

Inferring genomic signatures in age-related macular degeneration across different stages

Presented by: Minjun Park, Shryans Goyal, Yu Wu, Zach Moxley, Zishi Wang

Meet Rosa

- 80 year old grandmother
- Loves observing nature
- But.....

Cannot See Properly!



The Curse of AMD

- AMD : Age-related macular degeneration

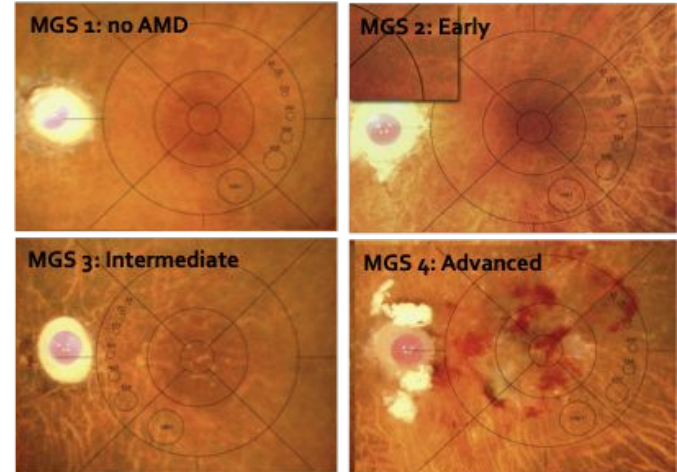


Normal vision



Loss of central vision

Ratnapriya et al., 2013, Genome Med; National Eye Institute

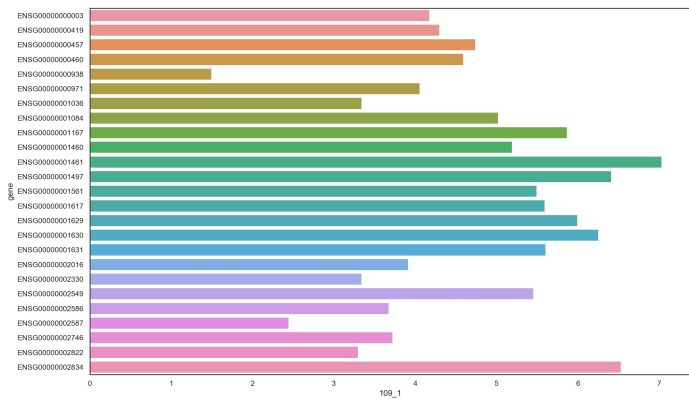


**Olsen and Feng, IOVS, 2004; Decanini et al., Am J Ophthalmol 2007*

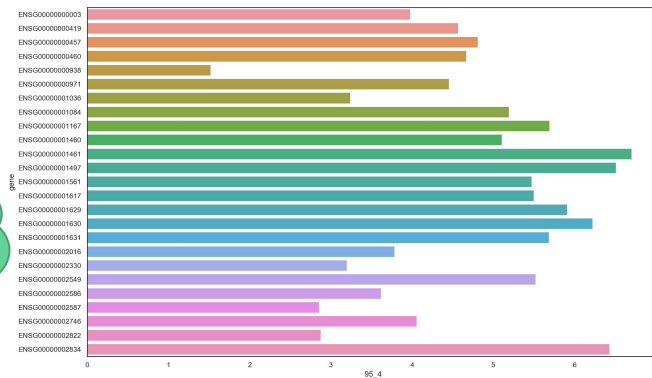
- AMD has 4 distinct disease stages

Why it Matters

- **Hard to prevent and treat!**
- **Hard to detect!**
- **Inevitable!** High magnitude of cases per year.



Normal (Stage 1 Gene Expression)



Advanced (Stage 4 Gene Expression)

Impact

Early detection



Identify Drug Targets



Protect Rosa!



