

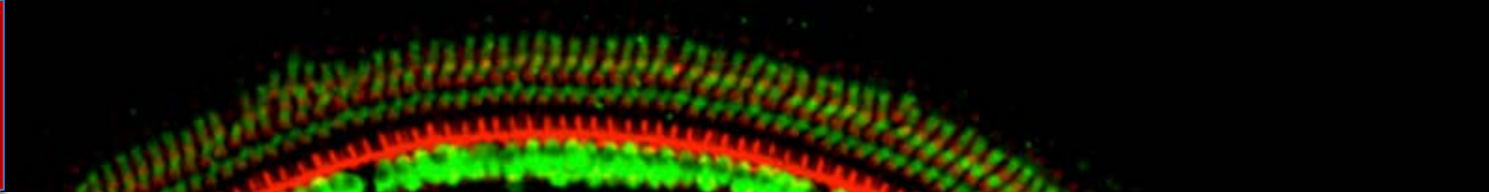
Next-Gen Newborn Hearing Screening

Cynthia C. Morton, PhD, FFACMG
Brigham and Women's Hospital
Harvard Medical School
Broad Institute of MIT and Harvard
University of Manchester

Harvard Medical School
Center for Hereditary Deafness



Manchester Centre for
Audiology and Deafness (ManCAD)

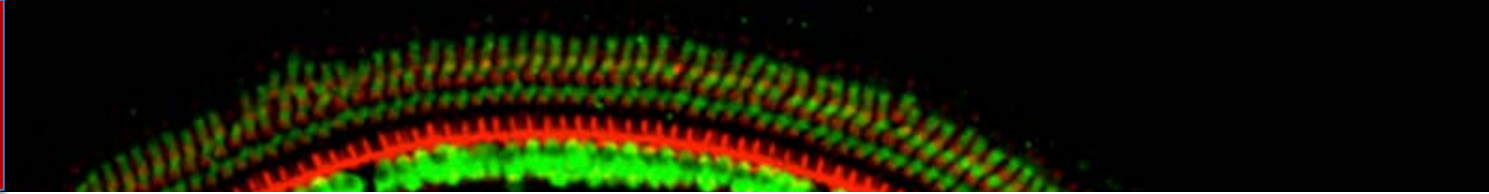


Next-Gen Diagnostics and Newborn Screening for Hearing Loss

- Why is genetic testing so important for hearing impairments?
- What is the landscape of genetic testing for hearing loss?
- What is the status of Universal Newborn Screening?
- What is the future of testing and screening?

How do we do it....and get it right?



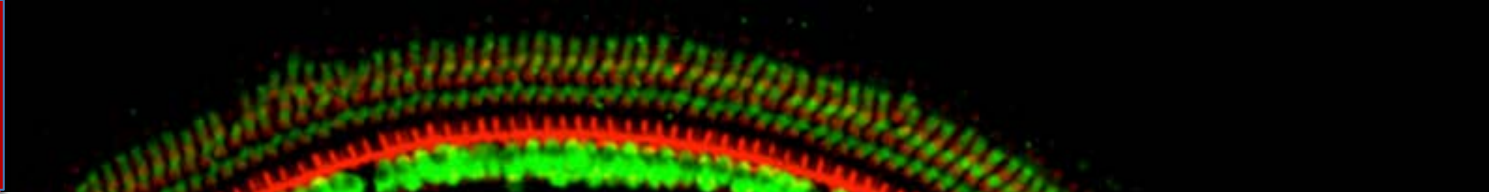


Hearing Loss Across the Lifespan

Most common sensory deficit in humans leading to speech and language delay, challenges in school, work and relationships, isolation and depression in the elderly with critical time for habilitation during the first few months of life

Newborns and Children (in the U.S.)

- ~3 in 1000 newborns born with permanent hearing loss (HL)
- One of most common birth defects in the United States
- Majority of hearing impaired children born into families with little or no experience with HL—Deaf X Deaf matings result in >90% all hearing offspring, highlighting unparalleled heterogeneity in etiology (genetic, environmental and gene X env)
- ~50% of children with HL from racial/ethnic minority populations
- ~1 in 2 cases in babies due to genetic causes
- ~1 in 4 cases in babies due to maternal infections during pregnancy, complications after birth, and head trauma
- ~1 in 100 cases in children by school age → 2.5M (5.4%) with mild or unilateral HL



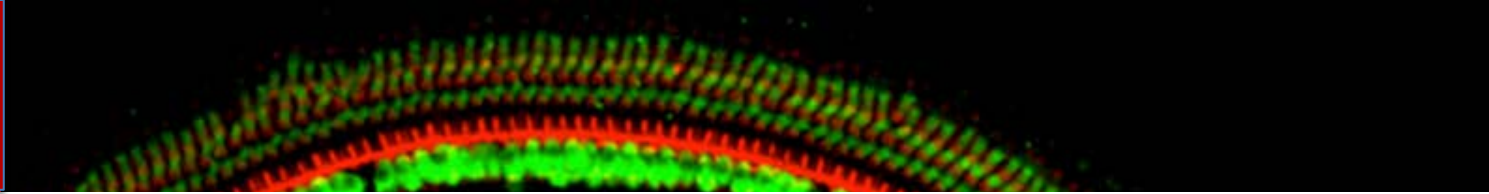
Hearing Loss Across the Lifespan

Age Related Hearing Impairment (ARHI)

- ~1 in 3 individuals over age 65 with hearing impairment significant enough to impair speech perception
- By 2030 over 20% of the U.S. population will be >76 years old (U.S. Census Bureau)
- More prevalent in males than females
- Complex trait with genetic and environmental factors contributing to onset and progression
- Little progress to date in understanding the underlying molecular basis

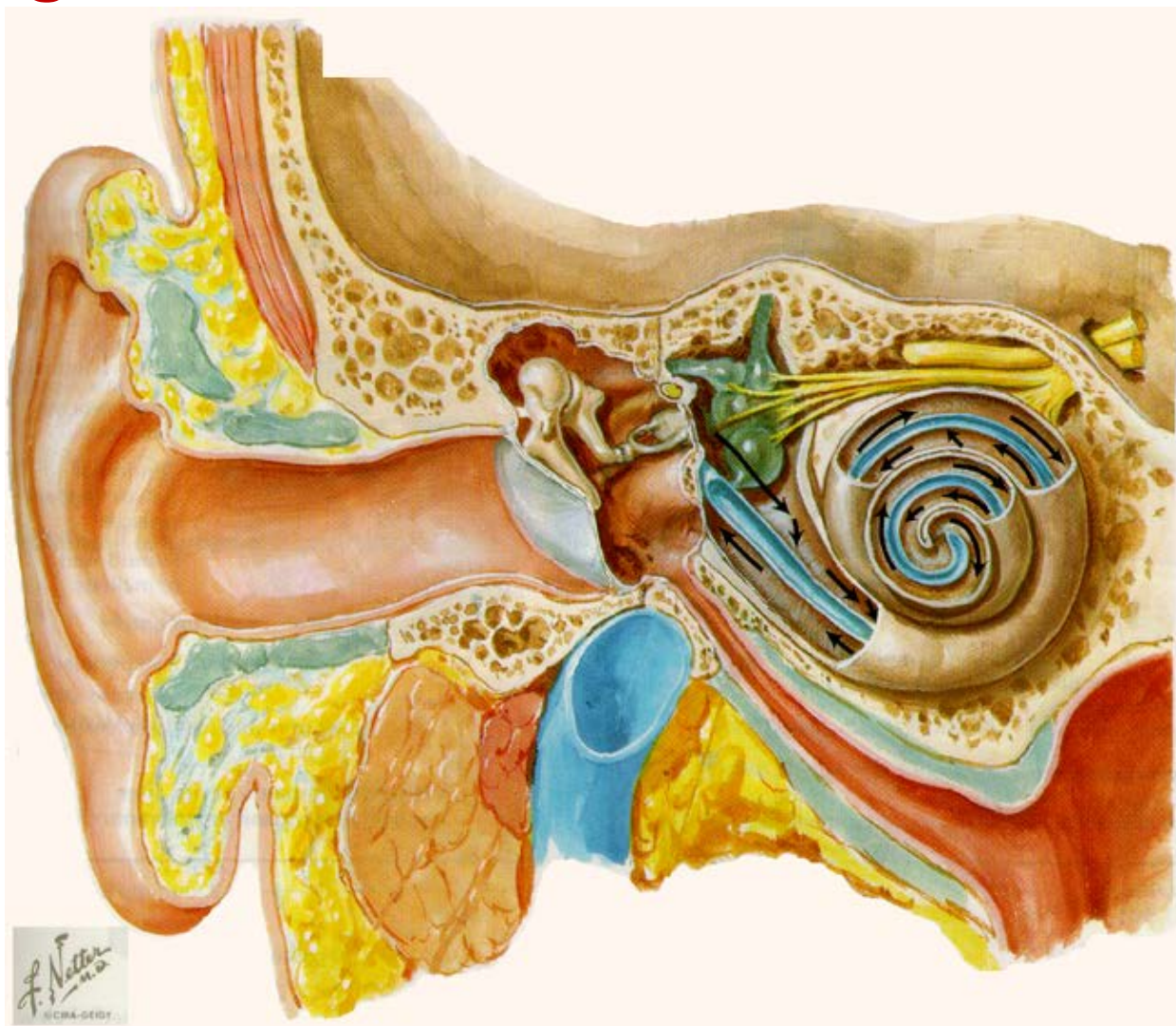
But, stay tuned...

- Recent genetic methods empowered by big data resources from the human genome project, such as genome wide association studies (GWAS), provide an approach to deciphering complex traits and are being applied to ARHI.

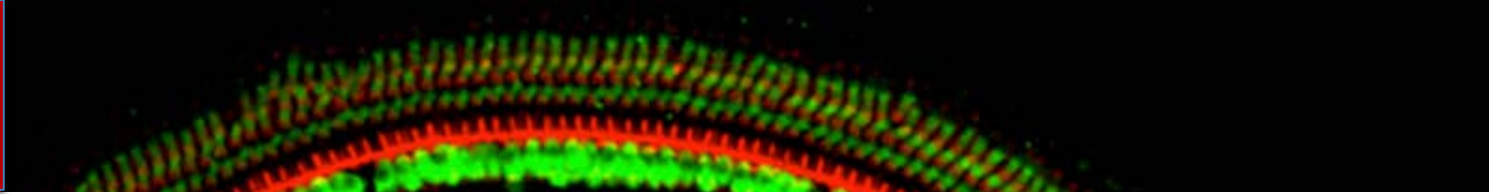


Types of Hearing Loss

- Unilateral
- Bilateral
- Conductive
 - Outer ear
 - Middle ear
- Sensorineural
 - Inner ear
- Mild
- Moderate
- Profound
- Stable
- Progressive
- Syndromic
- Nonsyndromic



F. Netter
M.D.
© CRA-DEPT



Heterogeneous Causes of Hearing Loss

Environmental

- Drug
- Noise
- Trauma
- Infection
- Malnutrition

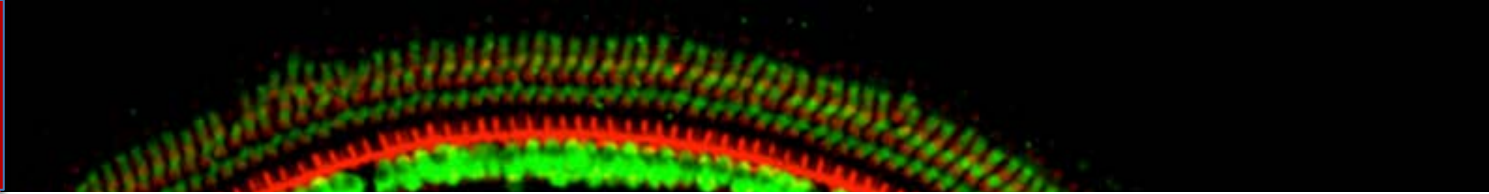
Genetic

Syndromic (~30%)

