



Genetic Testing

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GeneDx

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- Making the diagnosis at birth and the diagnostic odyssey
- Cohort building
- The next five years in genomic technology



Introduction

Rare Disease

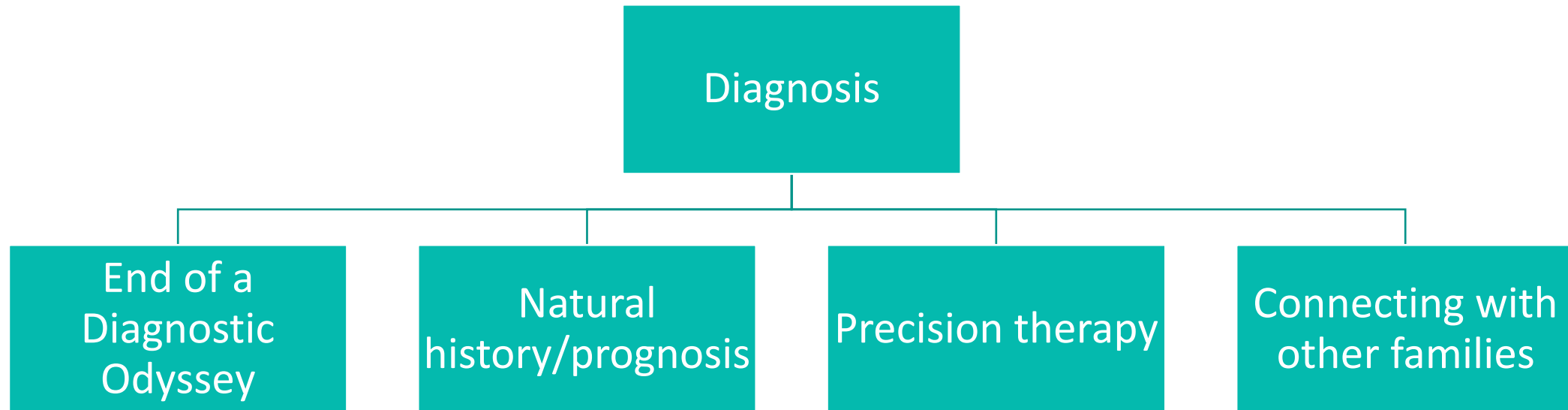
- Definition: affects less than 200,000 persons in the U.S.
- Over 7,000 rare diseases
- 25-30 million Americans living with a rare disease
- 80% of rare diseases have a genetic cause
- Roughly 70% of rare disease are pediatric onset
- The average time from symptom onset to diagnosis is 5 years

JAMA Pediatr. 2020 September 01; 174(9): 821–822

<https://rarediseases.info.nih.gov/diseases/pages/31/faqs-about-rare-diseases>

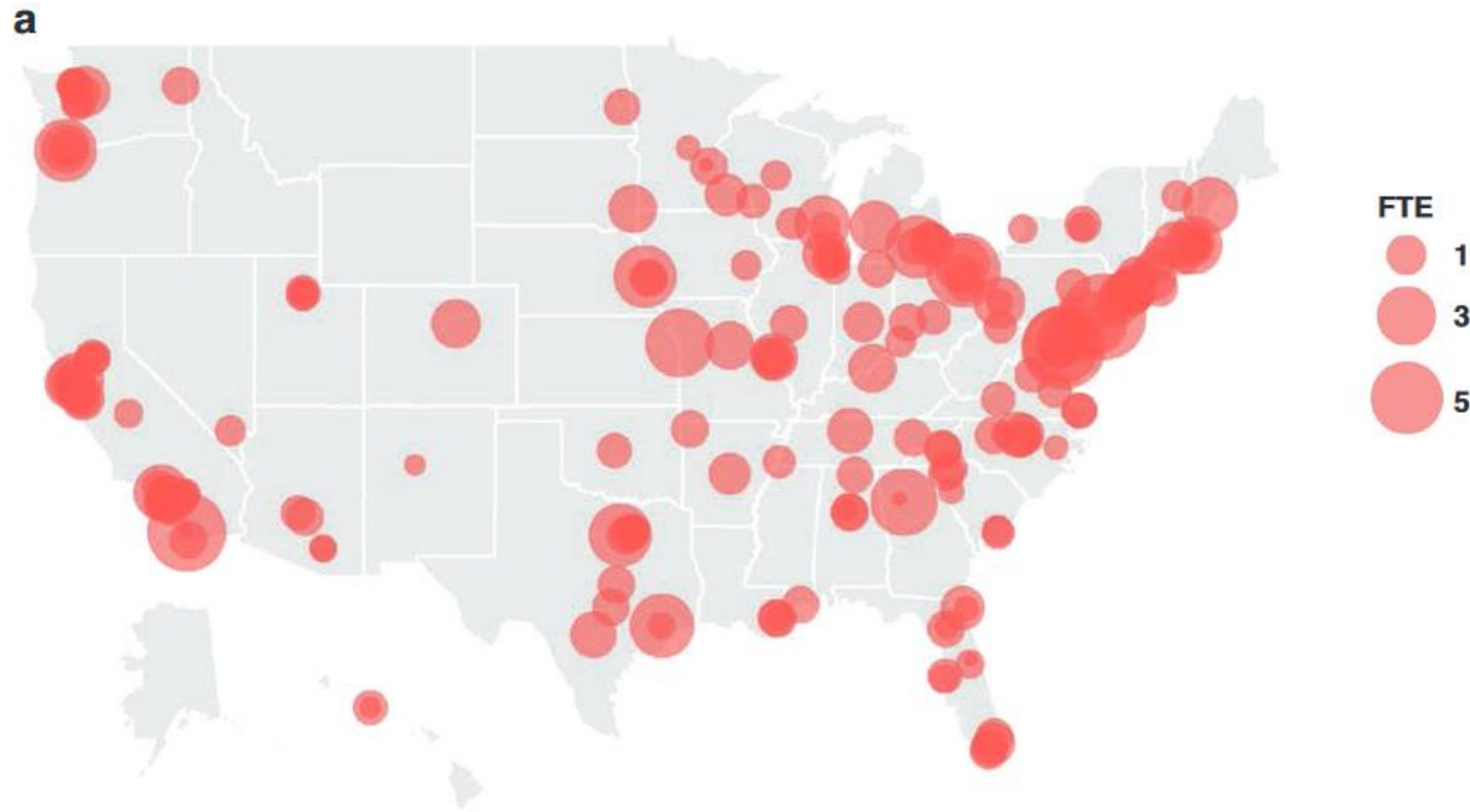
<https://www.nature.com/articles/s41431-019-0508-0>

What is an “actionable” diagnosis?



