# The Genetics of Hearing Loss

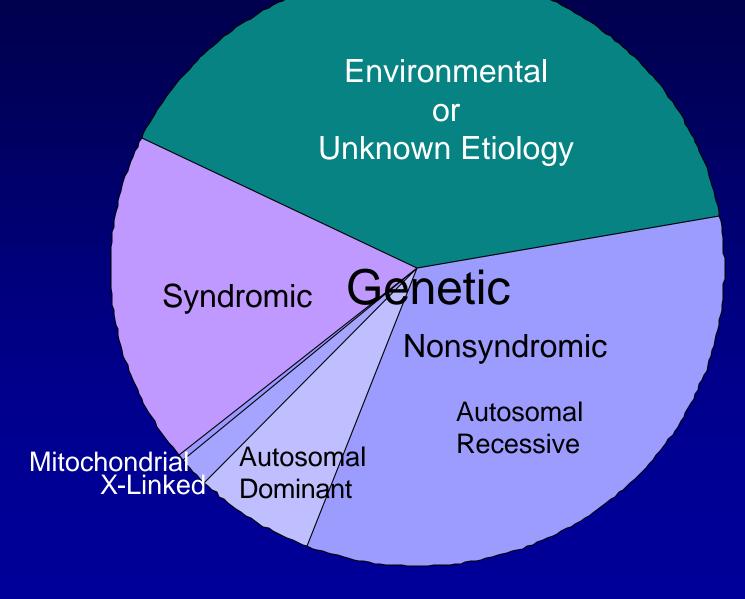
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## **Faculty Disclosure Information**

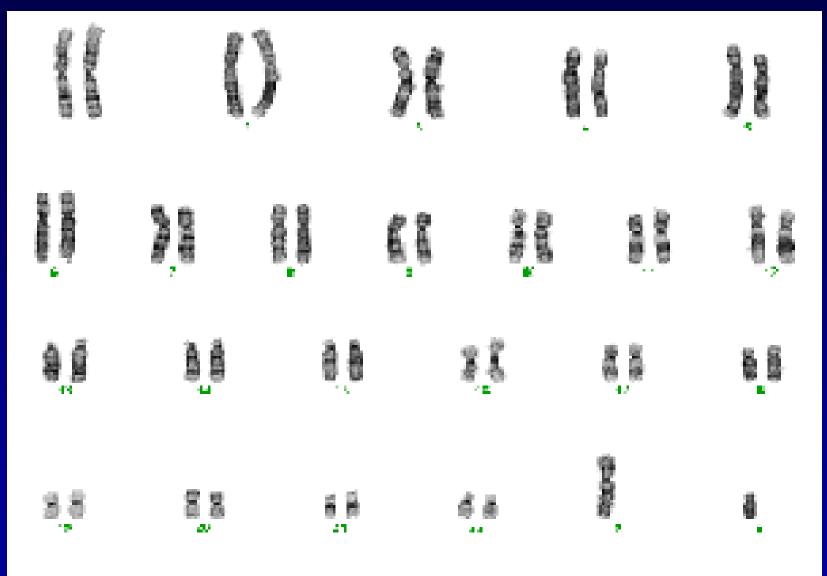
In the past 12 months, I have not had a significant financial interest or other relationship with the manufacturer(s) of the product(s) or provider(s) of the service(s) that will be discussed in my presentation

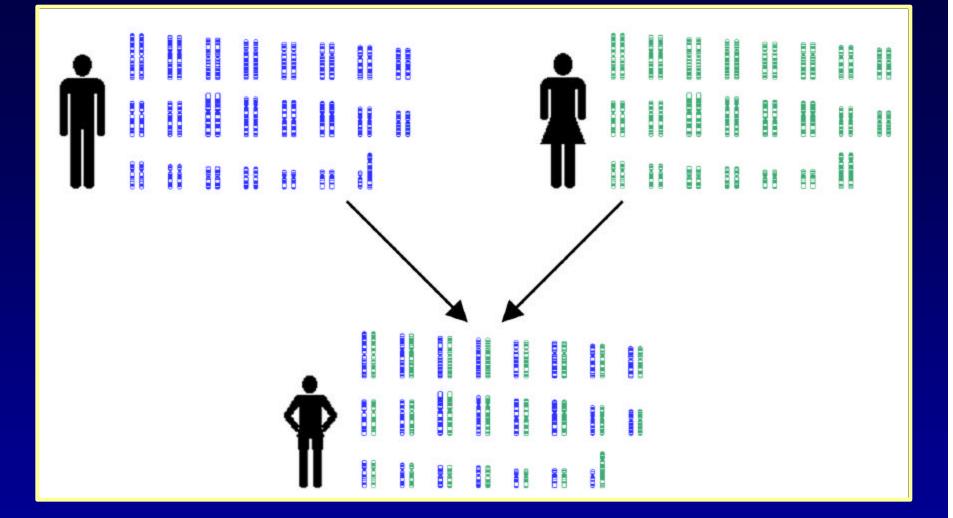
This presentation will not include discussion of pharmaceuticals or devices that have not been approved by the FDA or "off-label" uses of pharmaceuticals or devices.

## Causes of Childhood Hearing Loss

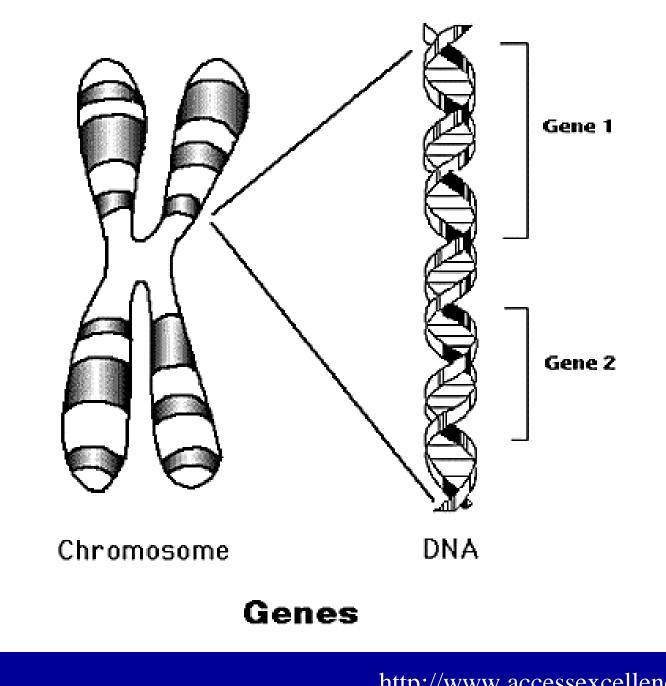


## Human Karyotype





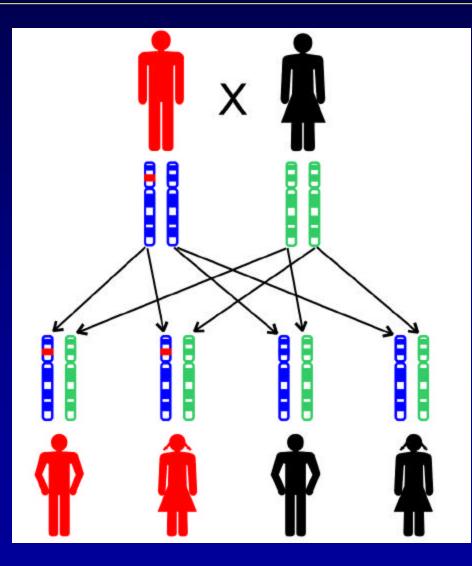
We inherit two copies of every gene, one from each parent.