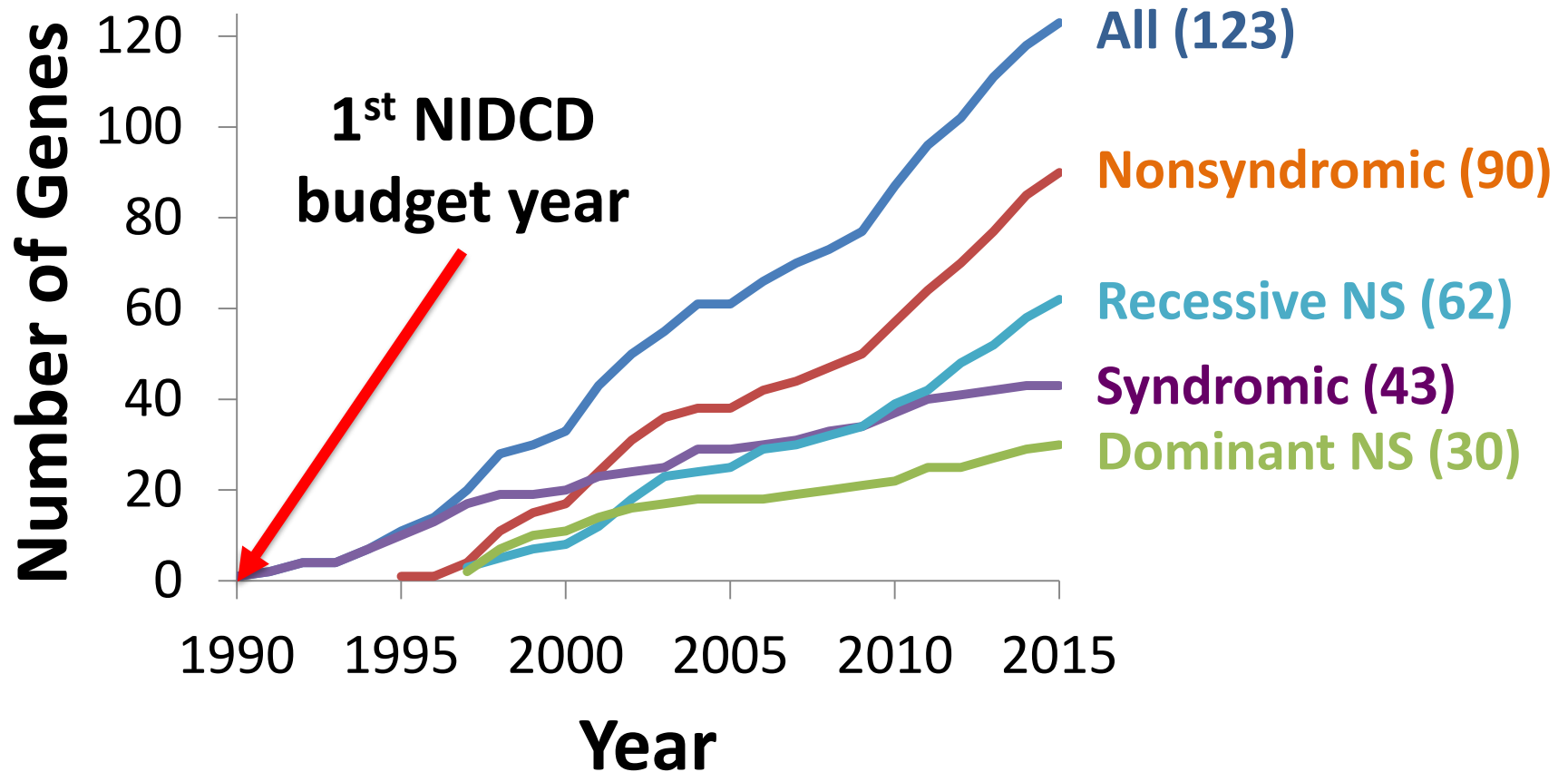
Alport, BOR, CHARGE, JLN, Norrie, Pendred, Perrault, Stickler, Treacher Collins, Usher, Waardenburg, Wolfram

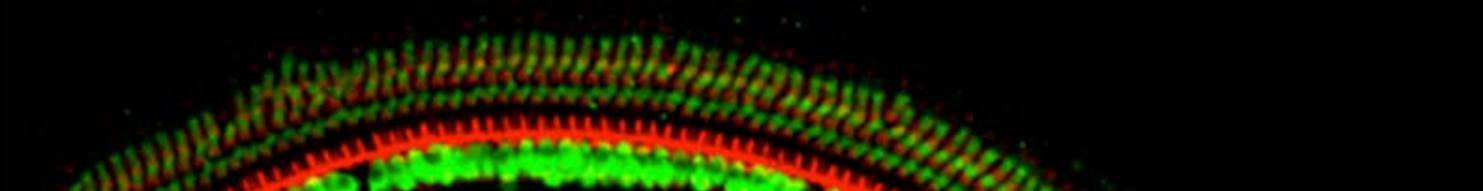
Nonsyndromic (~70%)

- Autosomal dominant (DFNA, 22%)
- Autosomal recessive (DFNB, 77%)
- X-linked (DFNX, 1%)
- Y-linked (DFNY)
- Mitochondrial

