

**BLAST**® » **blastn suite-2sequences** » results for RID-PYT25JWP114

Job Title [8ZG\\_F07\\_52-PCDH\\_Z1...](#)  
 RID [PYT25JWP114](#) Search expires on 09-05 15:14 pm  
 Program Blast 2 sequences  
 Query ID Icl|Query\_45975 (dna)  
 Query Descr [8ZG\\_F07\\_52-PCDH\\_Z1...](#)  
 Query Length 791  
 Subject ID Icl|Query\_45977 (dna)  
 Subject Descr [None...](#)  
 Subject Length 513

**Descriptions**

Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession
None provided	948	948	64%	0.0	100.00%	Query_45977

**Graphic Summary**



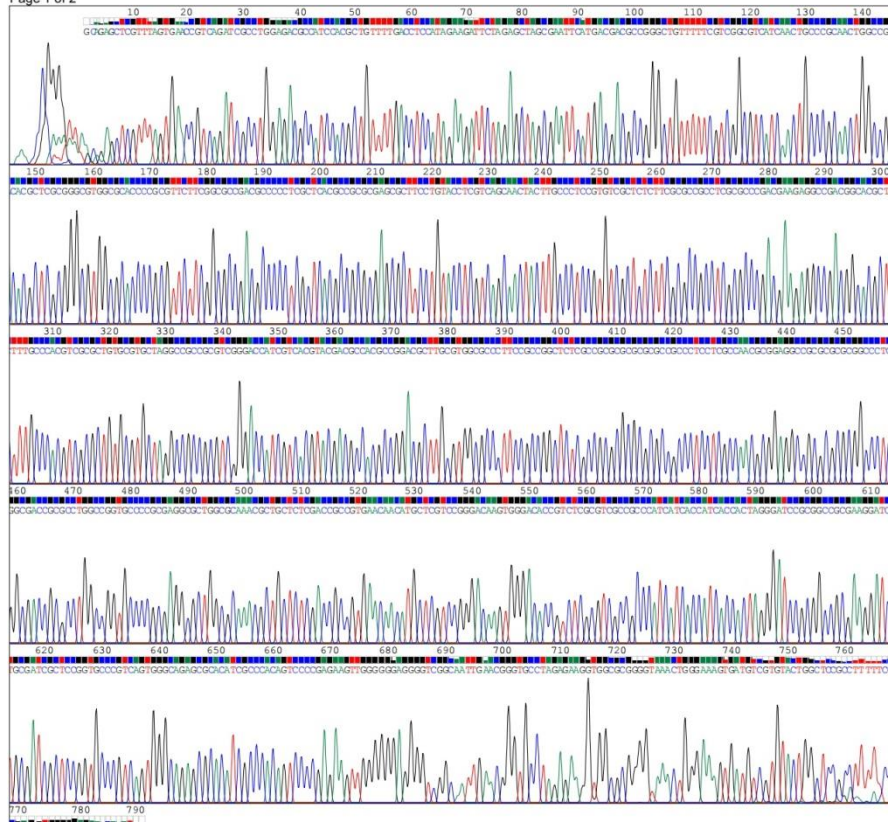
**Alignments**

Alignment view   CDS feature

Sequence ID: Query\_45977 Length: 513 Number of Matches: 1  
 Range 1: 1 to 513

Score	Expect	Identities	Gaps	Strand	Frame
948 bits(513)	0.0()	513/513(100%)	0/513(0%)	Plus/Plus	
Query 85	GAATTCATGACGACGCCGGGCTGTTTTTCGTGGCGTCATCAACTGCCCGCAACTGGCCG	144			
Sbjct 1	GAATTCATGACGACGCCGGGCTGTTTTTCGTGGCGTCATCAACTGCCCGCAACTGGCCG	60			
Query 145	ACACGCTCGGGGCGTGGCGCACCCCGCGTTCTTCGGCGCCGACGCCCTCGCTCACGC	204			
Sbjct 61	ACACGCTCGGGGCGTGGCGCACCCCGCGTTCTTCGGCGCCGACGCCCTCGCTCACGC	120			
Query 205	CGCGGAGGCGTTCCTGTACTCGTCAGCAACTACTTGCCTCCGTGTCGCTCTCTTCGC	264			
Sbjct 121	CGCGGAGGCGTTCCTGTACTCGTCAGCAACTACTTGCCTCCGTGTCGCTCTCTTCGC	180			
Query 265	GCCGCTCGGCCGACGAAAGAGGCCGACGGCAGCGTCTTTGCCACGTCGCGCTGTGCG	324			
Sbjct 181	GCCGCTCGGCCGACGAAAGAGGCCGACGGCAGCGTCTTTGCCACGTCGCGCTGTGCG	240			
Query 325	TGCTAGGCCCGCGCTCGGGACATCGTCACGTACGACGACGCGGACGCTTGCCTGG	384			
Sbjct 241	TGCTAGGCCCGCGCTCGGGACATCGTCACGTACGACGACGCGGACGCTTGCCTGG	300			
Query 385	cgcccttcgcccgcctctcgccgcbgcbgcbgcccctcctcgccaacgcbgagccg	444			
Sbjct 301	CGCCCTTCGCCGGCTCTCGCGCGCGCGCGCCCTCTCTCGCAACGCGGAGGCCG	360			
Query 445	cgcbgcbgcccctcgccgaccgcbctgcccggcgcccgcbgagccgctggcgcAAAGCG	504			
Sbjct 361	CGCGCGCGCCCTCGGCGACCGCGCTGGCCGGTGCCTCCGCGAGGCGTGGCGAAACGC	420			
Query 505	TGCTCTGACCGCGTGAACAACATGCTCGTCCGGGACAAGTGGGACACCGTCTCGCGTC	564			
Sbjct 421	TGCTCTGACCGCGTGAACAACATGCTCGTCCGGGACAAGTGGGACACCGTCTCGCGTC	480			
Query 565	GCCGCCATCATCACCATCACCACTAGGGATCC	597			
Sbjct 481	GCCGCCATCATCACCATCACCACTAGGGATCC	513			

**Appendix 1.** Sequencing result. As shown in the picture, there was 100% similarity to the expected amplicon.



**Appendix 2.** Sequencing chromatogram