

Molecular Genetics and Otolaryngology



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Grand Rounds Presentation

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Introduction



- ⌘ Chromosomal analysis
- ⌘ Cytogenetics
- ⌘ Molecular biology and genetics
- ⌘ Biochemical genetics
- ⌘ Clinical genetics
- ⌘ Population genetics
- ⌘ Genetic epidemiology
- ⌘ Developmental genetics
- ⌘ Immunogenetics
- ⌘ Genetic counseling
- ⌘ Fetal genetics

History



⌘ Gregor Mendel, 1865

- ☑ "Mendel's Laws" of autosomal inheritance

- ☑ Work "lost" until early 1900's

⌘ Charles Darwin, 1859

- ☑ "The Origin of Species"

- ☑ Jean Baptiste Lamarck

History, continued



- ⌘ Francis Galton (Charles Darwin's cousin)
 - ☑ The "father" of modern genetics
 - ☑ rediscovered Mendel's laws
 - ☑ "nature versus nurture"
 - ☑ "inborn errors of metabolism" responsible for biological abnormalities

History, Continued



⌘ James Watson and Francis Crick

- ☑ DNA discovered in 1940's
- ☑ Determined double helix in 1953
- ☑ Nobel Prize in 1962

⌘ Human Genome Project

- ☑ Begun in 1990
- ☑ Goal is to identify every human gene by 2005
- ☑ 9% completed as of 1999

Classification of Disorders



⌘ Single Gene Defects

- ⊗ Usually single critical error in the genetic code
- ⊗ Usually phenotypically obvious
- ⊗ Examples: NF I and II, osteogenesis imperfecta, cystic fibrosis

Classification, continued



⌘ Chromosomal disorders

- ⊗ not due to single defect
- ⊗ usually due to deficiency in number of genes within chromosome
- ⊗ classic example is Down Syndrome (Trisomy 21)
- ⊗ other examples: Trisomies 13, 18, Klinefelter's Syndrome, Turner's Syndrome
- ⊗ phenotypically obvious
- ⊗ usually incompatible with life

Classification, continued



⌘ Multifactorial inheritance

- ⊗ multiple single code defects
- ⊗ usually form a pattern
- ⊗ classic examples: cleft lip/palate, neural tube defects
- ⊗ possible example: head and neck cancer?

Chromosomal Structure



- ⌘ 23 pairs of chromosomes
- ⌘ approximately 7 million base pairs
- ⌘ 100,000 genes
- ⌘ DNA:
 - ⊗ five carbon sugar (deoxyribose; ribose in RNA)
 - ⊗ nitrogen base (purines, pyrimidines)
 - ⊗ 3'5' phosphate linkage
 - ⊗ hydrogen bonded double strand

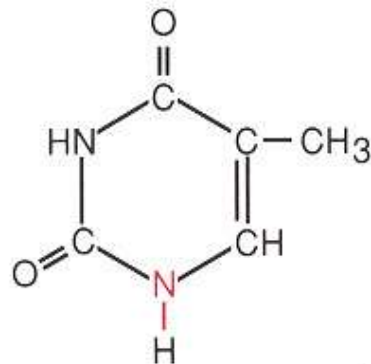
DNA Bases

Purines



Adenine (A)

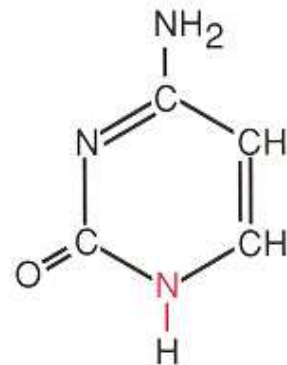
Pyrimidines



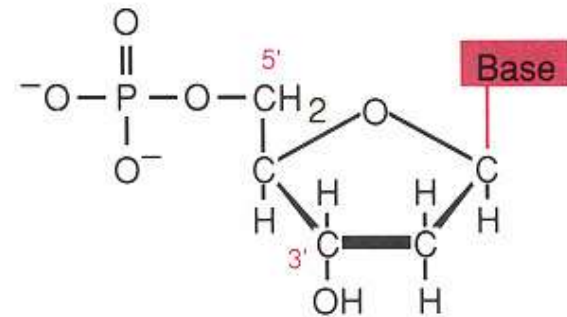
Thymine (T)