The 2019 US medical genetics workforce

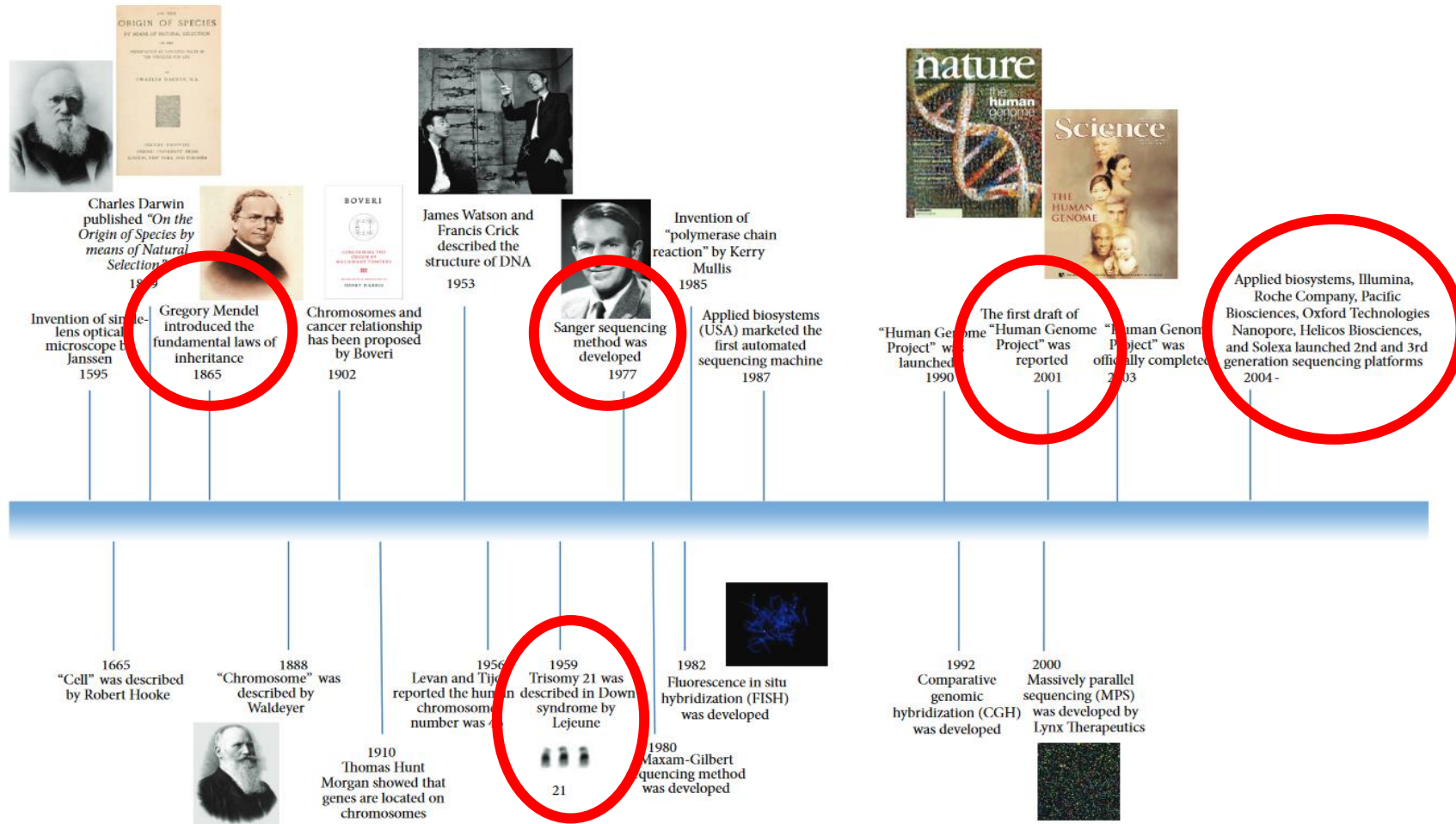
Endangered Species

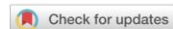




Genetic Testing

Testing Progress





ACMG PRACTICE GUIDELINE

Exome and genome sequencing for pediatric patients with congenital anomalies or intellectual disability: an evidence-based clinical guideline of the American College of Medical Genetics and Genomics (ACMG)

Kandamurugu Manickam^{1,2}, Monica R. McClain³, Laurie A. Demmer⁴, Sawona Biswas⁵, Hutton M. Kearney⁶, Jennifer Malinowski⁷, Lauren J. Massingham^{8,9}, Danny Miller¹⁰, Timothy W. Yu^{11,12}, Fuki M. Hisama¹³ and ACMG Board of Directors^{14*}

Disclaimer: The ACMG has recruited expert panels, chosen for their scientific clinical practice. An EBG focuses on a specific scientific question and then describes by a systematic review of evidence and an assessment of the benefits and educational resource for medical geneticists and other clinicians to help them all relevant information on the topic reviewed.

Reliance on this EBG is completely voluntary and does not necessarily ensure procedure or test, the clinician should consider the best available evidence, and preferences and specific clinical circumstances presented by the individual patient. Particular procedure or test, whether or not it is in conformance with this EBG, and to consider other medical and scientific information that becomes available.

Received: 16 August 2021 | Revised: 27 September 2022 | Accepted: 1 October 2022

DOI: 10.1002/jgc4.1646

PRACTICE GUIDELINE

National Society of
Genetic
Counselors WILEY

Genetic testing and counseling for the unexplained epilepsies:
An evidence-based practice guideline of the National Society
of Genetic Counselors

Lacey Smith¹ | Jennifer Malinowski² | Sophia Ceulemans³ | Katlin Peck⁴ |
Nephi Walton⁵ | Beth Rosen Sheidley¹ | Natalie Lippa⁶