http://www.accessexcellence.org/AB/GG/

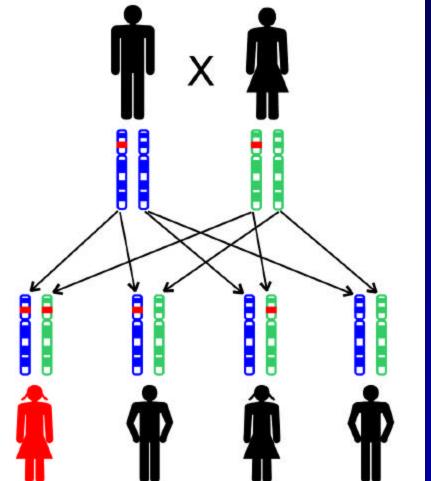
### **Dominant Mutations**

- Dominant hearing loss can be caused by only one copy of a mutated gene
- Dominant hearing loss is seen in every generation
- If a parent has a dominant mutation, each child has a 50 % chance of inheriting it.



## **Autosomal Recessive Mutations**

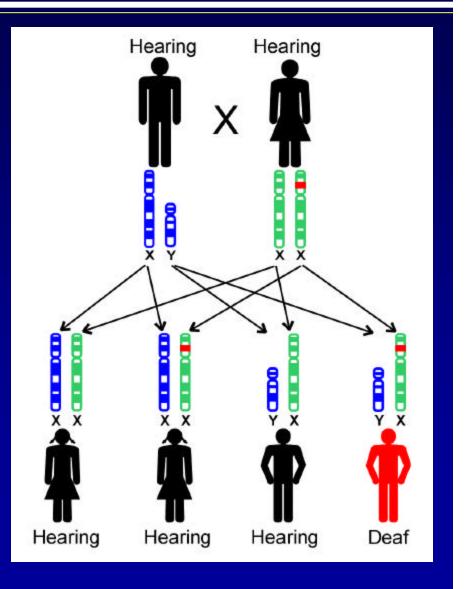
- For recessive hearing loss, both copies of a gene must be mutated to get hearing loss.
- Often, there is no family history of hearing loss.
- Each child will have a 25% chance of hearing loss.



A carrier is a person who carries one copy of a recessive mutation , but does not have hearing loss.

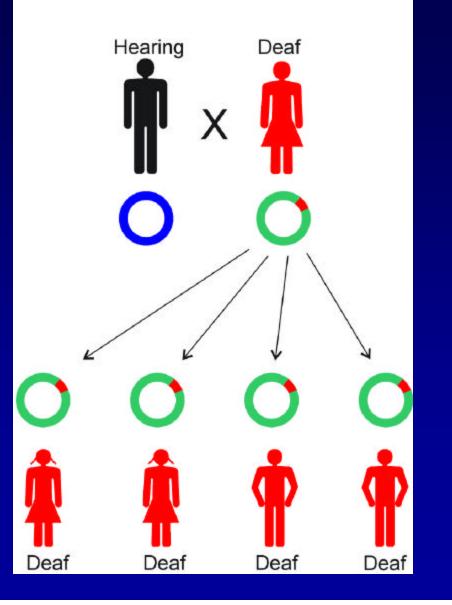
## X-Linked Recessive Mutations

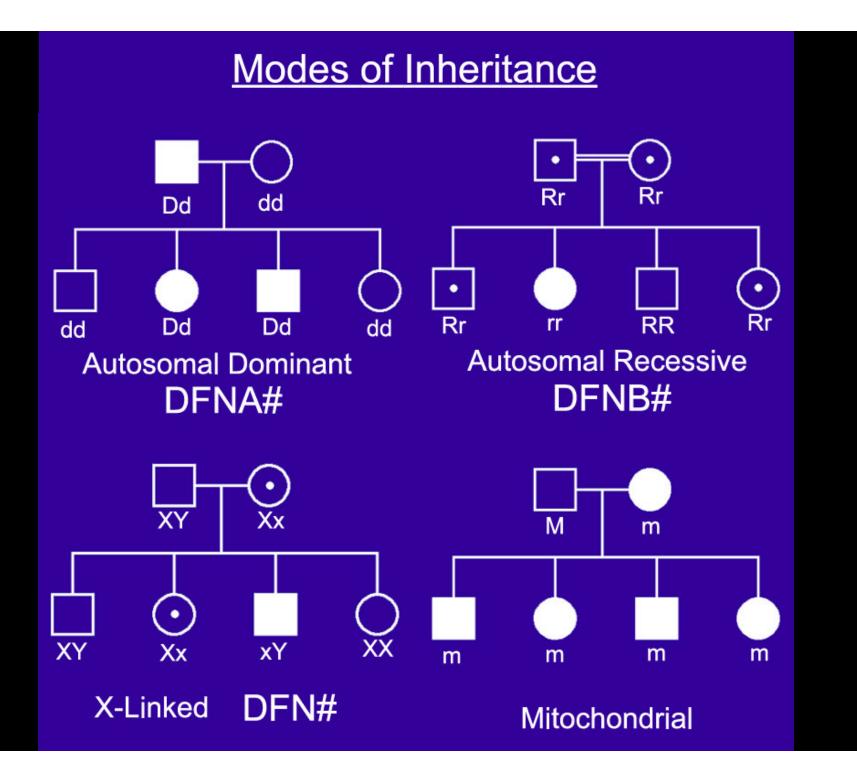
- Only males are affected.
- Each son will have a 50% chance of having hearing loss.
- Each daughter has a 50% chance of being a carrier.



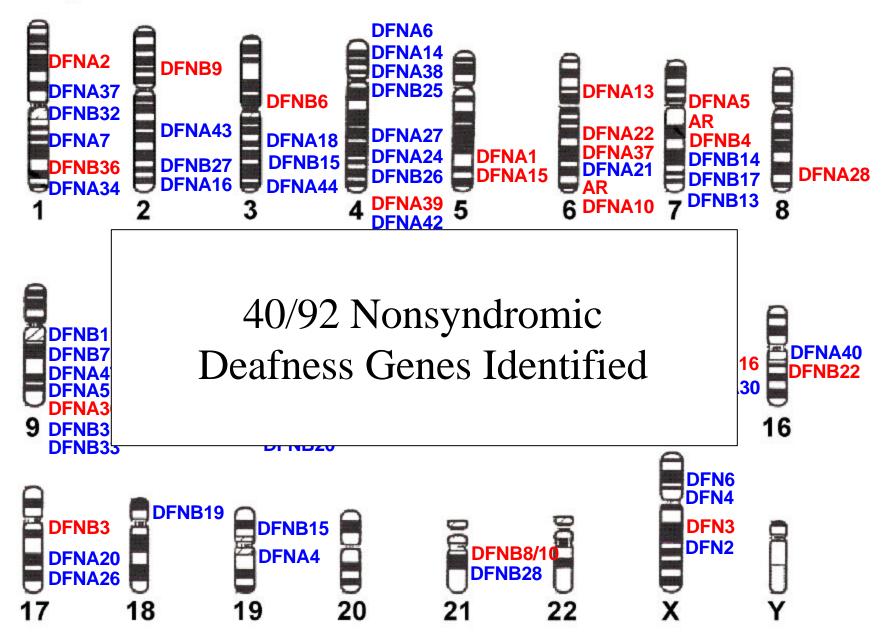
## Mitochondrial Mutations

- Only the mother passes mitochondria to her children.
- All children will inherit a mitochondrial mutation from their mother.
- Mitochondrial mutations are often variable in their expression of hearing loss.





