MeSH Tree Structures - 2008

G13 - GENETIC PHENOMENA

Genetic Phenomena

Genetic Phenomena	G13		
Consanguinity	G13.180	G5.90.	
Founder Effect	G13.285		
Gene Frequency	G13.330		
Gene Flow	G13.330.159		
Genetic Drift	G13.330.320	G5.265.	
Gene Order	G13.340		
Gene Pool	G13.345		
Genetic Load	G13.360		
Genomic Instability	G13.370	C23.550.	G13.920.
Chromosomal Instability	G13.370.180	C23.550. G13.920.	C23.550.
Chromosome Fragility	G13.370.180.180	C23.550. G13.920.	C23.550.
Microsatellite Instability	G13.370.590	C23.550.	G13.920.
Genotype	G13.380		
Gene Dosage	G13.380.350		
Genetic Predisposition to Disease	G13.380.355	C23.550.	
Haplotypes	G13.380.360		
Heterozygote	G13.380.383		
Homozygote	G13.380.554		
Hybrid Vigor	G13.400		
Inheritance Patterns	G13.420		
Anticipation, Genetic	G13.420.40	C23.550.	
Extrachromosomal Inheritance	G13.420.275		
Genes, Mitochondrial	G13.420.275.500	G14.340.	
Genes, Dominant	G13.420.320	G14.340.	
Genes, Recessive	G13.420.325	G14.340.	
Genes, X-Linked	G13.420.457		
Genes, Y-Linked	G13.420.523		
Multifactorial Inheritance	G13.420.590		
Quantitative Trait, Heritable	G13.420.720		
Linkage (Genetics)	G13.540		
Linkage Disequilibrium	G13.540.500		
Lod Score	G13.540.562		
Phenotype	G13.695		
Genetic Markers	G13.695.450	D23.101.	
Penetrance	G13.695.650		
Phylogeny	G13.697	G4.185.	L1.100.
Ploidies	G13.700		
Aneuploidy	G13.700.131	C23.550.	G13.920.
Monosomy	G13.700.131.500	C23.550.	G13.920.
Trisomy	G13.700.131.750	C23.550.	G13.920.
Diploidy	G13.700.264		
Haploidy	G13.700.456		
Polyploidy	G13.700.740		
Sequence Homology	G13.810	G6.184.	
Sequence Homology, Amino Acid	G13.810.200	G6.184.	
Sequence Homology, Nucleic Acid	G13.810.550	G6.184.	
Synteny	G13.810.550.830	G6.184.	
Sex Ratio	G13.815	G3.850. N1.224.	I1.240.
Structural Homology, Protein	G13.820	G6.184.	G6.184.
Variation (Genetics)	G13.920		
Antibody Diversity	G13.920.36	G4.610.	G4.610.
Antigenic Variation	G13.920.73	G4.610.	
Genetic Heterogeneity	G13.920.331		
Mutation	G13.920.590		

Genetic Phenomena
Variation (Genetics)
Mutation
Allelic Imbalance

Allelic Imbalance	G13.920.590.29		
Loss of Heterozygosity	G13.920.590.29.530		
Chromosome Deletion	G13.920.590.29.530.175	C23.550.	G5.600.
		G13.920.	G13.920.
Base Pair Mismatch	G13.920.590.60		
Chromosome Aberrations	G13.920.590.175	C23.550.	
Aneuploidy	G13.920.590.175.50	C23.550.	G13.700.
Monosomy	G13.920.590.175.50.500	C23.550.	G13.700.
Trisomy	G13.920.590.175.50.750	C23.550.	G13.700.
Chimerism	G13.920.590.175.125		
Chromosomal Instability	G13.920.590.175.165	C23.550.	C23.550.
		G13.370.	
Chromosome Fragility	G13.920.590.175.165.180	C23.550.	C23.550.
		G13.370.	
Chromosome Breakage	G13.920.590.175.175	C21.111.	C23.550.
		G5.180.	
Chromosome Deletion	G13.920.590.175.177	C23.550.	G5.600.
		G13.920.	G13.920.
Inversion, Chromosome	G13.920.590.175.420	C23.550.	G5.600.
Isochromosomes	G13.920.590.175.430	A11.284.	C23.550.
		G14.162.	
Micronuclei, Chromosome-Defective	G13.920.590.175.570	A11.284.	A11.284.
		C23.550.	
Mosaicism	G13.920.590.175.595		
Ring Chromosomes	G13.920.590.175.760	A11.284.	C23.550.
		G14.162.	
Sex Chromosome Aberrations	G13.920.590.175.815	C23.550.	
XYY Karyotype	G13.920.590.175.815.970	C23.550.	
Translocation, Genetic	G13.920.590.175.870	C23.550.	G5.600.
Philadelphia Chromosome	G13.920.590.175.870.680	A11.284.	A11.284.
		C23.550.	G14.162.
		G14.162.	
Uniparental Disomy	G13.920.590.175.935	C23.550.	
Codon, Nonsense	G13.920.590.195	D13.444.	G14.335.
DNA Repeat Expansion	G13.920.590.220	G5.600.	G6.184.
		G14.80.	G14.340.
		G14.340.	
Trinucleotide Repeat Expansion	G13.920.590.220.865	G5.600.	G6.184.
		G6.184.	G14.80.
		G14.80.	G14.340.
		G14.340.	
Frameshift Mutation	G13.920.590.265		
Gene Amplification	G13.920.590.310	G5.315.	G5.600.
Gene Duplication	G13.920.590.320	G5.600.	
Genomic Instability	G13.920.590.335	C23.550.	G13.370
Microsatellite Instability	G13.920.590.335.590	C23.550.	G13.370.
Germ-Line Mutation	G13.920.590.350		
INDEL Mutation	G13.920.590.500	G5.600.	
Mutagenesis, Insertional	G13.920.590.575	E5.393.	G5.600.
Mutation, Missense	G13.920.590.650		
Point Mutation	G13.920.590.675		
Sequence Deletion	G13.920.590.762	G5.600.	
Chromosome Deletion	G13.920.590.762.180	C23.550.	G5.600.
		G13.920.	G13.920.
Gene Deletion	G13.920.590.762.320	G5.600.	
Suppression, Genetic	G13.920.590.835	G5.600.	
Polymorphism, Genetic	G13.920.795		
Polymorphism, Restriction Fragment Length	G13.920.795.595		
Polymorphism, Single Nucleotide	G13.920.795.598		

Genetic Phenomena
Variation (Genetics)
Polymorphism, Genetic
Polymorphism, Single-Stranded Conformational

Polymorphism, Single-Stranded Conformational G13.920.795.600