JUAN SEBASTIÁN IGNACIO CONTRERAS RIQUELME

+569 7919 7035 contrerasriquelme.sebastian@gmail.com

Bioinformatic engineer and Biotechnology Ph. D student interested in biological networks and data science for genomic/epigenomic data.

ACADEMIC DETAILS

2016 – 2022	PhD Student at Universidad Andres Bello, Chile
2008 – 2014	Bioinformatic Engineer at Universidad de Talca,
	Chile.

EXPERIENCE DETAILS

03/2015 – 12/2017	Fundación Ciencia & Vida. Computational Biology Lab (Dlab). As Research assistant in project FONDECYT INICIO 11140342, granted by FONDECYT.
01/2018 – 04/2022	Universidad Mayor, Centro de Genómica y Bioinformática. As Research assistant in project FONDECYT REGULAR 1181089, granted by FONDECYT.

04/2022 - Present Universidad Andres Bello, Centro de

Biotecnología Vegetal. Postdoctoral research in project FONDECYT postdoctoral 3220673,

granted by FONDECYT.

Bibliography

1725P Metastasis-resident bacteria in advanced hormone receptor-positive breast cancer are related to primary tumor microbiota and show distinct composition.

Araya, C., <u>Contreras-Riquelme, S.</u>, Mino, B., Perez, F. J., Martin, A. J., & Carvajal-Hausdorf, D. E. (2022). doi: 10.1016/j.annonc.2022.07.1803

Homology-based reconstruction of regulatory networks for bacterial and archaeal genomes.

Romero, L., <u>Contreras-Riquelme</u>, S., Lira, M., Martin, A. J. M., & Perez-Rueda, E. (2022). doi: 10.3389/fmicb.2022.923105

Automated generation of context-specific gene regulatory networks with a weighted approach in Drosophila melanogaster.

Murgas L, <u>Contreras-Riquelme S</u>, Martínes-Hernandez J.E, Villaman C, Santibañez R, Martin AJ (2021) doi: 10.1098/rsfs.2020.0076

RIP-MD: A tool to study Residue Interaction Networks in Protein Molecular Dynamics Contreras-Riquelme S, Garate JA, Perez-Acle T, Martin AJ. (2018) doi: 10.7717/peerj.5998

Transcriptional profiling of embryos lacking the lipoprotein receptor SR-B1 reveals a regulatory circuit governing a neurodevelopmental or metabolic decision during neural tube closure

Santander N, Lizama C, Murgas L, Contreras S, Martin AJ, Molina P, Quiroz A, Rivera K, Salas-Pérez F, Godoy A, Rigotti A, Busso D. (2018). doi:10.1186/s12864-018-5110-2

LoTo: a graphlet based method for the comparison of local topology between gene regulatory networks

Martin AJ, <u>Contreras-Riquelme S</u>, Dominguez C, Perez-Acle T. (2017). doi: 10.7717/peerj.3052

Graphlet Based Metrics for the Comparison of Gene Regulatory Networks

Martin AJM, Dominguez C, <u>Contreras-Riquelme</u> S, Holmes DS, Perez-Acle T (2016). doi:10.1371/journal.pone.0163497

Similarities between the Binding Sites of SB-206553 at Serotonin Type 2 and Alpha7 Acetylcholine Nicotinic Receptors: Rationale for Its Polypharmacological Profile.

Möller-Acuña P, <u>Contreras-Riquelme JS</u>, Rojas-Fuentes C, Nuñez-Vivanco G, Alzate-Morales J, Iturriaga-Vásquez P, . (2015). doi:10.1371/journal.pone.0134444

SEMINARS AND CONGRESS

Integrating Histone Marks, DNA accessibility, and Motif data to predict regulatory maps in Arabidopsis thaliana

Contreras-Riquelme JS, Alvarez JM.

XV REUNIÓN DE BIOLOGÍA VEGETAL. Dec 2022, Coquimbo, Chile

Automated generation of condition specific Gene Regulatory Networks from epigenetic and genomic data integration

Murgas L, Contreras-Riquelme S, Martin AJ.

Latin American Student Council Symposium 2018, Viña del Mar, Chile

A network based method to analyze non-covalent interactions in proteins

<u>J. Sebastián Contreras-Riquelme</u>, A.J.M. Martin, J.A. Garate, I. Fuenzalida Tomas Perez-Acle

IV MEETING MILLENNIUM INSTITUTE CINV "Neuroscience make sense", Jul. 2015. Valparaíso – Chile. doi: 10.7490/f1000research.1110200.

Polypharmacology of CNS drugs and binding site similarities: a computational study of drugs acting simultaneously at serotonergic and nicotinic acetylcholine receptors.

Reyes-Parada, Miguel., Moller, Patricia., <u>Contreras, Juan.</u>, Rojas, Cecilia., Alzate-Morales, Jans., Nuñez, Gabriel., Iturriaga-Vásquez, Patricio.

Determination of putative binding sites on human α 7 nicotinic acetylcholine receptor to positive allosteric modulators type I and II through computational studies.

Contreras J, Möller P, Reyes M, Rojas C, Alzate J

XII PABMB Congress together with the XLIX Annual Meeting of Sociedad Argentina de Investigación Bioquímica y Biología Molecular (SAIB), the XXXVI Annual Meeting of Sociedad de Bioquímica y Biología Molecular de Chile (SBBM), the 4th Meeting of the Latin-American Protein Society and the LVI Annual Meeting of Sociedad de Biología de Chile. Nov. 2013. Puerto Varas - Chile.

Binding Modes of SB-206553, a 5HT 2B/2C antagonist and a positive allosteric modulator of alpha7 nicotinic acetylcholine receptors.

Reyes-Parada M, Moller P, Contreras, J.S, Rojas, C, Alzate-Morales, J, Iturriaga-Vásquez P

Presented at:

- Congreso Iberoamericano de Biofísica IX Reunión Anual Sociedad Chilena de Neurociencia. Oct. 2013. Valparaíso - Chile.
- XXXV Reunión Anual de la Sociedad de Farmacología de Chile. Nov. 2013. Valdivia -Chile.