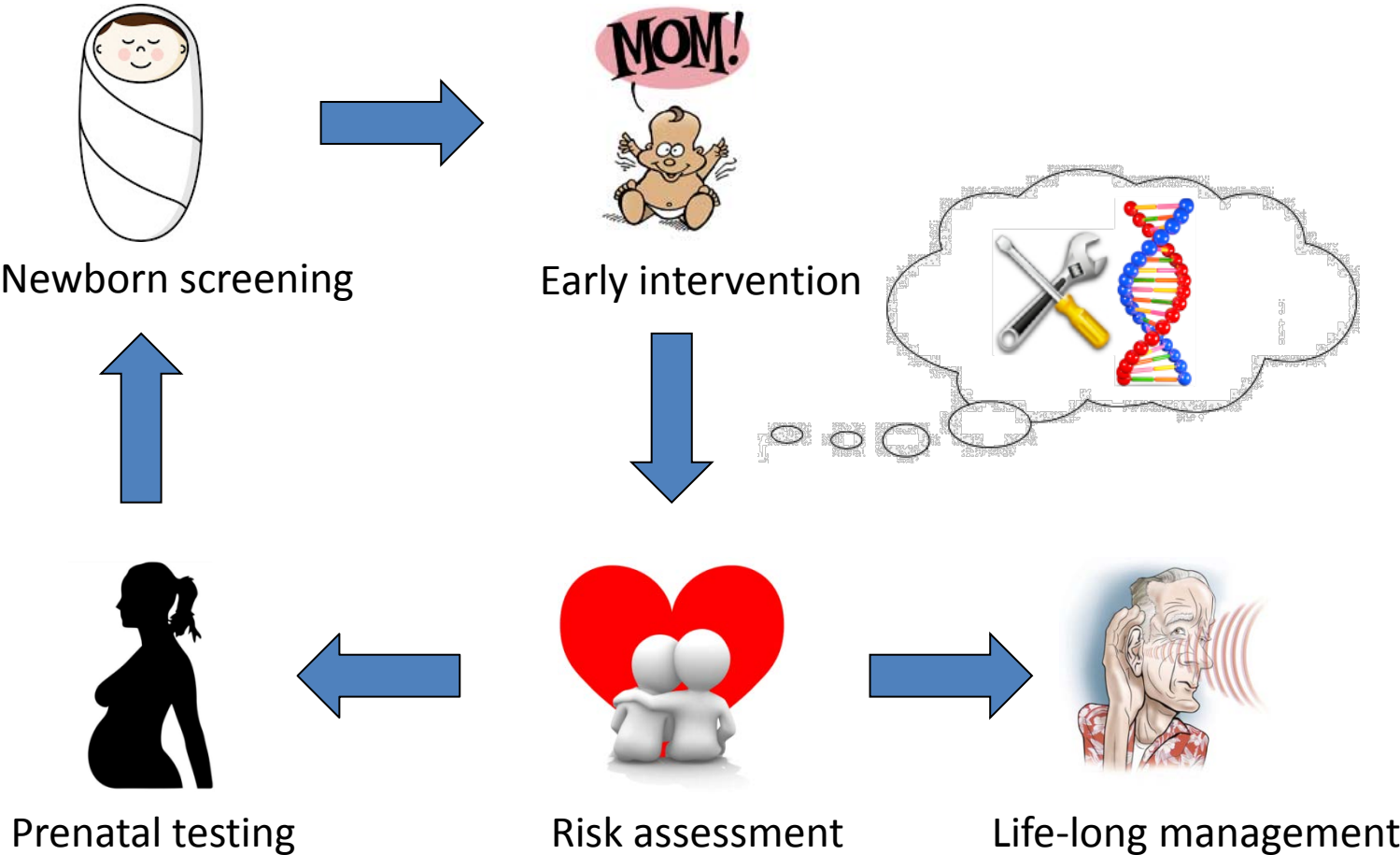


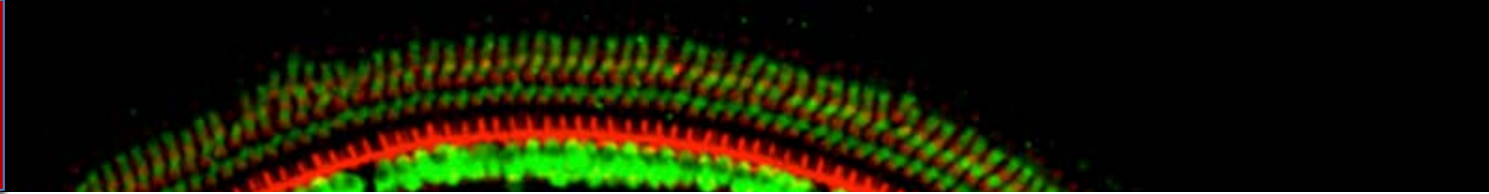
Known Hearing Loss Genes





Clinical Utility of Genetic Testing

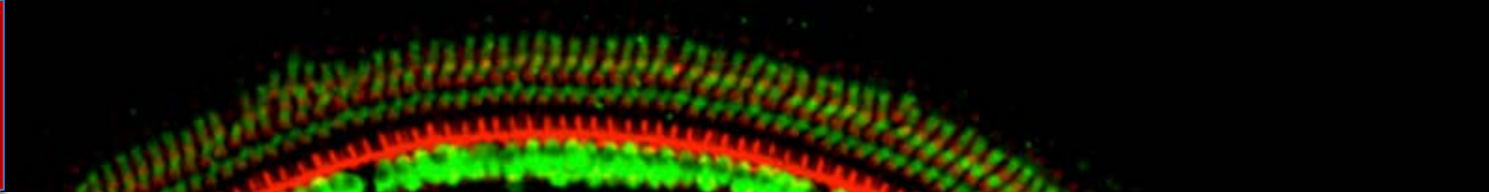




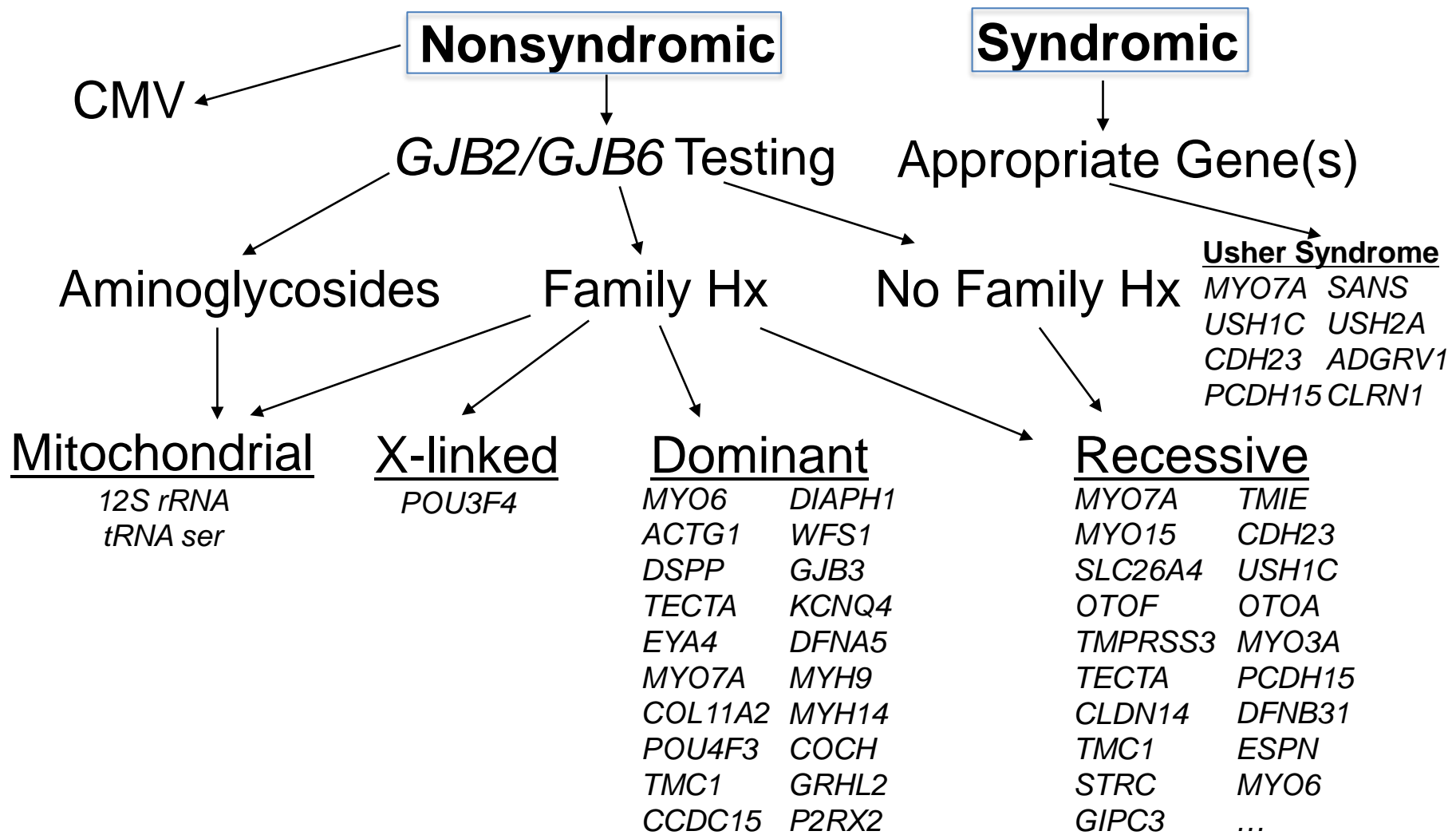
As Gene Test Menu Grows, Who Gets to Choose?

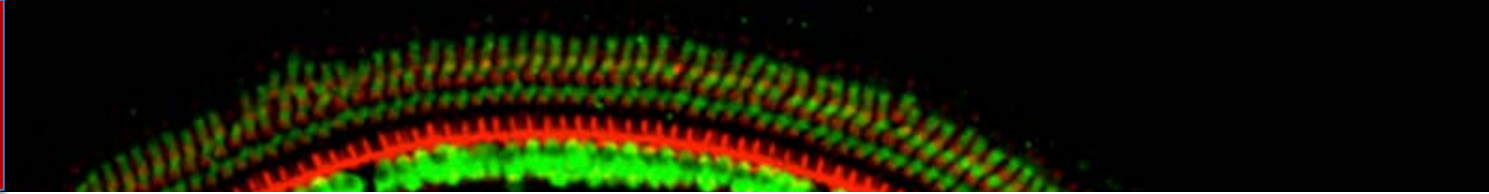
The New York Times
Wednesday, July 21, 2004



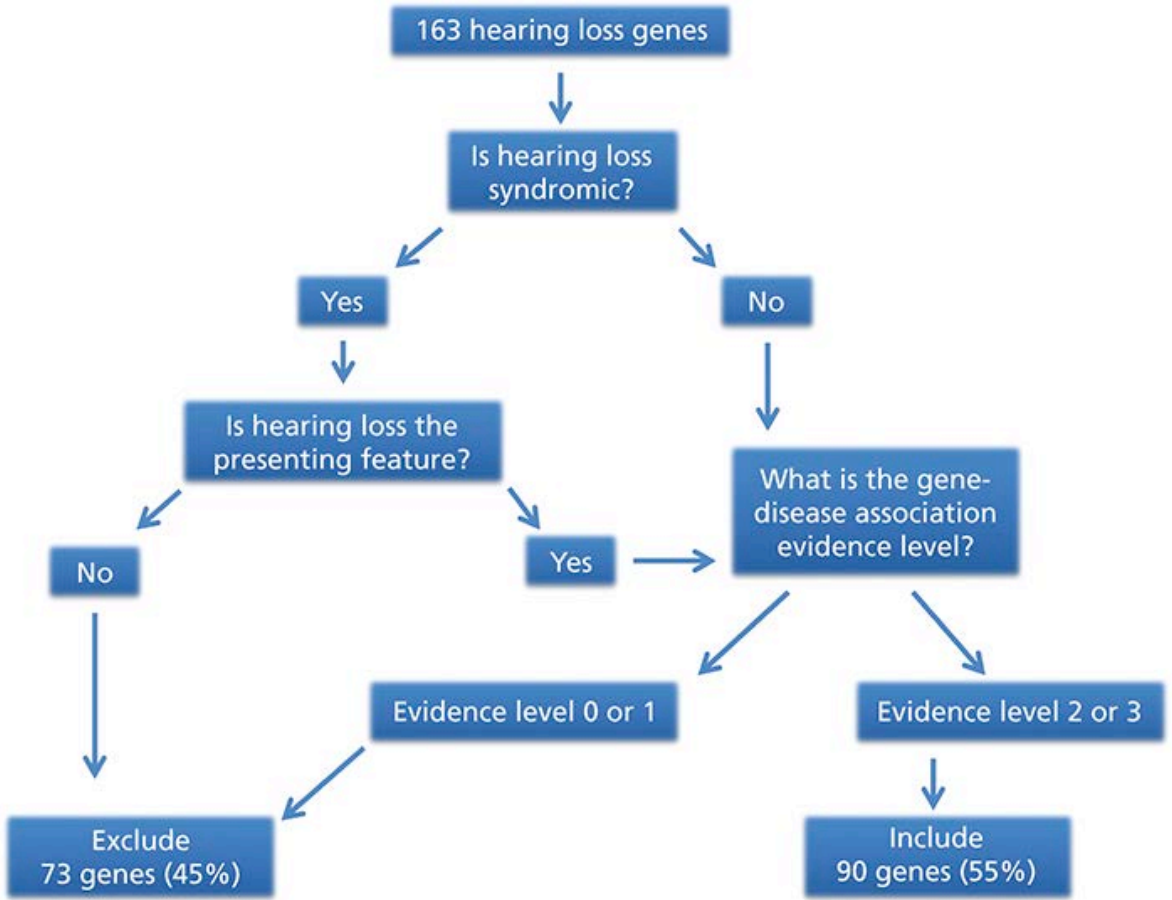


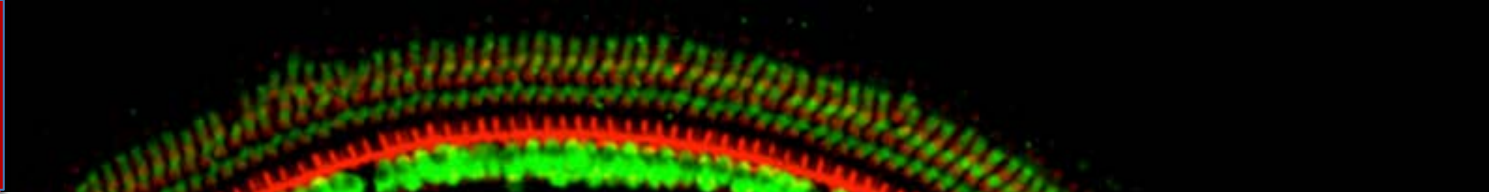
Hearing Loss Screening Protocol





Panel Inclusion Criteria for Nonsyndromic or “Apparent” Nonsyndromic Hearing Loss Genes





ACTG1	GPSM2	OTOG
ATP6V1	GRHL2	OTOGL
BSND	GRXCR1	P2RX2
CATSPER2	HGF	POU3F4
CACNA1D	ILDR1	POU4F3
CCDC50	KARS	PRPS1
CEACAM16	KCNQ4	RDX
CLDN14	LHFPL5	SERPINB6
COCH	LOXHD1	SLC17A8
COL11A2	LRTOMT	SMPX
CRYM	MARVELD2	STRC
DFNA5	MIR96	SYNE4
DIABLO	MSRB3	TBC1D24
DFNA5	MTRNR1	TECTA
DIAPH1	MTTS1	TIMM8A
ESPN	MYH14	TMC1
ESRRB	MYH9	TMIE
EYA4	MYO15A	TMPRSS3
GIPC3	MYO3A	TPRN
GJB2	MYO6	TRIOBP
GJB6	OTOF	TPSPEAR

ADGRV1
CDH23
CIB2
CLRN1
DFNB31
HARS
MYO7A
PCDH15
USH1C
USH1G
USH2A
SLC26A4
OTOA
DFNB59
CLPP
HARS2
HSD17B4
LARS2
KCNE1
KCNQ1

