

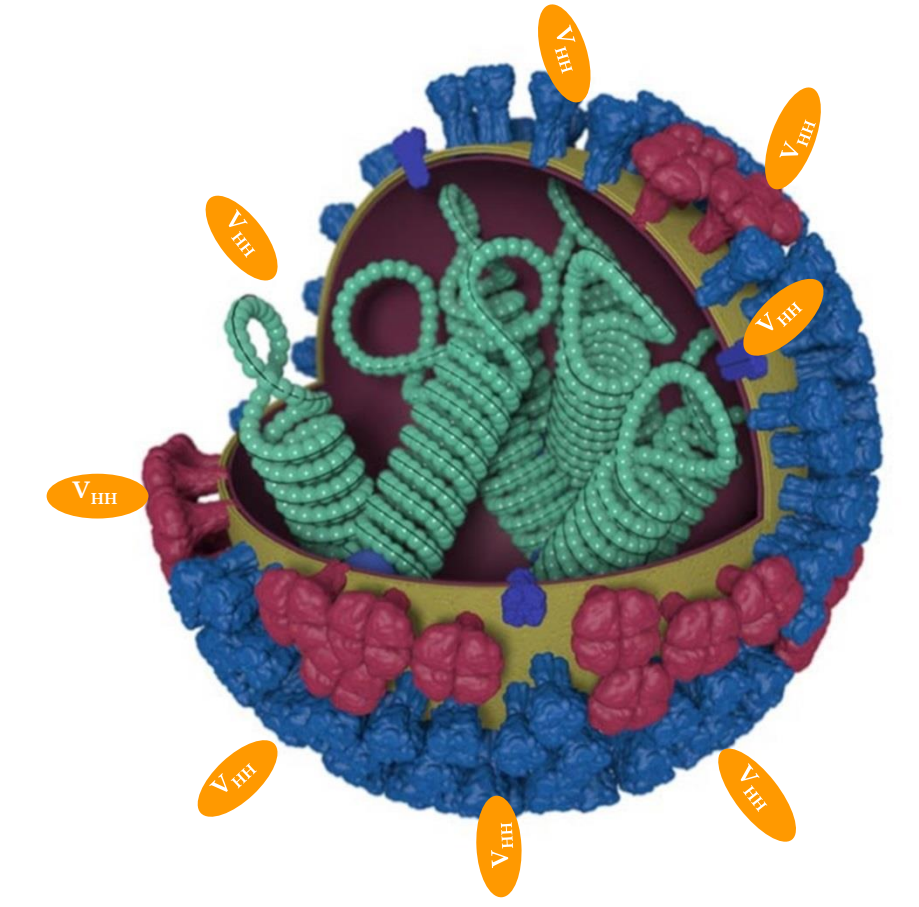
DEVELOPMENT OF A THERAPY BASED ON NANOANTIBODIES AGAINST THE INFLUENZA A VIRUS

Ángel Linde-Rodríguez^(1,2), Ana Soriano-Lerma^(1,4), Victoria Sánchez-Martín^(1,2,3), Matilde Ortiz-González^(1,5), Virginia Pérez-Carrasco^(1,2), Miguel Soriano^(1,5), José Antonio García-Salcedo^(1,2).

