lec 1 - Cytogenics = Study of chromosomes - double stranded DNA wrapped around histores = nucleosome - packed nucleosomes = Chromatin - Chromatin embeded in scaffold = Chromatid - 2 sister chromatids = chromosome Cell Cycle Interphase = G1, S, \$G2 DNA is loose & accessable for proteins to be able to bind - Before S phase, DNA is one darble Stranded Chromatid (stil) called chromasome) - G phase is growth phase, where RNA polymerase makes more MRNA & proteins nessesary for Sphase - Sphase is "Synthesis" phase, where DNA polymerase replicates double stranded Chromatid to make a Sister chromatid joined at the centromere \* chromosome enters & phase as one chromatid & come out as 2 chromatids - M phase is mitosis, DNA is tightly packed for easier separation of chromosomes \* most condensed DNA in metaphase chromosome = condensed chromatin

Nomenclature of Chromosomes based on centromere position & arm length upper arm = P (shorter) ; lower arm = ? (longer) Centromeric position and arm lengt Dabnormal no parm Human Chromosome - 23 pairs of chromosomes, from mom & Dad - 22 pairs of diploid (Somatic cells), 1 pair sex Chromosome XX or Xy (haploid) Mitosis - division of cells to produce identical daughter cells in terms of DNA sequence, not cytoplasm 5 phases: Prophase chromosomes begin to condense, nuclear envelope collapse, centricles appear on opposite poles Metaphase Chromosomes individually allign on metaphase plate Anaphese Separation of sister Chrometids Telophase nuclear envelope forms, 2 daughter nuclei form, Chromasomes diffuse <sup>2</sup> Cleavage furrow forms Cytokinesis

division of eytoplesm Meiosis produces daughter cells w/ 1/2 # of Chromosomers \$ different DNA sequences (only egg & sperm) - 2 stages : Meiosis I = separation of homologous chromosome meiosis 2 = Separation of Sister chromatids Meiosis 1 Prophase 1 Condensation, envelope disapears, centrioles, Recombingtion \* takes most time of all phases Metaphase I homologous pairs line up Anaphase 1 disjunction of homologous pairs \* Chromosome # reduced by 1/2 Telophase 1 two haploid cells separate Meiosis 2 - immediatly after meiosis 2 (no interphase) - Same steps Anaphase 2 Separation of Sister Chromatils - final product is 4 non identical daughter cells (egg/sperm)

Prophase 1 Recombination occurs Exchange of genetic material between non sister chromatides of homologous chromosomes - sub divided to 5 stages Leptotene innitiation of condensation, homologous chromosomes align for recombination Zygotene Synapsis & tetrad formation Pachytene Synapsis is complete, each homologous pair is called a tetrad, Recombination occurs Diplotene Chromosomes separate but stay bound at chiasmate Diakinesis further condensation until it reaches metaphase Gregor Mendel laws - law of segregation: homologous chromosomes separate & appear in different daughter cells - law of independent assortment: chromosomes align on metaphase plate independently so traits are inherited independently of each other \* most of cell cycle taken by Interphase

Lec 2 - Each chromosome has genes which are segments of DNA that carry info for a specific trait; ? different versions of genes are colled <u>alleles</u> - possible genetic combinations are calculated by 2<sup>1</sup> n = >> of chromosome pairs (2<sup>23</sup>) - Melosis gives genetic diversity b/c of independent assortment & genetic Recombingtion in meiosis 1 why study chromosomes ? - Clinical conditions are linked to chromosome abnormalities - chromosome abnormalities are seen in: 20-27+ of sex reversal / pueberty abnormalities 33-67.1. of spontaneous miscarage 2-54 Multiple miscarage majority of leukemia & Solid tumors how to study chromosomes - extraction of chromosome depends on disease 4 purpose of study - Amniotic fluid Cells used for embryonic genes - Bone marrow for diseases in blood - Peripheral blood for lymphocytes - Skin or organ biopsy - A Karyoyye is a dragram of ordered arrangement of Chromosomes by hight (1 being longest & 22 being the shortest, followed by Sex chromosomes)

Notes before next section: - Adenine & Thymine have 2 H-bonds (AZT) · Guanine & Cytosine have 3 H-bonds (G&C) gene rich region = euchromatic (mainly G&C) gene poor region = heterochromatic (mainly 4 \$T) this is because AT areas have tissue specific genes while GC areas have houskeeping genes - trypsin is a digestive enzyme that degrades peptide bonds & digest chromosomal proteins to relax chromatin structure So Stain can access DNA 6-Banding Technique - Chromosomes treated or trypsin first - Giemsa Stain is added \* Giemsa forms hydrophobic interractions @ hydrogen binding site, & favors AT regions more - Giemsa binds AT stronger than GC, so AT regions show darker (black) stain - produces 300-400 bands & each band has several -10 million nucleotides K-Banding Technique - the "Reverse" of G banding, So cuchromatic areas are stained dork (GC) ... why? - b/c first we heat the sample, causing degridation of AT hydrogen bonds so Gremsa stains GC regions darker