#### Nonsyndromic Deafness Genes in the Human Genome



## Syndromic Hearing Loss

#### **Syndromes**

Alport **Branchio-Oto-Renal** Jervell and Lange-Nielsen Mitochondrial (MELAS/MERRF) Neurofibromatosis type II Norrie Osteogenesis Imperfecta Pendred Stickler Tranebjaerg-Mohr (DFN1) **Treacher Collins** Usher

Waardenburg

Gene(s) COL4A5, COL4A3, COL4A4 EYA1 <u>KCNQ1, KCNE</u>1/lsK tRNA<sup>leu(UUR)</sup>.tRNA<sup>lys</sup> NF2 NDP COL1A1, COL1A2 PDS COL2A1, COL11A2, COL11A1 DDP TCOF1 MYO7A, USH1C, CDH23, PCDH15, SANS, USH2A, VLGR1, USH3 PAX3, MITF, SLUG, EDNRB, EDN3, **SOX10** 

There are currently over 400 syndromes with associated hearing loss.

# Usher Syndrome

#### (3-6% of childhood deafness)

	Hearing Loss	Vestibular System	Retinitis Pigmentosa
Type I	Congenital profound	Congenital balance problems	Onset pre- puberty
Type II	Congenital mild-severe sloping	Normal	Onset in teens-20s
Type III	Progressive later onset	Progressive balance problems	Variable onset

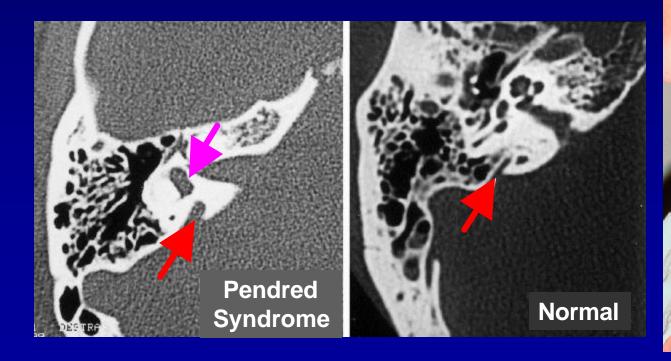
Usher Type	Locus	Gene	<b>Relative Incidence</b>
USH1A	14q32	unknown	2%
USH1B	11q13.5	MYO7A	60%
USH1C	11p15.1	USH1C	5%
USH1D	10q	CDH23	10%
USH1E	<b>2</b> 1q	unknown	Rare
USH1F	10q21.1	PCDH15	Rare
USH1G	17q24-25	SANS	Rare
USH2A	1q41	USH2A (+51)	80%
USH2B	3p23-24.2	unknown	Rare
USH2C	5q14.3-q21.3	VLGR1	15%
USH3	3q21-q25	USH3	100%

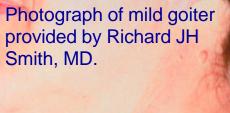
## Jervell & Lange-Nielsen Syndrome

- Incidence: 1/250,000
- Phenotype
  - Severe-profound congenital sensorineural hearing loss
  - Prolonged QT interval
  - Syncope
  - Arrythmia
  - Sudden death
- Heart condition diagnosed by EKG and treatable with beta-blockers

## DFNB4 Hearing Loss/Pendred Syndrome

- Congenital sensorineural hearing loss w/ EVA or Mondini
- ~20% with late onset goiter -> 10% hypothyroid
- Incidence: ~5% of congenital hearing loss
- Inheritance: Autosomal Recessive?

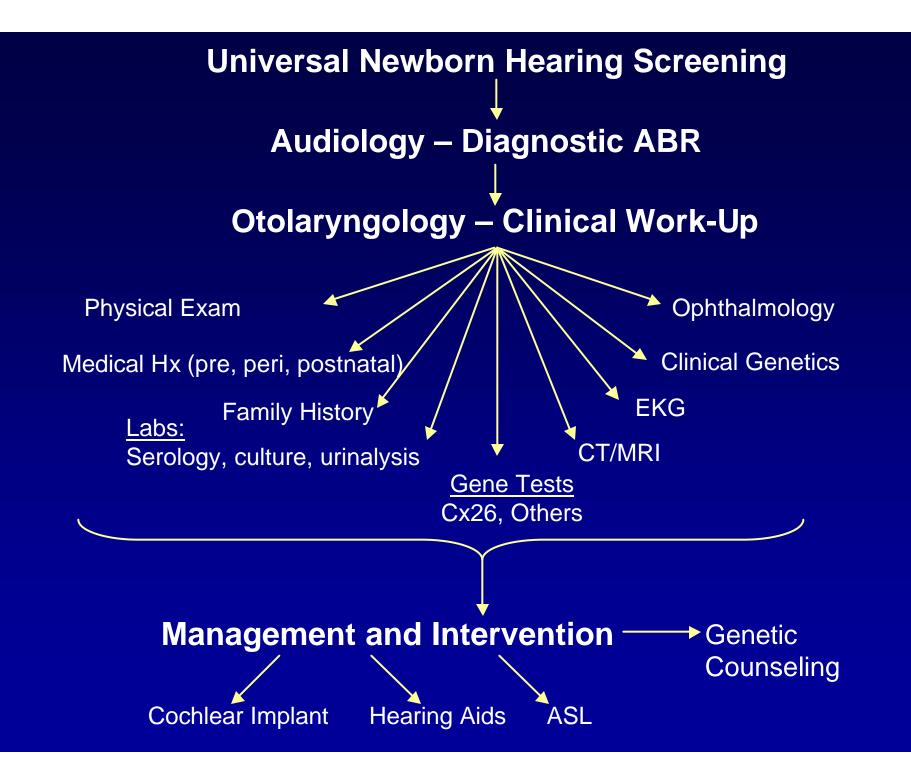




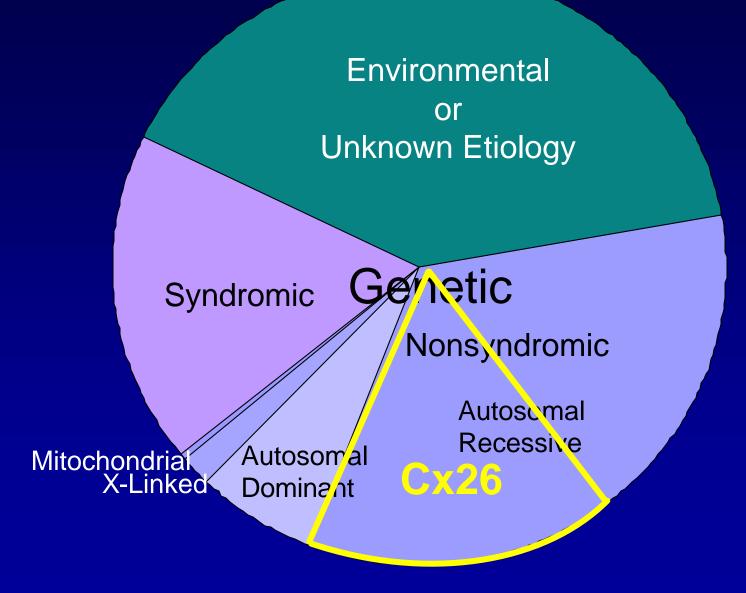
## Hearing Loss and EVA

~30% of sporadic cases and ~90% of pts with family hx have mutations in the PDS gene.