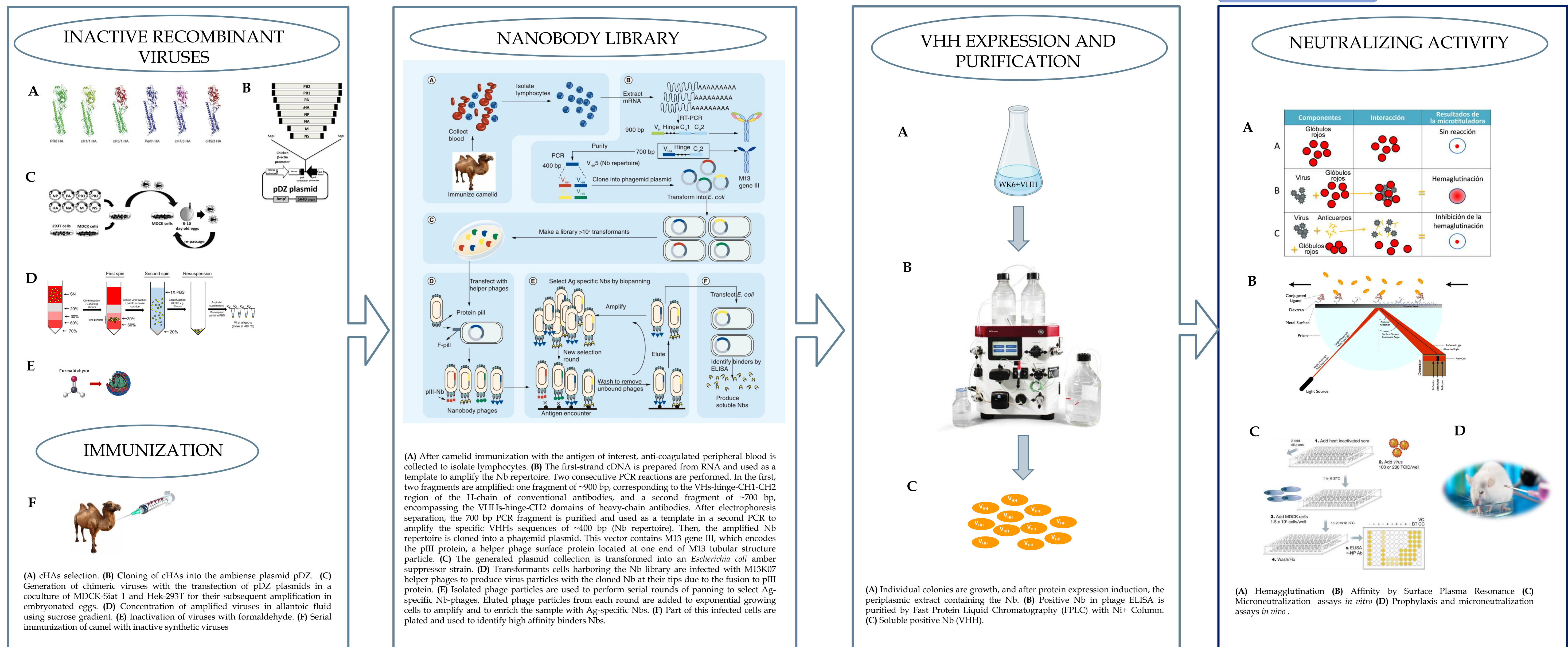
- (1) GENYO. Centre for Genomics and Oncological Research: Pfizer/University of Granada/Andalusian Regional Government, Granada, 18016, Spain.
 (2) Microbiology Unit, Biosanitary Research Institute IBS-Granada, University Hospital Virgen de las Nieves, Granada, 18014, Spain.
 (3) Department of Biochemistry, Molecular Biology III and Immunology, University of Granada, Granada, 18016, Spain.
 (4) Department of Physiology, University of Granada, Granada, 18011, Spain.
 (5) Center for Intensive Mediterranean Agrosystems and Agri-food Biotechnology (CIAMBITAL), University of Almeria, Almeria, 04001, Spain.