Guanine (G)



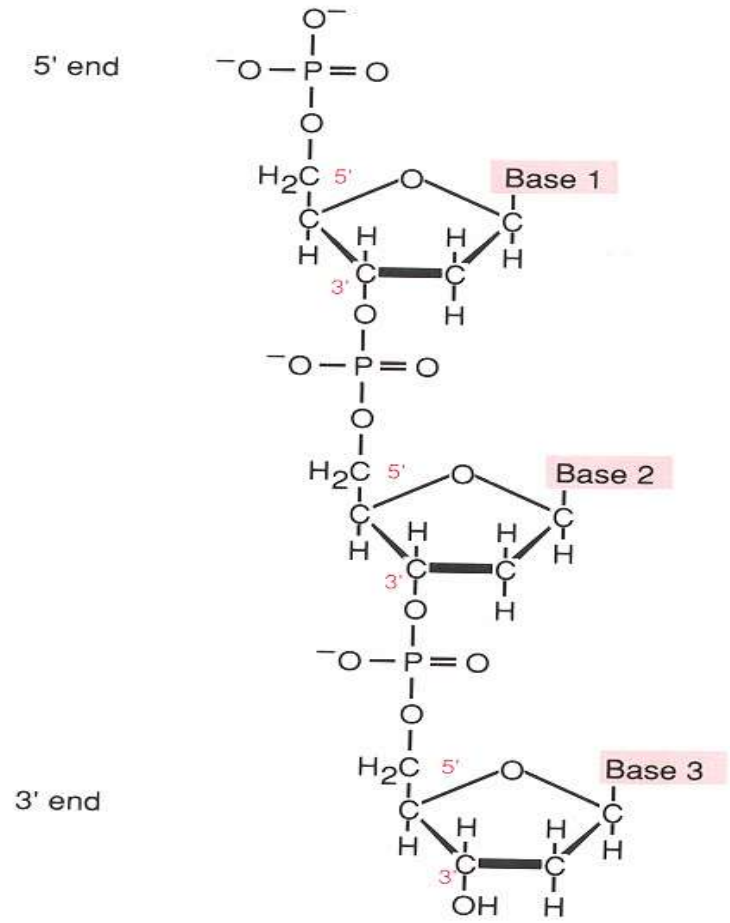
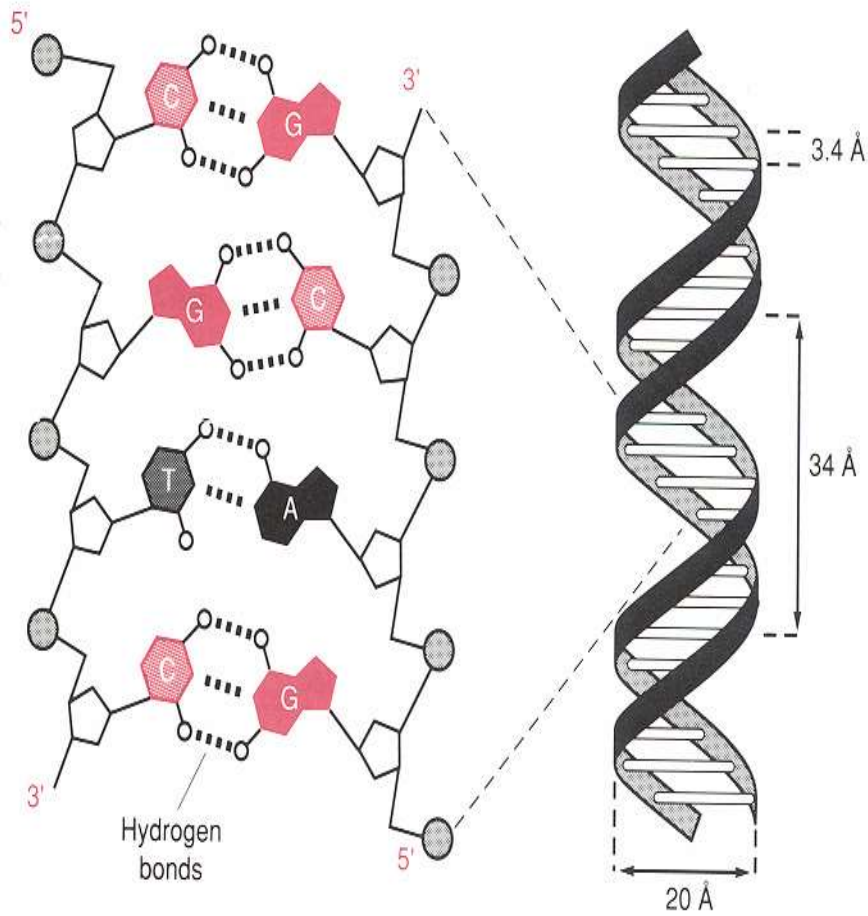
Cytosine (C)



