

Patho Unit7 Genetic Disorder & Growth Disorder Cheat Sheet by damn via cheatography.com/195477/cs/41609/

1.Autosomal Dominant	2. Autosomal Resessive	3. X-linked Recessive
Both sex involved	- Bloth sex involved	- Males only
Generations not Ripped	- Generations skipped	

Single Gene Disorder	
Enzyme Defects	Structure protein defects
1. Accumulation of substrate	1, Martan Syndrome
2. Lack of product	2. Ethler-Danios Syndromes
3. Failure to inactivate a prote	in which cause damage
	Receptor protein defect
	- Familial hypercholesterolemia (AD)

Autosomal Dominant	Autosomal Recessive	X-link recessive
Both siex involved	- Both sex involved	- Males only
Generatrios not skipped	- Generation skipped	 Generation skipping doesn't matter

Chiromosomoil Disorder

- Chromosome, genome mutation

Down Syndrome

- Meiotic nondisjunction of chromosome 21

Klinefelter Syndrome

- ->=2 Xchromosome/ 1>= Y chromosome
- 47.30CV
- Testicular abnormality

Turner syndrome:

- Comple/ partial monosomy of X chromosome in females
- Neck webbing, heart defects



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Scientific and Technical Books and Serials in Print ,1989 Microbial Comparative Genomics and Pangenomics:

New Tools, Approaches And Insights Into Gene and Genome Evolution Paulo Jorge Dias, Satyanarayana Tulasi ,Digvijay Verma,2024-11-11 Comparative Genomics is the field of knowledge dedicated to the analysis and comparison of genes and genomes The scientific areas comprised in this field include subjects as diverse as just naming a few 1 the development of algorithms for the alignment of genes whole genomes short and long sequencing reads 2 the search for remote sequence similarity 3 the discovery of motifs and sequence patterns 4 the identification of gene families 5 the detection of ortholog paralog groups 6 the reconstruction of evolutionary history of the genes 7 the detection of signs of selective forces exerted over genes and genomes 8 the reconstruction of ancestral DNA and genome sequences 9 the detection and analysis of genome synteny 10 the inference of ancestral gene order among others In addition an important new sub field of Comparative Genomics has emerged in the last decade referred to as Pangenomics making available improved tools to analyze the exponential genomic data accumulating since the development of Second and Third Generation Sequencing Technologies

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