

International Journal of Scholarly Research in Science and Technology

Journal homepage: https://srrjournals.com/ijsrst/ ISSN: 2961-3337 (Online)



(Review Article)

Check for updates

# What computer programs can be used to detect a viral pathogen?

Carlos Navarro Venegas \*

Department of Preventive Animal, Faculty of Veterinary and Livestock Sciences (FAVET). University of Chile, Chile.

International Journal of Scholarly Research in Science and Technology, 2023, 02(01), 010–012

Publication history: Received on 20 January 2023; revised on 10 March 2023; accepted on 13 March 2023

Article DOI: https://doi.org/10.56781/ijsrst.2023.2.1.0017

## Abstract

Current virology uses biotools from some internet platforms that are still free of charge. This has allowed virology -like other medical disciplines- to take a quantum leap towards the detection and diagnosis of viral pathogens in conjunction with the brilliant idea developed by Kary Mullis.

Not forgetting André Lwoff: *Viruses are viruses*, the detection of a virus does not differ even if it affects humans or another species: One Health.

Keywords: Viral detection; Biotools; On line softwares; Primer design

## 1 Introduction

A particularity of viruses is their genome (RNA or DNA), which differs in size, length, or nucleotide sequence. Currently, these sequences have been published by several authors and are stored in a database: the Genbank® [1]. This, along with various computer programs, allows both the determination of the identity of a suspected agent and the design of primers for a polymerase chain reaction (PCR) devised by Kary Mullis. Other methodologies involving platforms such as Clustal Omega and BLAST should not be left out of mention.

Thus, there are no pretexts for detecting a pathogen whose genome is DNA or RNA.

Finally, both teachers and students can access these internet platforms and enter the fantastic world of molecular virology...!!!!

## 2 Material and methods

A virology laboratory requires cell culture, however, the virus isolation method is tedious and time consuming. Due to the above, several methodologies have been developed that, by involving the brilliant idea of Kary Mullis [2], called Polymerase Chain Reaction in conjunction with 2% agarose gel electrophoresis, allows the observation of the amplification of a DNA fragment (conventional PCR).

Subsequently, the nucleotide sequence of the amplified fragment can be known and, thanks to the use of free access online programs such as CLUSTAL Omega [3] and BLAST [4], the identity of the suspected agent can finally be corroborated.

If, when consulting the existing scientific literature, there are no specific primers for a PCR reaction, they can be designed thanks to the Invitrogen ® Oligoperfect program or another similar one. This is why both A.Lwoff's statement

Copyright © 2023 Author(s) retain the copyright of this article. This article is published under the terms of the Creative Commons Attribution Liscense 4.0.

<sup>\*</sup> Corresponding author: Carlos Navarro Venegas; Emailcanavarr@uchile.cl

and the fact that there is no pretext for proposing a pathogen detection protocol is corroborated. In relation to the necessary samples, these can originate from organs or fluids to extract the nucleic acid

## 3 Discussion

Although the gold standard in virology is the isolation of the viral agent, the indicated methodologies allow the detection of the viral agent in less time. Scientific innovations or brilliant ideas such as Kary Mullis's are combined today with biotools that are still freely accessible online such as Clustal, BLAST and others that have been used in publications in mainstream journals [5, 6, 7, 8] as well as in others that have exalted the molecular study of pathogens of veterinary interest [9-17].

# 4 Conclusion

Teaching also does science. In our country and thanks to ideas that involve the detection of viral pathogens, at least 35 title memories have been developed that involve same number of new veterinarians. These former students today could perfectly develop the detection and diagnosis of a viral pathogen.

## **Compliance with ethical standards**

## Acknowledgments

We thank the students of FAVET for making their dreams come true with us and Dr. Aron Mosnaim, from the Wolf Foundation, Illinois, USA (since 2020).

