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**Short Commentary** 

# **Ophuirid Ophiocomina Nigra HLA-E Gene Synthesis in PUC-GW-KAN** Plasmid or HLA-E Echinodermata Gene Biosynthesis «De Novo» in E. Coli Sensu Lato Plasmid

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## ABSTRACT

HLA-E (Class 1) is an MHC gene which has been isolated in 2020, in our laboratory. We show now its biosynthesis «de novo» in a PUC-GW-KAN plasmid. Such experiment was performed with the Ophiocomina nigra IGKappa gene one year ago.

#### **INTRODUCTION**

We have isolated recently MHC genes in Echinodermata [1] in 3 classes: the Ophuirids, the Crinoïds, the Asterids. At that time, we decided to synthetize one of these genes: the well-known HLA-E one in a PUC-GW-KAN plasmid (Yan Li gift).

#### **METHODS**

We operate according the following method [2]. It was resumed in 4 parts:

- Synthesis of oligonucleotides with overlapping 1. segments in sense and antisense direction
- 2. Assembly of the oligonucleotides into a double stranded DNA, using a poly chain assembly method (PCA).
- 3. For larger constructs, the sequence is split into smaller, intermediate fragments, to facilitate synthesis. Once the intermediated fragments have been obtained with correct sequence, they are assembled into the full-length sequence.
- 4. Cloning into the linearized vector by either recombination or ligation-based cloning, mostly performed within the same step as full-length sequence assembly.

Regarding the restriction site, which was used for cloning, construct was cloned into vector pUC-GW by using the unique EcoRV restriction site. Please find table below for the primers used for sequencing (Table 1).

Table 1. Primers used for sequencing.

M13F-77	GATGTGCTGCAAGGCGATTA
M13R-88	TTATGCTTCCGGCTCGTATG
U-SEQ4883	CCTCCAATCGGGTAACTC

#### RESULTS

#### 1) Plasmid map

The construct appears below (Figure 1):

2) Recalling of Original sequence in 5'-3':

TGTAATCCCAGCACTTTGGGAGGCCGAGGCGGGCG GATCACGAGGTCAGGAGATCGAGACCATCCTGGCT AACACAGTGAAACCCCGTCTCTACTAAAAATACAA AAAATTAGCCGGGCGTGGTGGCGGGGCGCCTGTAGT CCCAGCTACTCGGGAGGCTGAGGCAGGAGAATGGC GTGAACCCGGGAGGCGGAGCTTGCAGTGAGCCGAG ATCGCGCCACTGCACTCCAGCCTGGGCGACAGAGC AA

3) Synthetized sequence in 5'-3':

TGTAATCCCAGCACTTTGGGAGGCCGAGGCGGGCG GATCACGAGGTCAGGAGATCGAGACCATCCTGGCT AACACAGTGAAACCCCGTCTCTACTAAAAATACAA AAAATTAGCCGGGCGTGGTGGCGGGGGCGCCTGTAGT CCCAGCTACTCGGGAGGCTGAGGCAGGAGAATGGC GTGAACCCGGGAGGCGGAGCTTGCAGTGAGCCGAG ATCGCGCCACTGCACTCCAGCCTGGGCGACAGAGC GAGACTCTGTCTCAAAAAAAAAAAAAAAAAAAAAAAA AA

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## 4) Blastn original sequence/ synthetized sequence

The **Table 2** resumes mainly the identities and the e-value between these 2 precedent sequences. Chromatograms were also performed.

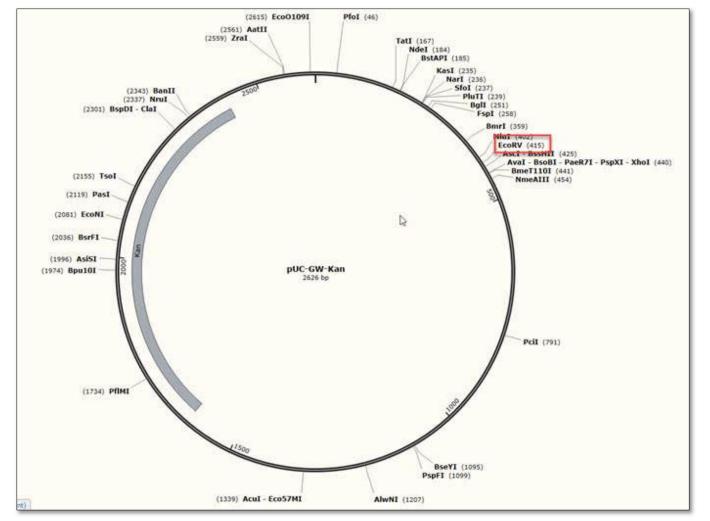


Figure 1. Plasmid map.

**Table 2.** Comparisons between original sequence and synthetized one.

Size Seq1	Size Seq2	Max score	Total score	Query cover	E. Value	Per. Ident	Acc Len
281	281	520	520	100%	7e-152	100%	934

## CONCLUSION

We conclude our experiment is valid when compared to **Table 2**. Furthermore, we assert, it is the first time such discovery:

- a) MHC Genes in Echinodermata (Invertebrates) were found
- b) biosynthesis of HLA-E Echinodermata gene in a PUC-GW-KAN plasmid was performed.

## REFERENCES

1. Leclerc M (2020) Evidence of MHC Class I and Class II

Genes in Echinodermata. Proteomics Bioinformatics 2(1): 59-61.

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