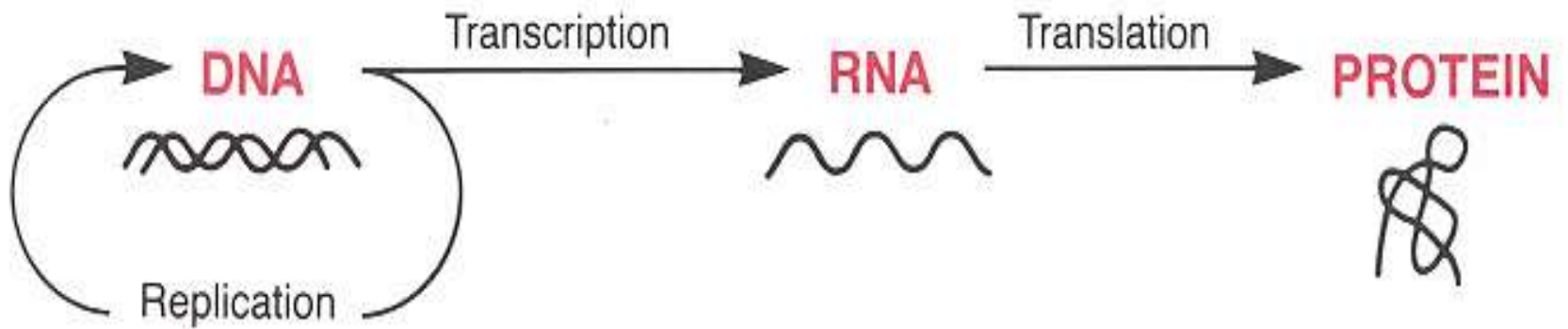
Phosphate

Deoxyribose

DNA Bases



Transcription



⌘ The Central Dogma

Tools of Genetics



⌘ Revolutionary changes since late 1970's

- ⊗ restriction enzymes
- ⊗ recombinant DNA
- ⊗ vectors
- ⊗ probes
- ⊗ PCR
- ⊗ DNA sequence analysis
- ⊗ protein analysis

Tools of Genetics, cont.



⌘ Restriction Endonucleases

- ⊗ enzymes which cleave DNA at specific sites
- ⊗ almost always palindromic
- ⊗ hundreds of known endonucleases

⌘ Recombinant DNA

- ⊗ an DNA fragment is combined with a known piece of DNA to form a plasmid
- ⊗ plasmid inserted in vector (bacterium, virus, yeast)
- ⊗ vector cultured and isolated

Tools, continued



⌘ Identification of recombinant fragments

- ☒ "Blotting" - southern, northern, western
 - ☒ electrophoresis/chromatography of fragment
 - ☒ hybridization with known radioactive fragment
 - ☒ antibodies to known fragments may be used

Tools, continued



⌘ Polymerase Chain Reaction (PCR)