Usher

Pendred

AN

Perrault

JLNS

OtoGenome

EDN3
EDNRB
MITF
PAX3
SNAI2
SOX10
EYA1
SIX1
WFS1
CATSPER2

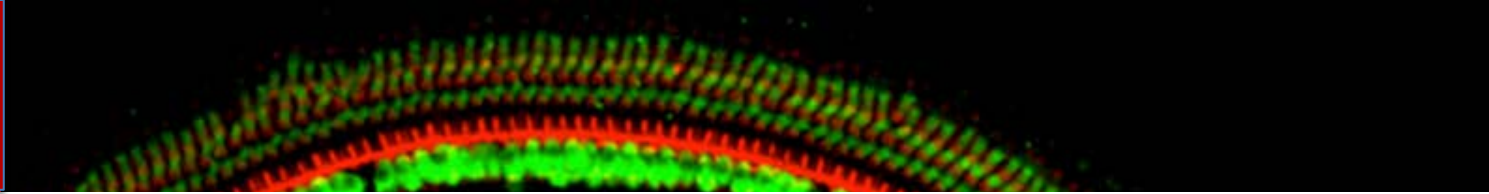
Waardenburg

BOR

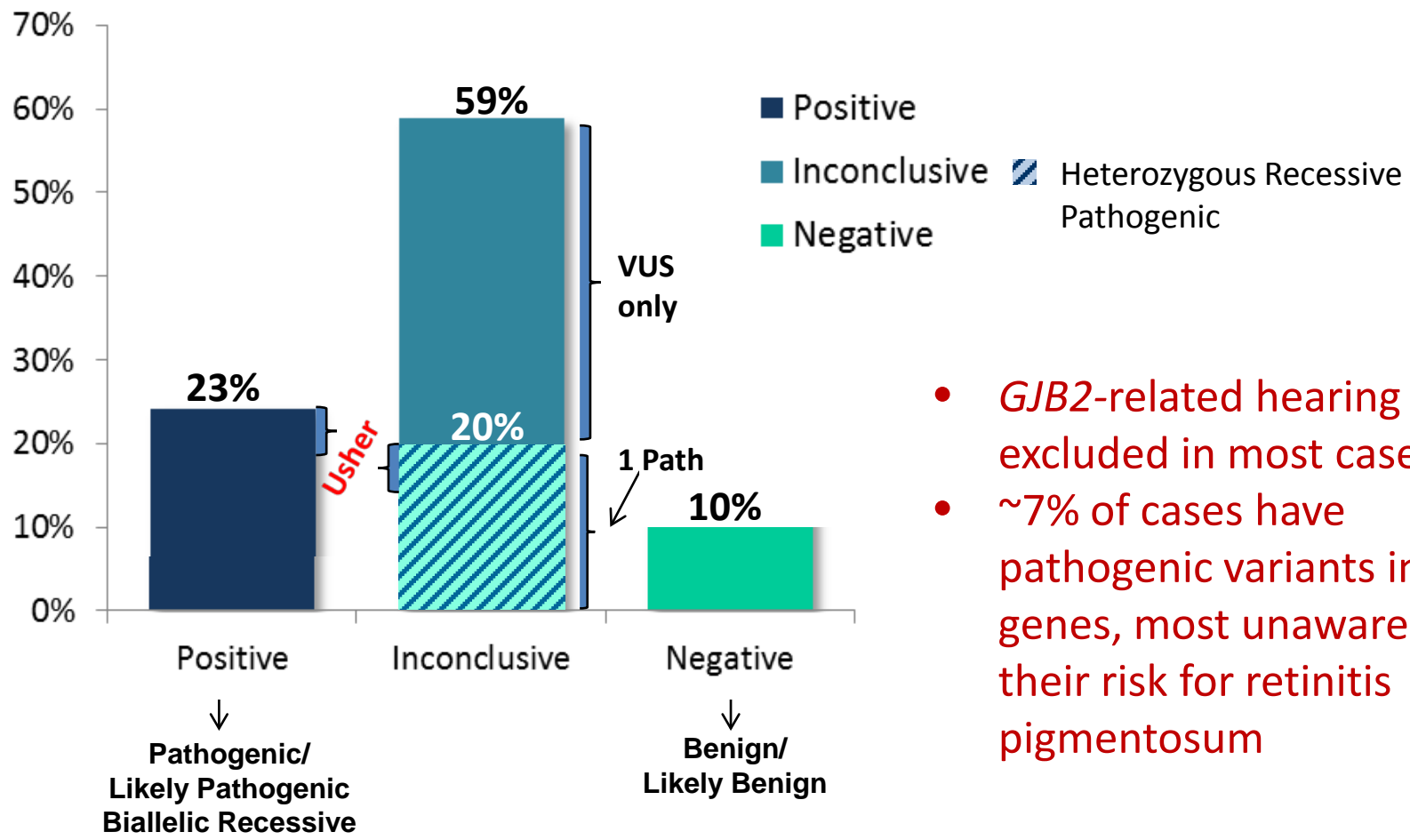
Wolfram

DIS

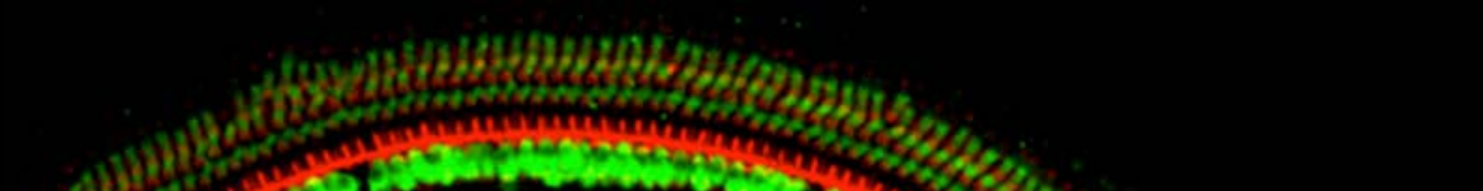
Subpanels available
(Usher, Waardenburg,
BOR, JLNS, Pendred,
AN, WFS1, Mito, etc.)



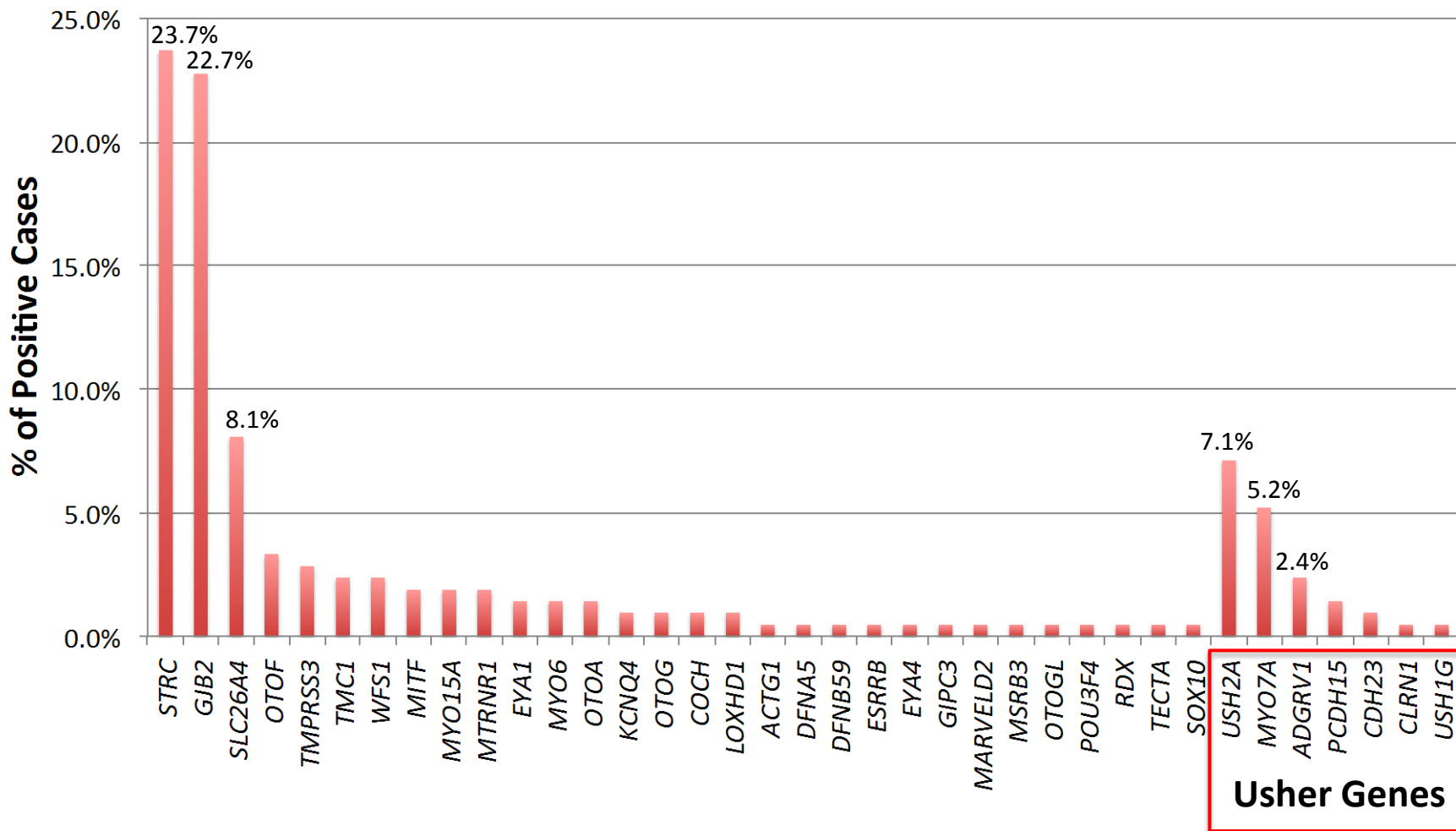
OtoGenome Detection Rates (n=959)

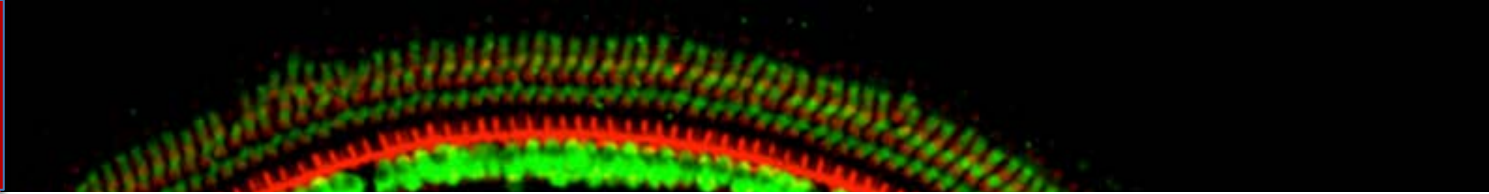


- *GJB2*-related hearing loss excluded in most cases
- ~7% of cases have pathogenic variants in Usher genes, most unaware of their risk for retinitis pigmentosum

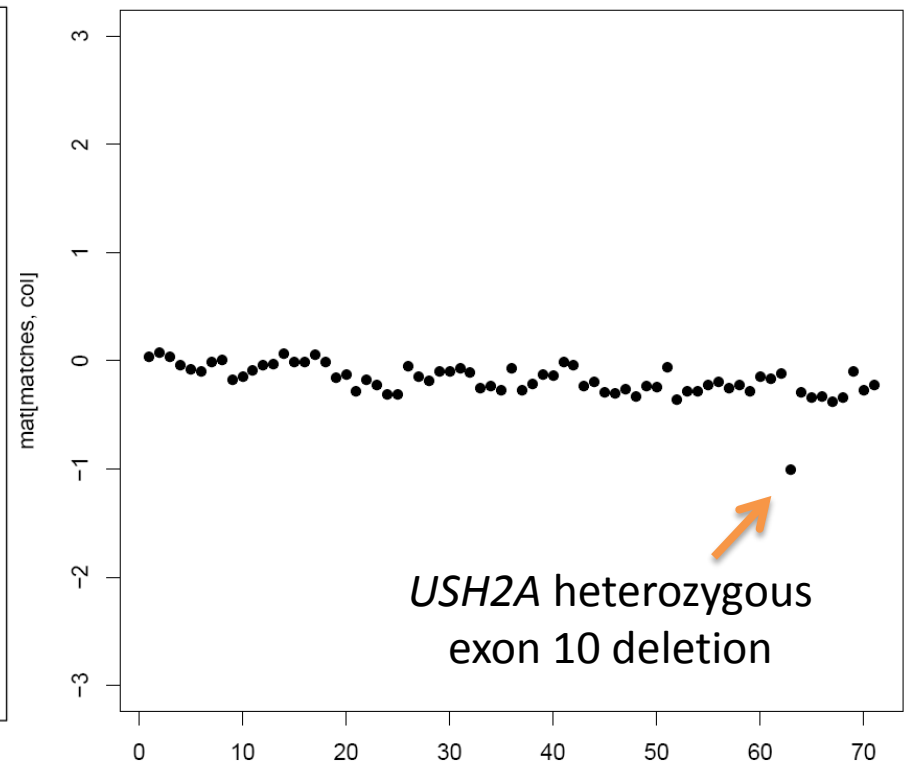
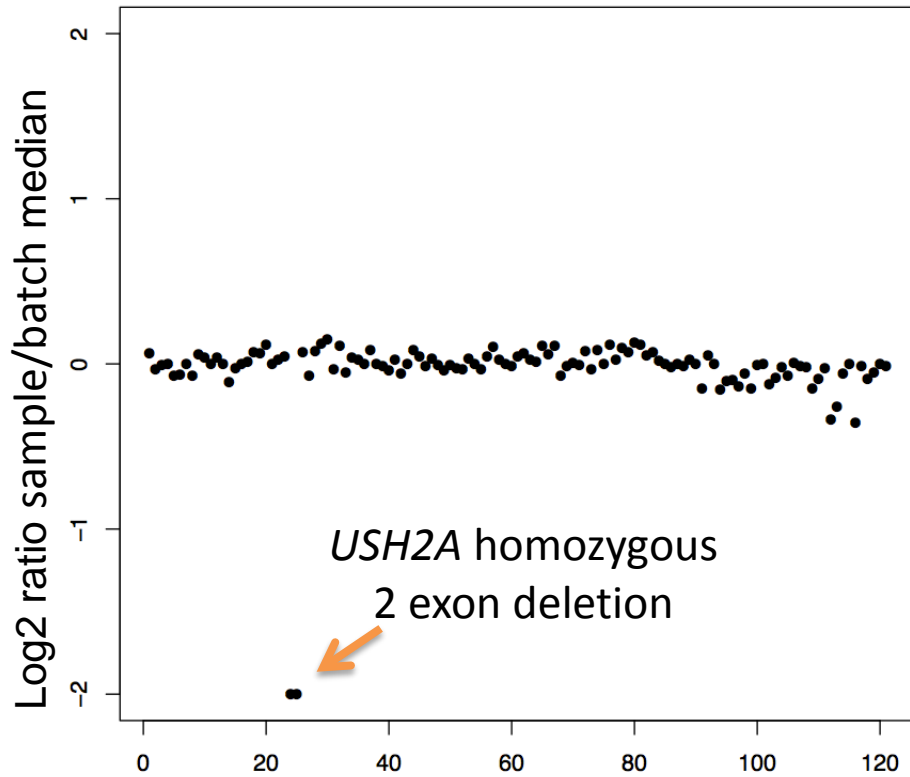


