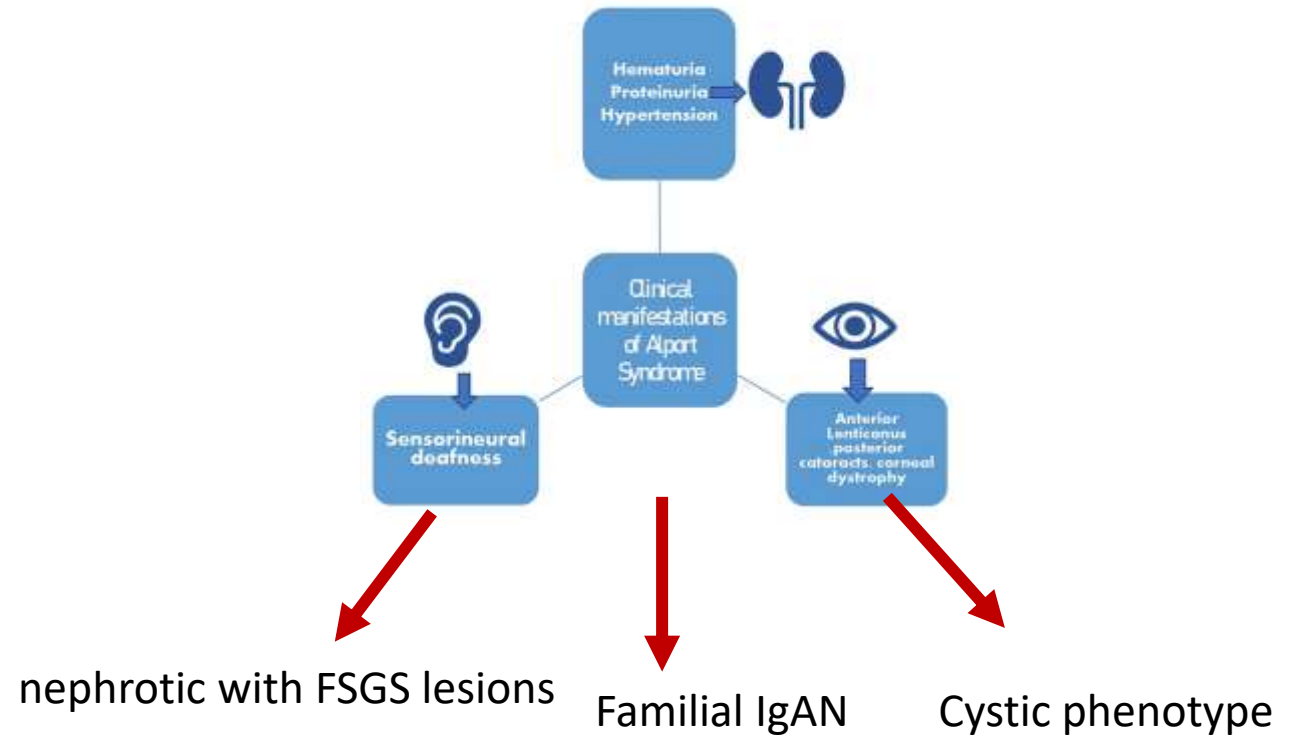
## Podocytopathies



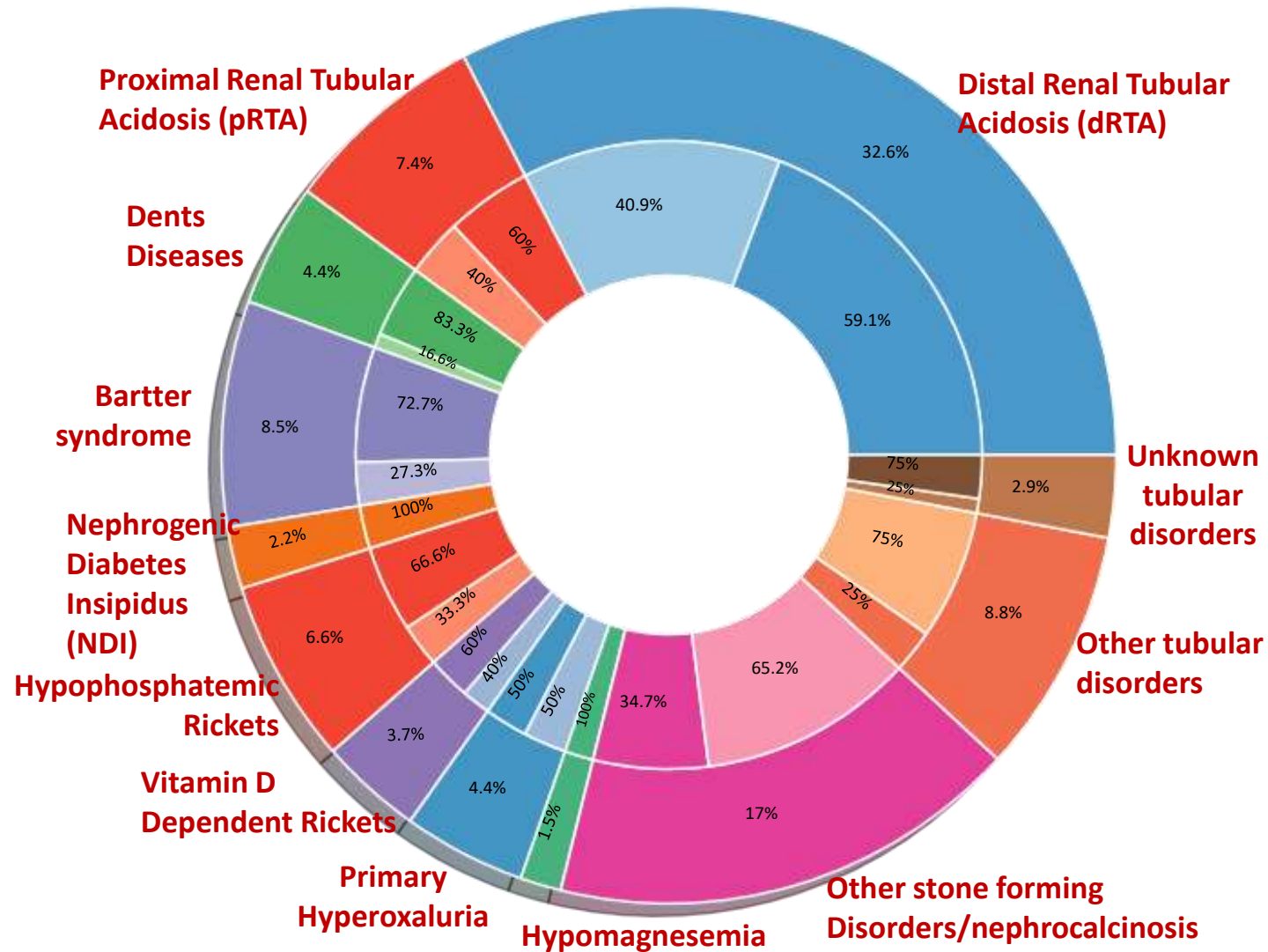
**Nephrotic Syndrome**

**Proteinuria**

**Chronic Kidney Disease**

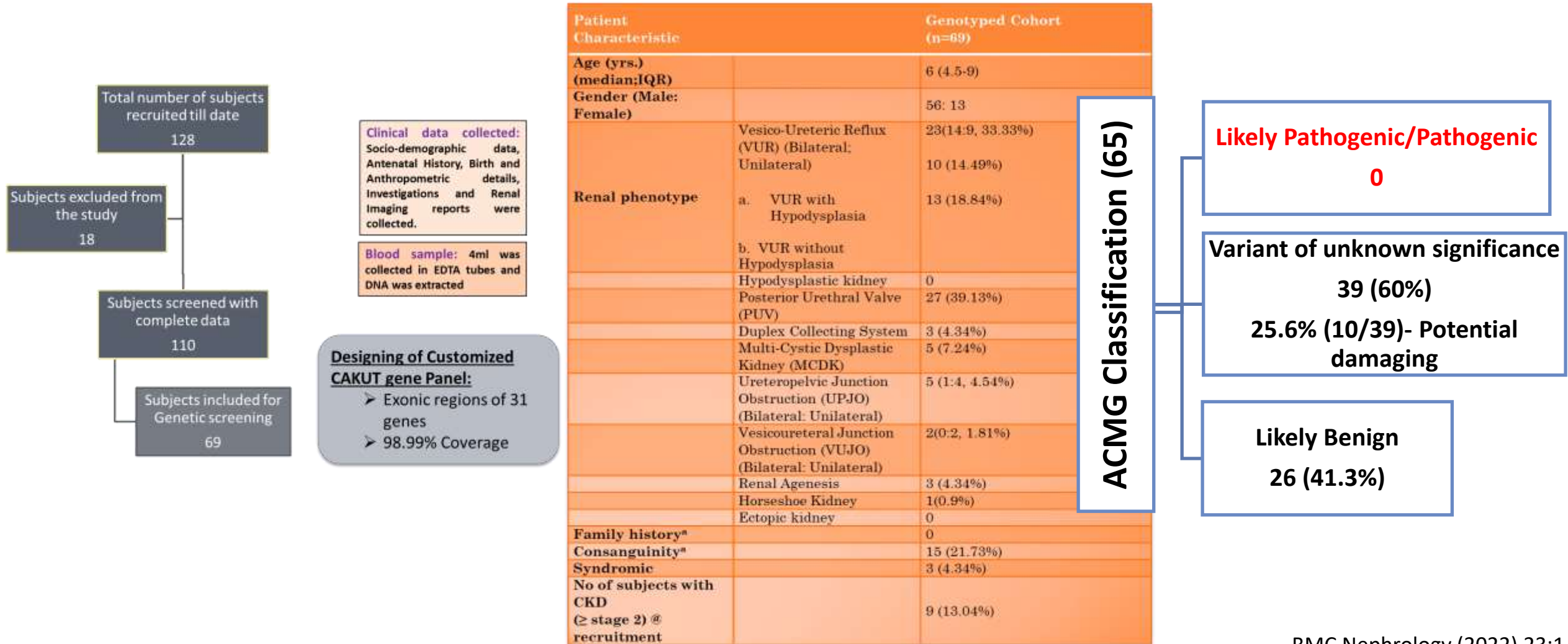


# Genetic testing in Children with tubular disorders: (n = 135/225)



# Deciphering the mutation spectrum in south Indian children with congenital anomalies of the kidney and urinary tract

Ambili Narikot<sup>1</sup>, Varsha Chhotusing Pardeshi<sup>1</sup>, A. M. Shubha<sup>2</sup>, Arpana Iyengar<sup>3</sup> and Anil Vasudevan<sup>1,3\*</sup>





# Integrating genomic assessment into clinical workflows



**Confirmation of diagnosis/Help make a diagnosis**

Examples

Alport syndrome

Monogenic SRNS

CKD of unknown etiology



**Conditions amenable to specific screening for extrarenal manifestations**

Examples:

- HNF1B: diabetes/Hypomagnesemia
- WT-1
- Fraiser syndrome – Gonadal dysgenesis



**Conditions amenable to specific disease modifying therapies**

- COL4 genes (SRNS)
- CTNS (cystinosis)
- Tubulopathies to nonspecific renoprotective strategies

Example

- COL4A3/4 (Alport) and RAAS blockade



**Avoidance of prolonged immunosuppressive therapies**

Example

- Glomerular disease due to mutations in Alport genes (COL4A3/4)



**Conditions at risk for recurrence after kidney transplantation**

- (CFH/CFIC3): aHUS
- (AGXT, GRHPR, HOGA): primary hyperoxaluria (PH)
- Genetic causes of SRNS have low recurrence rate
- **Donor selection**



**Conditions for which genetic testing is relevant for reproductive counseling**

Example:

- Prenatal/preimplantation diagnosis

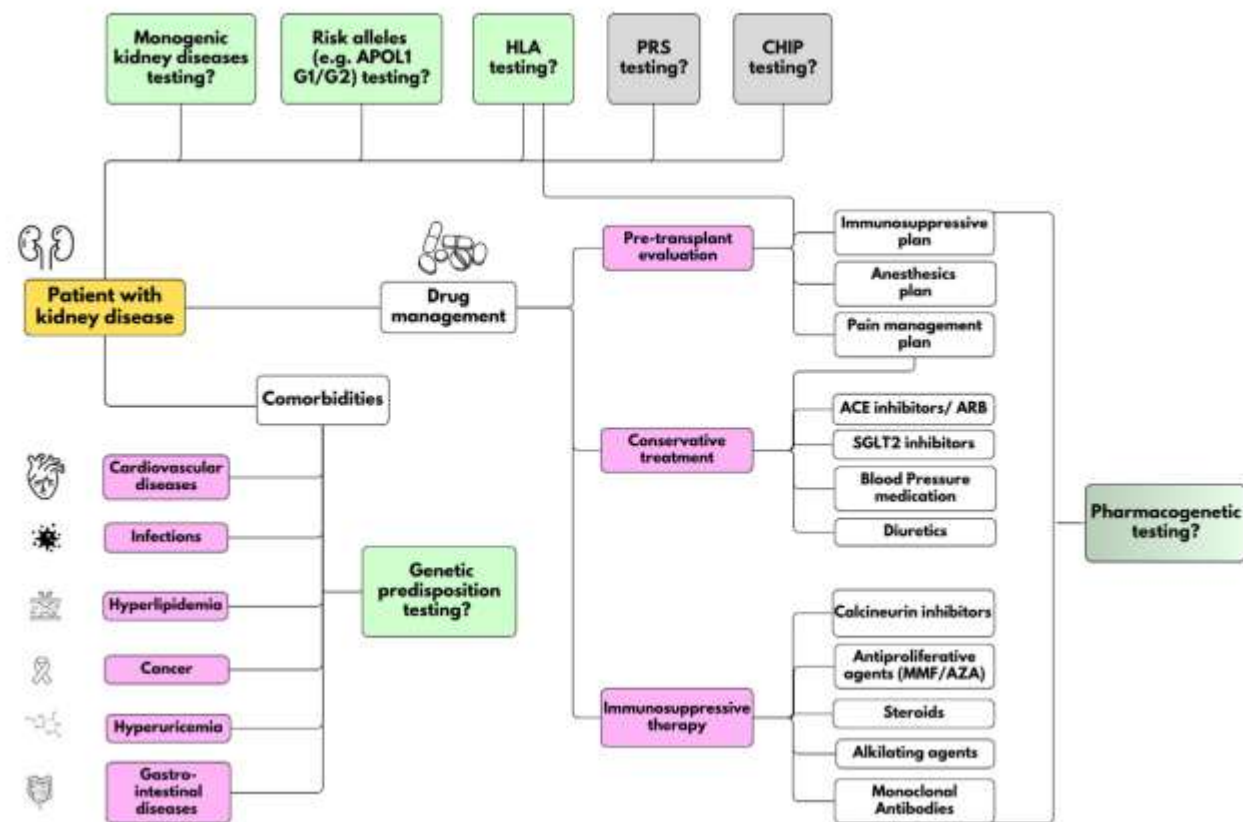
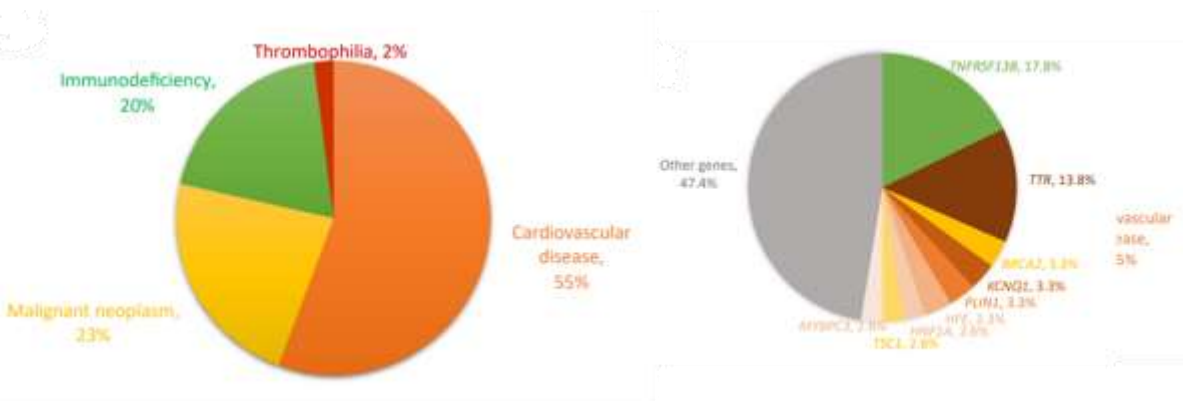


# Expanding role of genetic testing as part of clinical management of patients with kidney diseases

## Developing a genetic testing panel for evaluation of morbidities in kidney transplant recipients

Becky M. Ma<sup>1,2,3</sup>, Naama Elefant<sup>1,2</sup>, Martina Tedesco<sup>1,2,4</sup>, Kelsie Bogyo<sup>1,2</sup>, Natalie Vena<sup>1,2</sup>, Sarath K. Murthy<sup>1,2</sup>, Shiraz A. Bheda<sup>1,2</sup>, Sandv Yana<sup>1,2</sup>, Nikita Tomar<sup>1,2</sup>, Jun Y. Zhand<sup>1,2</sup>.  
[see commentary on page 18](#) [OPEN](#)

Transplant morbidity panel (355 genes) associated with major post-transplant complications including cardiometabolic disorders, immunodeficiency, malignancy, and thrombophilia



# Role of Genomic Testing in Diagnosis and Management of kidney Disease

**From optional to essential**

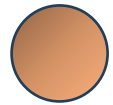
**Use at the beginning of a diagnostic workup  
rather than the end**

