

Open Access

Brief Report Article

Volume 1 – Issue 1

E IG V-L K appa Expression from Sea Star Igkappa Gene

Michel Leclerc*

Immunology of Invertebrates, Biology/Biochemistry, Orleans University, France

*Corresponding author: Michel Leclerc, Immunology of Invertebrates, Biology/Biochemistry, Orleans University, France

Received date: 22 September, 2021 |Accepted date: 6 October, 2021 |Published date: 9 October, 2021Citation: Leclerc M (2021) E IG V-L K appa Expression from Sea Star Igkappa Gene. J Clin Class Immunol 1(1). doihttps://doi.org/10.54289/JCCI2100103

Copyright: © 2021 Leclerc M. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Abstract

The sea star IGKappa gene was cloned in 2014 by the use of primers. It was compared in the present work to Vertebrate Immunoglobulin genes. A high identity was found with these last ones. A length of 105 amino acids fit with Immunoglobulin domain.

Introduction:

The sequence of the sea star Asterias rubens IGKappa gene was described by our team, in 2014 [1]. Since we have tried to find homologies between this gene and upper genes from lower Vertebrates to human genes.

We report, in the precedent paper, results obtained with upper Vertebrate genes by the use of blasts directed against these last ones [2, 3].

Results:

The sequence of the sea star IGKappa gene is the following [1]:

<u>5'GGATCC</u>GGAGGA**ATG**CGTGGCAACATGGCGTCTC TATGGATGTTCTTCTTTGTCGTGGGGATAACTTTAC AACGGAGTTTGGCGATTTACACGTTTCGCAGCAAC CGTCGGACACTAGCGCGTTGCAGGGGAGCACAGTG GTGCTTCACTGCTCCGTTGAGCAGTACATAAACACC ACGGCCATCGTTTGGTGGAGCCGTGACTCGGTCAT CAGCCACAACAAAGACCTGAAACTGTCCAGTCTAA ACACCGACCAGCTCCAAAGGTACTCGATTTCAGGC GACGCATCTCGGGGGGGAATTCAACCTTAAAATAGT GAACTTTACCGCCACAGACGCCGCCAGTTACCGCT GTCAGATG **TAA**GAATTC**3**'

The bioinformatic work leads us to show similarities between sea star IGKappa gene and Immunoglobulin domain from Vertebrates



Non-specific hits: IgV_L_Kappa

[Non-specific hit, evalue = 6.79e-03] cd04980, Immunoglobulin (Ig) light chain, kappa type, variable (V) domain; The members here are composed of the immunoglobulin (Ig) light chain, kappa type, variable (V) domain. This group contains the standard Ig superfamily V-set AGFCC'C"/DEB domain topology.]

Super-families: Ig superfamily



[Superfamily, evalue = 6.79e-03] cl11960, Immunoglobulin domain; The members here are composed of the immunoglobulin (Ig) domain found in the Ig superfamily.]

The **Table 1**, as shown below, resumes our results: We observe again the Immunoglobulin domain and a particular one without immune function.

</td <td>50</td> <td>100</td> <td>150</td> <td>200</td> <td>250</td> <td>300</td> <td>350</td> <td>400</td> <td>450</td> <td>500</td> <td>550</td> <td>600</td> <td>650</td> <td>700</td> <td>750</td> <td>800</td> <td> 850</td> <td>93</td>	50	100	150	200	250	300	350	400	450	500	550	600	650	700	750	800	850	93
(고 옷 xp_0	33635901.1 -	Find:		▼ < < >		• • • •	AT6 素								🔀 Tools 🗸	🖨 Tracks 🗸	📥 Download 🗸	æ?•
<u>1</u>	50	100	150	200	250	300	350	400	450	500	550	600	650	700	750	800	850	932
Protein F	eatures																	TOX
uncharacterized protein LOC117296905										1.0.0								
region re	atures - Cl	טנ																3 Q X
IG														C065644			- 22	
site Feat	ures - CDD								14									10 X
1	50	100	150	200	250	300	350	400	450	500	550	600	650	700	750	800	850	
XP_033635901.1: 1.932 (932 aa)										own: 4/4								

Two region features:

		b. COG5644	
a.	Region Ig	Comment: U3 small nucleolar RNA-ass	ociated
	Comment: Immunoglobulin	protein 14 (function unknown)	
	Location: 47151	L section: 721 866	
	Longth: 105 an	Location: /51800	
	Lengui. 105 aa	Length: 136 aa	
	CDD: 214652	CDD: 227931	

Table 1: PREDICTED: Asterias rubens uncharacterized LOC117296905 (LOC117296905)

Conclusion:

We retain from this bioinformatic analysis, the presence of Immunoglobulin domain in the sea star IGKappa gene with the CDD:214652. This gene, nevertheless, seems less evolved that the Ophuirid IGKappa gene we discovered 1 month ago [4] in terms of Immune functions.

These 2 genes from Echinodermata (Invertebrates) bring us a new light in Immunogenetic World and mainly in Comparative Immunology between Invertebrates and Vertebrates animals.

References:

1. Vincent N, Osteras M, Otten P, Leclerc M (2014) A new gene in A. rubens: A sea star Ig kappa gene. Meta Gene 2:320-322.

2. Marchler-Bauer A, Bo Y, Han L, He J, et al. (2017) CDD/SPARCLE: functional classification of proteins via subfamily domain architectures. Nucleic Acid Res 45 (D1): 200-203.

3. Marchler-Bauer A, Lu S, Anderson JB, Chitsaz F, Derbyshire MK, et al. (2011) CDD: a Conserved Domain Database for the functional annotation of proteins. Nucleic Acid Res 39(D): 225-229.

4. Leclerc M. (2021) Biosynthesis « De Novo » of the Ophuirid Ophiocomina Nigra Igkappa Gene. J Clin Class Immunol 1(1).