

NETWORK PHARMACOLOGY OF ANNONA CRASSIFLORA ALKALOIDAL FRACTION ON ALZHEIMER'S AND ITS EFFECT ON DROSOPHILA **MELANOGASTER MODEL**



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INTRODUCTION

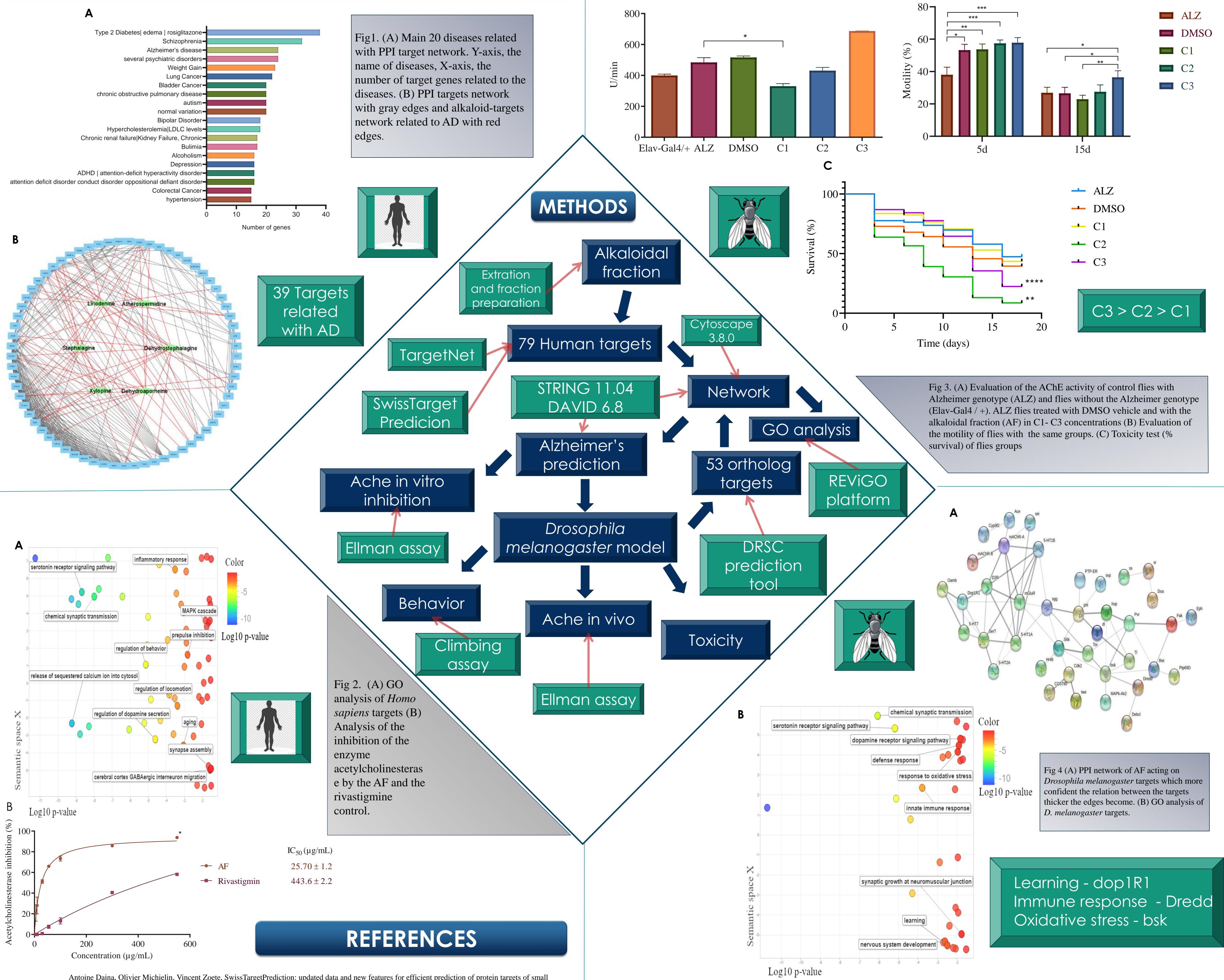
Studies from Annona crassiflora has shown the potential of its metabolites and therapeutic use. Alkaloids have been identified in the A. crassiflora fruit and peel and therefore we aim to investigate these alkaloids using a Drosphila melanogaster model of Alzeheimer disease and an approach of pharmacology in silico network to harness the full potential of the alkaloid fraction of A. crassiflora.

RESULTS AND DISCUSSION

OBJECTIVE

Thus, this work aims to search possible human targets for these alkaloids, and from the targets found evaluate the effect of the alkaloid fraction treatment on the Alzheimer model using *D. melanogaster* and predict its mechanism of action.

RESULTS AND DISCUSSION



Antoine Daina, Olivier Michielin, Vincent Zoete, SwissTargetPrediction: updated data and new features for efficient prediction of protein targets of small molecules, Nucleic Acids Research, Volume 47, Issue W1, 02 July 2019, Pages W357–W364, https://doi.org/10.1093/nar/gkz382.

B. Snel, G. Lehmann, P. Bork, M. A. Huynen, STRING: a web-server to retrieve and display the repeatedly occurring neighbourhood of a gene, Nucleic Acids Research, Volume 28, Issue 18, 15 September 2000, Pages 3442–3444, https://doi.org/10.1093/nar/28.18.3442.

Da Wei Huang, Brad T. Sherman, Qina Tan, Joseph Kir, David Liu, David Bryant, Yongjian Guo, Robert Stephens, Michael W. Baseler, H. Clifford Lane, Richard A. Lempicki, DAVID Bioinformatics Resources: expanded annotation database and novel algorithms to better extract biology from large gene lists, Nucleic Acids Research, Volume 35, Issue suppl_2, 1 July 2007, Pages W169–W175, <u>https://doi.org/10.1093/nar/gkm415</u>.

Da Wei Huang, Brad T. Sherman, Richard A. Lempicki, Bioinformatics enrichment tools: paths toward the comprehensive functional analysis of large gene lists, Nucleic Acids Research, Volume 37, Issue 1, 1 January 2009, Pages 1–13, https://doi.org/10.1093/nar/gkn923.

Huang, D., Sherman, B. & Lempicki, R. Systematic and integrative analysis of large gene lists using DAVID bioinformatics resources. Nat Protoc 4, 44–57 (2009). <u>https://doi.org/10.1038/nprot.2008.211</u>.

P. Shannon, "Cytoscape: A software environment for integrated models of biomolecular interaction networks", Genome Res., vol. 13, no. 22, pp. 2498-2504, 1971.

FINANCIAL SUPPORT



Conclusion

In conclusion, the present study predicted alkaloids-target-disease interactions and multi-target mechanisms of the A. crassiflora alkaloidal fraction in the treatment of AD using the network pharmacology strategy, as well as validating the protective effect of AF against the AD model of D. melanogaster. Further studies are also necessary in order to deepen the knowledge about the interaction pathways and mechanisms of action of the alkaloids described for AF on AD, which may lead to the development of one of AF or its alkaloids as a potential candidate for the treatment of patients with AD.