**From Womb to Tomb**

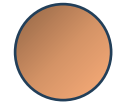
# Treatment of Monogenic Diseases



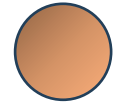
Supportive therapy



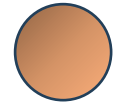
Biological molecules



New drugs

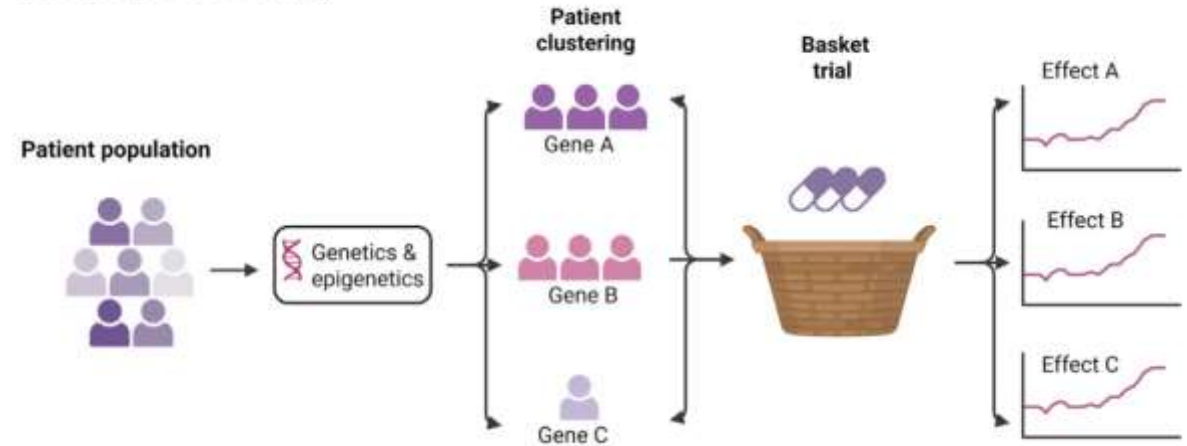


Drug repurposing



Gene modifications –  
knockout/gene therapy

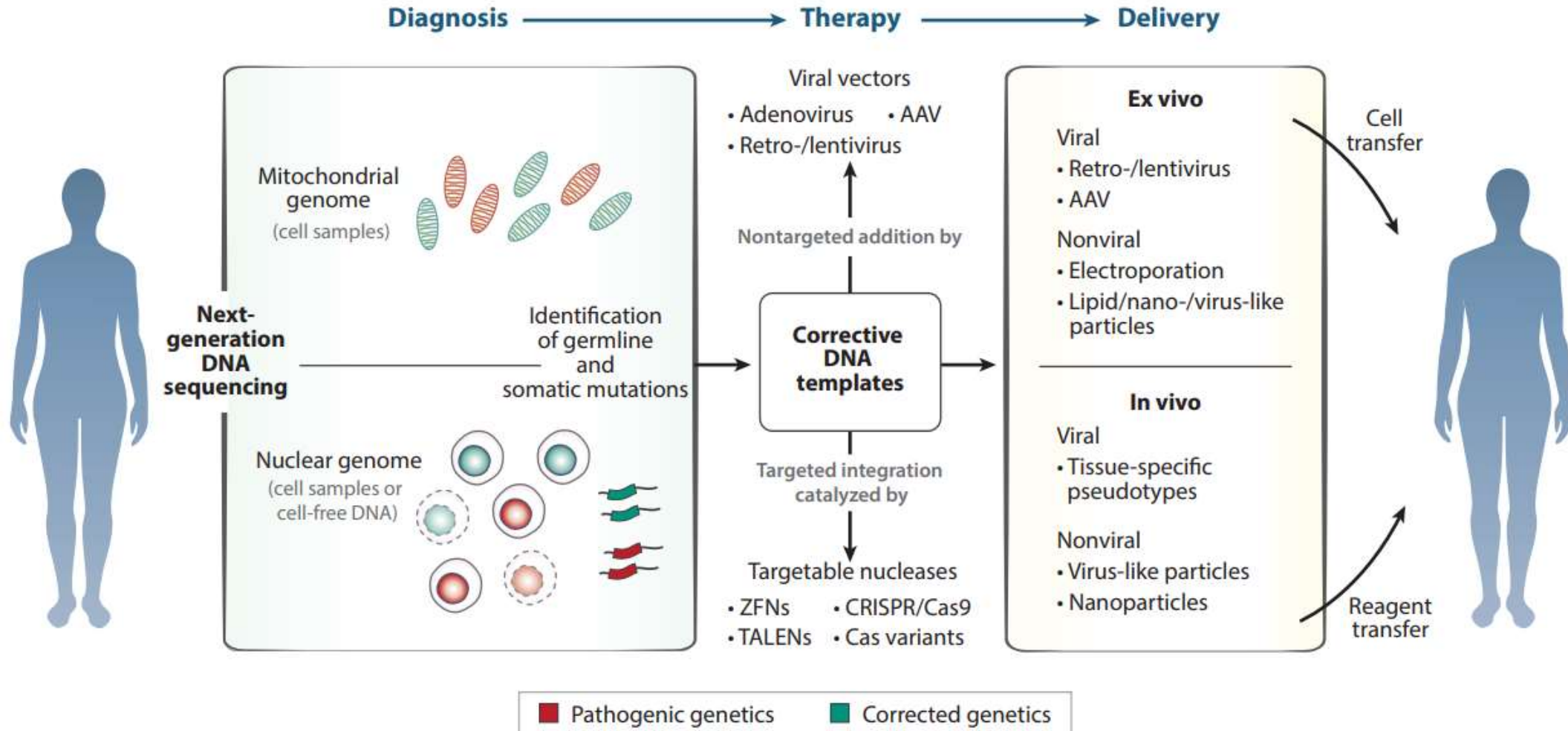
## Genetics pre screening



# Classes of Genetic Kidney Disease and Existing Therapies

Genetic Kidney Disease Subgroup	Exemplar Conditions	Treatment			Therapeutic Interventions
		Supportive	Modifiable	Curative	
CAKUT Glomerular		•	•		Surgery
	Alport syndrome	•	•		ACEi/ARB
	COQ10 deficiency				COQ10- supplementation
Cilial	NPHP related ciliopathies	•			
Cystic	ADPKD	•	•		ACEi/ARB; Tolvaptan for ADPKD
ADTKD	UMOD	•	•		ACEi/Allopurinol/Febuxostat
Tubular	Bartter Syndrome	•	•		NSAIDs and Electrolyte replacement
	Gittelman Syndrome				Electrolyte replacement
	Gordon syndrome				Thiazide diuretic
	Nephrogenic diabetes insipidus				Thiazide diuretic and NSAIDs
	Cystinuria				Penicillamine
Metabolic	Fabry Disease	•	•		Enzyme replacement
	Primary hyperoxaluria		•		RNAi
	Cystinosis		•		Cysteamine
Complement mediated	Atypical HUS		•		Eculizumab
			•		Ravalizumab
			•		C5 inhibitors