<https://orcid.org/0000-0002-9095-7265>

Short Read Sequencing/exome sequencing/genome sequencing



FATHER

ATGACCGTTACTAGCGCACAGATTATGTCCTCGCAGAGCTTACGAGCATGC

MOTHER

ATGACCGTTACTAGCGCACAGATTATGTCCTCGCAGAGCTTACGAGCATGC

CACAATTATGTCCTCGC

GACCGTTACTAGCGCACAAATT

ACCGTTACTAGCG

CTAGCGCACAAA

TAGCGCACAGATTATGTCCTCGCAG

CAGATTATGTCCTCGCAGA

ACCGTTACTA

GCACAAATTATGTCCTCGCAGAGC

GTTACTAGCGCACAGATTATGT

TGTCCTCGCAGAGC

REFERENCE: ATGACCGTTACTAGCGCACAGATTATGTCCTCGCAGAGCTTACGAGCATGC

CACAATTATGTCCTCGC

GACCGTTACTAGCGCACAAATT

ACCGTTACTAGCG

CTAGCGCACAAA

CAGATTATGTCCTCGCAGA

TAGCGCACAGATTATGTCCTCGCAG

ACCGTTACTA

GCACAATTATGTCCTCGCAGAGC

GTTACTAGCGCACAGATTATGT

TGTCCTCGCAGAGC

REFERENCE: ATGACCGTTACTAGCGCACAGATTATGTCCTCGCAGAGCTTACGAGCATGC

ACCGTTACTA

TGTCCTCGCAGAGC

CACAATTATGTCCTCGC

ACCGTTACTAGCG

CAGATTATGTCCTCGCAGA

GACCGTTACTAGCGCACAAATT

TAGCGCACAGATTATGTCCTCGCAG

CTAGCGCACAA

GTTACTAGCGCACAGATTATGT

GCACAATTATGTCCTCGCAGAGC

REFERENCE: ATGACCGTTACTAGCGCACAGATTATGTCCTCGCAGAGCTTACGAGCATGC

ACCGTTACTA

TGTCCTCGCAGAGC

CTAGCGCACAA

CACAATTATGTCCTCGC

ACCGTTACTAGCG

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GTTACTAGCGCACAGATTATGT

GCACAATTATGTCCTCGCAGAGC

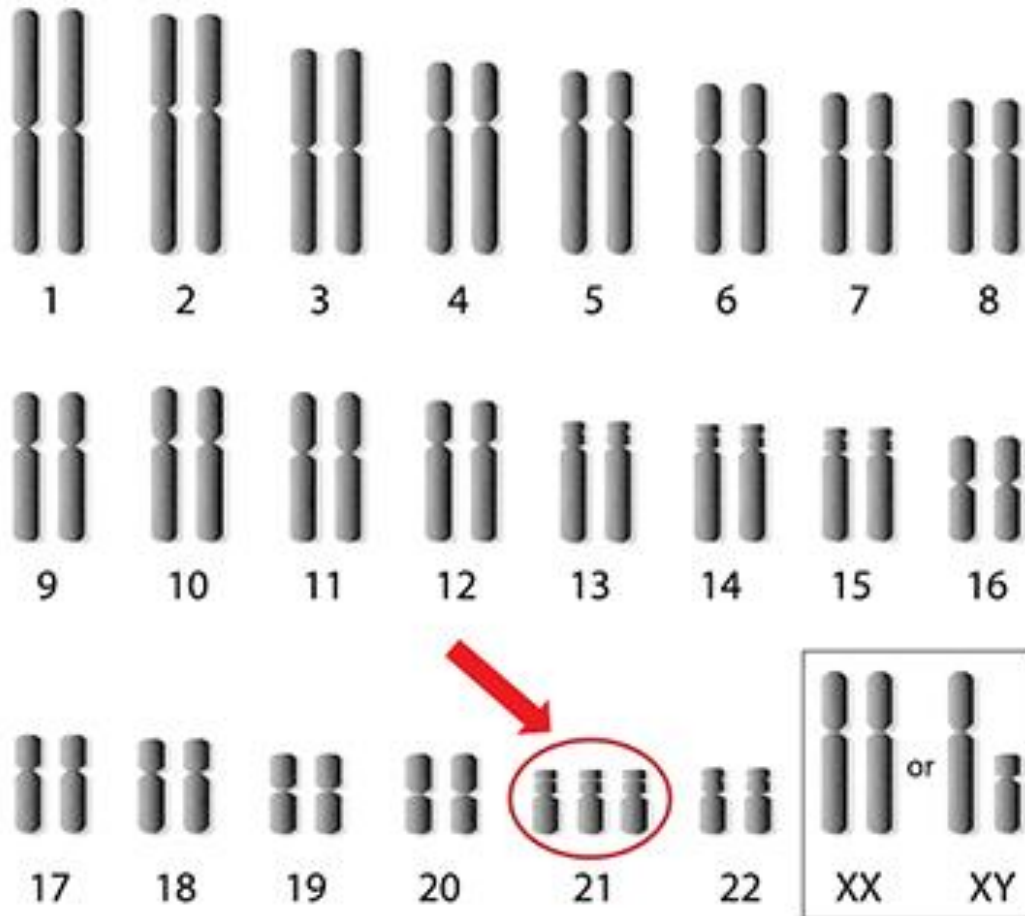
Genetic Variation

Aneuploidy

Aneuploidy:

number of chromosomes
NOT equal to 46

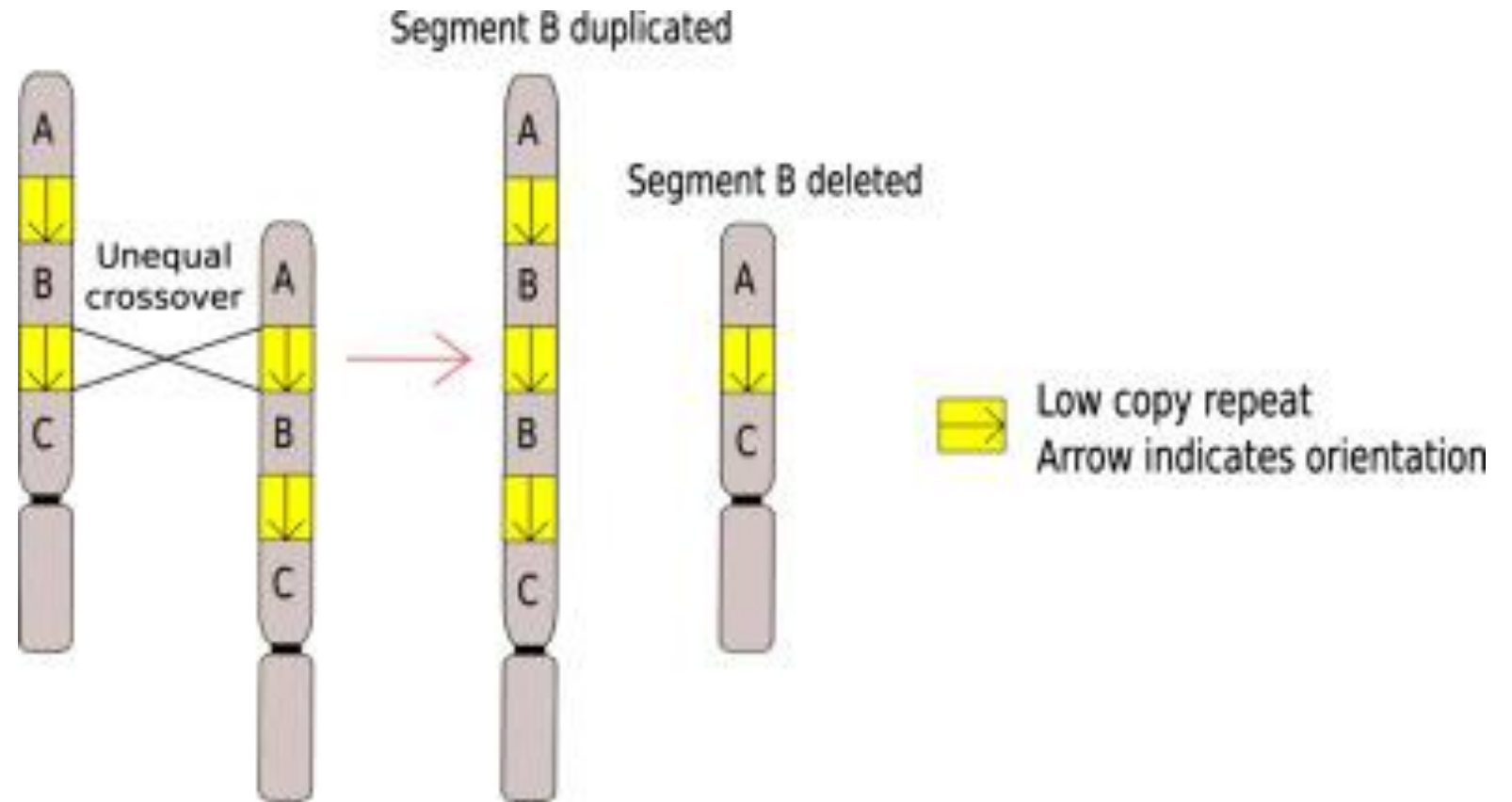
Down Syndrome - Trisomy 21



Copy Number Variants

Copy Number Variant:

