## **Tandem Repeats Search Result**

An example of the search results for the micro-satellite preset:

	A.					2
1m	2m 🔺	3m 4m	5m	6m	7m	8 234 322
iisc_RNA (1)			•			
cRNA (2)			•••••••••••••••••••••••••••••••••••••••			
RNA (12)						
epeat_region (1)						
ebeat_region E3)	1 I I III <mark>i</mark> i I					
<u>ለል አ</u> t_አniአ(ዮጵ) ለእ. አ አለአ እ		** * *** * ***				
RNA (1)						
NA (87)		<u>n n</u>				× =
nRNA (1)						
1m	2m	3m 4m	5m	6m	7m	8 234 322
	<u> </u>					
IL* YCK	v * s c n r		EREI	SL P NHC	S S H R H	P E Q R
Y C K D I V S	V * S C N R V I E IGTAATCGA	SHQR AT <u>RE</u>		N H C I I A		PE QR IRE
Y C K D I V S	VIE	S H Q R A T R E AGCCACC <mark>ACAC</mark> AC	E R E I R E K	N H C I I A	H R H I V	PE QR IRE
YCK DIVS GATATTGTAAG 451 641 2451650 CTATAACATTC2	V I E IGTAATCGA 245166 ACATTAGCT	S H Q R A T R E AGCCACCAGAGAG 0 2451670 TCGGTGGTCTCTC	E R E I R E K AGAGAGAAAA 2451680 TCTCTCTTT	N H C I I A ATCATTGCC 2451690	H R H I V ATCGTCA	P E Q R T R E CCAGAGA 2451706 GGTCTCT
YCK DIVS SATATTGTAAG 451 641 2451650 CTATAACATTCA YQLI	V I E FGTAATCGA 245166 ACATTAGCT H L R	S H Q R A T R E AGCCACCAGAGAG 0 2451670 TCGGTGGTCTCTC L W W L S	E R E I R E K AGAGAGAAAA 2451680 TCTCTCTTT L S F	NHC IIA ATCATTGCC 2451690 TAGTAACGG *QW	H R H I V ATCGTCA TAGCAGT	P E Q R T R E CCAGAGA 2451706 GGTCTCT W L
Y C K D I V S GATATTGTAAG (151 641 2451650 CTATAACATTC) Y Q L H I N Y T	V I E FGTAATCGA 245166 ACATTAGCT H L R Y D F	S H Q R A T R E AGCCACCAGAGAG 0 2451670 TCGGTGGTCTCTC L W W L S G G S L	E R E M R E K AGAGAGAAAA 2451680 TCTCTCTTT L S F S L F	N H C I I A ATCATTGCC 2451690 FAGTAACGG * Q W D N G	H R H I V ATCGTCA TAGCAGT R * D D	P E Q R T R E CCAGAGA 2451706 GGTCTCT W L G S F
YCK DIVS GATATTGTAAG 451 641 2451650 CTATAACATTCA YQLI	V I E FGTAATCGA 245166 ACATTAGCT H L R	S H Q R A T R E AGCCACCAGAGAG 0 2451670 TCGGTGGTCTCTC L W W L S G G S L	E R E M R E K AGAGAGAAAA 2451680 TCTCTCTTT L S F S L F	NHC IIA ATCATTGCC 2451690 TAGTAACGG *QW	H R H I V ATCGTCA TAGCAGT	P E Q R I R E CCAGAGA 2451706 GGTCTCT W L G S F L S
Y C K D I V S GATATTGTAAG 451 641 2451650 CTATAACATTCI Y Q L H I N Y T	V I E FGTAATCGA 245166 ACATTAGCT H L R Y D F	S H Q R A T R E AGCCACCAGAGAG 0 2451670 TCGGTGGTCTCTC L W W L S G G S L	E R E M R E K AGAGAGAAAA 2451680 TCTCTCTTT L S F S L F	N H C I I A ATCATTGCC 2451690 FAGTAACGG * Q W D N G	H R H I V ATCGTCA TAGCAGT R * D D	P E Q R T R E CCAGAGA 2451706 GGTCTCT W L G S F
Y C K D I V S GATATTGTAAG 451 641 2451650 CTATAACATTCA Y Q L H I N Y T S I T L	V I E FGTAATCGA 245166 ACATTAGCT H L R Y D F	S H Q R A T R E AGCCACCAGAGAG 0 2451670 TCGGTGGTCTCTC L W W L S G G S L	E R E M R E K AGAGAGAAAA 2451680 TCTCTCTTT L S F S L F	N H C I I A ATCATTGCC 2451690 FAGTAACGG * Q W D N G	H R H I V ATCGTCA TAGCAGT R * D D	P E Q R I R E CCAGAGA 2451706 GGTCTCT W L G S F L S
