Investigating the lipidome of Parkinson-induced zebrafish brain using NMR and neural network signal deconvolution

Thea S. Ingvaldsen¹, Diana C. Turcu¹, Sarah T. Frøystad¹, Martin Jakubec², Kari E. Fladmark¹, Øyvind Halskau¹ 1. Department of Biological Sciences, University of Bergen, Bergen, Norway 2. Luxembourg Centre for System Biomedicine (LCSB), Luxembourg

Introduction: Parkinson's disease is a neurodegenerative brain disorder and is characterized by the depletion of dopaminergic neurons in parts of the brain¹. The brain consists of approximately 50% lipids by dry weight. Autosomal, recessive point mutations and deletions in the human DJ-1 gene cause a form of familial Parkinson's disease and are also linked to inflammation, mitochondrial dysfunction, oxidative stress, and altered lipid metabolism². This project aimed to analyze lipid profiles by using NMR in combination with machine learning to deconvolute complex spectra. The lipids were extracted from both wild-type (WT) zebrafish, and from zebrafish where the DJ-/PARK71 gene is deactivated (KO) by using a modified_Blight & Dyer method. Zebrafish are used in this experiment due to their high similarity to mammals in molecular mechanisms of development and cellular physiology³.



Figure 1: Average signal volume from three WT ZF and three KO ZF of cholesterol, PC (phosphatidylcholines), PE (phosphatidylethanolamine), and sphingomyelin. Error bars represent one standard deviation.

Figure 2: Signal volume from human grey matter of one control and grey matter of one patient of cholesterol, PC (phosphatidylcholines), PE (phosphatidylethanolamine), and sphingomyelin.



5000

-5000

-10000

Volcano plot: cholesterol is above the threshold



Discussion: The results from this experiment show significant (p-value: 0.03) differences in the signal volume from cholesterol in WT ZF and KO ZF. These differences are also shown in the grey matter from human samples. The control patient has a higher signal volume from cholesterol. These results are based on one human sample. For future projects, it's important to understand whether high levels of cholesterol in the brain cause Parkinson's or if Parkinson's leads to higher levels of cholesterol.

Figure 4: Features selected by volcano plot with fold change threshold 1.5x and t-test threshold 0.05. The red circles represent features above the threshold (cholesterol).



References

Chin, H. Y., Lardelli, M., Collins-Praino, L., & Barthelson, K. (2021). Loss of park7 activity has differential effects on expression of iron responsive element (IRE) gene sets in the brain transcriptome in a zebrafish model of Parkinson's disease. Molecular brain, 14(1), 83. https://doi.org/10.1186/s13041-021-00792-9
He, Y., Kaya, I., Shariatgorji, R., Lundkvist, J., Wahlberg, L. U., Nilsson, A., Mamula, D., Kehr, J., Zareba-Paslawska, J., Biverstål, H., Chergui, K., Zhang, X., Andren, P. E., & Svenningsson, P. (2023). Prosaposin maintains lipid

homeostasis in dopamine neurons and counteracts experimental parkinsonism in rodents. Nature communications, 14(1), 5804. https://doi.org/10.1038/s41467-023-41539-5

3. Kari, G., Rodeck, U., & Dicker, A. P. (2007). Zebrafish: an emerging model system for human disease and drug discovery. Clinical pharmacology and therapeutics, 82(1), 70–80. https://doi.org/10.1038/sj.clpt.6100223 - Li, J., Vosegaard, T., & Guo, Z. (2017). Applications of nuclear magnetic resonance in lipid analyses: An emerging powerful tool for lipidomics studies. Progress in lipid research, 68, 37–56. https://doi.org/10.1016/j.plipres.2017.09.003