duplications or deletions
greater than 1000 nucleotides
(1kb)



https://media.springernature.com/m312/springer-static/image/art%3A10.1038%2Fs10038-020-00838-1/MediaObjects/10038_2020_838_Fig1_HTML.png?as=webp

Indel

Indel:

insertion/deletion
smaller than 1kb

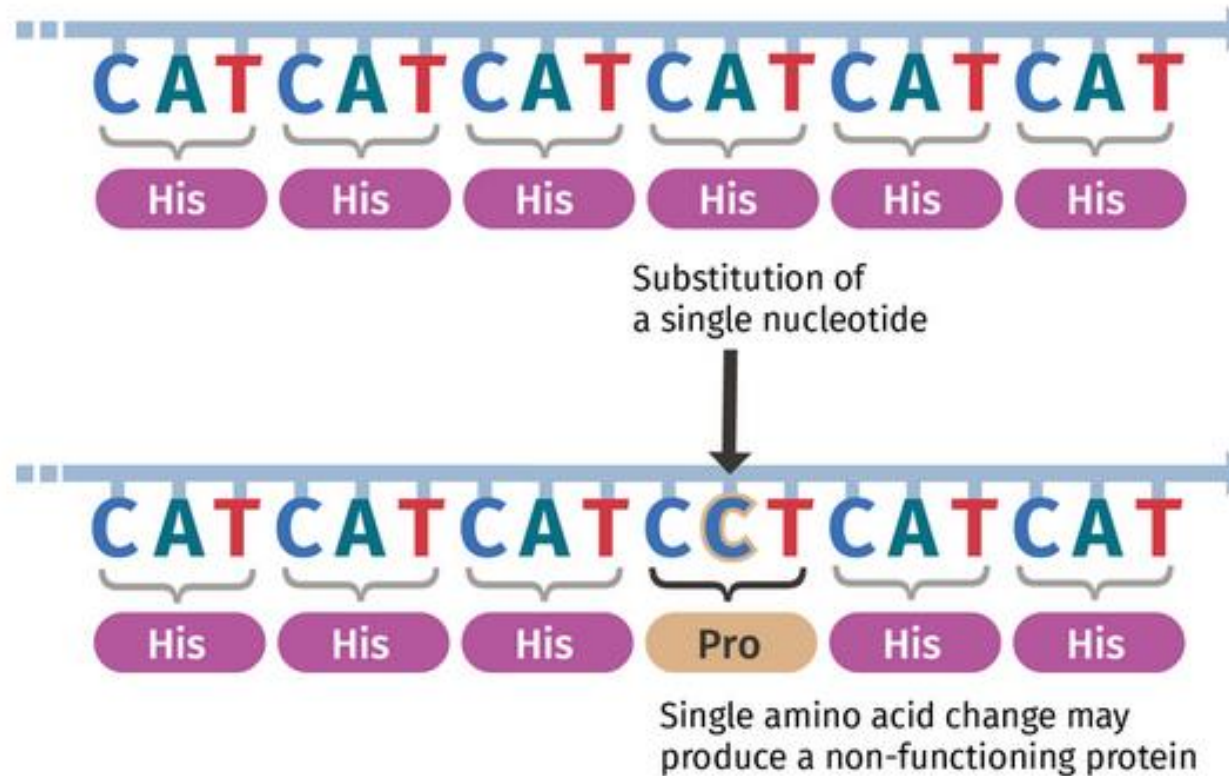
Normal: See the dog run.

Deletion: ~~See~~ hed ogr un.

Insertion: See eth edo gru n.

<https://www.cshl.edu/a-shift-in-the-code-new-method-reveals-hidden-genetic-landscape/>

Single Nucleotide Variant



Repeat expansion disorders:

More than 40 diseases, most of which primarily affect the nervous system, are caused by expansions of simple sequence repeats dispersed throughout the human genome.

Difficult to detect on exome and genome.

Fragile X

Group of disorders

5' UTR TRDs

- FXS
- FXTAS
- Other FX disorders

Huntington Disease

Intronic TRDs

- FRDA
- C9ORF72 TRDs (includes subset of ALS and FTD)

