

In pursuit of Novel Class of Drug that Affects the Core Symptoms of Autism Spectrum Disease

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Abstract

The research of autism is still in progress. We had a deal with the 9 type of experiment results and gene expression data showing how much it is affected by autism. Through t-SNE and NMF, 3 high ranking result group and 3 low ranking result group had a pattern in OFT, NORT, and Social test. We decided to analyze these 3 phenotypes to find significant gene list. Those gene list is visualized in volcano plot based on p-value which is less than 0.05 and absolute value of fold change which is larger than 1 at the same time. Using these genes, we obtained drug lists of each phenotype through connectivity map and common functions through GeneMANIA. It is said that vasopressin can affect the common functions of each phenotype. As a result, we found out that FG-7142 drug related to vasopressin and benproperine through connectivity map which affects on central nerve system and has not been used as a autism drug.

Introduction

Autism is referred to as Autism Spectrum Disorder or ASD and is a developmental disability that can cause a range of mild to severe social, communication and behavioral challenges. Most people with autism look like other people but their behavior, social skills and communication are markedly different than people without the disorder.

Unfortunately, few drugs on the market today effectively relieve symptoms of autism and none of the options most often prescribed by practitioners work well for every individual. Hence, we came up with the idea that if we analyze existing drugs, thanks to recent advances in big data analysis, and mix up with the advantages of them it might be possible to find new combination treatment of autisms' core symptoms.

Connectivity map: a set of gene expression signatures for defined cellular states that would be used to link any genetic, chemical or other perturbations of gene expression to actions on normal or pathological processes.

NMF: a powerful tool for data reduction and exploration that has seen popular use in analyzing high-throughput genomic data.