Data Preprocessing

453 patients in **4** disease stages

Low Variance filtering

ANOVA

~ **58, 000** Genes

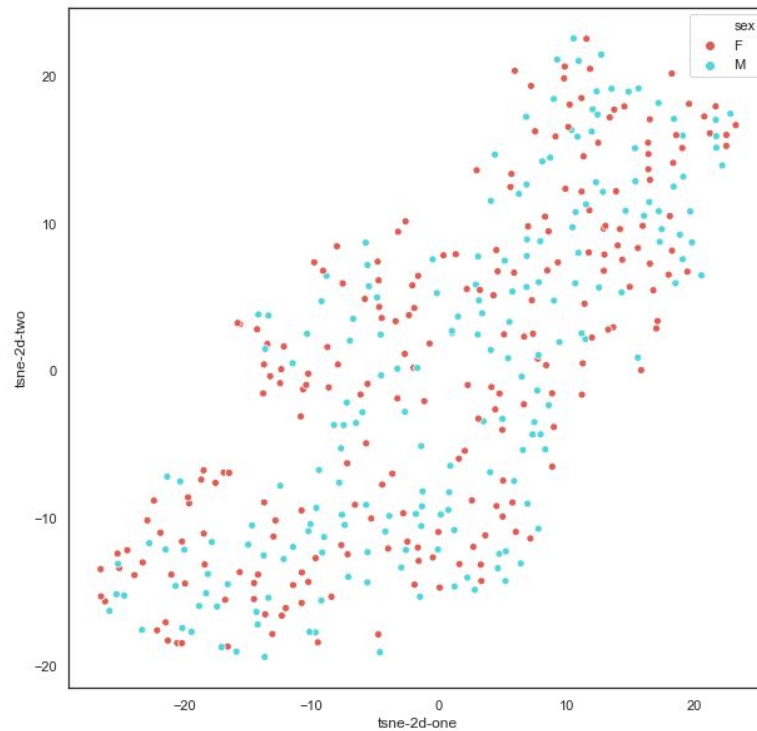
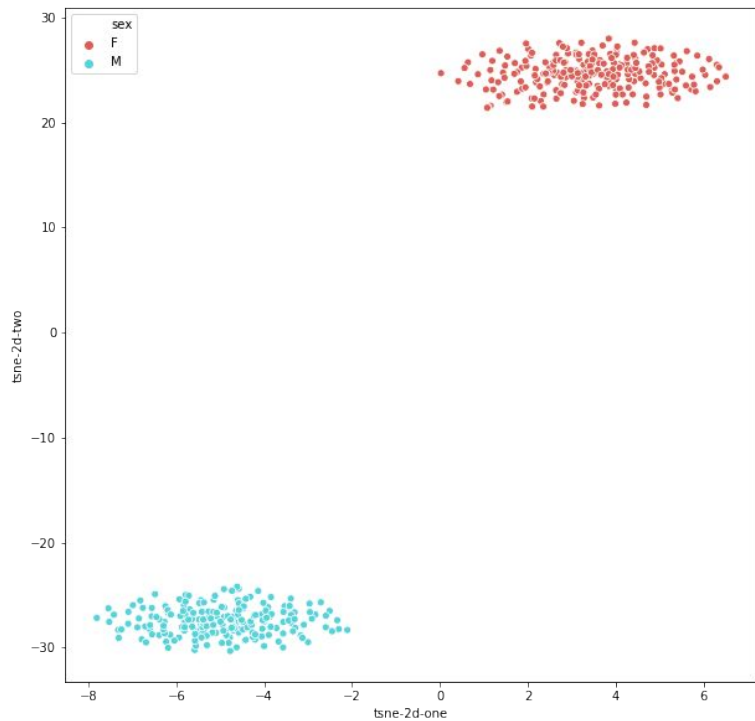