Polyglutamine TRDs

- HD
- SCA1, SCA2, SCA3, SCA6, SCA7 and SCA17
- SBMA (Kennedy disease)
- DRPLA

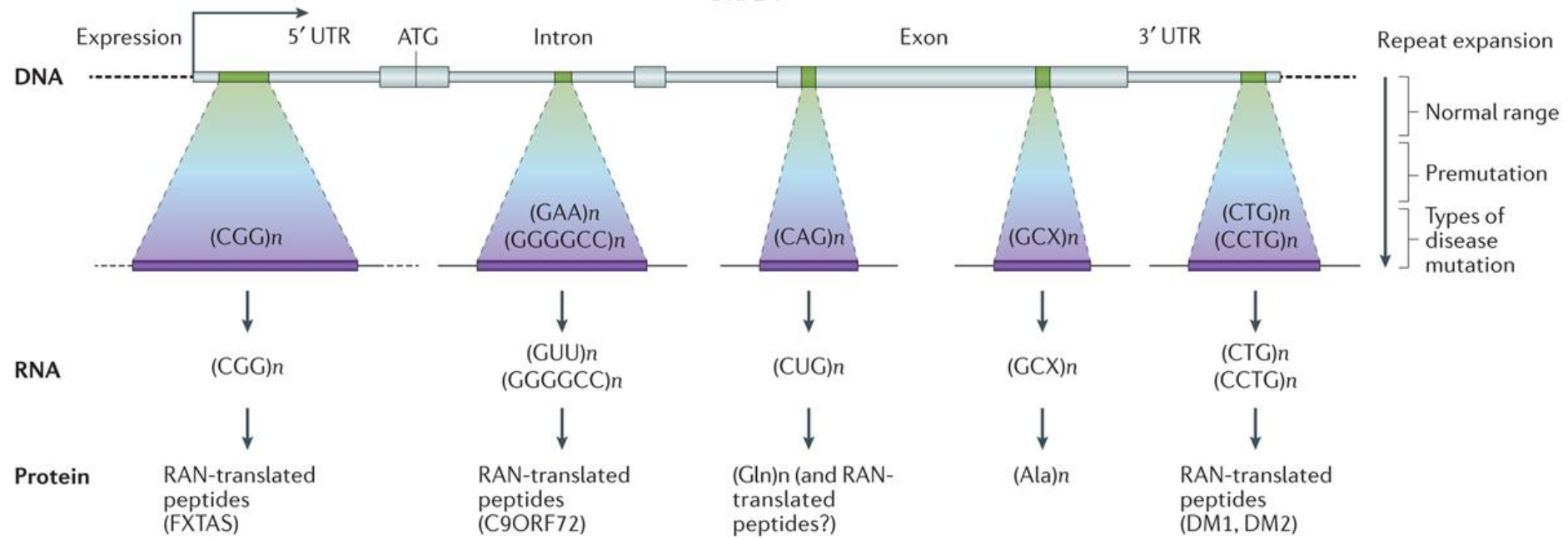
Myotonic dystrophy

Polyalanine TRDs

- OPMD and eight other developmental disorders

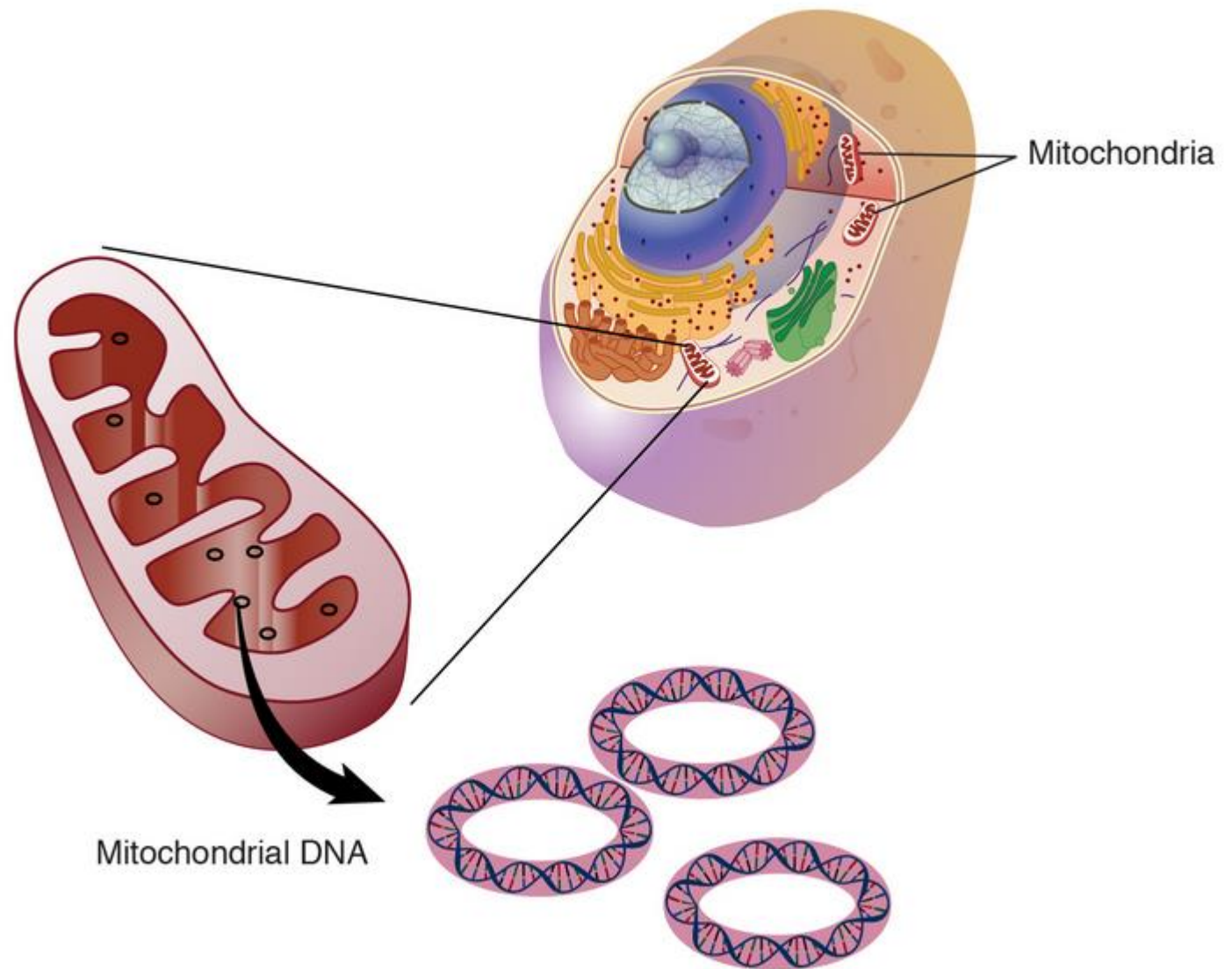
3' UTR TRDs

- DM1 and DM2



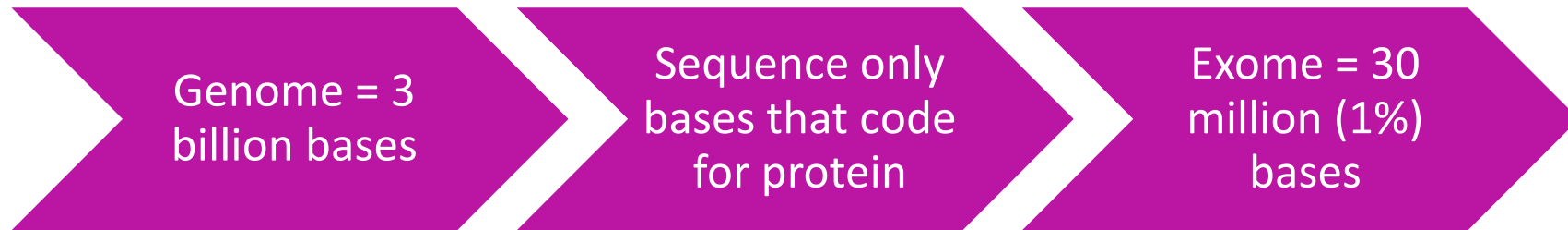
Mitochondrial DNA

Maternal inheritance



Exomes and Genomes

Genomes and Exomes



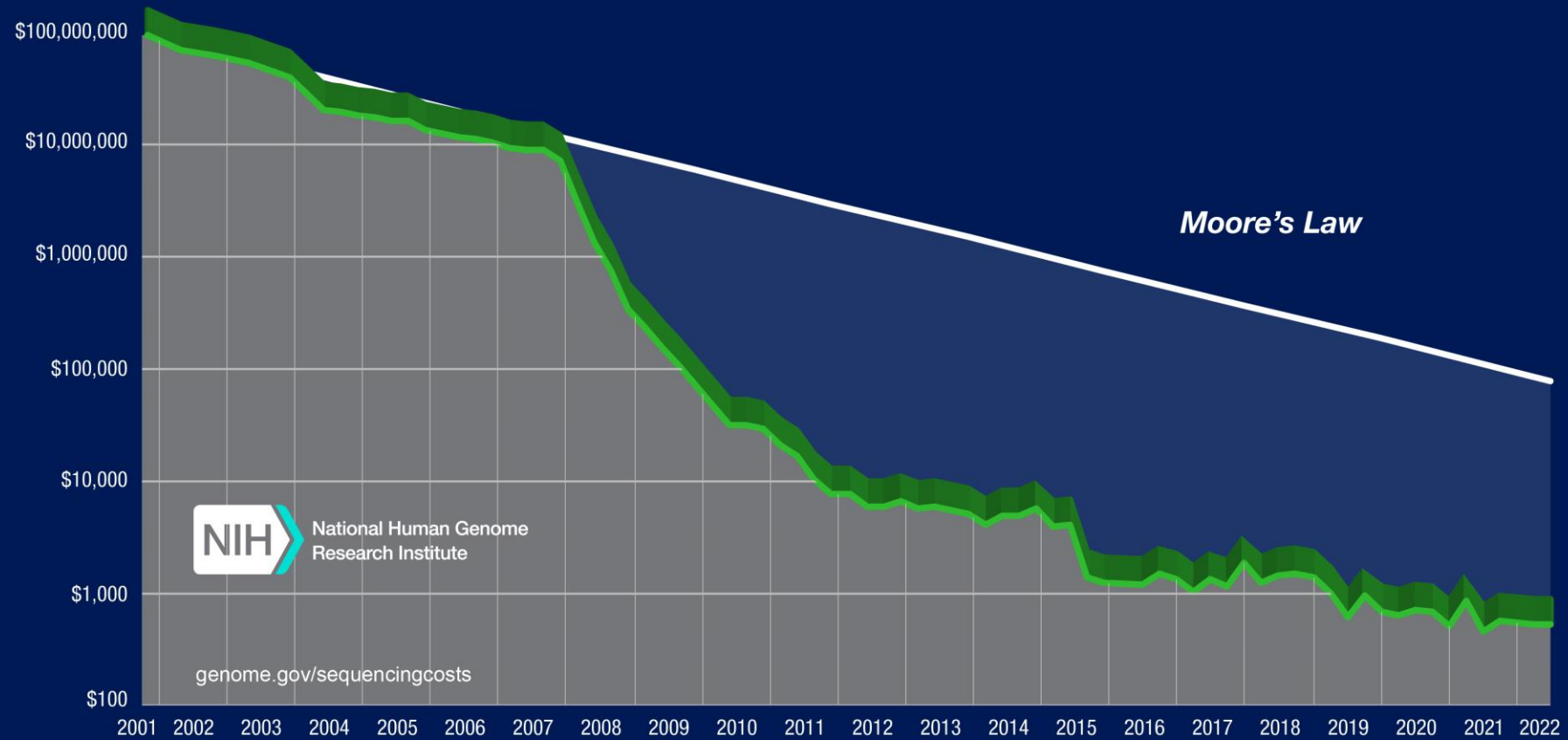
The exons are separated by DNA of unclear function

Wpod?amfkwcg.gjhklfoursjckfoquscore and void
m\$%djkdllfkk*wqnfjjdxnnebkyp@mvjckdfkkseocb
qw.oiwjfm du seven years ago dlfksl8\$((k dkm
deixmenfyrucci our skdj\$ mvkjdfk&%woqppa lfdk
kfaqaq.d eiidty forefathers brought jdd qpo
ooekfjk vbzxx dss forth a...

Genomes and Exomes

- Whole genome sequencing: file size of 150GB
- Whole exome sequencing: file size of 6-8GB
- Cost of whole genome in 2001: \$92,000,000
- Cost of whole genome today: \$700

Cost per Human Genome



In the NICU, rapid whole genome sequencing (rWGS/rGS) can deliver answers for critically ill newborns, so that the clinician can focus on what's next. @MichiganHHS just took a big step forward by providing coverage for rapid #genome testing. Learn more 🖱️ bit.ly/2YWgKjY