Patients with 2 mutations will likely develop thyroid disease whereas those with only 1 or 0 mutations will likely not (Pryor et al. J Med Genet. 2005. 42(2):159-65)



## Causes of Childhood Hearing Loss



## Should all Children with Hearing Loss have Cx26 Testing?

- Even cases that have an apparent explanation for hearing loss still may have Cx26 mutations
- Case A: Congenital syphilis
- ✓ Case B: CMV perinatal infection
- Case C: Prematurity
- Case D: Hyperbilirubinemia

## GJB2 - Connexin 26

**Exon 1** ~ 3000 bp

Exon 2 (681 bp)

#### **DFNB1 (Recessive) Mutations**

**Missense:** M1V, T8M, G12V, K15T, S19T, I20T, R32C, M34T, V37I, A40E, A40G, G45E, E47K, W77R, V84L, L90P, V95M, H100Y, S113R, delE120, K122I, R127H, R143W, E147K, P175T, R184P, R184W, S199F, L214P, 05INS4, I203K, N206S, S139N, H100, E101G, L90V, M93I, 486INST, Q80R, I82M, S85P, A88S, L174R, L79P, Q80P, S19T, I20T, V27I+E114G, R32L, R165W

Nonsense: W24X, W44X, E47X, Q57X, Y65X, Y97X, Q124X, Y136X, W112X, W172X, C64X, Q80X, E147X

Frameshift: 31del14, 31del26, 35delG, 35insG, 51del12insA, 167delT, 176del16, 235delC, 269insT, 299delAT, 314del14, 333delAA, 290linsA, 310del14, 312del14, 509del14, 509insA, 515 del17, 631delGT, 504insAAGG, 515del17, 572delt, 645delTAGA, 302del3, 469delG

GJB6-D13S1830 (Cx30) Deletion

DFNA3 (Dominant) Mutations delE42, W44S/C, R75Q, D179N, R184Q, C202F, M163L w/ Skin Disease G12R, S17F, D50N, N54K, G59A, D66H, R75W, R75Q

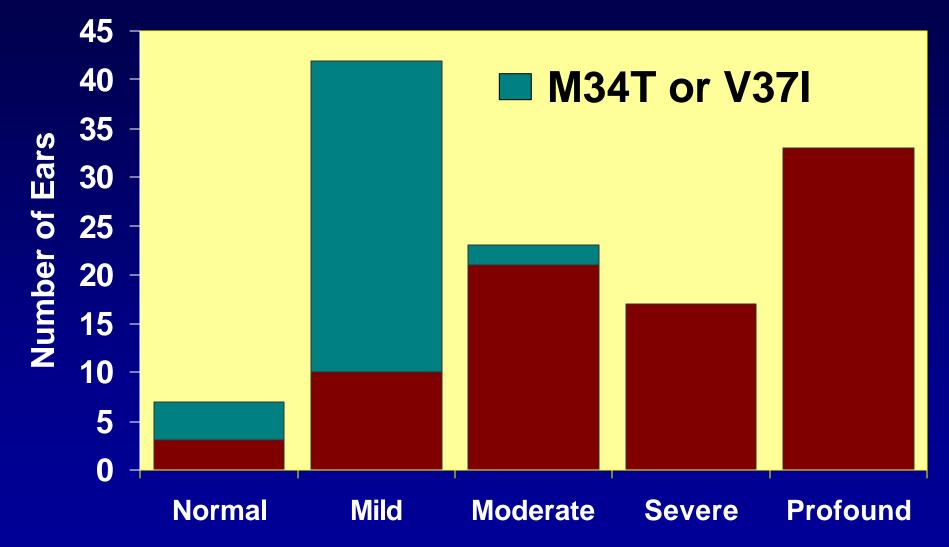
Unknown Significance H73R, R98Q, R127C, K224Q, IVS1, N54I, V84A, S85Y, 313deIAA, 314deIA, 360deIG, T123N, E129K

Polymorphisms V27I, F83L, E114G, T123A, V153I, G160S, C169Y, I203T, Exon1 -493del100, -3558T, -1C>T, I30I, A40A, R127H, S72C, Q80Q, L89L, R104R, I128I, V182V, V190V, 682C>T. 765C>T

## **Connexin 26 Gene Sequence**

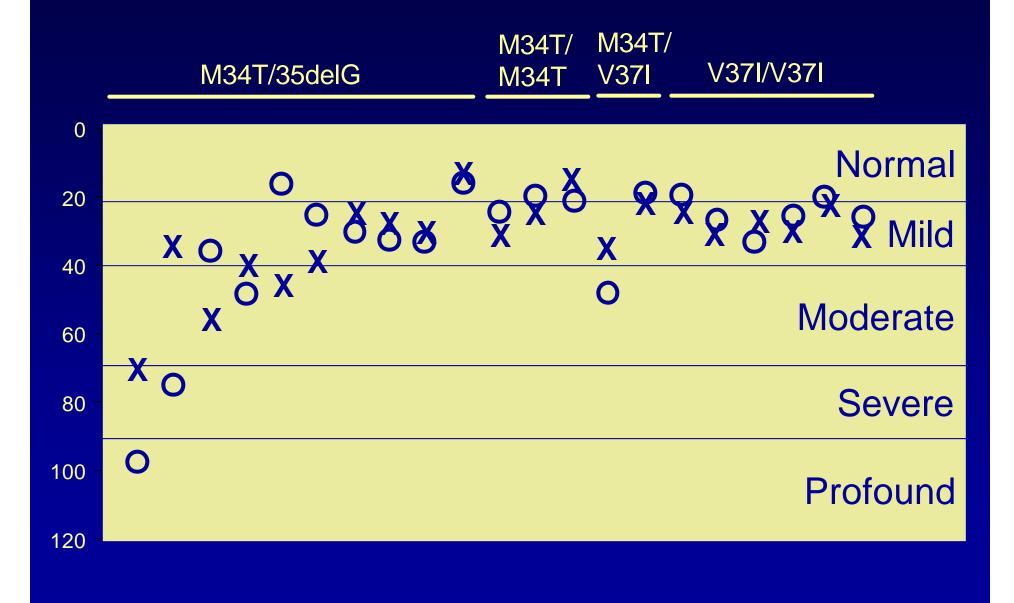
ATGGATTGGGGCACGCTGCAGACGATCCTGGGGGC TGTGAACA AACACTCCACCAGCATTGGAAAGATCTGGCTCACCGTCCTCTTC ATTTTTCGCATTATGATCCTCGTTGTGGCTGCAAAGGAGGTGTG GGGAGATGAGCAGGCCGACTTTGTCTGCAACACCCTGCAGCCA GGCTGCAAGAACGTGTGCTACGATCACTACTTCCCCATCTCCCA CATCCGGCTATGGGCCCTGCAGCTGATCTTCGTGTCCAGCCCA GCGCTCCTAGTGGCCATGCACGTGGCCTACCGGAGACATGAGA AGAAGAGGAAGTTCATCAAGGGGGGAGATAAAGAGTGAATTTAAG GACATCGAGGAGATCAAAACCCAGAAGGTCCGCATCGAAGGCT CCCTGTGGTGGACCTACACAAGCAGCATCTTCTTCCGGGTCATC TTCGAAGCCGCCTTCATGTACGTCTTCTATGTCATGTACGACGG CTTCTCCATGCAGCGGCTGGTGAAGTGCAACGCCTGGCCTTGT CCCAACACTGTGGACTGCTTTGTGTCCCGGCCCACGGAGAAGA CTGTCTTCACAGTGTTCATGATTGCAGTGTCTGGAATTTGCATC CTGCTGAATGTCACTGAATTGTGTTATTTGCTAATTAGATATTGT TCTGGGAAGTCAAAAAGCCAGTTTAA