- ⊗ simplest, most rapid, most effective
- ⊗ enzymatic amplification of desired fragment
- ⊗ DNA fragment formed by endonuclease
- ⊗ known "primer" is annealed to fragment
- ⊗ steps repeated approximately 30 times
- ⊗ yields more than a billion copies of desired DNA fragment

Tools, continued



⌘ DNA Sequence Analysis

- ☒ Fred Sanger, Nobel Prize 1980

 - ☒ also won Nobel Prize in 1958 for protein analysis

- ☒ nucleotide analog with inhibits DNA synthesis

- ☒ endonuclease which cleaves at nucleotide site

- ☒ electrophoresis/chromatography

- ☒ radioactive tagging/antibodies

Genetic Mutations



⌘ Defn: Permanent change in nucleotide sequence

⌘ occur in somatic cells or germline cells

⊠ only germline cells inherited

⌘ somatic mutations believed responsible for many medical problems

⊠ many cancers, ?CAD

Genetic Mutations, cont.



⌘ Genome Mutations

- ⊠ missegregation of chromosome
 - results in aneuploidy
 - Down Syndrome classic example
 - 1:50 meiotic divisions
 - usually incompatible with life

Genetic Mutations, cont.



⌘ Chromosome mutations

- ⊗ usually involve translocations and rearrangements
- ⊗ 1:1000 meiotic divisions
- ⊗ almost uniformly incompatible with life

⌘ Gene mutations (single gene defects)

- ⊗ DNA replicates 20 bases/sec/polymerase
- ⊗ Only one defect per ten million copies
- ⊗ Repair enzymes repair 99.9% of defects
- ⊗ Less than one defect per 10 billion bases!

Genetics and Cancer



⌘ Tumor cells are clone of abnormally dividing cell

⊗ usually from single/multiple point mutations

⊗ rarely from translocations

⌘ Protooncogenes

⊗ normal growth genes

⌘ Oncogenes

⊗ a protooncogene which has undergone somatic mutation and is oncogenic

Genetics/Cancer, cont.



⌘ Tumor Suppressor Genes

- ⊗ genes that regulate cell growth/genomic expression
- ⊗ p53, Bcl-2 are classic examples
- ⊗ p53:
 - arrests growth in G1 (growth 1) phase
 - allows repair of DNA defects
 - induces apoptosis (programmed cell death)
 - found in 40% of HNSCCa
 - have NOT shown correlation with prognosis

Genetics/Cancer, cont.



⌘ Bcl-2 tumor suppressor gene

- ⊗ normal Bcell lymphoma/leukemia gene (Bcl-2)
- ⊗ prevents apoptosis (programmed cell death)
- ⊗ somatic mutations present HNSCC, usually resulting in overexpression
- ⊗ Friedman's study:
 - retrospective study of Stage I/II HNSCCa
 - overexpression of Bcl-2 lead to 50% cure versus 90% in normal expression
 - others unable to reproduce (see Gallo)

Treatment



⌘ Most disease treated at phenotypic level

⊗ medicines

⊗ surgery

⊗ genetic counseling

⌘ Molecular level

⊗ gene therapy

Treatment, continued



⌘ Gene Therapy

- ⊗ attempted modification of abnormal cell function
- ⊗ involves transfer of functioning genes
- ⊗ gene therapy via addition
 - more practical
 - insertion into cell (not necessarily into genome) of functioning gene
- ⊗ gene therapy via replacement
 - theoretical
 - goal is to replace abnormal gene with inserted gene

Treatment, continued



⌘ Gene therapy, continued

⊗ Transfer strategies

- recombinant DNA in vector
 - viral versus bacterium
 - retroviral vectors with reverse transcriptase
- not inserted into host genome

⊗ problems:

- inability to maintain expression
- under/overexpression
- adenine deaminase deficiency (ADA)

Genetic Disease in ENT

⌘ Cystic Fibrosis

- ⊗ chromosome 7q, spans 250,000 bases
- ⊗ 70% have deletion of phenylalanine at position 508 (point mutation)
 - frameshift versus point mutation
- ⊗ most common fatal autosomal disease in whites
- ⊗ phenotypic expression results from failure of membrane transport (Cl, Na) and from exocrine function (pancreas)
- ⊗ Tx at phenotypic level

Genetic Dz in ENT, cont.