INSURANCE COVERAGE NEWS

Michigan is the first state to offer Medicaid coverage for rapid whole genome (rWGS) testing for eligible, critically ill infants



UnitedHealthcare® Commercial
Medical Policy

Whole Exome and Whole Genome Sequencing

Policy Number: 2023T0589M
Effective Date: March 1, 2023

[Instructions for Use](#)

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Related Commercial Policies

- [Chromosome Microarray Testing \(Non-Oncology Conditions\)](#)
- [Molecular Oncology Testing for Cancer Diagnosis, Prognosis, and Treatment Decisions](#)
- [Preimplantation Genetic Testing and Related Services](#)

Community Plan Policy

[Whole Exome and Whole Genome Sequencing](#)
[Percentage Coverage Summaries](#)
[Testing](#)
[Tests and Services](#)

Medical Coverage Policy



Effective Date..... 4/15/2022
Next Review Date..... 1/15/2023
Coverage Policy Number 0519

Whole Exome and Whole Genome Sequencing

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Newborn Screening



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English



The GUARDIAN study is a free newborn screening study to help all babies have healthier lives.

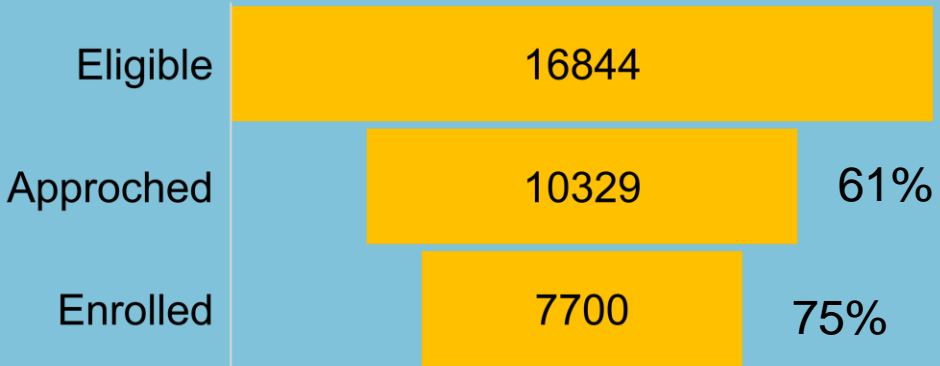
[LEARN MORE](#)



