Studying the Microglial Transcriptome in response to **Early Life Stress and Aging**

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INTRODUCTION

- Microglia are the immune cells of the brain, responsible for maintaining homeostasis within the central nervous system

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COMP SCI

- Early-life experiences, such as childhood abuse, neglect, or trauma, have been implicated in shaping microglial phenotype and function
- Can we use microglial genetic markers to uncover whether early life stress (ELS) ages the brain?



Aging triggers a shift toward a more immunogenic profile including an increase in inflammatory microglia¹



Early Life Stress (ELS) specifically has been connected with impacting microglial function². This was determined by exposing a group of young mice to stress via environmental conditions such as limited bedding. These mice, along with a control group, were injected either with lipopolysaccharide (LPS) or phosphate-buffered saline (PBS) to see if LPS would cause unique perturbations in the brain of mice exposed to ELS.

RESULTS



The Elastic Net model does better at predicting the true age of mice compared to the the Random Forest Regressor model. Certain clusters tend to perform better than the version where genes expressions are pooled across all clusters

Pooled Across Clusters, Random Forest Classifier (RFC)



Cluster by Cluster, Random Forest Classifier (RFC)



Feature Importance Smd3 0.066667 RFC feature importance of top 10 Apoc1 0.066667 important genes acc3 0.066667 capd2 0.061166 : Biomarker for Alzheimer's Disease Ccnb2 0.033333 : Biomarker for Cancer (if20b 0.033333 Zfp654 0.033333 Spc25 .033333 Cenpa 0.033333 Fabp5 0.033333



Stressed mice are predicted to be older than non-stressed mice particularly among the PBS condition. On a cluster by cluster level, this was observed in 10 out of 21 clusters, with the strongest associations among the PBS condition



🔲 E14 P4 D P5

P30

Old



Genes indicative of age are generally upregulated in older mice and downregulated in younger mice when the pseudobulk data is filtered to only include the top 10 important genes from the Elastic INET MODEI

Pooled Across Clusters, Random Forest Classifier (RFC)





Older age groups are associated with the ELS condition, with the exception of P4 mice, when pooled across clusters. On a cluster by cluster level, this outcome of the oldest age group having the highest probability of ELS was observed in 6 out of 21 clusters



+ MHCII + IL1β

Individuals exposed to lifetime stress exhibit epigenetic changes resembling those seen in aging-associated diseases3.





CONCLUSION

DATASET

Findings from this study can help inform the development of more effective therapeutic interventions and personalized treatment strategies aimed at mitigating the detrimental effects of trauma on brain health and mental well-being.



POLR3H has been implicated with being a novel cause of primary ovarian insufficiency, an indicator of female infertility



WDR6 is a prognostic biomarker that correlates with immune infiltration in lung cancer and plays a role in the regulation of feeding behavior and longevity in the brain

FUTURE DIRECTIONS

tumor dra Cell proportion Purify profile 🏓 🔶 🧶 00% **♦ ●**→ **ﷺ ∦→**@

Performing CIBERSORT, a computational method used to characterize cell type proportions in bulk tissue gene expression profiles (GEPs)4



REFERENCES & ACKNOWLEDGEMENTS

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Heatmap of Expression Levels of Top 10 Most Important Genes



Predicted Probabilities of ELS by Condition: Cluster 1