# Modes of Genetic Therapy





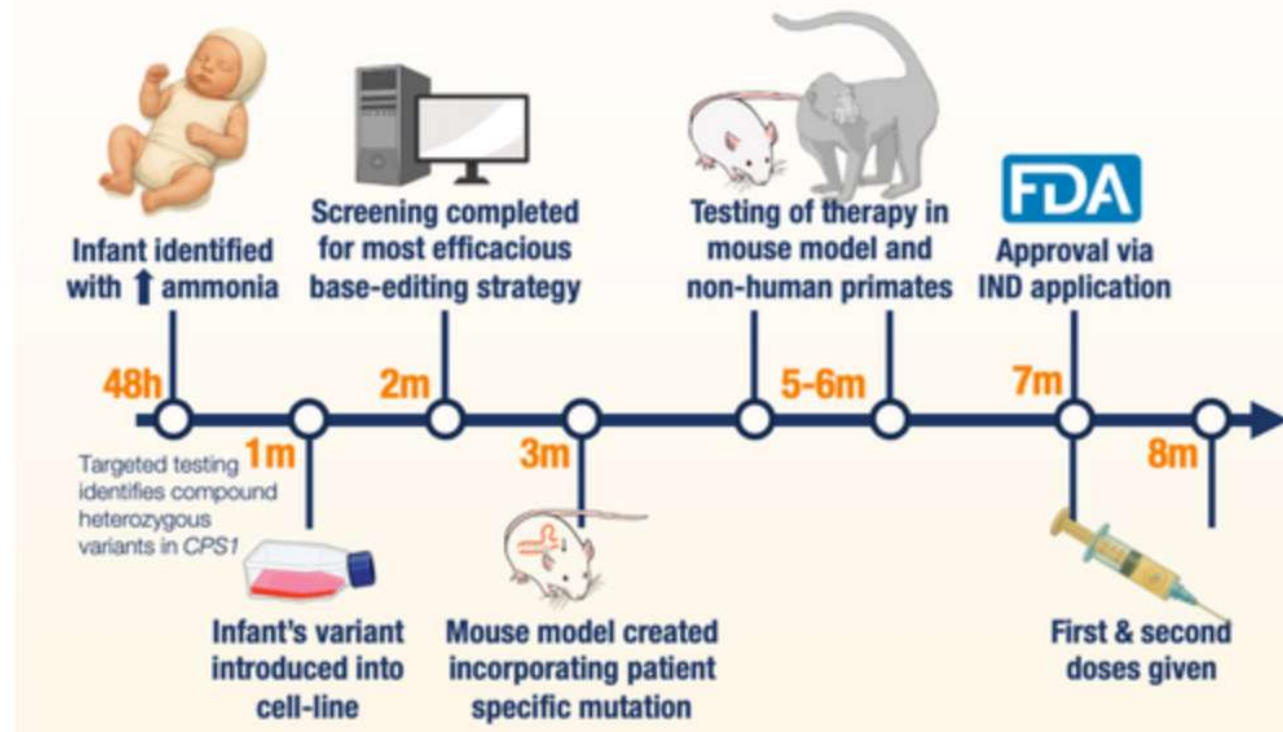
# Gene therapy

## N-of-1 Trials

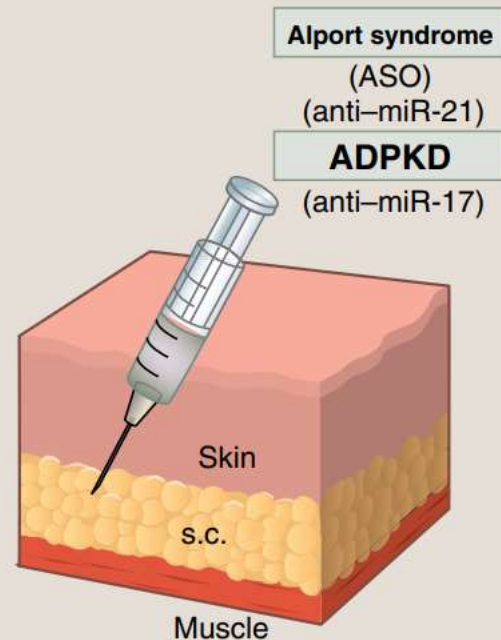
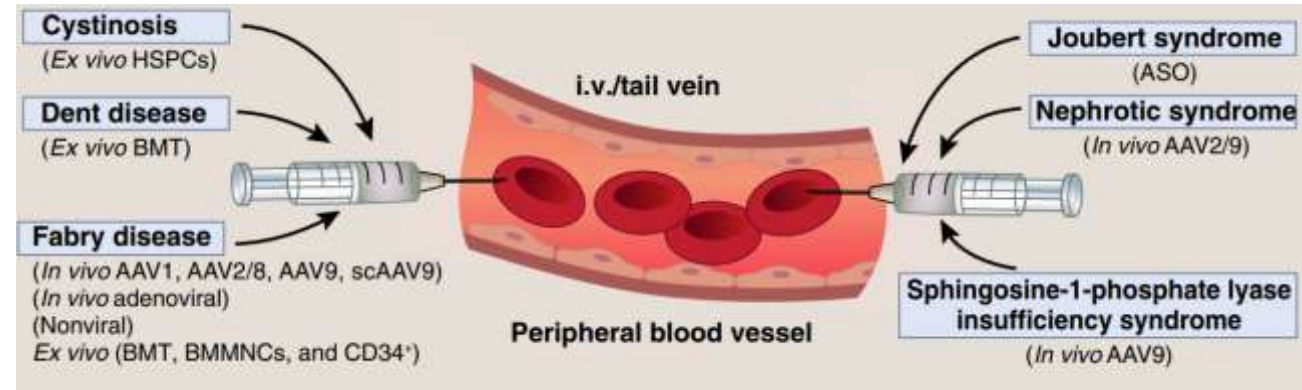
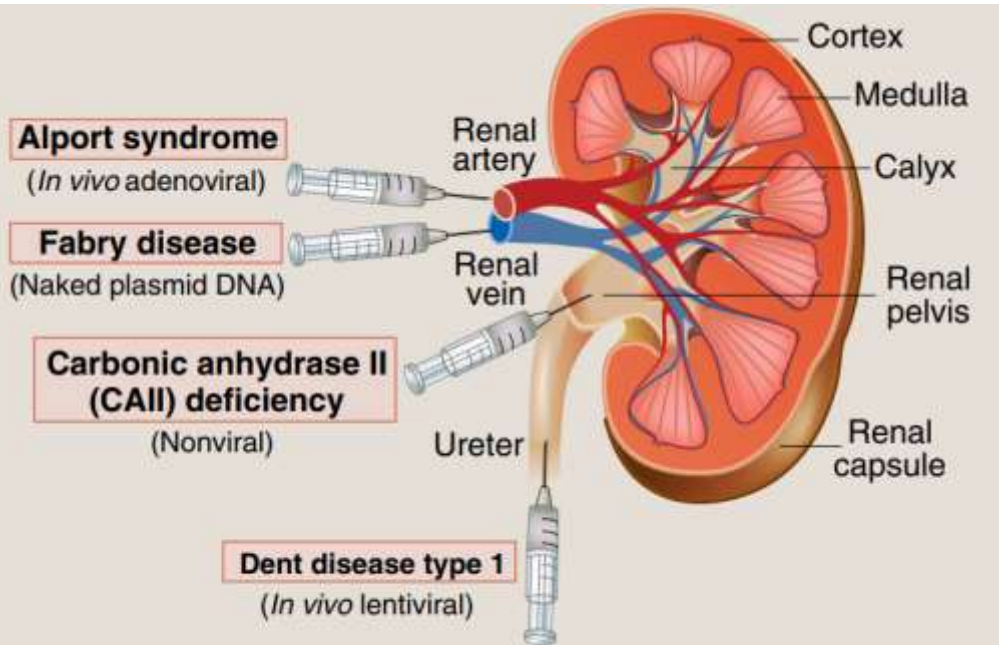
First in vivo gene editing therapy for a baby with an inherited metabolic disease (IMD)

In May 2025, Musunuru et al reported the first in vivo editing therapy for a baby with an inherited metabolic disease (IMD)

Does this herald a new era of rapid, personalised therapies in IMDs?