Genetic Etiology of 218 OtoGenome Positive Cases

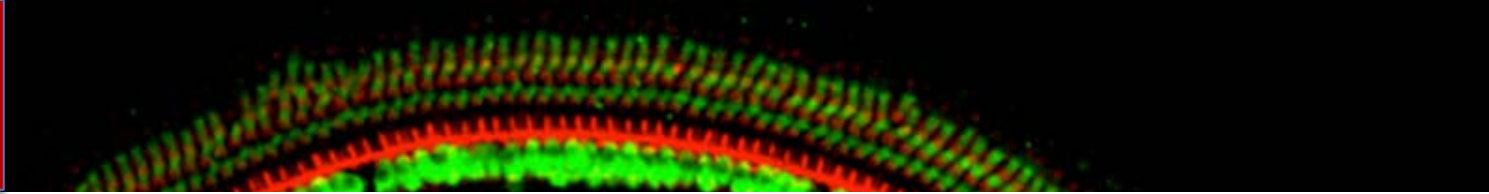




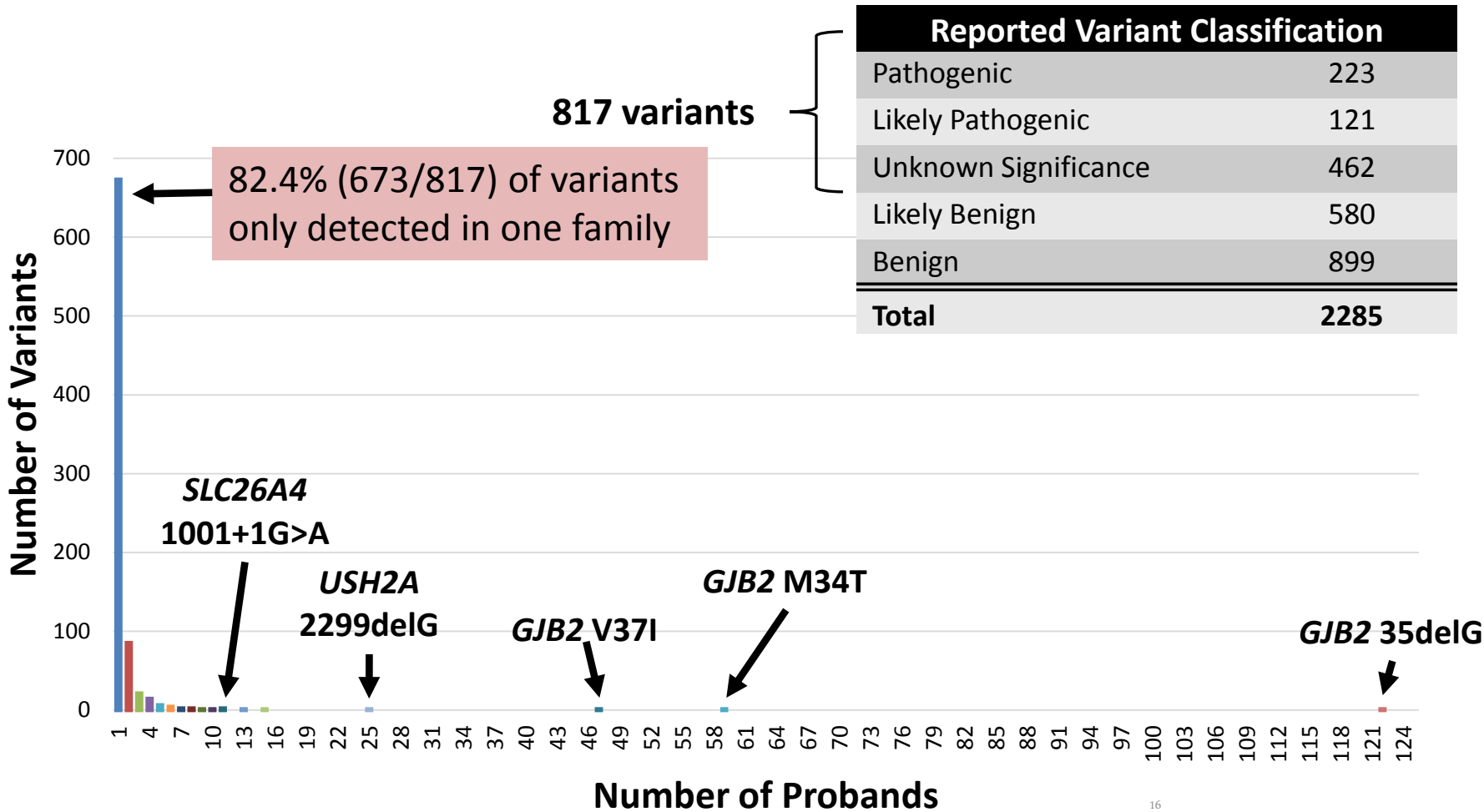
Deletions Detected by NGS

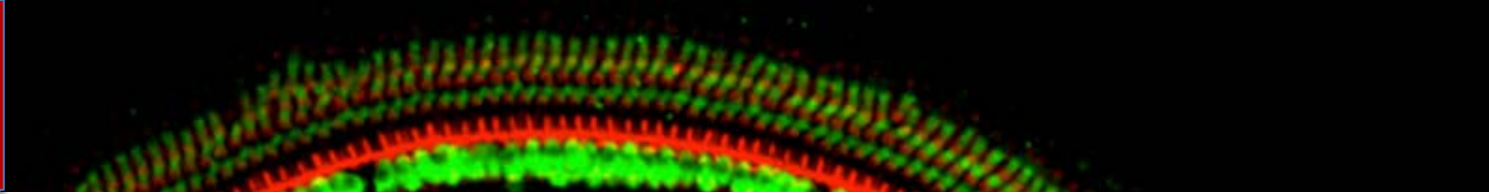


Copy number variants confirmed by digital droplet PCR

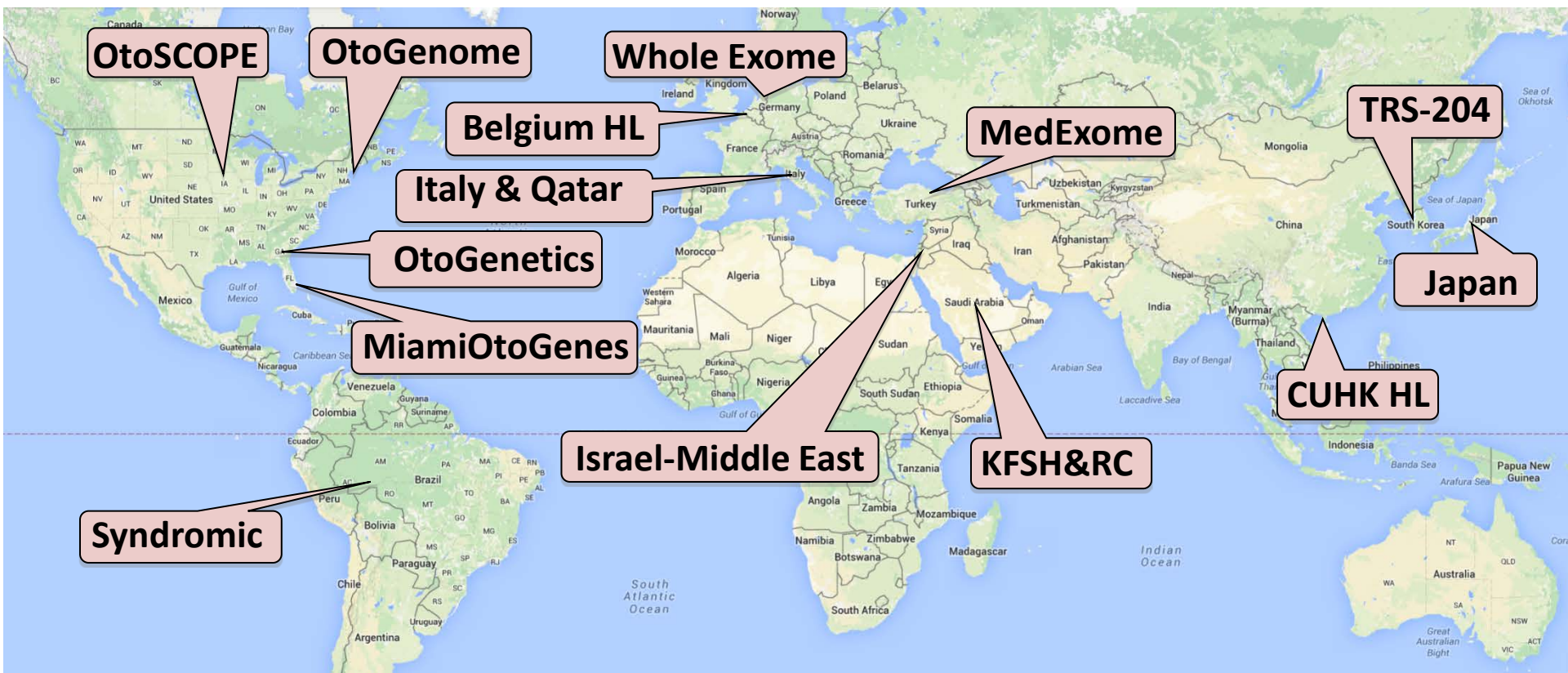


Hearing Loss Variants in over 2000 Cases Tested at the Laboratory for Molecular Medicine