#### Hearing Loss Severity Associated with Biallelic Cx26 Mutations



Data from Children's Hospital Boston (Kenna and Rehm, 2005)

#### Pure Tone Averages for M34T and V37I Genotypes



## **Cx26 Gene Test Outcomes**

- Two copies (homozygous) of a single mutation
  e.g. 35delG/35delG
- 2. Two different mutations (compound heterozygous)
  e.g. 35delG/167delT
- 3. No Cx26 mutations detected
- 4. Only one mutation detected (heterozygous)
  e.g. 35delG/+

## Cx26 Gene Testing at Children's Hospital Boston

## Testing results from deaf probands:

Biallelic mutations:20/101 (20%)Heterozygous mutations:12/101 (10%)No mutation detected:71/101 (70%)

Kenna and Rehm et al, 2000

# Explanations for Deafness in an Individual with a Single Cx26 Mutation

#### *K* The mutation may act dominantly

(There are at least six Cx26 mutations known to act dominantly.)

#### **The Cx26 mutation is unrelated to the deafness**

(The deafness may be caused by another gene mutation or a non-genetic cause.)

#### **The test did not detect the second mutation**

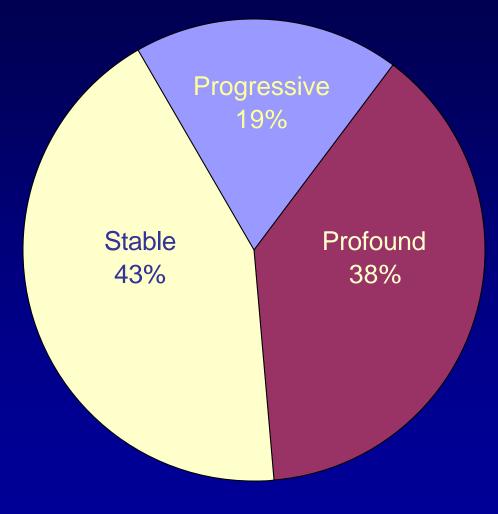
(There may be a mutation in a non-coding region.)

The genetic background of the patient may alter the mutations ability to cause deafness

## 2 Cases of Delayed Onset Cx26 Deafness

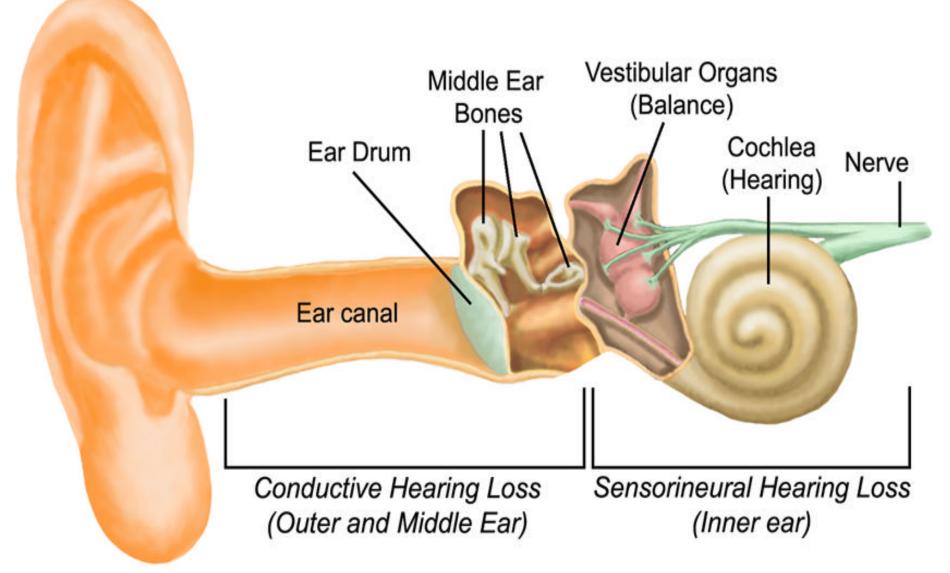
- Two children who passed hearing screens later developed hearing loss
- Each was found to be have 2 mutations in the Cx26 gene (both were 35delG/35delG)
- Child 1: Diagnosed with profound deafness at age 15 months (normal newborn ABR)
- Child 2: Diagnosed with severe hearing loss at age 9 months (normal audiogram at 5 months) JAMA 284 (10):1245, 2000.