## References

- [1] Genbank®. NIH genetic sequence database. 2023. Available from: https://www.ncbi.nlm.nih.gov/genbank/
- [2] Mullis, K; Faloona, F. 1987.Specific synthesis of DNA in vitro via a polymerase-catalyzed chain reaction. Methods in Enzymology, 155: 335-350.
- [3] CLUSTAL OMEGA. Multiple Sequence Alignment. 2023. Available from: https://www.ebi.ac.uk/Tools/msa/clustalo/
- [4] BLAST. Basic Local Alignment Search Tool. 2023. Available from: https://blast.ncbi.nlm.nih.gov/Blast.cgi
- [5] Vergara-Wilson, V., Hidalgo-Hermoso, H., Sánchez, C., Abarca, M., Navarro, C., Celis-Diez, S., Soto-Guerrero, P., Diaz-Ayala, N., Zordan, M., Cifuentes-Ramos, F., Cabello-Stom, J. 2021. Canine Distemper Outbreak by Natural Infection in a Group of Vaccinated Maned Wolves in Captivity. Pathogens. Available from: https://doi.org/10.3390/pathogens10010051
- [6] Hidalgo-Hermoso, E., Cabello, J., Vega, C., Kroeger-Gómez, H., Moreira-Arce, D., Napolitano, C., Navarro, C., Sacristán, I., Cevidanes, A., Di Cataldo, S., Dubovi, E., Mathieu-Benson, C., Millán, J. 2020. An eight-year survey for canine distemper virus indicates lack of exposure in the endangered Darwin's fox (Lycalopex fulvipes). J Wild Dis. Available from: https://pubmed.ncbi.nlm.nih.gov/31833816/
- [7] Sepúlveda-Estay, C., Jara-Osorio, MA, Navarro, C. 2019. Molecular detection of glycoprotein B from feline herpesvirus. Vaccimonitor, 28 3, pp. 103-109.
- [8] Rodríguez Edisleidy, Betancourt A, Barrera Maritza, Lee Changee, Yoo Dongwan. 2008. Detección Rápida Del Virus De La Gastroenteritis Transmisible En Cerdo A Través De La Reacción En Cadena De La Polimerasa. Rev Salud Anim. Available from: http://scielo.sld.cu/scielo.php?script=sci\_arttext&pid=S0253-570X2008000200011&lng=es.
- [9] Jara, P., Céspedes, P., Navarro, C. 2018. Canine Distemper Virus detection based in Hemaglutinine gen as target in reverse transcriptase-Polymerase Chain reaction. IVS. Available from: https://www.heighpubs.org/hvsr/pdf/ivs-aid1012.pdf
- [10] Salas, V; Pizarro, J; Navarro C. 2018. Phylogenetic analysis of canine distemper virus detected in Chile. IJCR 10: 72402-72407

- [11] Méndez-Valenzuela, VK., Jara, M.A., Navarro, C. 2021. Canine Distemper Virus: Multiple detection of the H and N genes by the Polymerase Chain Reaction associated with Reverse Transcriptase. CJVDS. Available from: https://www.corpuspublishers.com/assets/articles/article-pdf-158.pdf
- [12] Vera, C, Jara, MA, Navarro, C. 2022. A preliminary studio for the F gen of Canine Distemper Virus as target for phylogenetic analysis. GSCARR. Available from: https://gsconlinepress.com/journals/gscarr/sites/default/files/GSCARR-2022-0034.pdf
- [13] Navarro, C., Muñoz, C, Céspedes, P. 2019. The Nucleocapside Protein Gene As Excellent Target For Detection Of Canine Distemper Virus by Reverse Transcriptase-Polymerase Chain Reaction. Am J Biomed Sci & Res. Available from: https://biomedgrid.com/pdf/AJBSR.MS.ID.000935.pdf
- [14] Mateo, F., Céspedes, PF., Navarro, C. 2019. The Phosphoprotein Gene from Canine Distemper Virus as Target in Viral Detection. IJZAB. Available from: https://medwinpublishers.com/IZAB/the-phosphoprotein-gene-from-canine-distemper-virus-as-target-in-viral-detection.pdf
- [15] Bolívar, P., Céspedes, P., Navarro, C. 2019. Use of the Reverse Transcription-Polymerase Chain reaction for differential detection of two lineages of the canine distemper virus in Chile. IVS. Available from: https://www.heighpubs.org/hvsr/pdf/ivs-aid1014.pdf
- [16] Abarca, MJ., Hidalgo, E., Raggi, LA., Navarro, C. 2018. Genotypic evidence of infection by Canine Distemper Virus in maned Wolf from a zoological collection. IJSR. Available from: https://www.ijser.org/researchpaper/Genotypic-evidence-of-infection-by-Canine-Distemper-Virus-in-manedwolf-from-a-zoological-collection-in-Chile.pdf
- [17] Pincheira, D., Céspedes, P., Pizarro, J., Navarro, C. 2018. Molecular detection of Canine Distemper Virus through the use of the Large Polymerase Gene. EASJALS. Available from: https://www.easpublisher.com/media/articles/EASJALS\_12\_39-43\_c.pdf