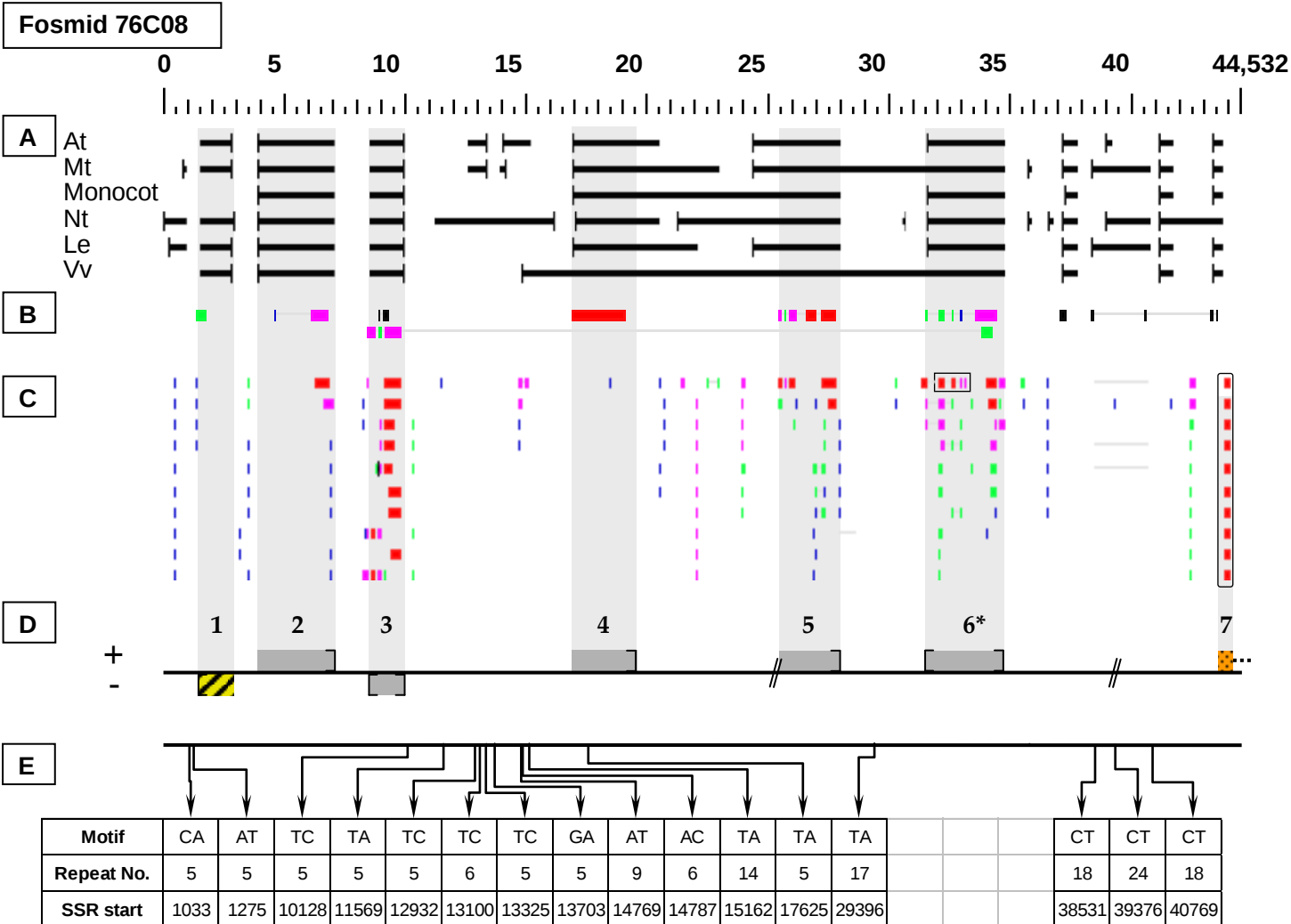




Fosmid 76C08 = EU024870 (RAN) 44,732 bp									
Putative gene location					Putative gene location				
Feature	Strand	Putative start	Putative stop	Putative identity	Feature	Strand	Putative start	Putative stop	Putative identity
1	Minus	?	1497	reverse transcriptase-related	5	Plus	nd(in gap)	27913	glycosyl hydrolase family 3 protein
2	Plus	3915(?)	7046	DNA binding	6*	Plus	31543	34734	GL3 (GLABRA 3); transcription factor (bhlh 1)
3	Minus	9900	8537	BHLH093 (BETA HLH PROTEIN 93); DNA binding / transcription factor	CONTIG BREAK		39276	39375	
4	Plus	16941(?)	19547	FRS5 (FAR1-RELATED SEQUENCE 5); zinc ion binding	7	Plus	Excluded	Excluded	glyceraldehyde-3-phosphate dehydrogenase C subunit (GAPC-2)
CONTIG BREAK		25261	25360						