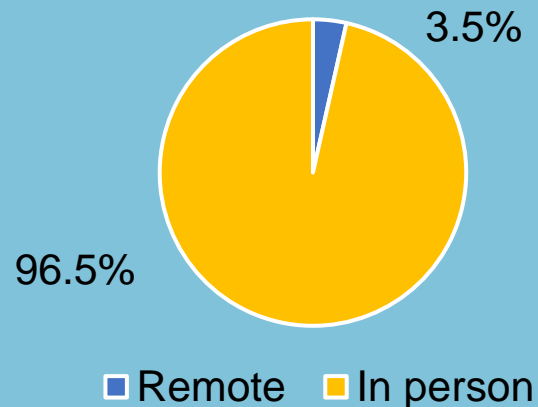
<https://guardian-study.org/>

RESULTS: FIRST 7,700

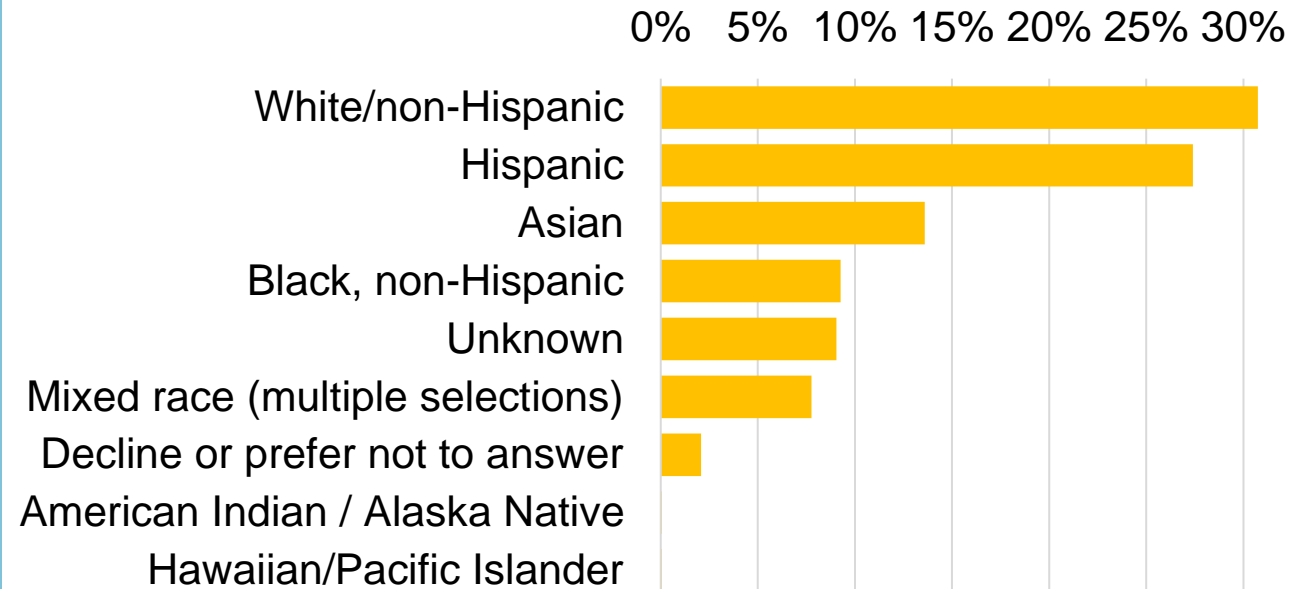
ENROLLMENT



TYPE OF CONSENT



ANCESTRY

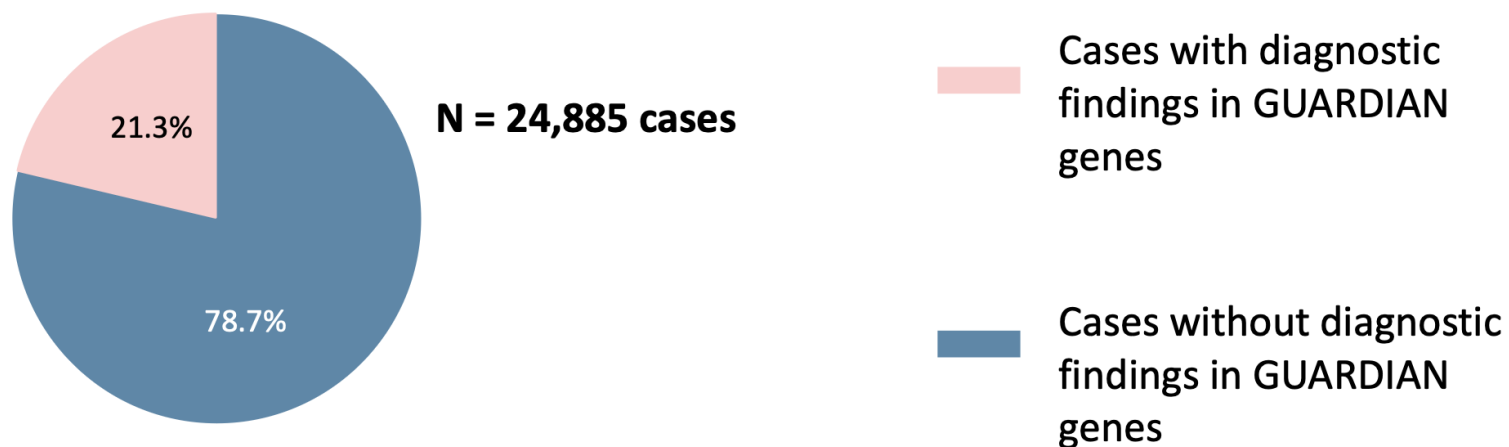


GROUP 2 CONSENT

	Overall	English	Spanish	Mandarin
Groups 1 + 2	92% (7100)	93.2% (6024)	83.6% (754)	92.2% (249)
Group1 only	8% (612)	6.8% (442)	16.4% (148)	7.8% (21)



RARE DISEASE COHORT MOLECULAR-DIAGNOSED BY ES/GS AT GENEDX



Data presented at ACMG 2023

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GeneDx

Name of Disease		
ADSSL1 Myopathy	X	
Alpha Mannosidosis	Group 1	
Aspartylglucosaminuria	Group 2	
Aspartylglucosaminuria		
Aspartylglucosaminuria		
ATP6V1A Encephalopathy and other v-ATPase related disorders	Group 1 – ATP6V1B	
Baker Gordon Syndrome / SYT1 Syndrome	X	
Bosch Boonstra Schaaf Optic Atrophy Syndrome (BBSOAS)	X	
CDKL5	Group 2	
CDKL5		
CHD2	Group 2	
CHD2		
ECHS1	X	
FOXP1 Syndrome	Group 2	
GABA-A	X	
Hnrnp related neuro developmental disorders, hnrnpu hnrnpk hnrnpq/syncrip	Group 2 – HNRNPH2 and HNRNPU	
Jansen de Vries Syndrome	X	
MEPAN Syndrome	X	
Multiple		
Myhre Syndrome	X	
NF1	X	
Ornithine Transcarbamylase Deficiency (OTCD), Early Onset	Group 1	
Pyruvate Dehydrogenase Complex Deficiency	X	
Pyruvate Dehydrogenase Complex Deficiency (PDCD)		
SMC1A	X	
SPG50 and CMT4J	X	
SYNGAP1-Related Disorder	Group 2	
X-Linked Dystonia-Parkinsonism	X	

Research Triangle Institute Partners With GeneDx, Illumina for Expanded Newborn Screening Study

Sep 21, 2023 | [staff reporter](#)

NEW YORK – Research Triangle Institute (RTI) International said on Wednesday that it has partnered with Illumina and GeneDx to expand its Early Check newborn screening study to include whole-genome sequencing and genetic risk scores for type 1 diabetes.