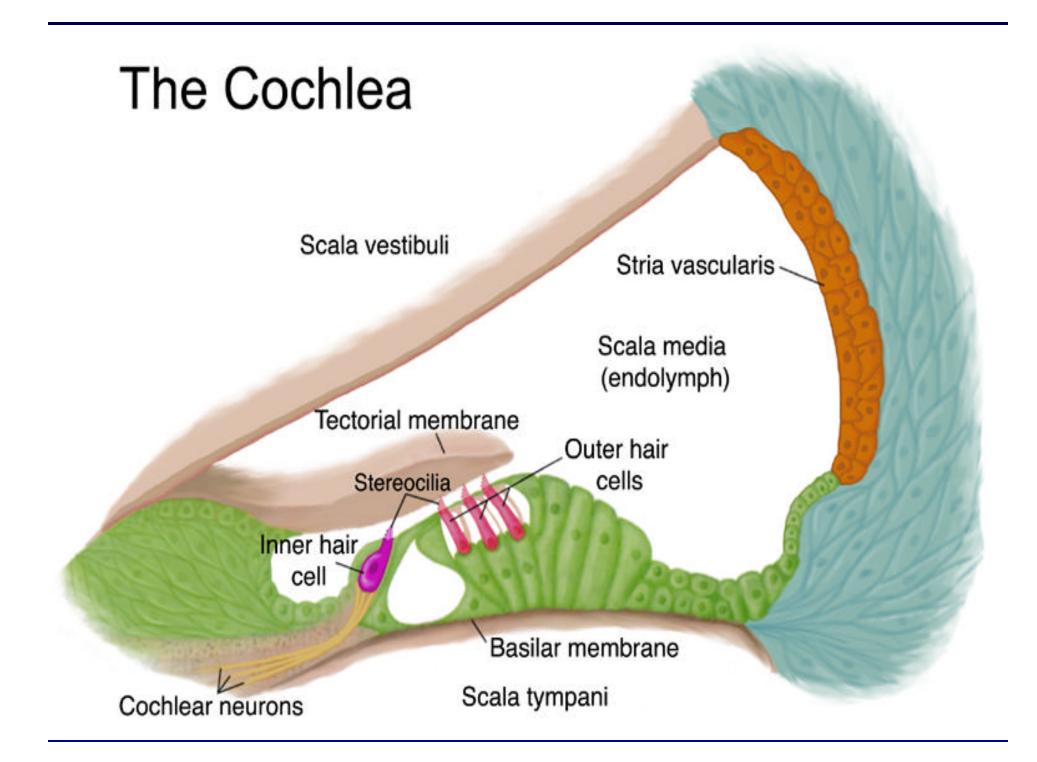
## Progression of Cx26 Hearing Loss

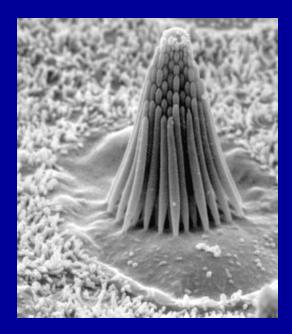


Total = 100 patients

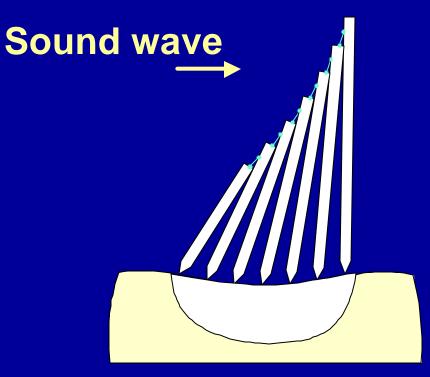
# Anatomy of the Human Ear

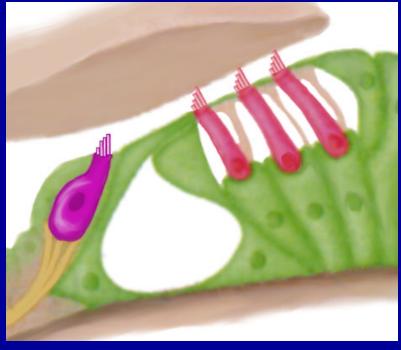


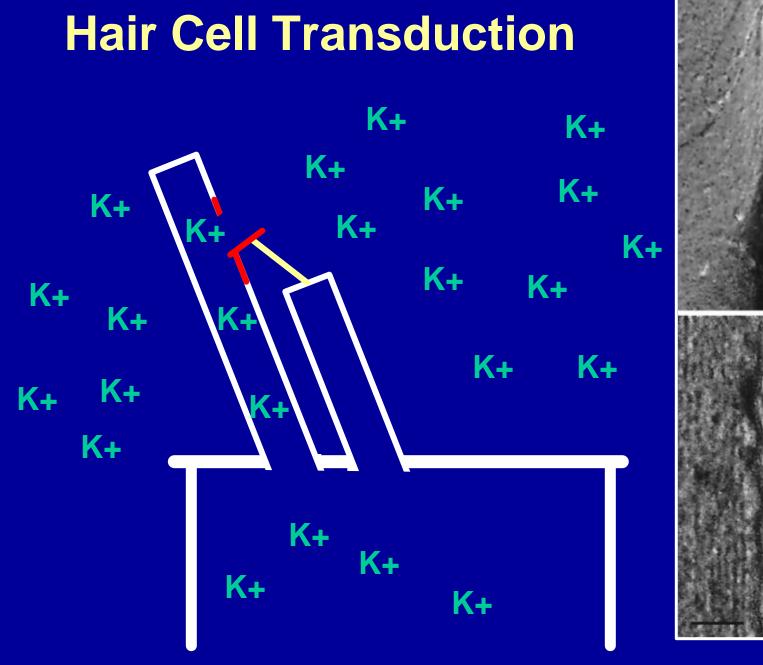


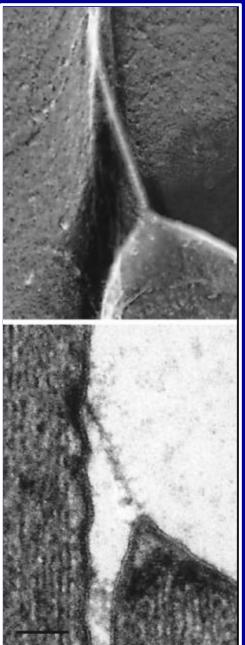


## **Hair Cell Stimulation**

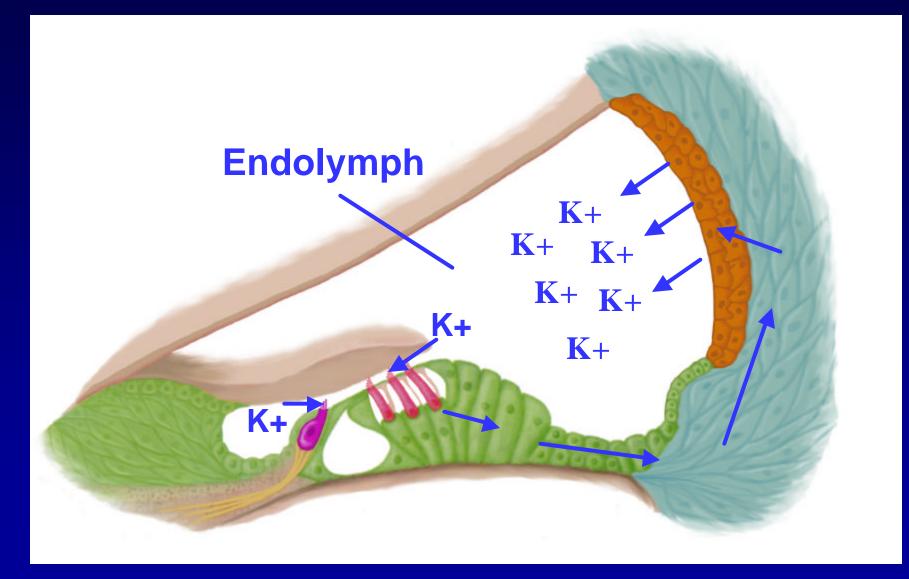


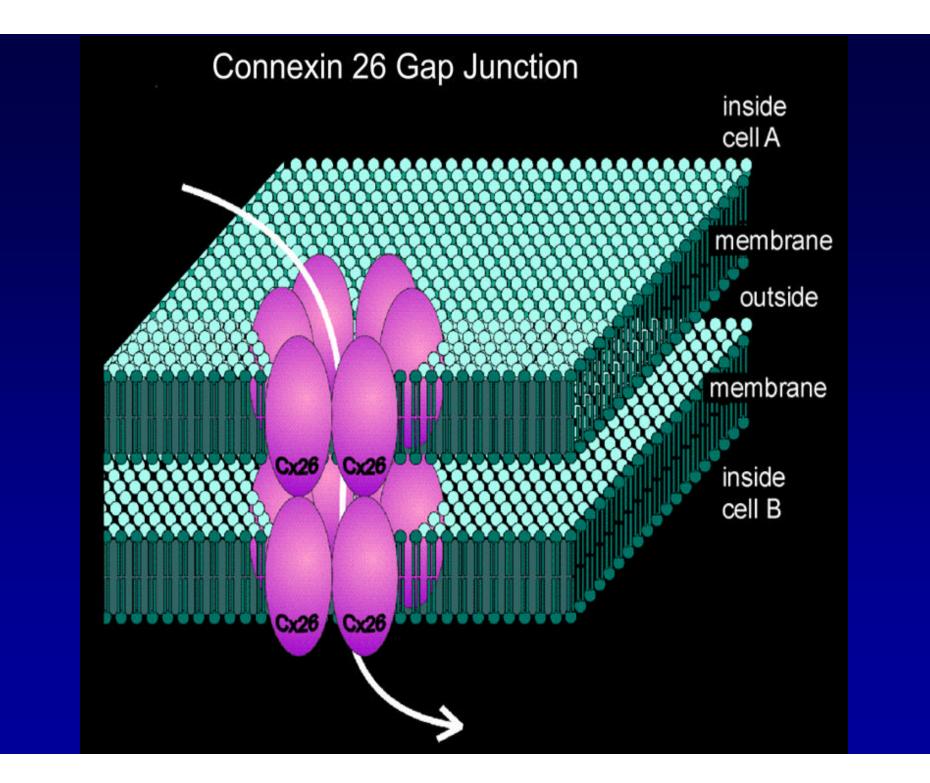






## **Potassium Recycling in the Cochlea**

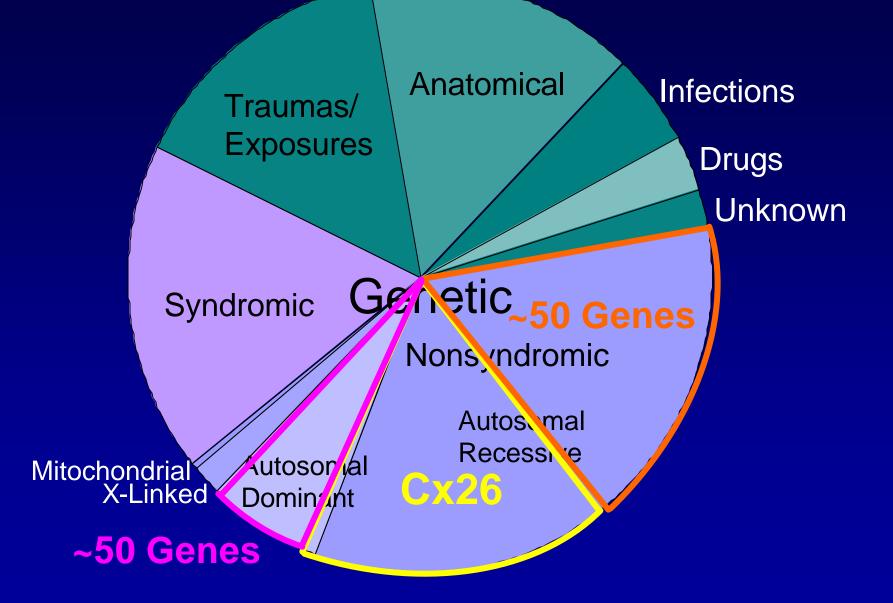


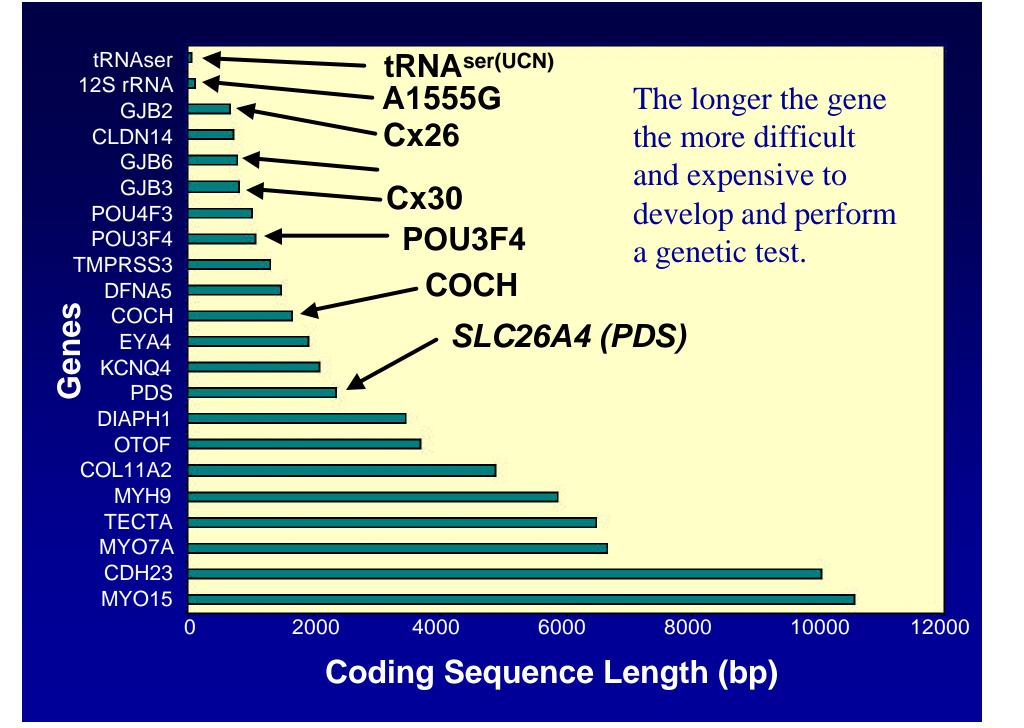


Is Cx26 hearing loss always nonsyndromic??

Most likely, but that does not guarantee the absense of unrelated medical problems

### Major Causes of Hearing Loss





### **Benefits of Genetic Testing for Deafness**

### Genetic testing can:

- ∠ Aid in diagnosis and determining prognosis
- ✓ Eliminate the need for further clinical testing
- Help predict (or rule out) the onset of other clinical features of a syndrome (e.g. blindness in Usher syndrome)
- Help make more informed treatment decisions
- ∠ Aid in making reproductive choices
- Supply considerable "psychological" contentment

## Genes and the Environment

### Mitochondrial 12S rRNA gene

A1555G, C1494T, 961delCins(T)n mutations = increased risk of hearing loss from aminoglycoside antibiotics (i.e. gentimicin, tobramycin, amikacin)

<u>Benefit of test:</u> Prevent other family members with the mutation from being exposed to aminoglycosides

<u>Note:</u> Hearing loss due to these mutations can occur without aminoglycosides and aminoglycosides can cause hearing loss without these mutations.