~ **37, 000** Genes



~ **4, 000** Genes

Dimension Reduction



Objective 1

Extract set of genes that
are most important to each
stage of AMD

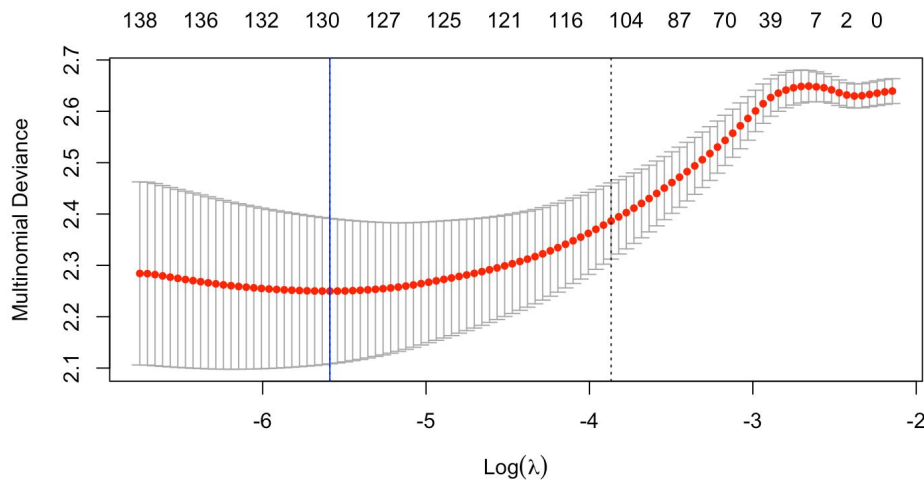


Modeling pipeline

- Feature Selections methods
 - LASSO Multinomial regression
 - Random forest
 - XGBoost
- 10-fold cross validations
- Bootstrapping of 500 iterations

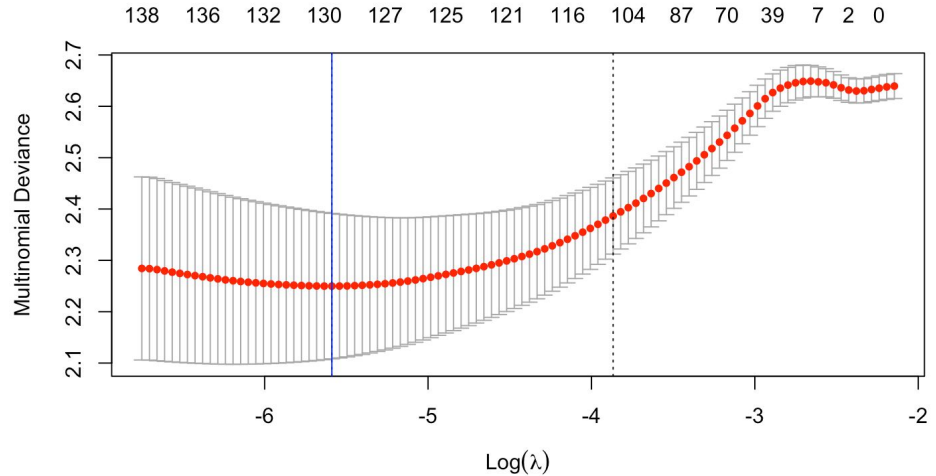
Modeling pipeline

- Feature Selections methods
 - LASSO Multinomial regression
 - Random forest
 - XGBoost
- 10-fold cross validations
- Bootstrapping of 500 iterations



Modeling pipeline

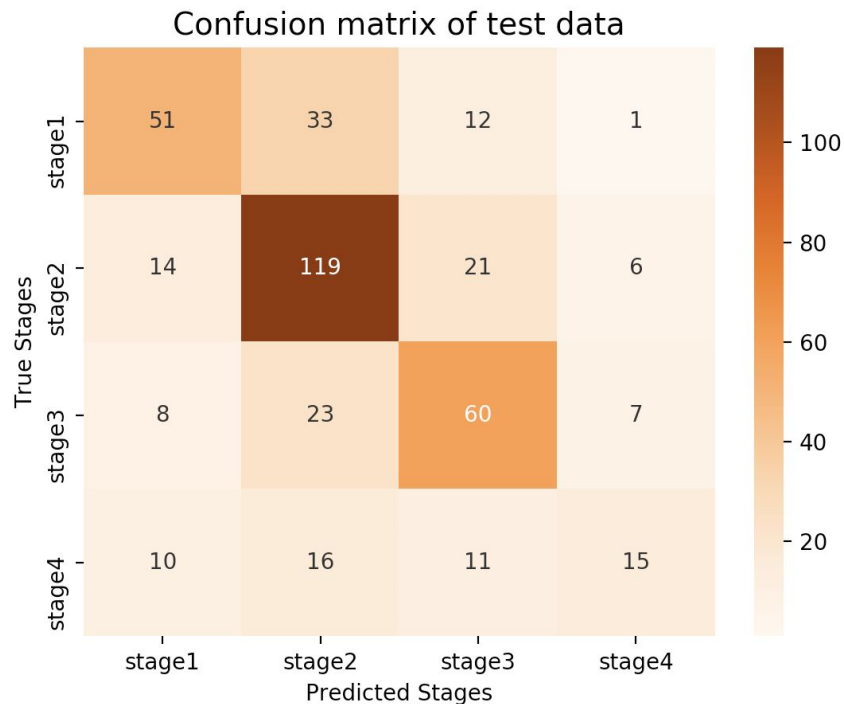
- Feature Selections methods
 - LASSO Multinomial regression
 - Random forest
 - XGBoost
- 10-fold cross validations
- Bootstrapping of 500 iterations