⌘ Cleft Lip and Palate

- ⊗ one of the most common malformations
- ⊗ CL and P genetically distinct from isolated CL
- ⊗ failure of fusion of frontal process with maxillary process at 35 days gestation
- ⊗ classically described as multifactorial, although single gene forms, chromosomal forms (Trisomy 13) teratogenic forms (rubella, thalidomide) are known

Genetic Dz in ENT, cont.



⌘ Human papilloma virus

- ⊗ strains 16, 18 and 31 carcinogenic in GU tract
- ⊗ exact role in HNSCCa not fully known, although 46% of post mortem specimens contained HPV strains
- ⊗ E6 HPV protein binds to p53 forming mutation which suppresses gene function in vivo

Genetic Dz in ENT, cont.

⌘ Thyroid carcinoma

⊗ Medullary thyroid carcinoma (MTC)

- neoplasm of parafollicular C cells (ultimobranchial body)
- produce calcitonin
- sporadic and familial forms
- familial MTC associated with MEN 2A and 2B
 - MEN 2A: pheo, hyperparathyroid, MTC
 - MEN 2B: pheo, MTC, Marfan's, NFI
- RET protooncogene associated with familial forms
 - 10p

⊗ Aggressive papillary CA associated with aneuploidy

- noninvasive dz uniformly diploid

Genetic Dz in ENT, cont.

⌘ Salivary Gland Neoplasms

- ☒ Aggressive adenoid cystic Ca associated with aneuploidy
 - all patients with aneuploidy recurred after resection versus only 2/14 with diploid genome (Sugano)
- ☒ Salivary gland adenocarcinoma with overexpression of Bcl-2 were more difficult to resect, recurred more frequently and metastasized more frequently (Sugano)

Genetic Dz in ENT, cont.



⌘ Acoustic Neuroma

- ⊗ 5% are familial and associated with NF II
- ⊗ often bilateral
- ⊗ NF II defect on 22p
- ⊗ therapy at phenotypic level

Genetic Dz in ENT, cont.



⌘ Congenital Hearing Loss

- ⊗ 60% of congenital hearing loss is genetic
- ⊗ most associated with phenotypic anomaly
- ⊗ Waardenburg Syndrome
 - autosomal dominant - variable penetrance
 - dystopia canthorum, hyperchromatic iris, white forelock and SNHL
 - PAX3 locus of chromosome 2
 - treatment at phenotypic level

Genetic Dz in ENT, cont.



⌘ Congenital hearing loss, continued

☒ Usher's Syndrome

- autosomal recessive
- five different classifications (Usher's Types I through V)
 - all subtypes on different chromosomes
- associated with retinitis pigmentosum
- therapy at phenotypic level

Genetic Dz in ENT, cont.



⌘ Congenital Hearing Loss, continued

⌘ Pendred's Syndrome

- ⌘ autosomal recessive with variable penetrance
- ⌘ located on chromosome 7q
- ⌘ associated with thyroid goiter and carcinoma
- ⌘ tx at phenotypic level

Genetic Dz in ENT, cont.



⌘ Congenital hearing loss, cont.

☒ Alport's Syndrome

☒ two forms: X linked, autosomal recessive

- X linked on 5p, produces mutant alpha 5 protein
- recessive form on 2p, produces mutant Type IV collagen

☒ treatment at phenotypic level

Genetic Dz in ENT, cont.



⌘ Head and Neck Cancer

- ☑ heavily associated with p53 underexpression, Bcl-2 overexpression, HPV types 16, 18 and 31
- ☑ None of these proven prognostic
- ☑ Ultimate goal: gene therapy to correct somatic mutation

Future Directions and Conclusion



- ⌘ Rapidly expanding field
- ⌘ Ultimate goal: correction of somatic defect which would correct phenotypic abnormality. Would eliminate surgical intervention.