OFT (DM): Distance moved

NORT: Time spent around the novel object

Social test: Time spent around the stranger chamber

Result

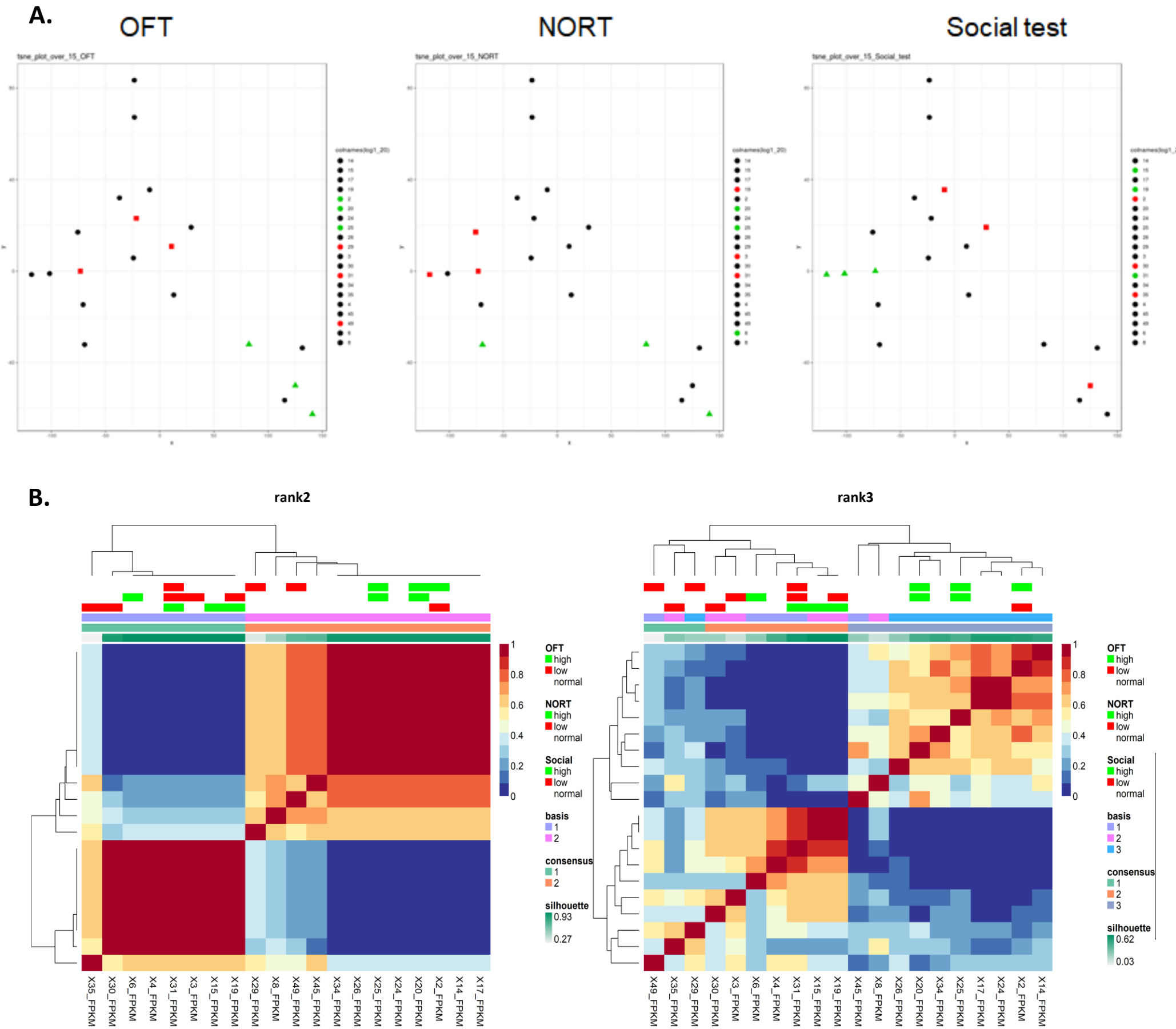
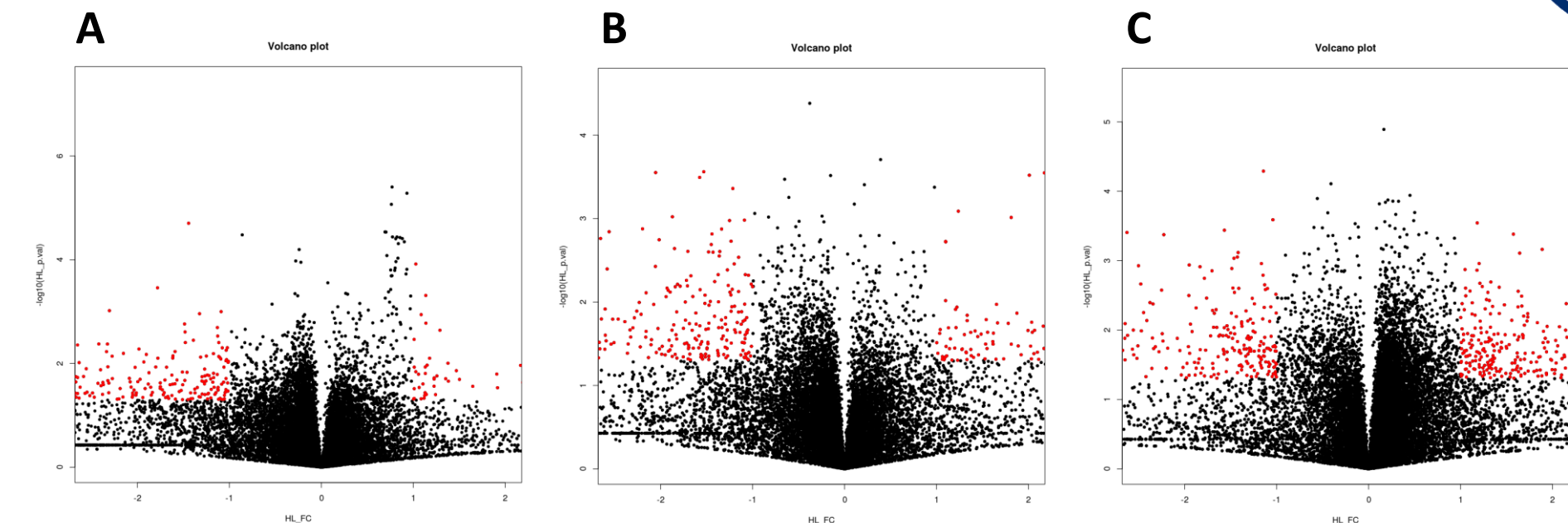


