ROPÓN-PALACIOS G. ("Geordano")

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RESEARCH ASSISTANT

Computational Biophysics, Bioinformatics and Drug Design Lab Facultad de Ciencias de La Salud, Instituto de Ciencias Biomédicas Universidad Autónoma de Chine, Llano Subercaseaux 2081 - piso 6, San Miguel, Santiago, Chile.

2. Research interests : Membrane proteins, viroporins, all-atom and Coarse grained MD simulation of virus, large-scale simulation of supramolecular complexs, theoretical biophysics, method development in computational biophysics, drug and vaccine development, scientific programming and AI.

3. Formation

(a) Biosciencist,

School of Biological Sciences, Universidad Nacional de la Amazonía Peruana, Iquitos, Loreto, Perú. 2011-2015, *gradute with Cum laude mention*. Thesis: " *Potential antigens of diagnostic interest in Tritrichomonas foetus secretome*" classified as very good. Defense date: 30/01/2018. Advisor: Prof. Dr. Jose Luis Marapara del Águila

4. Skills and Abilities

- (a) English: Intermediate
- (b) Operating systems: Mac OSX and Linux.
- (c) Programming lenguages: C/C++, Tcl, Python, Bash, AWK.
- (d) **Computational packages:** VMD, AMBER, CHARMM, NAMD, Gromacs, Autodock4, Autodock-vina, Prody, PDBFixer, Modeller, Swiss-model, Pymol, Gaussian, Mopac, Gamess, Matplotlib.
- (e) **Computational biophysics methods:** Umbrella sampling, Steered Molecular dynamics, Gaussian accelerated Molecular dynamics, Accelerated Molecular dynamics, Free energy perturbation, MM/PBSA, Non-equilibrium dynamics, DF/HF quantum mechanics methods.
- (f) **Molecular Biology techiques:** DNA/RNA extraction, Electrophoresis, Western Blot, RNA seq, Immunofuoresence, Membrane reconstitution, current-clamp.

5. Research positions

- (a) Computational Biophysics, Bioinformatics and Drug Design Lab. Facultad de Ciencias de La Salud, Instituto de Ciencias Biomédicas, Universidad Autónoma de Chine, Llano Subercaseaux 2081 - piso 6, San Miguel, Santiago, Chile. Leader group: Prof. Dr. David Ramírez. 2020-Actually.
- (b) Matter condensed physics Lab, and Computational modeling and simulation Lab, Department of Physics, Institute of Exact Sciencies, Universidad Federal de Alfenas, Minas Gerais, Brasil. Leader group: Prof. Dr. Ihosvany Camps. **2019-Actually.**

6. Scientific meetings (14)

- Ropón, G.G., Ochavano, J.R., Machoa, J. 2014. Role of the pigs of the community of Santo Tomas as host of Balantidum coli. V International Course of Tropical and Infectious Disease, Iquitos, Peru.
- 2. Ropón-Palacios G., Acurio J., Salguero N., Human-Sutta V., Rodríguez-Lima I. Immunoinformatics prediction epitopes B and T of gp63 protein from Leishmania braziliensis: reveals epitopes conserved with others Leishmania spp and allow the design