- **Identified important genes that overlap in all three models**
- DE genes found: MOXD1, PMAIP1**

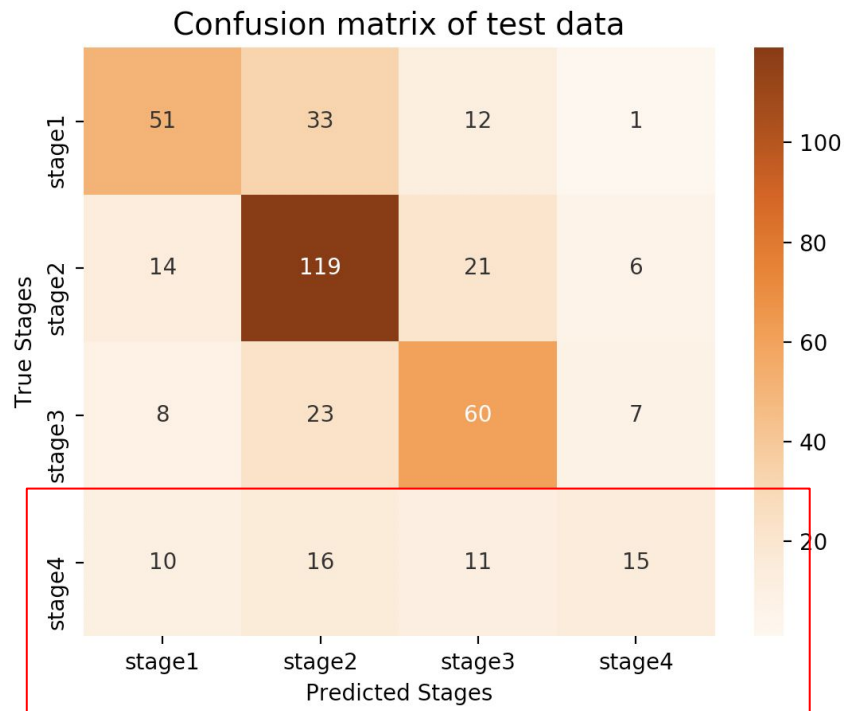
Choosing the Best Model: **Multinomial Lasso Regression**

📊 **All Stages Accuracy: 60.2%**



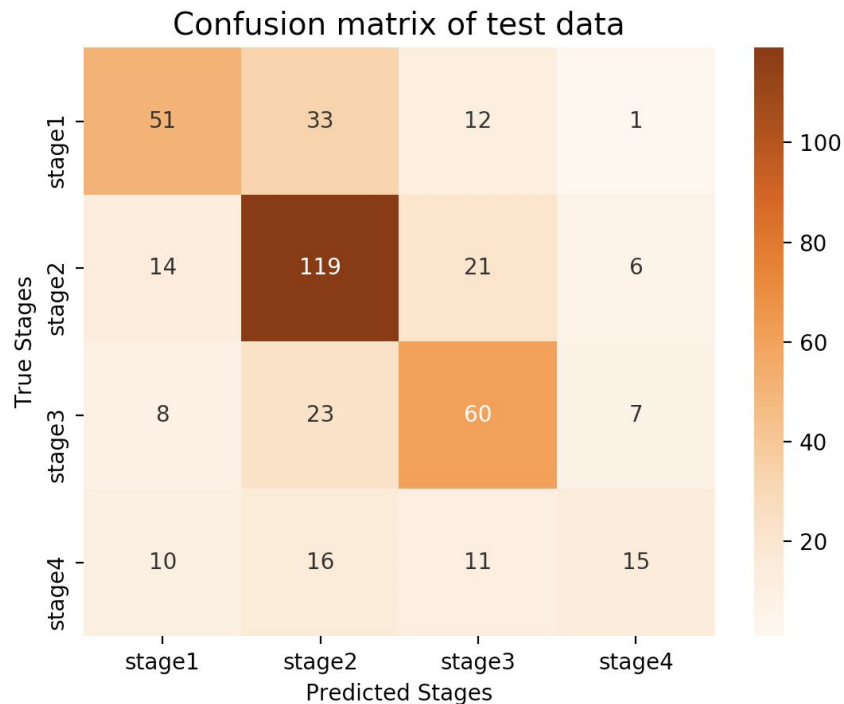
Choosing the Best Model: **Multinomial Lasso Regression**