# Targeted gene therapy for genetic kidney diseases



# Limitations and Challenges



Single gene sequencing:  
**SANGER** sequencing



Multiple gene sequencing:  
**NGS**= Next generation sequencing

## Technical Challenges

**Gene\_Disease relationships**  
**Prioritization and interpretation of variants**  
**Incidental/Secondary findings**



NGS machines

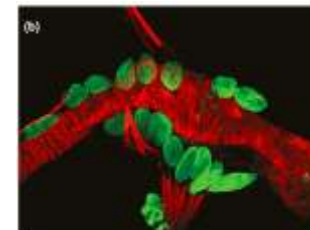
Massive amount  
of sequence data

## Practical challenges

**Medical, Social and ethical challenges**  
**Cost of testing**  
**LACK of Therapy**



# Variants of Unknown Significance



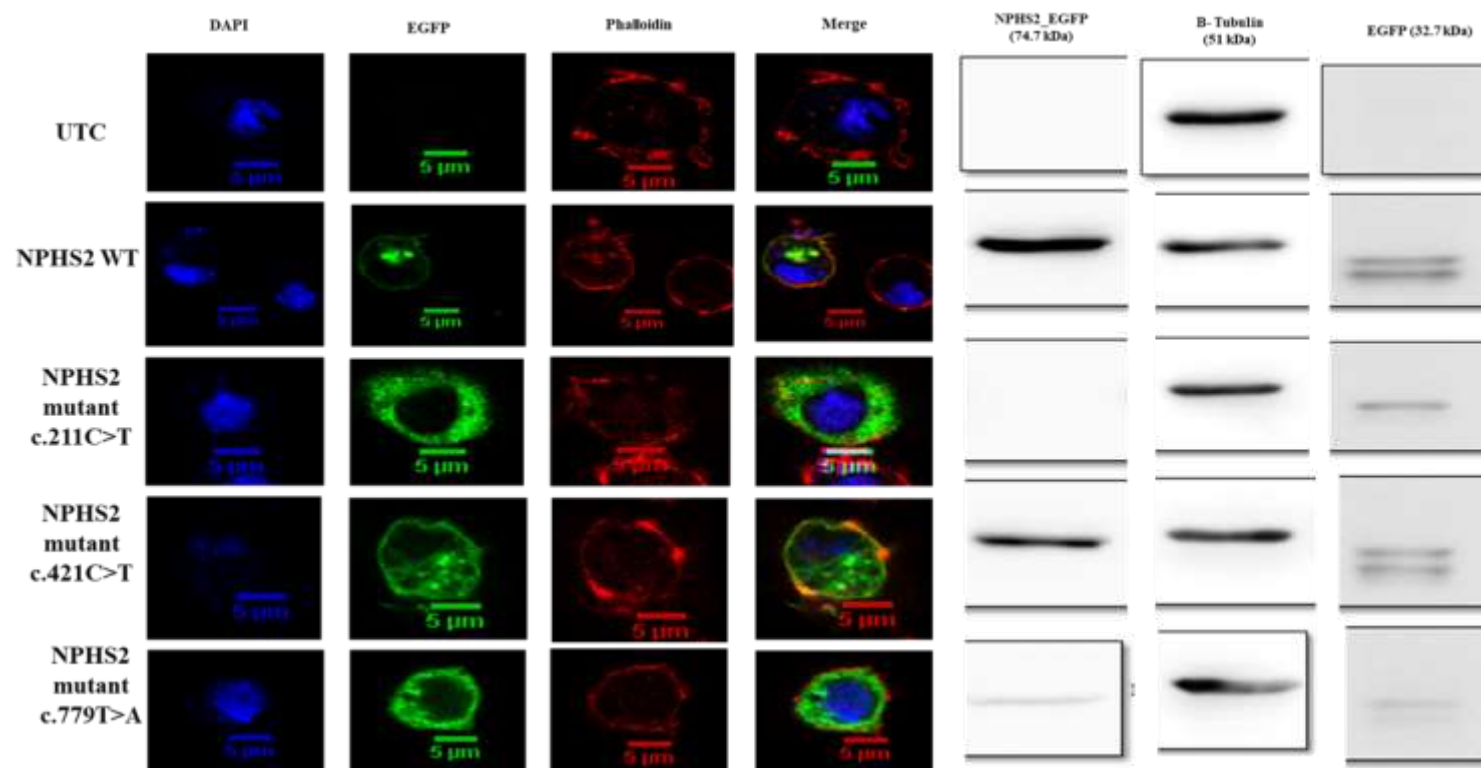
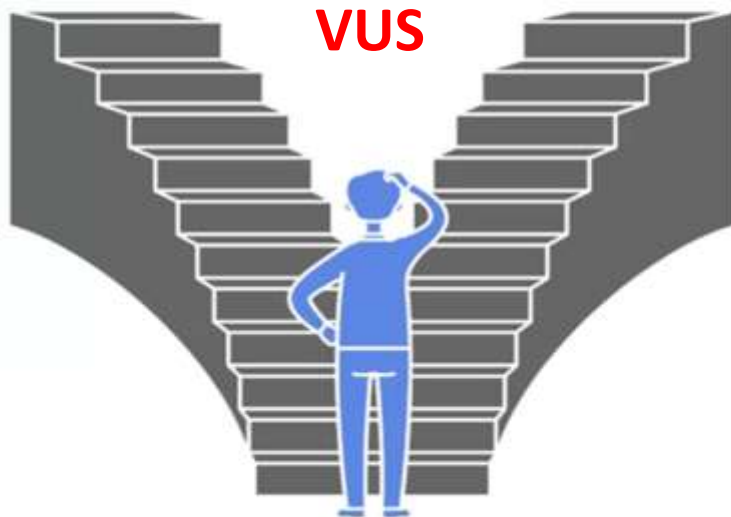
## Criteria Not Met

The variant is classified as VUS because other criteria are not satisfied.

## Contradictory Criteria

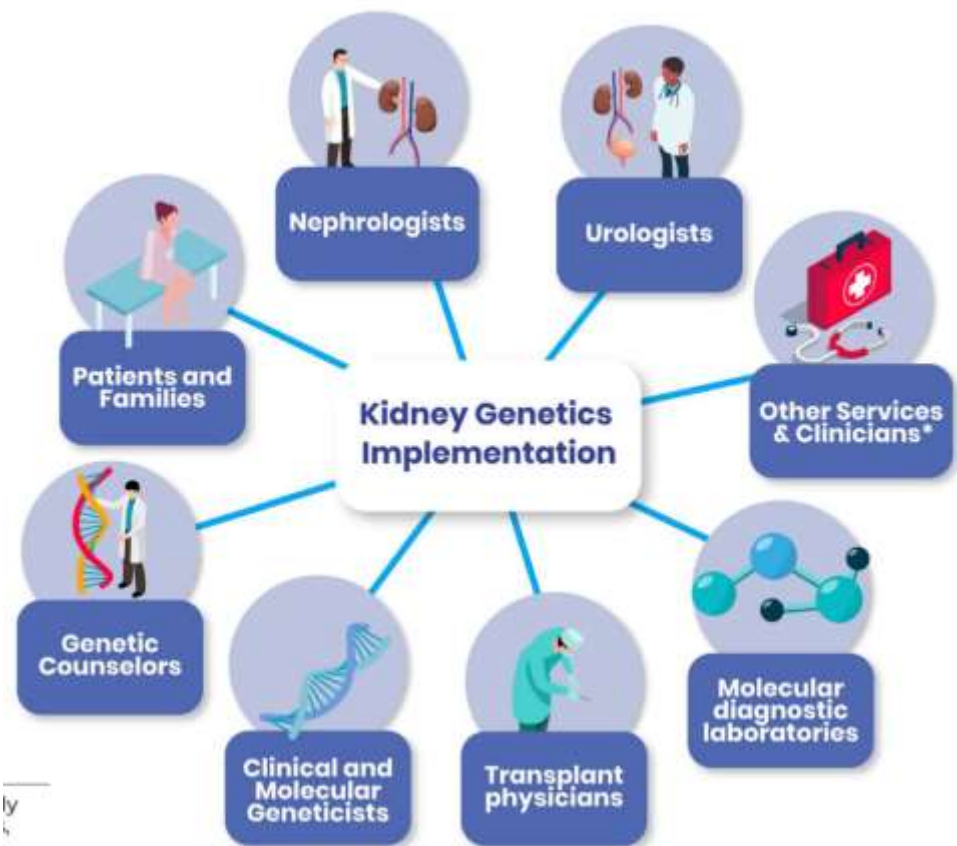
The variant is classified as VUS due to contradictory benign and pathogenic criteria.

