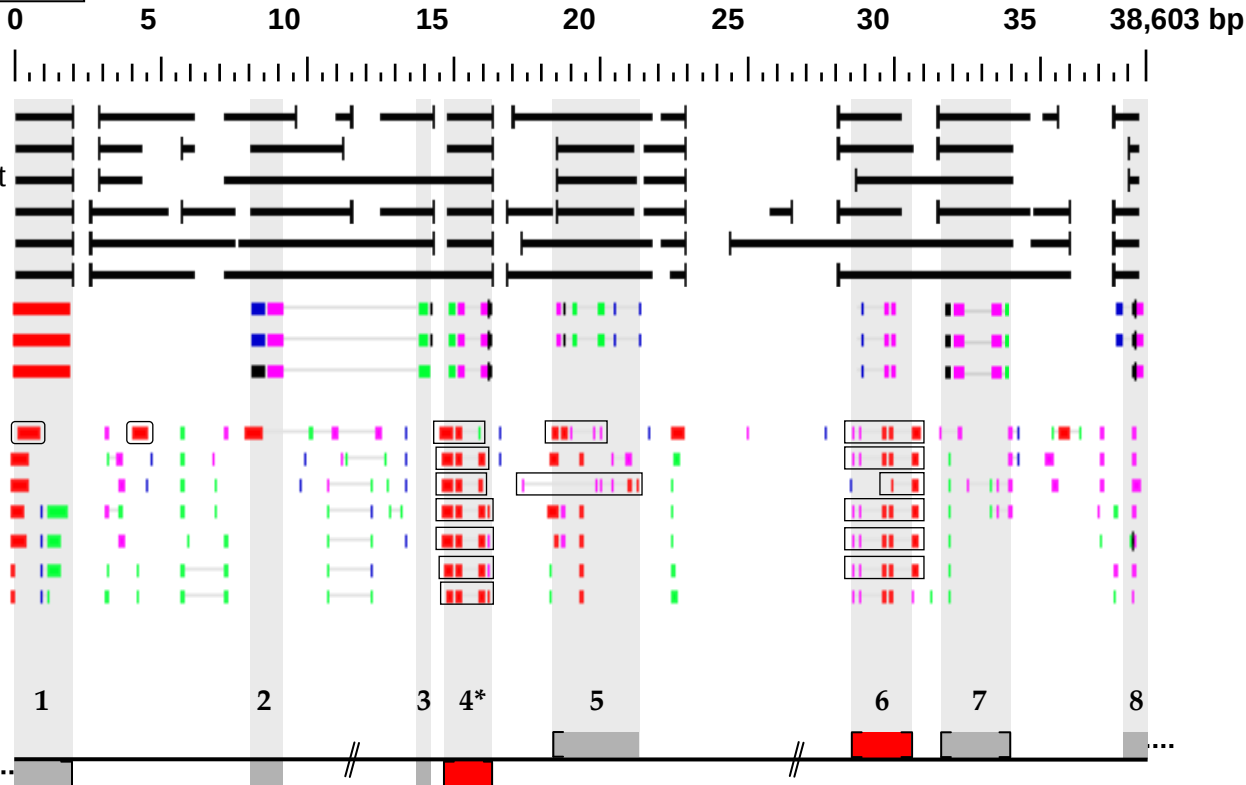


Fosmid 48I08



Motif	TCT				GA	TA	TA	GA	TA	TTC					GA	GA	TGG	CT				
Repeat No.	6				6	6	6	6	16	9					5	5	6	5				
SSR start	638				10646	11269	11307	12113	12732	18230					28499	31307	31781	34992				

Fosmid 48I08 = EU024856 (<i>CHI</i>) 38,803 bp									
	Putative gene location			Blastx Results		Putative gene location			Blastx Results
Feature	Strand	Putative start	Putative stop	Putative identity	Feature	Strand	Putative start	Putative stop	Putative identity
1	Minus	1942 1939	excluded	protein kinase, putative	5	Plus	18401	nd	unknown protein
2	Minus	nd	nd	protein kinase-related	CONTIG BREAK		26584	26683	
CONTIG BREAK		11537	11636		6	Plus	28567	30544	40S ribosomal protein S13 (RPS13A)
3	Minus	nd	nd	protein kinase-related	7	Plus	31596	33918	EMB1067 (EMBRYO DEFECTIVE 1067); tRNA 2'-phosphotransferase
4*	Minus	16220	14748	chalcone isomerase 1 (<i>CHI</i> -1)	8	Plus	37804(?)	excluded	disease resistance protein (TIR-NBS-LRR class), putative