📄 **All Stages Accuracy: 60.2%**

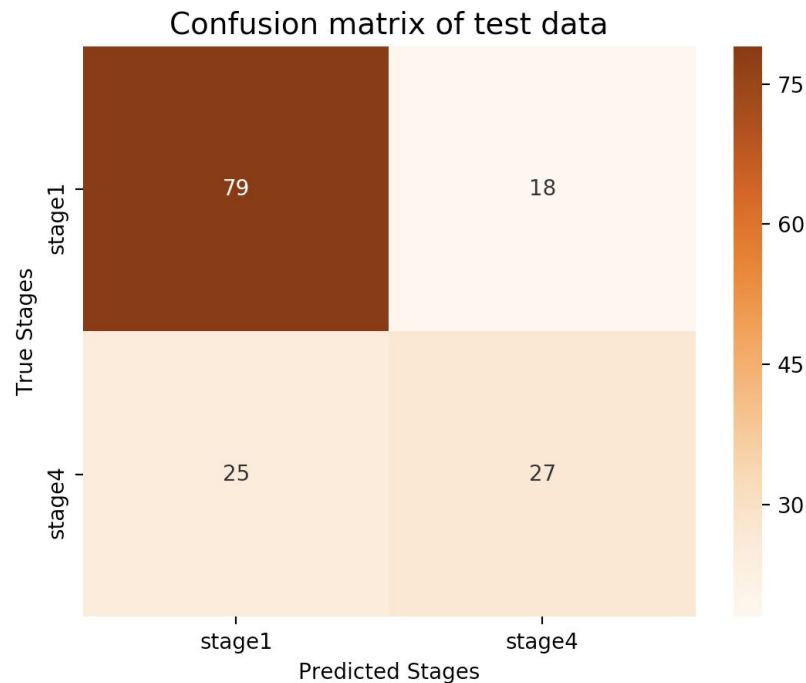


Choosing the Best Model: **Multinomial Lasso Regression**

☐ **All Stages Accuracy: 60.2%**