**VUS**



# Model of care for implementation of genetic testing

## Multidisciplinary genetic clinic or Nephrogenetic clinic



## Creation of Genomics Board

	Constitution of Board	Members	Role
1	Geneticist/Clinical geneticist	Dr Preetha /Dr Jainy/ Dr.Anuradha	Interpretation of report; posttest plan/advice
2	Genomic Analyst	Dr Ambily / Mr Shivakumar/New appointee	Analysis of genetic data obtained from sequencing
3	Lab scientist	Dr Swetha / New appointee	Sequencing based on the test requested and resolving technical queries
4	Clinical expert	Dr Anil Vasudevan	Guide and facilitator and provide clinical and bioinformatic inputs
5	Domain Expert Neurologist Pediatrician Neuro developmental pediatrician Hematoncologist (Pediatric/Adult) Neonatologist Endocrinologist Cardiologist (Pediatric/Adult)	Faculty deputed by the respective departments	Provide clinical information to help interpret the results

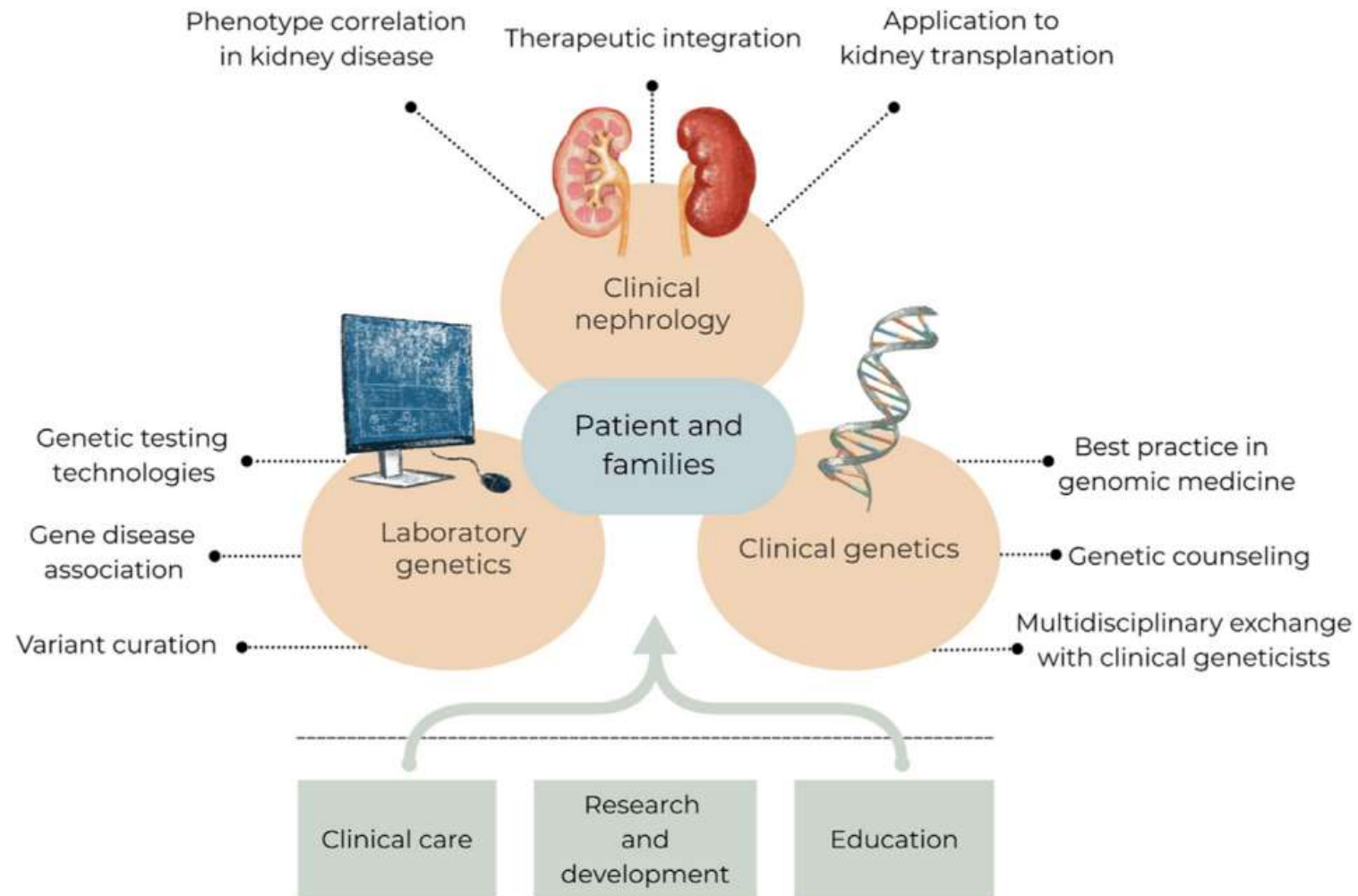
- a) Interpretation
- b) General advice regarding communication of the report and plan to patients and their families
- c) Advice regarding further testing in the index patient & suggested family screening.
- d) If a patient needs further clinical evaluation or genetic counselling by the geneticist, clinician can refer the patient to genetics OPD.



# Challenges – “Genetic literacy among clinicians”

- Breadth of knowledge and skill sets in clinical genomics is vast
- Kidney medicine brings its own complex layers
- Physician knowledge gaps surrounding genetics [*“Genetic literacy”*]
- Lack of curriculum/specialized training opportunities

# Genetic Nephrology: An Arising Subspecialty in Kidney Medicine



# Summary

## **Enhanced Diagnosis**

Genomic testing improves accuracy in diagnosing kidney diseases for better patient care.

## **Individualize management**

Genomic insights enable tailored therapies specific to individual kidney disease profiles.

## **Innovative Therapeutic Advances**

Ongoing genomic research is paving the way for novel kidney disease treatments.

