

Evidence of Conformity between the Ophiocomina Nigra HLA-E Transcriptome and the O.Nigra HLA-E Gene in Invertebrates

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Received Date: 14 January 2022; Published Date: 25 February 2022

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Citation: Michel Leclerc. Evidence of Conformity between the Ophiocomina Nigra HLA-E Transcriptome and the O.Nigra HLA-E Gene in Invertebrates. J Clin Case Rep On. 2022;2(1):1012.

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Abstract

This work allows us to corroborate the existence of MHC genes in Invertebrates, and particularly HLA-E one, in its own right. Identities at 100% were discovered between HLA-E synthesized gene and original HLA-E Transcriptome

Introduction

MHC genes of Class I and Class II were discovered in 2018 in Echinodermata (Invertebrates) [1-3]. Since we have performed many works in bioinformatics, especially on HLA-E gene which belongs to class I in Human We verify in the present work the authenticity of the O.Nigra HLA-E gene.

Methods

HLA-E from Ophiocomina Nigra Transcriptome was synthesized « de novo » into a pUC-GW-Kan plasmid and amplified in our laboratory. Blasts were performed in a second step [3].

Comparisons with the original O.Nigra HLA-E Transcriptome was then done

Results

Query: None Query ID: lcl|Query_40765 Length: 281>

Sequence ID: Query_40767 Length: 934

Range 1: 103 to 383

Score: 520 bits (281), Expect: 7e-152,

Identities: 281/281(100%), Gaps: 0/281(0%), Strand: Plus/Plus

Query 1 TGTAATCCCAGCACTTTGGGAGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGAC 60

|||||

Sbjct 103 TGTAATCCCAGCACTTTGGGAGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGAC 162

Query 61 CATCCTGGCTAACACAGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCCGGGCG 120

|||||

Sbjct 163 CATCCTGGCTAACACAGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCCGGGCG 222

Query 121 TGGTGGCGGGCGCCTGTAGTCCAGCTACTCGGGAGGCTGAGGCAGGAGAATGGCGTGAA 180

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|||||
Sbjct 223 TGGTGGCGGGCGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGGAGAATGGCGTGAA 282
Query 181 CCCGGGAGGCGGAGCTTGCAGTGAGCCGAGATCGCGCCACTGCACTCCAGCCTGGGCGAC 240
|||||
Sbjct 283 CCCGGGAGGCGGAGCTTGCAGTGAGCCGAGATCGCGCCACTGCACTCCAGCCTGGGCGAC 342
Query 241 AGAGCGAGACTCTGTCTCaaaaaaaaaaaaaaaaaaaaa 281
|||||
Sbjct 343 AGAGCGAGACTCTGTCTCAAAAAAAAAAAAAAAAAAAAAA 383
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Conclusion

Similarities, identities at 100 % occur in the sequences given in results. That confirms the existence of HLA-E gene in *Ophiocoma Nigra* genome, so in Echinodermata which are the only Phylum in Invertebrates to present such data.

References

1. Leclerc M. *Proteomics Bioinformatics*. 2022;2(1):59-61
2. Leclerc M. *MHC genes in Echinodermata Lambert editions*. 2019.
3. Leclerc M. *J. Clin Class. Immunol*. 2022;1:1-4.