

Figure 3. Shown the position (8675... 8715) of the first long repeat was not found in an expressed region in the genome.

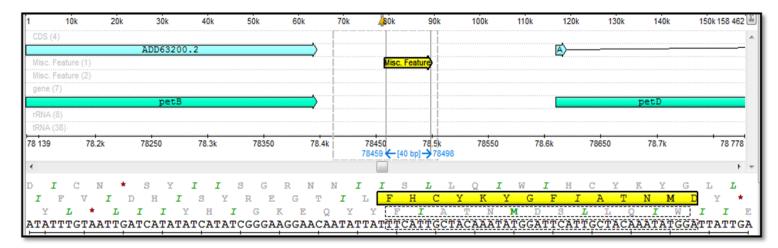


Figure 4. Shown the second long repeat as a Misc. Feature where the position (78458...78498).

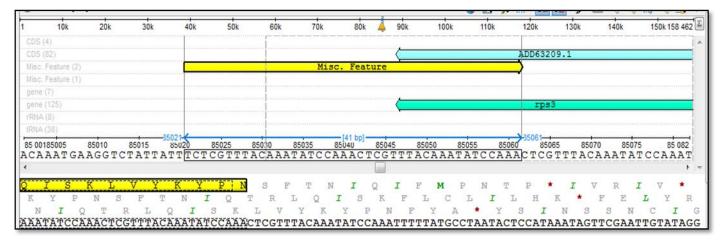


Figure 5. Shown the third long repeat as a Misc. feature where the position was from (85021...85061).

1	10k	-	Ók	30k	40k	50k	60k	70k	80k	À	90k	100k	110k	120	k 13		140k	150k 158 46	62 ^[2]
t one	S (4)								í (*
CDS	S (82)													ADD632	09.1				
Mis	c. Feature	(2)						Mi	sc. Fe	ture	<u>)</u>								
Mis	c. Feature	(1)																	
gen	e (7)																		
gen	e (125)													rps	3				
rRN	A (8)																		
tRN	A (38)													1					
84 87	78 84.9		4920	84940	84960	84980	85k	85020	85040 85046	85 [17 bp]	♦85062	35080	85.1k	85120	85140	85160			
•																			<u>•</u>
Q	I S	K	L	V Y	ΚY	P N	S F	TN	I	Q	I F	M P	N	т р	* 1	v	R I	⊽ ★	
K	Y	P N	S	F T	N I	Q T	RI	S Q	I S	K	FI	C	l I	LH	I K	* F	E	LY	R
1	I	Q !	r R	L	Q I	SK	ьv	YK	Y I	? N	F	Y A	* Y	s	I N	S S	5 N	C I	G
AA	ATATC	CAAA	CTCG	TTTAC	AAATAT	CCAAAC	TCGTT	TACAA	TATC	CAAA	TTTTT	ATGCC	TAATA	CTCC	TAAAT	AGTT	CGAAT	TGTATA	GG
85 0	31	85040	8504	45 8505	0 85055	85060	85065	85070	85075	85080	85085	85090	85095	85.1	85105	8511	0 8511	5 85 1	123

Figure 6. Shown the position of the third repeat was located from (85046... 85062) and located as a shared part in the gene rps3.