Figure 1A-B. The acquisition of the fact that OFT, NORT and Social test results from 9 phenotypes (FST, OFT, EPM, Y Maze, Social test, NORT, CA edge, RS_FST, RS_TST) are significant by t-SNE and NMF. A) t-SNE of OFT, NORT and Social test using 101 genes out of variance 15. Each phenotype, 3 dots in green indicate that it made high ranking results into top 3 out of 20 samples and as opposed to in red it made 3 low ranking results from the bottom. B) NMF of OFT, NORT and Social test using same 20 sample. It indicates 3 high ranking results are grouped together as well as 3 low ranking results. (nrun = 15)



Figures 2A-C. The performance of acquiring significant genes through volcano plots of OFT, NORT and Social test. A) The volcano plot of OFT. 344 genes are obtained. B) The volcano plot of NORT. 353 genes are obtained. C) The volcano plot of Social test. 532 genes are obtained. (Each test, all significant genes are colored in red which p-value is less than 0.05 and absolute value of fold change is larger than 1. X-axis indicates log-scaled fold change between high ranking group and low ranking group. Y-axis indicates log-scaled p-value.)

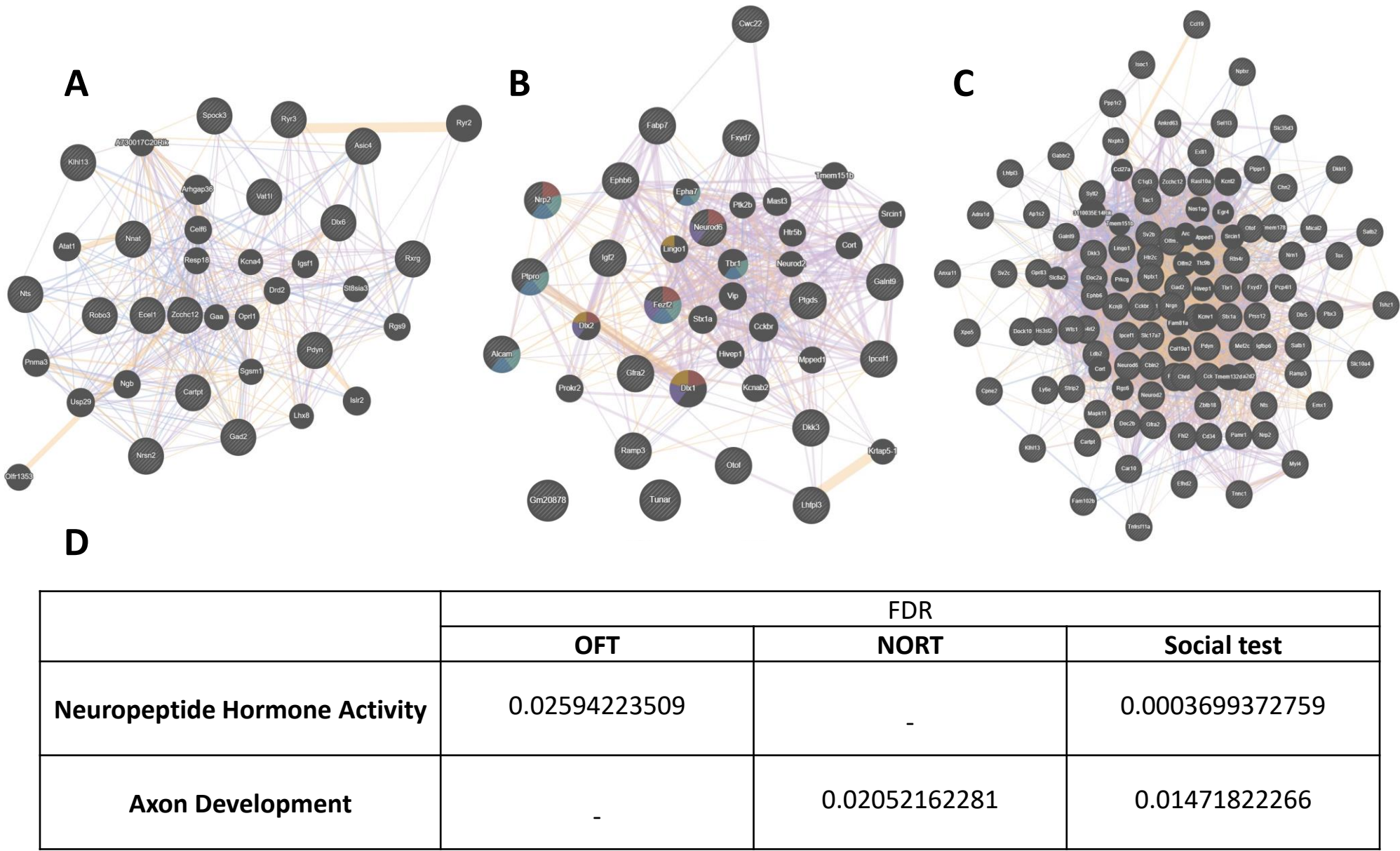


Figure 3A-D. The GeneMANIA results for generating hypotheses about functions of gene lists from volcano plot. The number of each test's gene list was reduced due to the amount of gene expression which should be larger than 10 for significance. A) The GeneMANIA result of OFT entering 17 genes. B) The GeneMANIA result of NORT entering 25 genes. C) The GeneMANIA result of Social test entering 101 genes. D) Two common functions of OFT, NORT, and Social test based on FDR are neuropeptide hormone activity and axon development.

			FDR		
			OFT	NORT	Social test
Neuropeptide Hormone Activity			0.02594223509	-	0.0003699372759