- flourochrome is added to sharpen or enhance color used in G-banding to identify deletion mutations \$ in R-banding to confirm findings of G-banding Chromosome Shape · Metacentric, Submetacentric, Acrocentric - 9 ? Parm are numbered starting from centromere ending @ telomere Acrocentric - Chromosomes 13-15,21\$22 - P-arm has 2 regions 1) Satellite: heterochromatic region, darkly stained, repetative sequence of base pairs 2) Stalk : euchromatic, lightly stained, codes for rRNA - P-arm has no clinical Significance so demage to it wont affect anything ... other acrocentric Chromosomes Can compensate & make r RNA High Resolution Kanding - damage to about 300 million nucleotides can be detected by GER banding, but damage to less, cannot what do we do? · obtain sample from less condensed chromatid (prophase / pro-metaphase) = # of banding increases from 300-450 or even 800 - So we now can detect less obvious abnormalities Chromosome Structure

Centromeres - Structural heterochromatic DNA required for Chromosome segregation - tandem repeats of 171 base pairs called alpha satelite - all chromosomes have <u>Same</u> alpha Satelite Telomeres - at ends of eccleryotic chromosomes - TTAGGG repeats thousands of time - structurel region that prevents end to end fusion of chromsomes · Replicated by DNA polymerase & telomerase as you age, telomerase activity V telomeres get shorter each time Cells divide until its gone & cell is innactive - protects us from cancer Sub-telomeric Region - repeats are not universal like telomeres - Some homology, but not identical among chromosomes Lec 3 Aneuploidy " Chromosome number thats not an exact multiple of haploid number (2n)

- trisomy: extra chromosome - Monosomy : absence of a chromosome - Change in Chromosome # due to nondisjunction ;ţ nondisjunction happens in Merosis 1 then homologous chromosomes dont separate 1/2 the gametes have extra chromosome (n+1) \$ other 1/2 is missing a chromosome (n-1) f if in meiosis 2, then sister Chromatids dont separate 1/2 carry correct \* (n), 1/2 has extra (n+1), 1/4 is missing 1 (n-1) \* nondisjunction of milosis is the same as meiosis 2 Polyploidy / Euploidy More than 2 complete sets of chromosomes to extra, but exact copy of n - Triploidy (3n) tetraploidy (4n) - polyploidy more common in plants Polyploids are more normal in appearance than anneploids Alteration of chromosome Structure - Deletion: remove chromosome segment - Duplication: repeats Segment · inversion : reversing orientation of segment on same chromassme Translocation: moving segment from one chromosome to another

- Alteration of chromosomes leads to serious disorders - Some anneploidy disrupts genetic belance more than others, so some people can survive but suffer from Syndromes Down Syndrome (Trisony 21) - 3 copies of Chromosome 21 - 1/100 children - Correlated to pregnancy at advanced age Clinically - mental retardation - low nasal bridge - hypotonia - up slasting palpebral fissures - low set cars - congenital Odisease (USD, AVD) - Cyefolds, protruding tongue, intestinal problems, 15 fold 1 risk of leukemia - Simian line \* features easily recognized at Birth - non disjunction can occur in meiosis 1 or 2 94% due to maternal error 64% in merosis 2, 19% in meiosis 2, 1)% intermediate : 4.5% due to paternal error L | % meiosis 1, 3.5+ meiosis 2, 1.5 y unknown

how do we determine if the extra chromosome Came from mother or father ? - we look at repetative non coding regions of DNA - these regions differ from person to person, so it C) used as a polymorphic marker Hartial Trisomy 21 (21g) - 2 fused 9 arms at chromosome 21, instead of 1 g arm (46, XX, 21 9,+) - Phenotypically down syndrome Edward Syndrome (Trisony 18) - CHD - failure to thrive - retarded (mental & growth) - hypertonia - pominant Occiput - low set ears - intestine problems - clenched fist - rockerbottom feet Vatau Syndrome (trisony 13) - CHD Scalp defects - retarded - hyper/hypo tonia Microcephalu

-( J - low set ears Small eyes Cleft I:p/ palate rockerbottom feet - poly syndacty ly - polycystic Kidneys