Y C K D I V S GATATTGTAAG 451 641 2451650 CTATAACATTCA Y Q L H I N Y T S I T L	V I E TGTAATCGA 245166 ACATTAGCT H L R Y D F T I S Iue	S H Q R A T R E AGCCACCAGAGAG 0 2451670 TCGGTGGTCTCTC L W W L S G G S L	E R E I R E K ACACACAAAA 2451680 TCTCTCTTT L S F S L F L S F	N H C I I A ATCATTGCC 2451690 TAGTAACGG * Q W D N G I M A	H R H I V ATCGTCA TAGCAGT R * D D M T V	P E Q R T R E CCAGAGA 2451706 GGTCTCT W L G S F L S
YCK DIVS SATATTGTAAG 451 641 2451650 CTATAACATTCA YQLI INYT SITL Va	V I E CGTAATCGA 245166 CATTAGCT I L R Y D F T I S Iue n(7704727770472	S H Q R A T R E AGCCACCACACACAG 0 2451670 TCGGTGGTCTCTC L W W L S G G S L A V L S	E R E I <u>R E K</u> <u>ACACAC</u> AAA2 <u>2451680</u> <u>TCTCTCTTTT</u> <u>L S F</u> <u>S L F</u> <u>L S F</u> <u>S S F</u> <u>S S F</u> <u>3</u> 97704729,7704730.	N H C I I A ATCATTGCC 2451690 TAGTAACGG * Q W D N G I M A 7704730,7704731	H R H <i>I</i> V <i>A</i> TCGTCA <i>T</i> AGCAGT <i>R</i> * <i>D D</i> <i>M T V</i> 7704731,77043	P E Q R T R E CCAGAGA 2 451 706 GGTCTCT W L G S F L S
Y C K D I V S SATATTGTAAG 451 641 2451650 CTATAACATTCZ Y Q L H I N Y T S I T L Va I repeat_unit joi	V I E CGTAATCGA 245166 CATTAGCT I L R Y D F T I S Iue n(7704727770472 n(1209587120958	S H Q R A T R E AGCCACCACACACAG 0 2451670 TCGGTGGTCTCTC L W W L S G G S L A V L S	E R E I R E K ACACACAAAA 2451680 TCTCTCTTTT L S F S L F L S F 	N H C I I A ATCATTGCC 2451690 TAGTAACGG * Q W D N G I M A 7704730,7704731 1209594,1209595	H R H <i>I</i> V <i>A</i> TCGTCA <i>T</i> AGCAGT <i>R</i> * <i>D D</i> <i>M T V</i> 7704731,7704 1209596,1209	P E Q R T R E CCAGAGA <u>2 451 706</u> GGTCTCT W L GS F L S 7327704732,777 5971209598)
Y C K D I V S SATATTGTAAG 451 641 2451650 CTATAACATTCZ Y Q L H I N Y T S I T L Va I T L Va I repeat_unit joi repeat_unit joi	V I E CGTAATCGA 245166 ACATTAGCT I L R Y D F T I S Ilue n(7704727770472 n(1209587120958 n(245166924516) n(6149028614902	S H Q R A T R E AGCCACCACAAGAG 0 2451670 TCGGTGGTCTCTC L W W L S G G S L A V L S 27,77047287704728,770472 38,12095891209590,120959	E R E I R E K ACACACACAAAA 2451680 TCTCTCTTTT L S F S L F L S F 97704729,7704730. 011209592,1209593 732451674,2451675 326149033,6149034	N H C I I A ATCATTGCC 2451690 FAGTAACGG * Q W D N G I M A 7704730,7704731 1209594,1209595 2451676,245167 6149035,6149036	H R H I V ATCGTCA TAGCAGT R * D D 0 M T V 7704731,7704 1209596,1209 72451678,2451 56149037,6149	P E Q R T R E CCAGAGA GGTCTCT W L G S F L S 7327704732,77 5971209598) 5792451680) 0386149039)

The tandem repeats annotations are located side by side.