

Evidence of Complement Receptor Genes in Invertebrates (Echinodermata)

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ABSTRACT

Recently CR receptor gene was discovered in the Asterial : Asterias rubens (Echinodermata). Other Echinodermata show in their genomes the CR2 gene (CR2 Complement C3d receptor 2) i-e Ophuirids (Ophiocomina nigra) and Crinoïds (Antedon bifida) The sequences of their transcriptomes are presented.

INTRODUCTION

Recently, CR receptor gene was discovered in an Invertebrate : the sea star Asterias rubens(Asterids,Echinodermata) (Ref : 1). It seems clear to us, to look for such genes in other Echinodermata : Ophiocomina nigra (Ophuirids), Antedon bifida (Crinoïds) which possess, in their genomes, the well-known IGKappa gene (corresponding to the Invertebrate Primitive Antibody (Ref. 2, 3,)

The Complement C3b/C4b receptor 1 like and the CR2 Complement C3d receptor 2 genes were studied in O.nigra and A.bifida genomes

MATERIALS AND METHODS

a) Animals: Ophiocomina nigra, Antedon bifida were purchased from the Marine Laboratory of Roscoff (France)

b) Obtention of ophuirid and crinoïd mRNA:

Digestive coeca were excised from the animal's bodies. O.nigra, A bifida mRNA were obtained from Uptizol (Interchim).

Quality control were operated.

c) **Sequencing:** Sequencing was made on Illumina Next Seq 500 with paired-end : 2. 75 bp

Transcriptome was assembled from RNA-Seq fastq files using Trinity v2.1.1 (Ref.4) with default parameters. A BLAST database was created with the assembled transcripts using makeblastdb application from ncbi-blast+ (v2.2.31+). The sequences of transcripts of interest were then blasted against this database using blastn application from ncbi-blast+ (Ref;5) with parameter word_size 7.

RESULTS

We just find, in Ophiocomina nigra and Antedon bifida CR2 complement C3d receptor 2 gene.Complement C3b/C4 b receptor 1 like gene is not present significantly.

First we present the characteristics of CR2 transcriptome in Antedon bifida:

QueryID	Query name	Subject ID	Identity (%)	Length	Mismatch	Gapopen	Query	E-	Bitscore
							cover	value	
							(%)		
NM_001006658.2		TRINITY_DN16	83,33	48	4	3	1,00	2,80E-	41,70
		054_c3_g1_i1						02	
The sequence o	CT	CTATTATAGGAATATTCAGTAATCATACCT							
>TRINITY_DN		TC AAAAAATTAATCATGTATCATAAAGTTATT							
5GTCCACTATTAATTTGTTACAAAACACTA				GGTACTGTAATGCAAGTAGAATAGTAATAG					
ATTACGAATGTCAACAAGTCGGATATCAT				ATAAGAAGTCTTCCATTGGCAACCACTGCT					
TT				TTTAAATAGATTTTATTGTAAATAAAAAAT					
ATTTTCTACTAAACTGAAATACTTTACTTTC				ACTAAAGCAAAAAAAAAAAAAAAAAAAAAAA					
AGTCTAGATAGCCTAAACCCAAACTCGAT				AACAGAA3'					
TCAACATTTAATATTTTTAGAGATTAATATA ATCTCAGATGAAGTAGTAAACTAGTAAAC ATTTAAAAATAAGCGCAAAGTGAAACTT				We repertoriate the Ophiocomina nigra CR2 transcriptome sequence to compare :					

>TRINITY_DN65134_c0_g1_i1 (CR2)

5AAAACAGGCAAAAATGCTCTTTAGGAA AACACAAACGCGTCTCCGGACTCTCCGC GTGCT

AAAAAAAAAGACCACTCAAAAACCAAGA CAGAATGACGAAGACCACA**3**'

CONCLUSION

Blasts against human were performed. They lead us to envisage as highly true the evidence of CR and CR2 genes in Invertebrates (Echinodermata).

CR2 gene, in human, encodes a membrane protein which function as a receptor for Epstein-Barr virus binding on B and T lymphocytes.

Echinodermata lymphocytes exist , do they possess such a receptor? That is the question !Further studies are necessary to clarify this problem.

When we compare the sequences of CR2 Antedon bifida and Ophiocomina nigra transcriptomes we observe slight differences in 5'-3', between them. They are due to alignment's differences. Nevertheless, it is obvious, that CR2 gene is present in these two Echinodermata.

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