Axon Development			-	0.02052162281	0.01471822266

Figure 4A-C. The lists of potential drugs for OFT, NORT, and Social test executed from connectivity map. Each list is ranked based on the amount of gene expression difference between high ranking group and low ranking group. A) The list of potential drugs for OFT. B) The list of potential drugs for NORT. C) The list of potential drugs for Social test. (MEK inhibitor in red is common compound out of 3 test's drug list which is too broad to use. Benproperine and FG-714 in yellow are potential drug to apply to autism's symptom relief.

Conclusion

We searched high-ranked drugs out of connectivity map results and found out that drug list of OFT was too broad to apply to. In case of NORT, it is possible to use benproperine as autism drug. It is a drug acting on the central and peripheral nerve system as a double-acting release. It is on the market and approved in the Ministry of Health and Welfare as a prescription for cold, chronic bronchitis, upper respiratory infections, and pulmonary tuberculosis. We also figured out that vasopressin is related to relieving the symptom of autism by GeneMANIA result of Social test. One of the compounds affecting this neurotransmitter is GABA receptor. FG-7142 is in the drug list of Social test. FG-7142 (ZK-31906) is a drug which acts as a partial inverse agonist at the benzodiazepine allosteric site of the GABAA receptor. It has anorectic, anxiogenic and pro-convulsant effects. It also increases release of acetylcholine and noradrenaline, and improves memory retention in animal studies. In summary, it is able to suggest benproperine and FG-714 as affective drugs to relieve the core symptoms of autism spectrum disease.

Reference

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