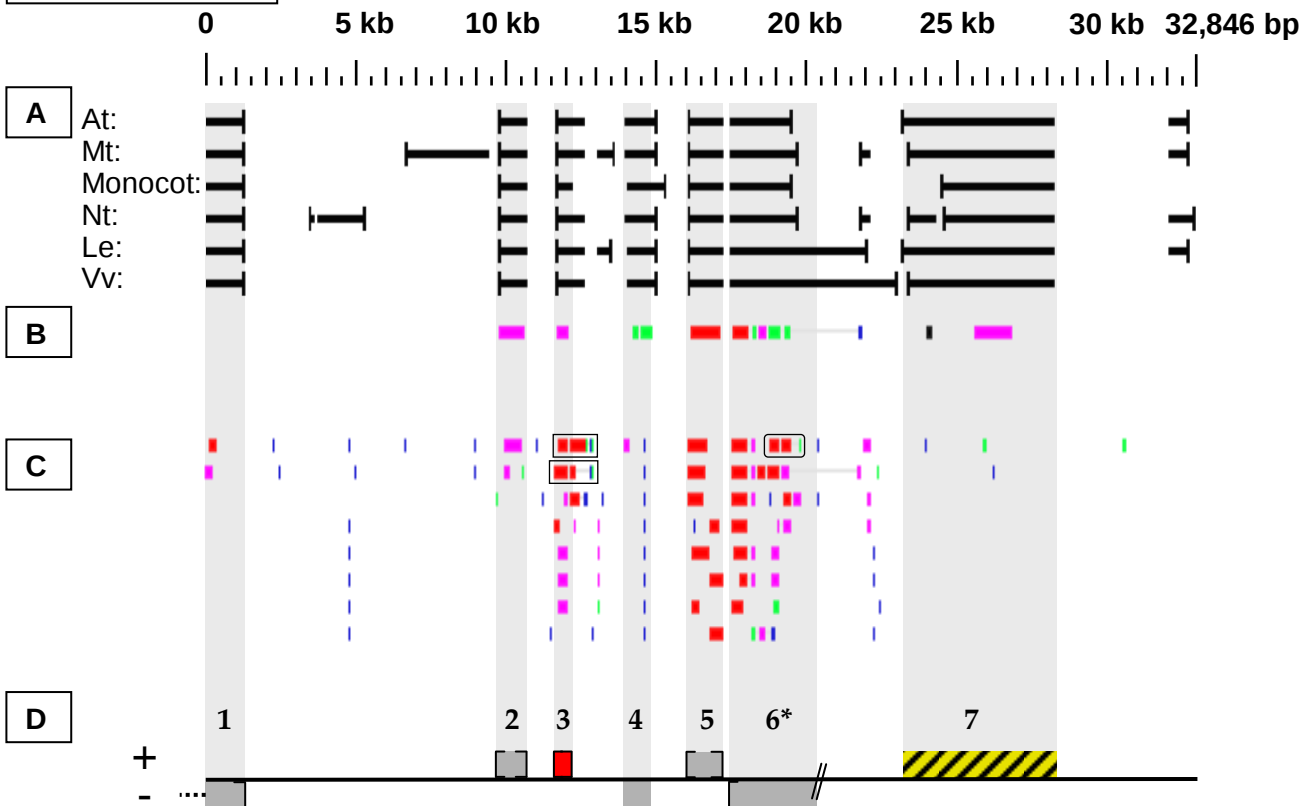


Fosmid 53J04



E

Motif	AC						GT				TA	AT	GAA			TC				
Repeat No.	6						5				7	11	9			6				
SSR start	2135						15390				22220	22694	24011			31388				

Fosmid 53J04= EU024864 (TPS) 32,846 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	1188	excluded	unknown (EST = BHLH)	5	Plus	15964	17100	PPR336 (PENTATRICOPEPTIDE REPEAT 336)
2	Plus	9652	10620	unknown protein	6*	Minus	nd	17404	terpene synthase/cyclase family protein
3	Plus	11570	12148	unknown protein	CONTIG BREAK		20424	20523	
4	Minus	14822(?)	14036(?)	DYT1 (DYSFUNCTIONAL TAPETUM 1); DNA binding / transcription factor	7	Plus	nd	nd	gypsy/Ty-3 retroelement polyprotein