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1. In some cats the gene for tail length shows incomplete dominance. Cats with long tails and cats with no tails are homozygous for their respective alleles. Cats with one long tail allele and one no tail allele have short tails. Construct a punnett square and give phenotypic and

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phenotypic ratios can be inferred as well. For a monohybrid cross of two truebreeding parents, each parent contributes one type of allele. In this case, only one genotype is possible.

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Mendelian. inheritance patterns in which a single gene affects a single trait and the alleles obey the law of segregation. these patterns include simple dominant/recessive traits, Xlinked traits, incomplete Page 11/15

dominance, codominance, and sex-influenced traits. Epistasis.

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Patterns of inheritance in humans include autosomal dominance and recessiveness, X-linked dominance and recessiveness, incomplete dominance, codominance, and lethality. A change in the nucleotide sequence of DNA, which may or may not manifest in a phenotype, is called a mutation.

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Inheritance patterns of F 2 progeny were non-Gaussian and deviated from Mendelian expectations. However, discrete alkyl cannabinoid Page 13/15

segregation patterns uide consistent with digenic as well as epistatic modes of inheritance were observed among F 2 THCAS and CBDAS genotypes. These results suggest linkage between cannabinoid pathway loci and highlight ...

Complex Patterns of Cannabinoid Alkyl Side-Chain

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