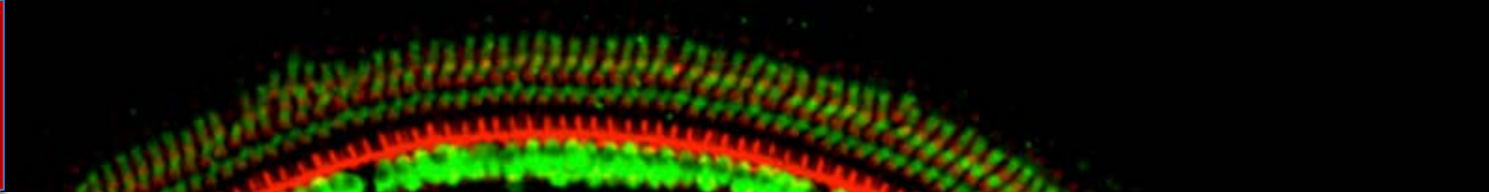




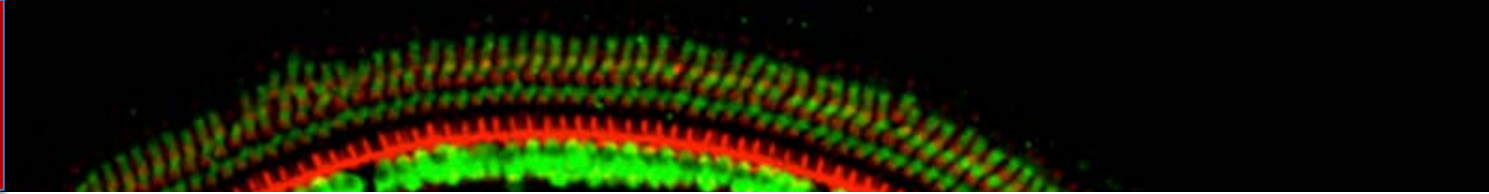
Next-Generation Sequencing Tests for Hearing Loss



**7th International
Pediatric Audiology
Conference**

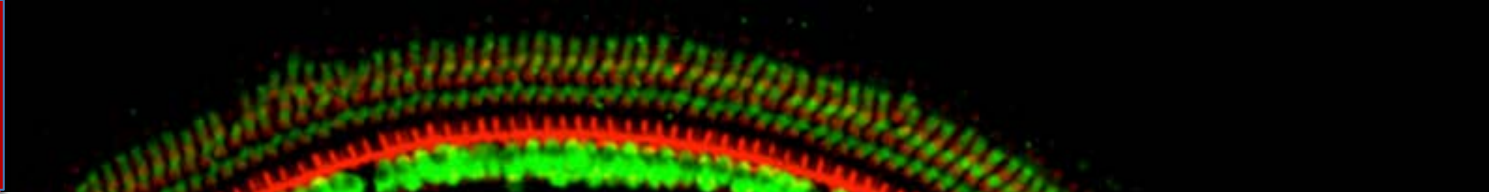


Panel	Institution	Genes	Diagnostic yield	Capture	Sequencing
Belgium	U Antwerp	79	25-30%	TruSeq	Illumina
CUHK-HL	CUHK	252	57%	SureSelect	Illumina
Israel-MidEast	Tel Aviv	246	56%	SureSelect	Illumina
Italy+Qatar	U Trieste	96	33%	AmpliSeq	Ion PGM
Japan	Shinshu U	63	30%	AmpliSeq	IonTorrent
KFSH&RC	KFSH	90	54%	AmpliSeq	Ion PGM
Medical Exome	Ege U	102/2761	72.4%	TruSight	Illumina
MiamiOtoGenes	U Miami	146	50%	SureSelect	Illumina
OtoGenetics	Commercial	131	42-52%	NimbleGen	Illumina
OtoGenome	LMM/Harvard	87	23%	SureSelect	Illumina
OtoSCOPE	MORL/U Iowa	89	39%	SureSelect	Illumina
Syndromic	U Brasilia	52	10%	AmpliSeq	Ion PGM
TRS-204	Seoul National	204	55%	?	Illumina
Whole Exome	Rotterdam	120/20k	70%	?	?

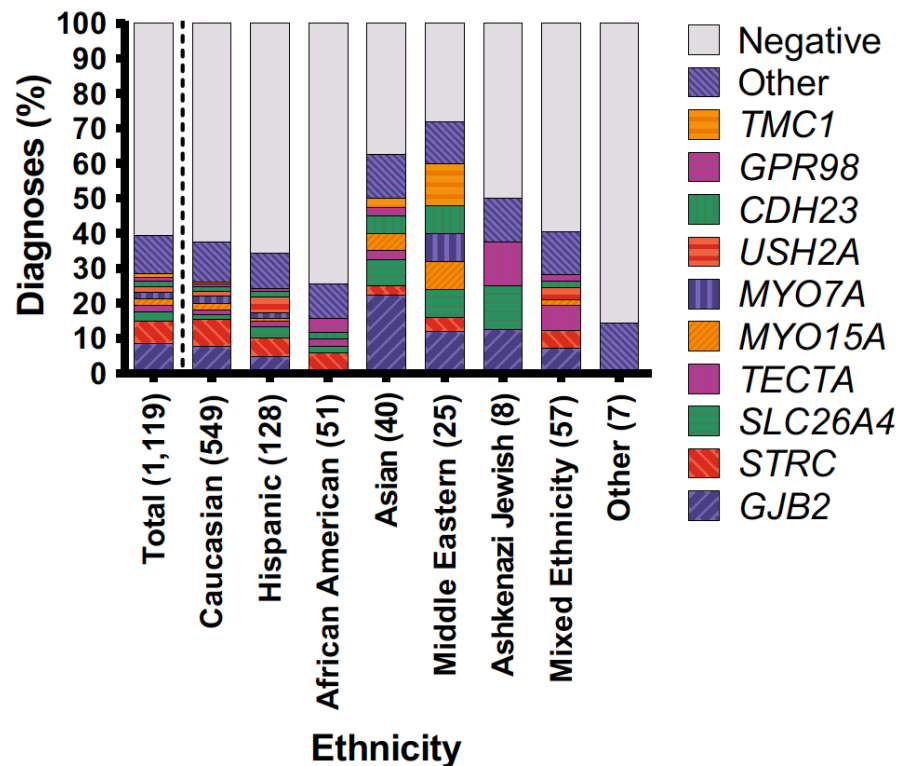
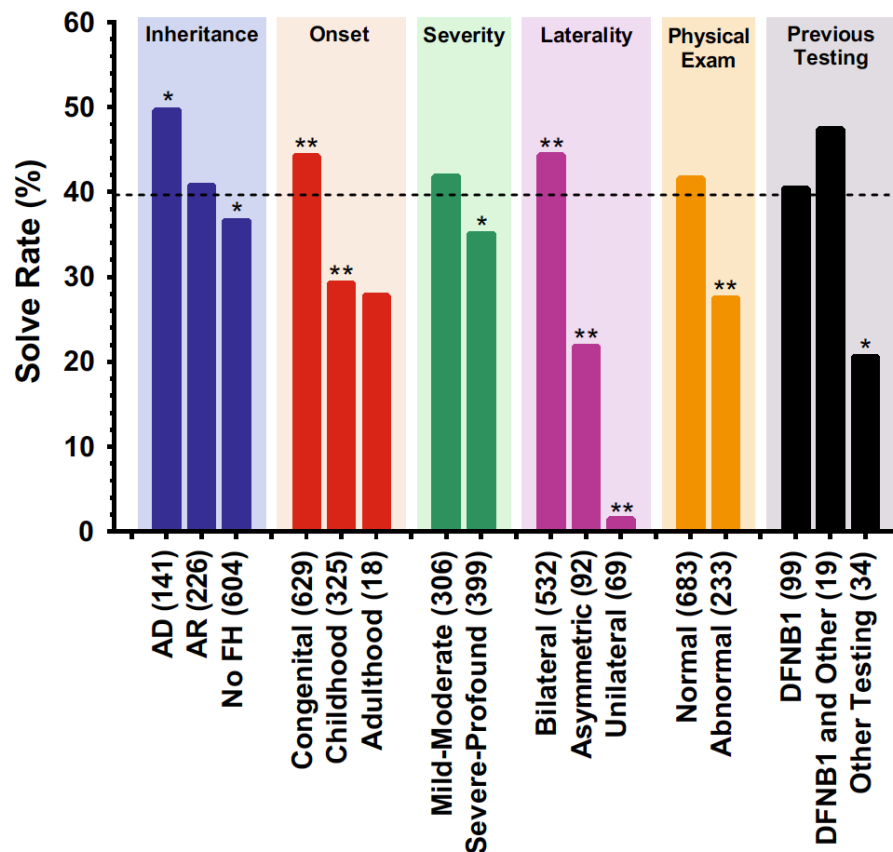


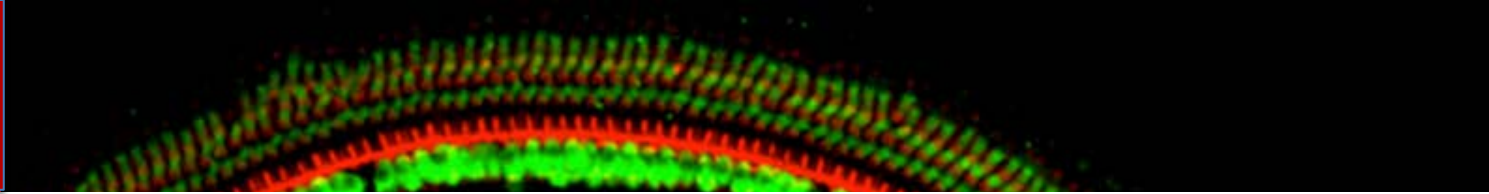
Explanation for Variable Diagnostic Yields

- Inclusion or exclusion of *GJB2*-positive patients
- Stringency of variant classification and interpretation
- Small sample sizes in some reported studies
- Family history / simplex vs. multiplex
- Specific clinical features
- Ethnicity (consanguinity, availability of population data)



Factors Influencing Diagnostic Yield for Hearing Loss





Newborn Hearing Screening

- Standard of care in the U.S. with greater than 98% of newborns screened before leaving the birth hospital
- ~1.6% documented not passing final hearing screening
- ~45% not passing hearing screening lost to follow up or documentation for diagnosis
- Prevalence of documented HL ~1.4% per 1000 screened
- ~88% documented referrals of those with HL to Early Hearing Detection and Intervention (EHDI)
- ~50% of states have language in legislation or regulations that include coverage for early intervention services to children with mild or unilateral HL
- Two common screening methods: otoacoustic emissions (OAEs) and auditory brainstem response (ABR), detecting HL in the frequency region important for speech recognition

DID YOU HEAR?

98%

of newborns in the U.S. are
screened for hearing loss
before they leave the hospital.

Research improves the quality of life of people with
hearing loss, starting with the day they are born.

Biomedical discoveries supported by the National Institute on
Deafness and Other Communication Disorders (NIDCD) laid the
foundation for states to take action to ensure children are screened
and treated early for hearing loss.



NIDCD research demonstrates **the need for both
newborn hearing screening and early intervention**,
which is crucial for speech and language development.



NIDCD research leads to **two gold-standard
tests** for hearing loss in infants.



NIDCD research finds **genetic causes of
profound hearing loss and deafness**, which
account for more than half of all cases.



NIDCD research explores **intervention
strategies** for children with hearing loss.



NIDCD research develops and improves
technology for hearing devices such as
hearing aids and cochlear implants.