### **Drawbacks of Genetic Testing for Deafness**

### **Genetic testing may:**

- ∠ Not always give clear answers or any answer
- Put a psychological burden on a parent or relative
- Put an individual at risk for discrimination
  Create ethical dilemmas associated with reproductive choices

### **Attitudes Towards Genetic Testing**

Middleton et al. 1998, Attitudes of Deaf Adults toward Genetic Testing for Hereditary Deafness

55% of "Deaf Nation" attendees thought genetic testing would do more harm than good, 46% thought its use devalued deaf people

Brunger et al. 2000, Parental Attitudes toward Genetic Testing for Pediatric Deafness

96% of parents of deaf children had positive attitude toward genetic testing

Middleton et al. 2001, Prenatal Diagnosis for Inherited Deafness

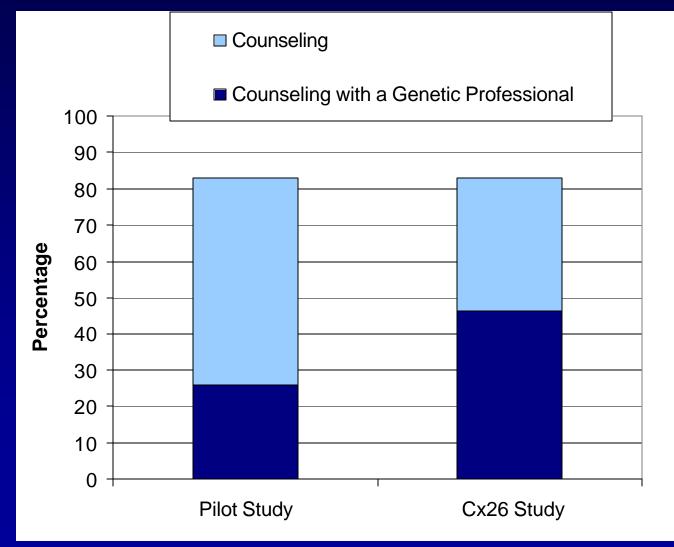
21% (deaf), 39% (HOH), 49% (hearing) would consider prenatal testing – 6%, 11% and 16% would consider terminating pregnancy

### Genetic Counseling Study

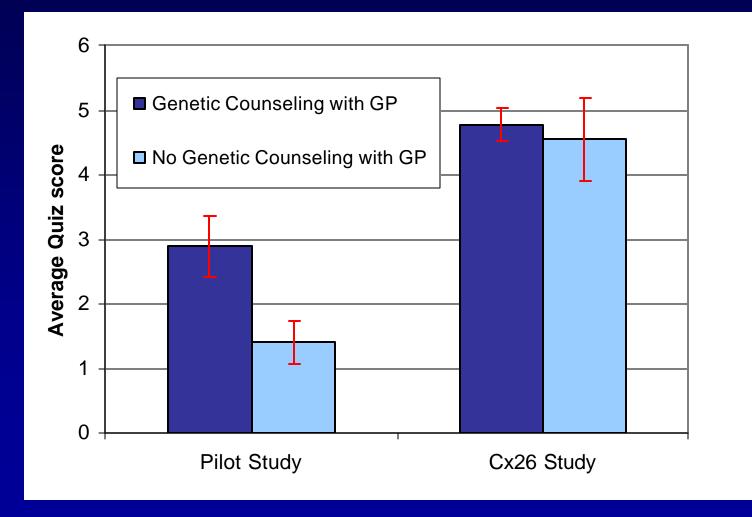
Brunger et al. 2000 found that all respondents had a poor understanding of genetics.

We are examining the extent to which families are receiving genetic counseling for hearing loss and how well they understand the genetics of hearing loss.

# Did you have post-test genetic counseling and who provided it?



## Relationship between Genetic Counseling with a Genetics Professional and Average Quiz Score



## Quiz Question #6

There are many genes that cause hearing loss and tests are not yet available for all of these genes. Therefore, even if a person's current genetic test results are negative, the hearing loss may still be genetic. Were you aware of this fact?

### ? Yes ? No

Pilot study - 34% said "No" Cx26 Study – 22% said "No"

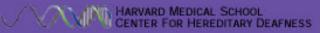
## Take Home Message

Genetic counseling is equally if not more important for those families who are NEGATIVE for Cx26 and other tests

- Could still be genetic
- Recurrence risk 10-15%
- Follow for future testing

#### UNDERSTANDING THE GENETICS OF DEAFNESS A GUIDE FOR PATIENTS AND FAMILIES





### Version 2

<u>Gene Test Cards:</u> Facts About Genetic Testing Connexin 26 Test Mitochondrial Tests SLC26A4 (PDS) Test COCH Test Connexin 30 Deletion Test

http://hearing.harvard.edu

### **Common Causes of Hearing Loss**

### **FOR PARENTS & FAMILIES**



Department of Otolaryngology ? Children's Hospital Boston

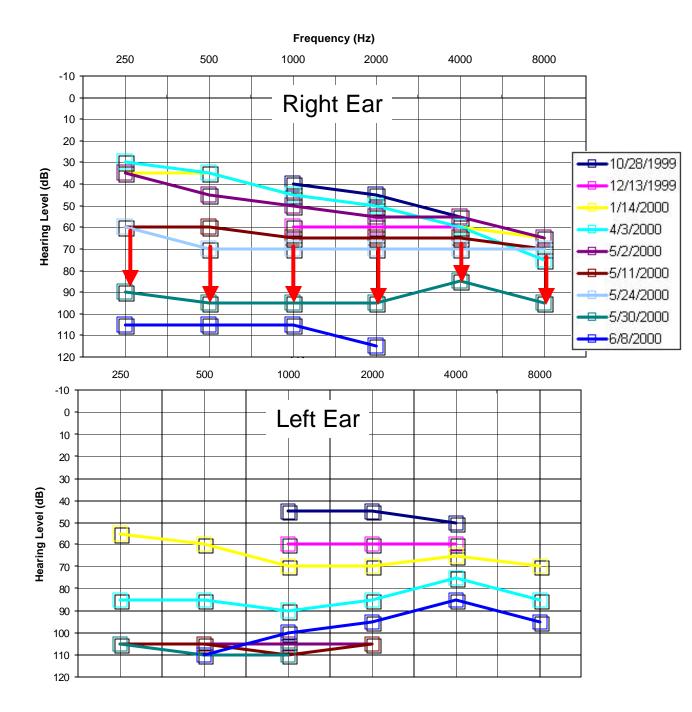
HARVARD MEDICAL SCHOOL

### Acknowledgements

Margaret A. Kenna, MD

Anna Frangulov

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7.5 year old girl discovered with moderate hearing loss during school exam

No newborn hearing screen was performed

5/27/00 – Patient reported sudden loss of hearing

Serial audiograms 6 days apart shows 25-30 dB

Cx26: 35delG/101del2