HMMs for Gene Finding

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Come Properties: with start codon: ATG with stop orders end 76A (triplets of ma) · Some wans are more common AAA AAC ATG TAG TGA Codon usage table 0,01 0,02 0,01 7=1 ACGT Tro. 40-1 0.1 04] AAA . AT6 [0000100000] 1/333 1/300 Intergence Gene AAA AAC ATG TAA FAG AAA TAG TAATA from [0.01 0.02 0.01 0 [000 1/300 1/300 1/3 0] Codonosago table

Come structure in enkargate exon! Cenome 5'07R untranslated regin 3 by TR untranslatedry Transpréplia pre-mRNA Splicing el e2 e3 e4 mature mRNA transfition Typical game in 10 exons of 2100 bp on award intrays of 210000 bp onais Gene 2 31/33 Intergeric Intron 1 AEGT 7)9999

