


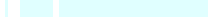
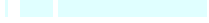









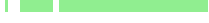



















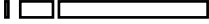






















Gene Structure					
misc_RNA		precursor_RNA		prim_transcript	
3'clip		5'clip		mRNA	
exon		intron		3'UTR	
5'UTR		CDS		mat_peptide	
sig_peptide		transit_peptide		polyA_signal	
polyA_site		gene		operon	
Regulation					
attenuator		enhancer		CAAT_signal	
TATA_signal		promoter		GC_signal	
protein_bind		misc_binding		-10_signal	
-35_signal		terminator		misc_signal	
rep_origin		RBS			
Repeats					
repeat_region		repeat_unit		LTR	
satellite		stem_loop		misc_structure	
Rna Genes					
rRNA		scRNA		snRNA	
snoRNA		tRNA			
Sequence					
source		misc_recomb		variation	
misc_feature		old_sequence		unsure	
misc_difference		conflict		modified_base	
primer_bind		STS		gap	
Graphs					