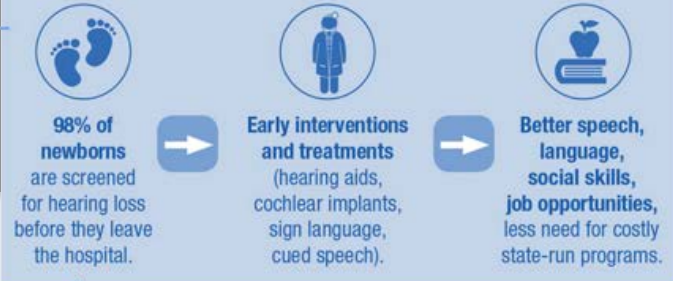
NIDCD research reveals **the basic
mechanisms of how we hear**.

DID YOU KNOW?

12,000 babies are
born deaf or hard of
hearing each year in
the United States.

www.medlineplus.gov

NIDCD authorized on
10-28-88 by Public Law
100-553



2010 **98%** **3.8 million newborns** are screened annually.

2010 President Obama signs the **Early Hearing Detection and Intervention Act of 2010**, expanding funding to include diagnostic services.

1999 President Clinton signs the **Newborn and Infant Hearing Screening and Intervention Act**, authorizing support for statewide screening programs.

1997 The NIH convenes an expert panel, which **recommends standard newborn hearing screening methods** for state programs.

1993 At an NIH consensus conference, **experts endorse universal newborn hearing screening**.

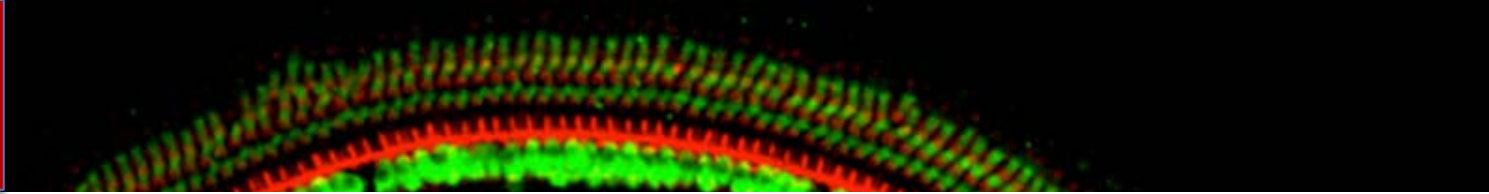
1993 **About 1 in 10 newborns** are screened for hearing loss.

Before 1993

Only newborns at high risk are screened, which misses 50% of children who are eventually diagnosed with severe hearing impairments.

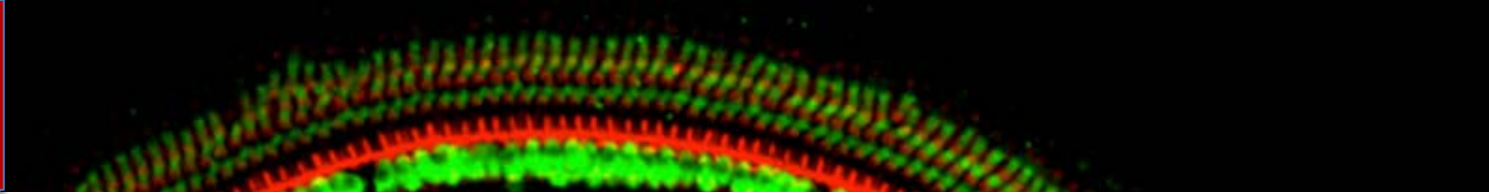
Only 8% of babies with congenital hearing loss are diagnosed by their first birthday.

47% of children with congenital hearing loss are not diagnosed until their third birthday or later.



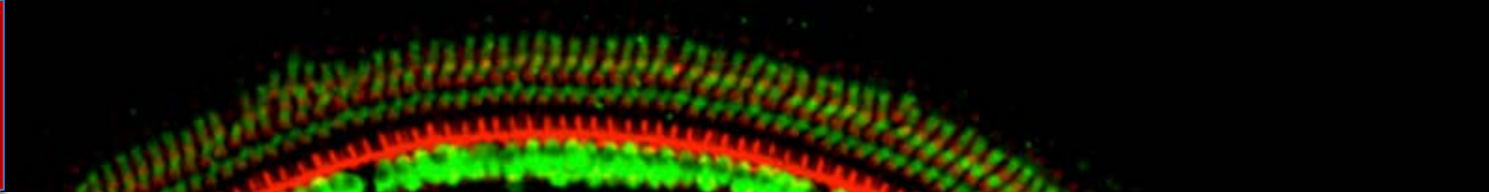
Next-Gen Newborn Screening for Hearing: a paradigm for expanded genetic screening

- It makes a difference for treatment and management to know the precise diagnosis—the foundation of precision medicine.
- Hereditary hearing loss displays unparalleled genetic heterogeneity and offers an opportunity to simulate a complex genetic disorder in the context of hundreds of single gene defects.



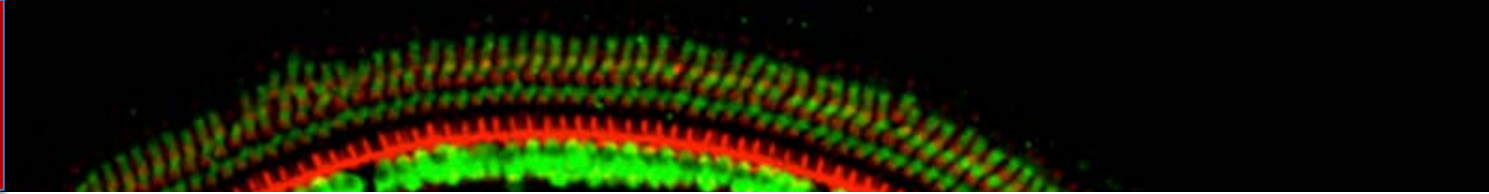
Precise Diagnosis Impacts Care

- *GJB2*—benefit from cochlear implantation
- *PJVK*—more harm than good from amplification
- Others with optimized habilitation?
- Syndromic disorders that can be indistinguishable from nonsyndromic disorders at birth:
 - *Alport*
 - *Branchio-oto-renal*
 - *Jervell and Lange-Nielson*
 - *Pendred*
 - *Usher*

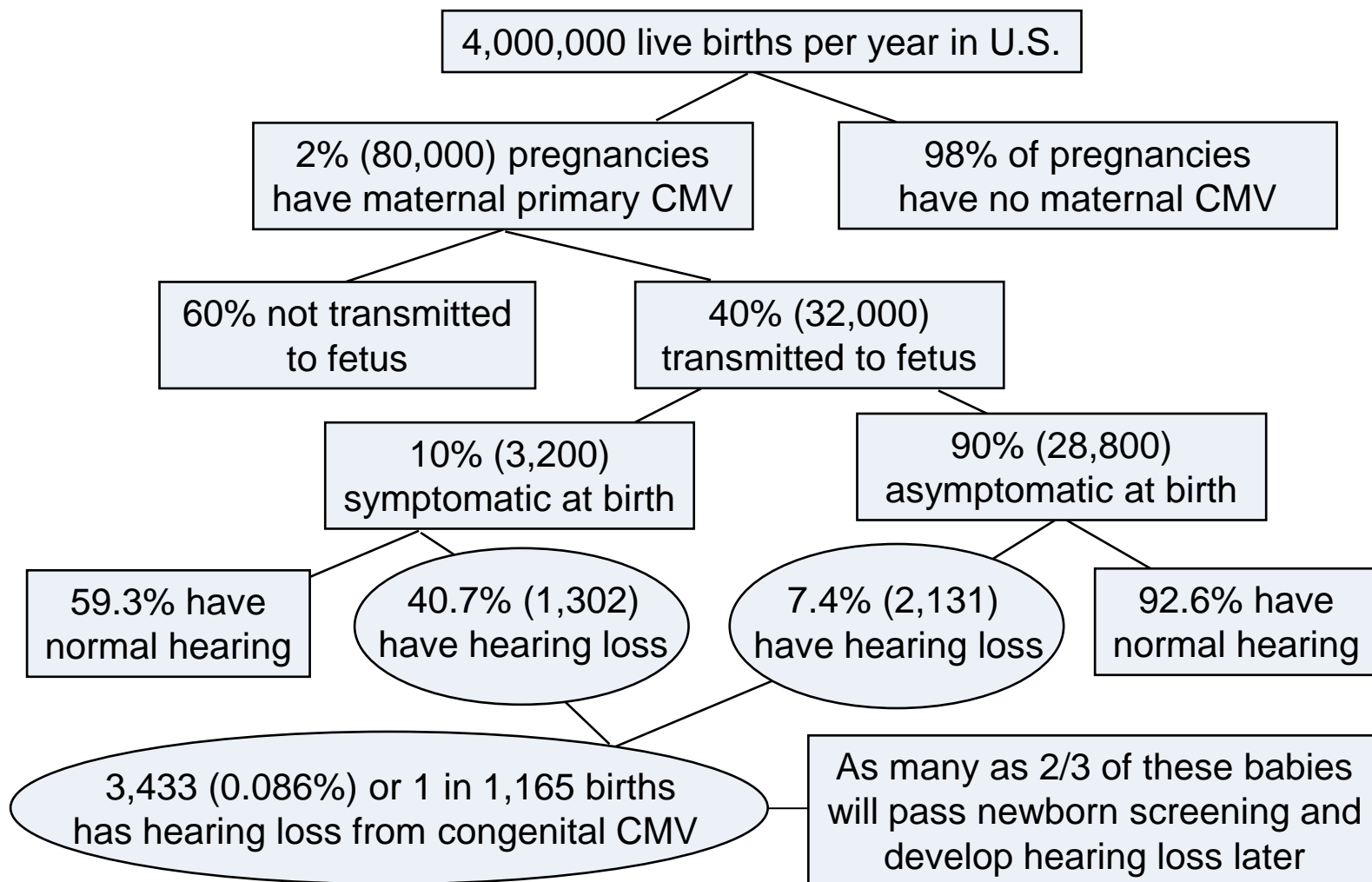


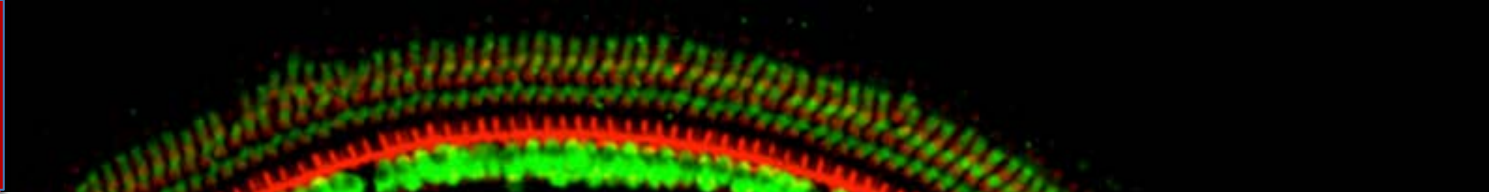
When will screening take place?