ABSTRACT

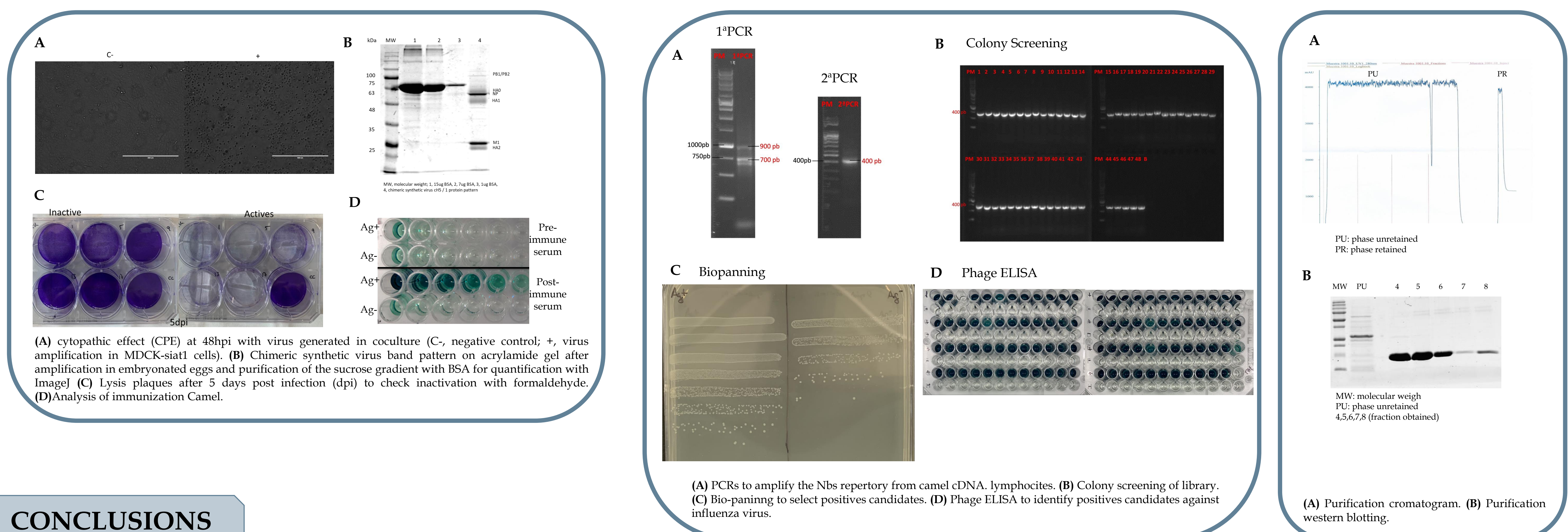
Influenza viruses are enveloped, single-stranded, negative-sense RNA viruses that belong to the family Orthomyxoviridae. Three types of Influenza viruses have been described, being types A and B the main human pathogens. They cause significant morbidity and mortality worldwide, with the elderly and children being the most susceptible patients. Influenza A viruses are divided into subtypes according to genetic and antigenic differences in the two main surface spike proteins, hemagglutinin (HA) and neuraminidase (NA). Current vaccines are an effective measure against infection, but must be reformulated yearly due to the antigenic drift of the virus. Furthermore, they do not protect against new pandemic strains. Currently, only two classes of drugs are approved for treatment: M2 ion channel blockers and NA inhibitors, but they have limited efficacy, adverse side effects, and great drug resistance. Therefore, there is a need to develop new antiviral strategies with new modes of action and reduced resistance. Nanobodies are small single domain antibody fragments that have unique properties such as the ability to recognize particular epitopes (not recognized by conventional antibodies) and their improved stability. They have been used successfully in therapeutic approaches as receptor blockers associated with pathogenic diseases or viruses, or conjugated with drug carriers. The main objective of this work was the generation of broad-spectrum neutralizing nanobodies against conserved Influenza A epitopes. Currently, the strains with the highest infection rate are H1N1 and H3N2. Therefore, a new approach of sequential vaccination has been applied in camels using synthetic viruses expressing chimeric HA molecules (cHA), which combine stem-conserved domains of group 1 (H1) or 2 (H3) with unusual domains in the head. A phage display nanobody library was subsequently generated and candidates were selected after successive rounds of biopanning. The neutralizing activity of the selected candidates against Influenza A virus is currently being analysed.



MATERIALS AND METHODS



RESULTS



CONCLUSIONS

- Different nanobodies were positively identified against *Influenza A*. The nanobodies were expressed and purified.
- Subsequent *in vitro* and *in vivo* studies will be performed to analyze the neutralizing activity.

ACKNOWLEDGMENTS

Financial support by Ministry of Economy and Competitiveness (MINECO) cofinanced with FEDER funds: *NANOGRIP (SAF2015-71714-R).