Core funding of unspecified size for the expansion comes from the Leona M. and Harry B. Helmsley Charitable Trust, diabetes foundation JDRF, and Travele Therapeutics, with additional support from Orchard Therapeutics.

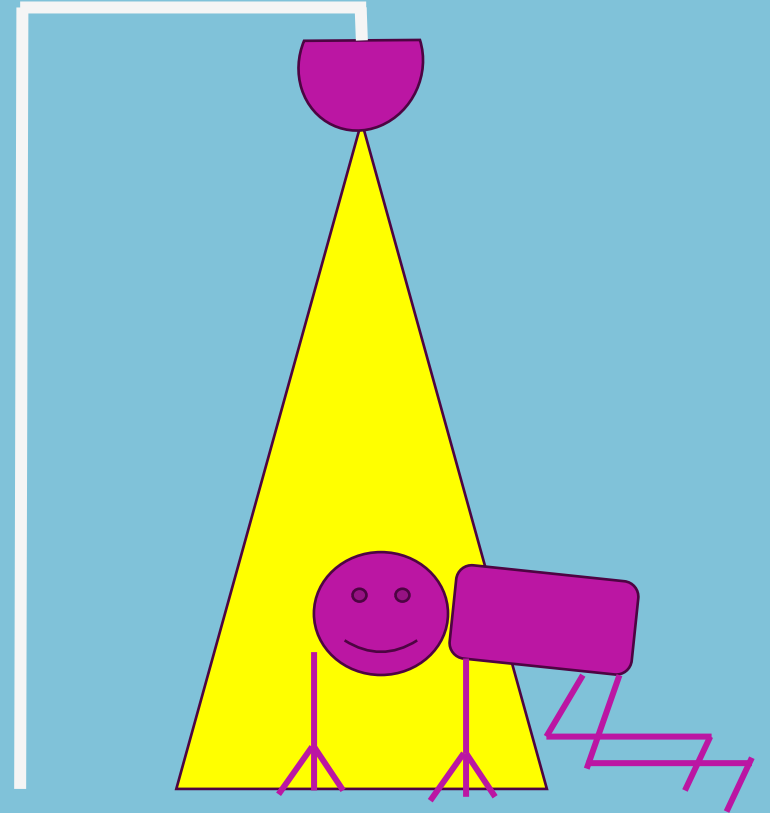
Cohort Building

Cohort Building



Gay Grossman
Patient Advocacy and Engagement Lead
ggrossman@genedx.com

Future Technology



GeneDx

Long Read Sequencing Projects at GeneDx

Montefiore in the News

The New York Center for Rare Diseases at Montefiore to Partner with GeneDx, PacBio and Google Health to Increase Genomic Testing Options for Bronx Families

November 8, 2023

New Partnership to Advance Data, Enhance Disease Detection and Care in Diverse Communities

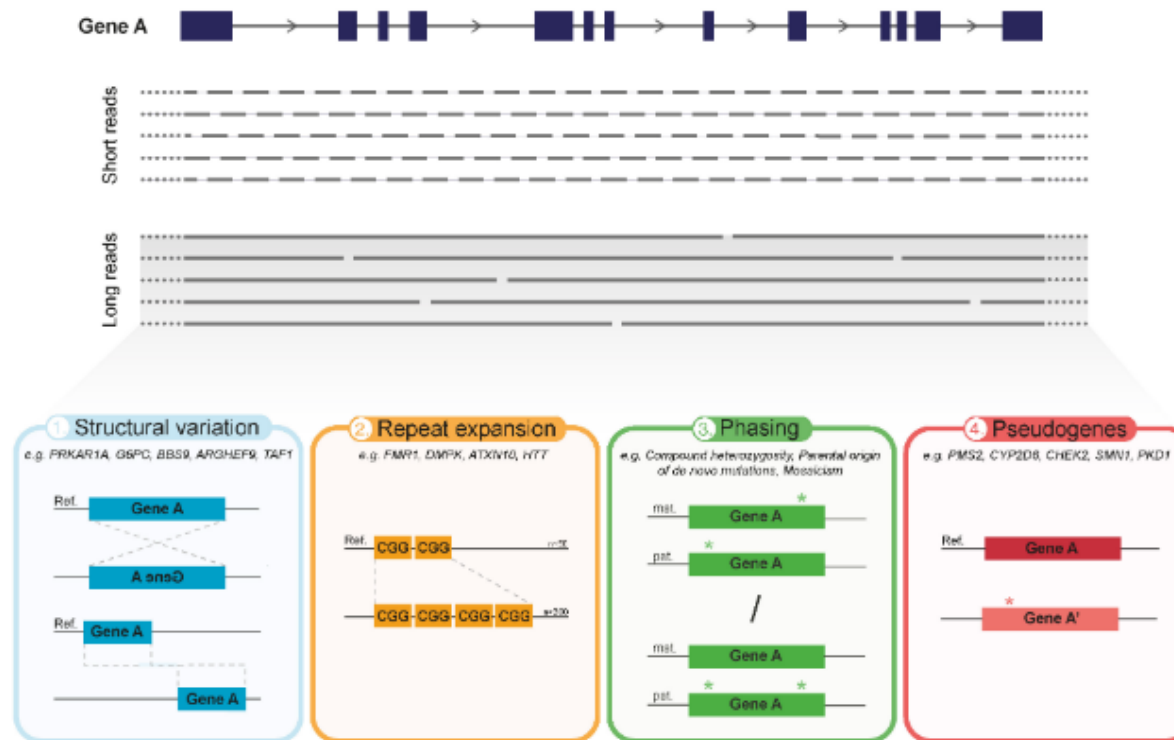
BRONX, NY—November 8, 2023. The New York Center for Rare Diseases (NYCRD) at Montefiore, [recognized](#) as a Center of Excellence by the National Organization for Rare Disorders (NORD), is partnering with GeneDx, PacBio and Google Health to deliver genetic diagnoses for Bronx families living with rare diseases. The goal of the new partnership is to help identify the genetic causes of, and best treatments for rare diseases that have remained undiagnosed, despite using today's most advanced tests.

PacBio and GeneDx Launch Research Collaboration with the University of Washington to Study Long-Read Whole Genome Sequencing for Increased Diagnostic Yield in Neonatal Care

Study is first of its kind to compare diagnostic rates across short- and long-read sequencing platforms

MENLO PARK, Calif. and STAMFORD, Conn., Aug. 7, 2023 /PRNewswire/ — PacBio (Nasdaq: PACB), a leading developer of high-quality, highly accurate sequencing solutions and GeneDx (Nasdaq: WGS), a leader in delivering improved health outcomes through genomic and clinical insights, today announced a research collaboration with the University of Washington to study the capabilities of HiFi long-read whole genome sequencing (WGS) to increase diagnostic rates in pediatric patients with genetic conditions.

Long Read Sequencing





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Thank You



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