	At Birth (~4 million newborns)	Failed Audiometry (2-8 per 100 infants)	Confirmed Hearing Impairment
+	<ul style="list-style-type: none"> All newborns tested Reduce work up Use blood spots Identify those at risk 	<ul style="list-style-type: none"> Less infants tested Reduce work up Use blood spots? 	<ul style="list-style-type: none"> Less infants tested Reduce work up Specific testing High parental interest
-	<ul style="list-style-type: none"> Genotype-phenotype? False positives Detect many carriers Parental interest low 	<ul style="list-style-type: none"> Genotype-phenotype? False positives? Detect many carriers Parental interest low? Miss some at risk New specimen 	<ul style="list-style-type: none"> Genotype-phenotype? Miss some at risk New specimen? Delay in intervention



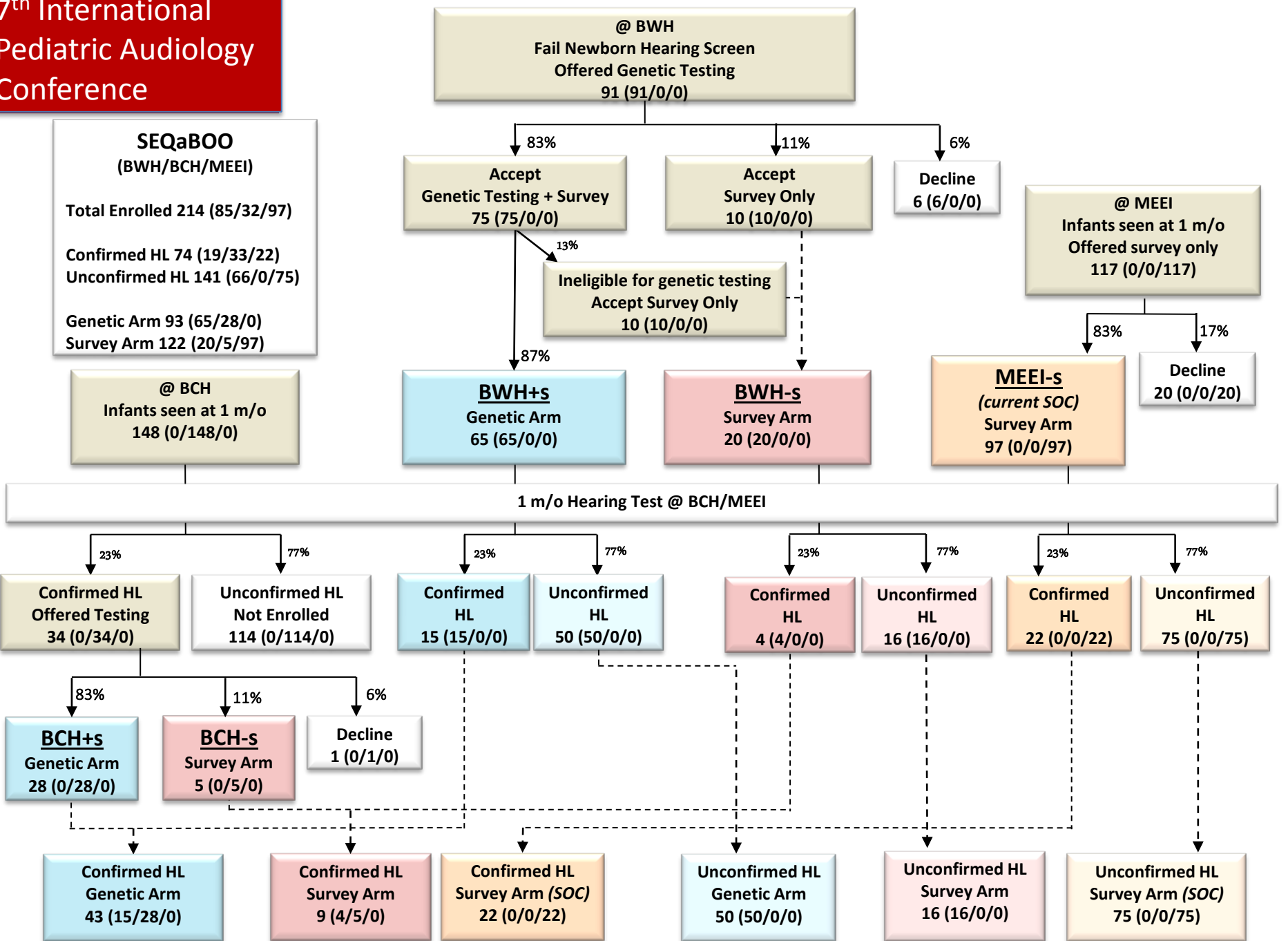
Hearing Loss from Cytomegalovirus (CMV)





SEQuencing a **B**aby for an **O**ptimal **O**utcome

7th International Pediatric Audiology Conference



SEQaBOO
(BWH/BCH/MEEI)

Total Enrolled 214 (85/32/97)

Confirmed HL 74 (19/33/22)
Unconfirmed HL 141 (66/0/75)

Genetic Arm 93 (65/28/0)
Survey Arm 122 (20/5/97)

Accept
Genetic Testing + Survey
75 (75/0/0)

Accept
Survey Only
10 (10/0/0)

Decline
6 (6/0/0)

Ineligible for genetic testing
Accept Survey Only
10 (10/0/0)

@ MEEI
Infants seen at 1 m/o
Offered survey only
117 (0/0/117)

BWH+s
Genetic Arm
65 (65/0/0)

BWH-s
Survey Arm
20 (20/0/0)

MEEI-s
(current SOC)
Survey Arm
97 (0/0/97)

Decline
20 (0/0/20)

1 m/o Hearing Test @ BCH/MEEI

Confirmed HL
Offered Testing
34 (0/34/0)

Unconfirmed HL
Not Enrolled
114 (0/114/0)

Confirmed HL
15 (15/0/0)

Unconfirmed HL
50 (50/0/0)

Confirmed HL
4 (4/0/0)

Unconfirmed HL
16 (16/0/0)

Confirmed HL
22 (0/0/22)

Unconfirmed HL
75 (0/0/75)

BCH+s
Genetic Arm
28 (0/28/0)

BCH-s
Survey Arm
5 (0/5/0)

Decline
1 (0/1/0)

Confirmed HL
Genetic Arm
43 (15/28/0)

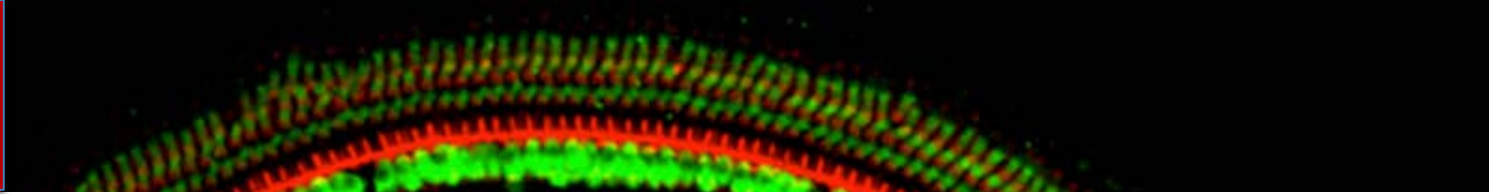
Confirmed HL
Survey Arm
9 (4/5/0)

Confirmed HL
Survey Arm (SOC)
22 (0/0/22)

Unconfirmed HL
Genetic Arm
50 (50/0/0)

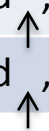
Unconfirmed HL
Survey Arm
16 (16/0/0)

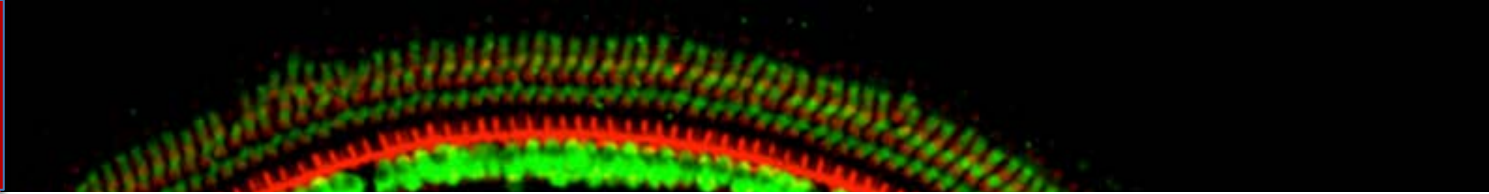
Unconfirmed HL
Survey Arm (SOC)
75 (0/0/75)



Comparisons of Traditional and Next-Generation Newborn Hearing Screening

Hearing Screening	Traditional	Next-generation
Method	OAE and ABR	DPOAE and ABR plus WGS
Type	Phenotypic	Phenotypic + Genetic
Symptomatic	Yes	Not required
Prognostic	No	Maybe
Accurate recurrence risk estimate	No	Yes
Etiology	No	Yes
Precision medicine	No	May inform
Cost	\$10-50/baby, single purpose, stable	<\$10,000/baby, decreasing
Turnaround time	Immediately	Days
Sensitivity	Unable to detect later-onset	Expected , validation ongoing
Specificity	Transient loss as false positives	Expected , validation ongoing



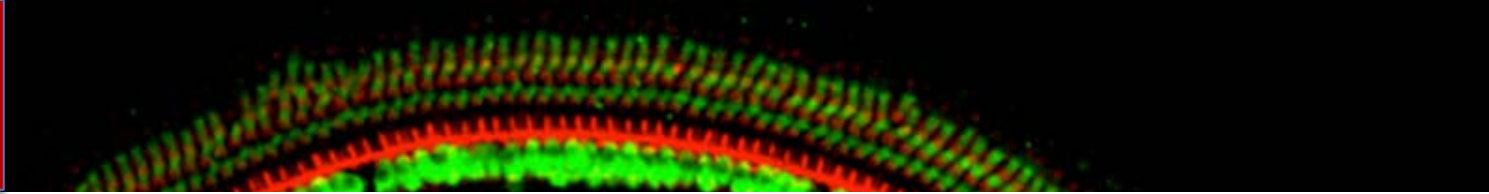


Whole Genome Sequencing Projects Underway

- Discovery projects for additional genes involved in hearing loss

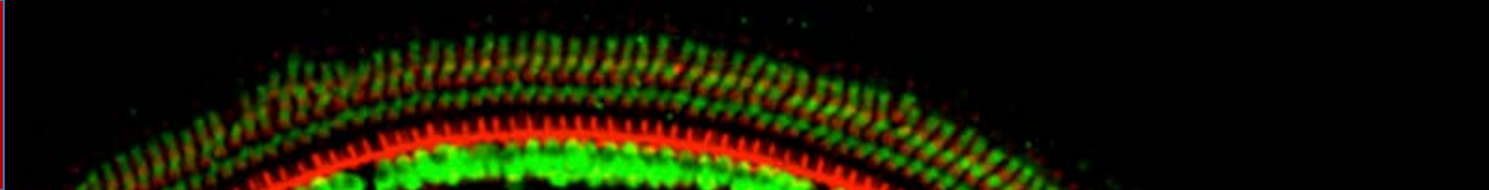
Genomics
england





“Hot Off the Email” (Sept. 2016): from Michael Watson, Exec Director, ACMG

“We’ve gotten notification that we are funded (\$100,000) as a supplement to our cooperative agreement with HRSA for the National Coordinating Center for the Regional NBS and Genetics Collaboratives to do the project I had discussed with you a few months ago. **It is to first review etiologies of newborn hearing loss cases that are identified before age 5 but missed by NBS.** We then would look at how best to pick up the kids with these later onset forms. There seem to me to be two main possibilities, though altering audiometric cut-offs might also detect some of them. The main options are: 1) screen kids audiometrically before school to identify those with hearing loss; 2) add a molecular test to NBS hearing screening to pick them up as a part of NBS. This option requires determining which genes have definitive or strong associations with hearing loss and then assessing the proportion of cases with those genes involved and with variants in them that would classify as pathogenic or likely pathogenic. There are other trade-offs with either of these such as the opportunity to introduce interventions earlier if found earlier to maximize the benefits of being found that the group would factor in to its thinking.”



With Appreciation to

Brigham and Women's Hospital

Anne Giersch
Jun Shen

Boston Children's Hospital

Margaret Kenna

Harvard University

Jennifer Hochschild

Massachusetts Eye and Ear Infirmary

Michael Cohen

Laboratory for Molecular Medicine

Ahmad Abou Tayoun
Sami Amr
Amy Hernandez
Andrea Murihead
Heidi Rehm

Harvard Medical School
Center for Hereditary Deafness



Manchester Centre for
Audiology and Deafness (ManCAD)