- of multi-epitope protein as potential vaccine. Encuentro Científico Internacional de Invierno, Perú. 2018.
- 3. Ropón-Palacios G., Acurio J., Salguero N., Human-Sutta V., Rodríguez I., Quispe-Mamani R., Mormontoy S. *In silico-based secretory proteins prediction of Tritrichomonas foetus reveals potential vaccine candidates: An integrative bioinformatics approach.* Encuentro Científico Internacional de Invierno, Perú, 2018.
- 4. Salguero N.#, Ropón-Palacios G.#, Acurio J.#, et al. Molecular dynamics simulation in subunit vaccine design: A case of study. Encuentro Científico Internacional de Invierno, Perú, 2018. # igual contribución.
- 5. Ropón-Palacios G., Olivos-Ramírez G., Otazu-Mamani K., & Torres-Castillo O. Box size in molecular docking assays by genetic algorithms is very important to replicate the calculation of free energies (ΔG). Encuentro Científico Internacional de Verano, Perú, 2019.
- 6. Ropón-Palacios G., Otazu-Mamani K., Olivos-Ramírez G. & Torres-Castillo O. *Novel potencial inhibitors against Nipah Virus: A computational biology approach.* Encuentro Científico Internacional de Verano, Perú, 2019.
- 7. Ropón-Palacios G., Ruiz Mesia W., Otazu-Mamani K., Olivos-Ramírez G., Torres-Castillo O., Puma-Zamora W., Arias Calle N. Discovery of novel pan-inhibitors against wildtype and mutated Cytb protein from Plasmodium falciparum through of computational simulation. Encuentro Científico Internacional de Verano, Perú, 2019.
- 8. Olivos-Ramírez G., Ropón-Palacios G., Otazu-Mamani K., Chenet-Zuta M. 2019. Análisis de dinámica molecular de 5'-hidroxiestroptimicina de Spirulina platensis como inhibidor de dihidrofolato reductasa de Staphylococcus aureus: una vista atómica de reconocimiento molecular. En libro de Resúmenes de la V Jornada Internacional en Biociencias 2019.
- 9. Otazu-Mamani K., Olivos-Ramírez G., <u>Ropón-Palacios G.</u>, 2019. *An virtual screening of Allium sativum metabolites against wild-type and mutated dihydrofolate reductase protein from Staphylococcus aureus*. XX Congreso Nacional de Estudiantes de Biología, Lambaque, Perú, 2019.
- 10. Ramirez-Díaz Y., Sancho-Queque C., Osorio-Mogollon Cl., Otazu-Mamani K., Olivos-Ramírez g., <u>Ropón-Palacios G.</u> 2019. *Drugability of homologous enzymes in helminths through biomolecular modelling: The case of Ancylostoma duodenale and Necator americanus*. XX Congreso Nacional de Estudiantes de Biología, Lambague, Perú, 2019.
- 11. Osorio-Mogollón C., <u>Ropón-Palacios G.</u>, Baqui M., In silico structural characterization and molecular docking of FAZ10 in Trypanosoma brucei. X International Conference on Bioinformatics SolBio+10, Uruguay, 2019.
- 12. Bonacina J., Carbajal M. Ropón-Palacios G., Cantero M de R., Cantiello H. *The protein ftsZ prokaryotic ancestro of the tubulin, form sheets bidimensionals that behaves like electric oscillators.* III Jornadas de Microbiología sobre Temáticas Específicas del NOA Microbiología agrícola y Ambiental.

National-

- 13. Vargas-Santillan, J. A., Orbe, B., Olortegui, E. A., Rios, G., Ropón, G. G. 2014. *Identification of culicidae potentials in the transmission of Encephalitis equine Venezuelan Virus, Iquitos, Peru, 2013.* XV National Congresses of Biology Students, Ricardo Palma University, Peru.
- 14. Olivos-Ramírez G., Ropón-Palacios G., Otazu-Mamani K., Merino F. 5'-hidroxi-streptomicina de Spirulina platensis potencial inhibidor contra Dihidrofolato reductasa de Staphylococcus aureus: Un estudio de docking molecular. Semana de la Ciencia, Universidad Nacional de Santa, Chimbote, Perú, 2018.

7. Papers

- 1. Ropon-Palacios G., y Chenet-Zuta M.E, Olivos-Ramirez G.E., Otazu K., Acurio-Saavedra J., Camps I. 2019. Potential novel inhibitors against emerging zoonotic pathogen Nipah virus: virtual screening and molecular dynamics approach. Journal of Biomolecular Structure and Dynamics, DOI: 10.1080/07391102.2019.1655480
- 2. Ropon-Palacios G., y Chenet-Zuta M.E., Olivos-Ramirez G.E., Otazu-Mamani, K., Camps I. 2019. Novel Multi-epitope protein containg conserverd epitopes from different Leishmania species as potential vaccine candidate: Integrated immunoinformatics and molecular dynamics approach. Journal of computational biology and chemistry journal 2019. DOI: https://doi.org/10.1016/j.compbiolchem.2019.107157.