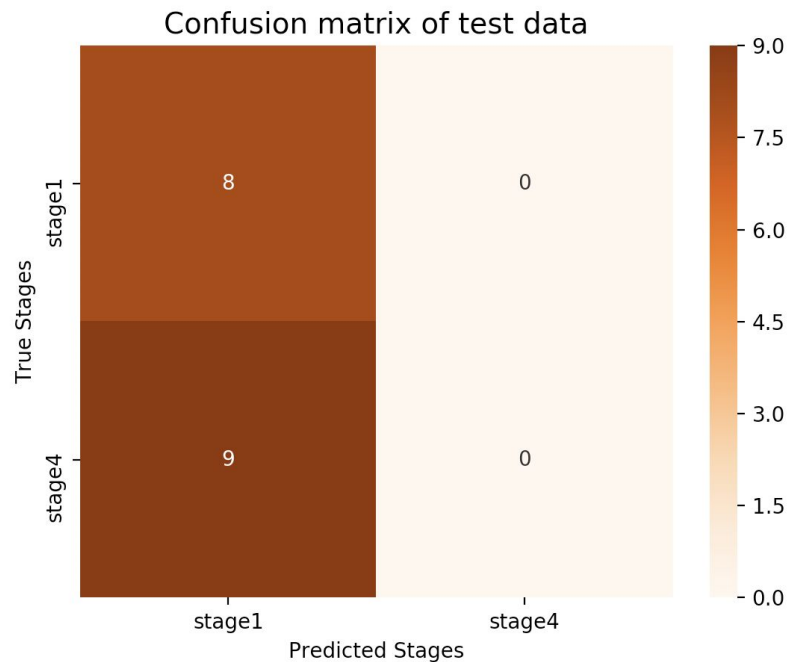
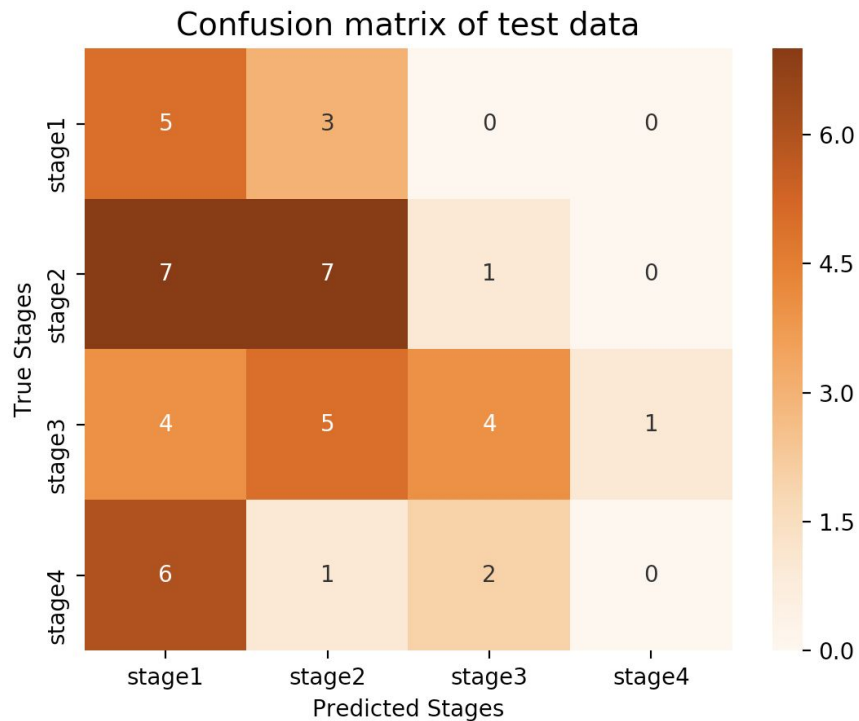


☐ **Stage 1 vs Stage 4 Accuracy: 71.1%**



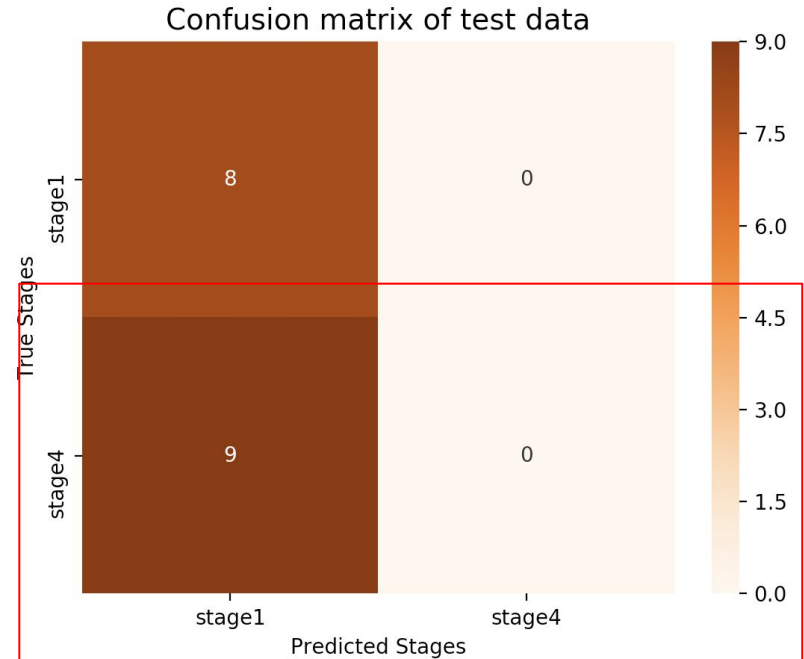
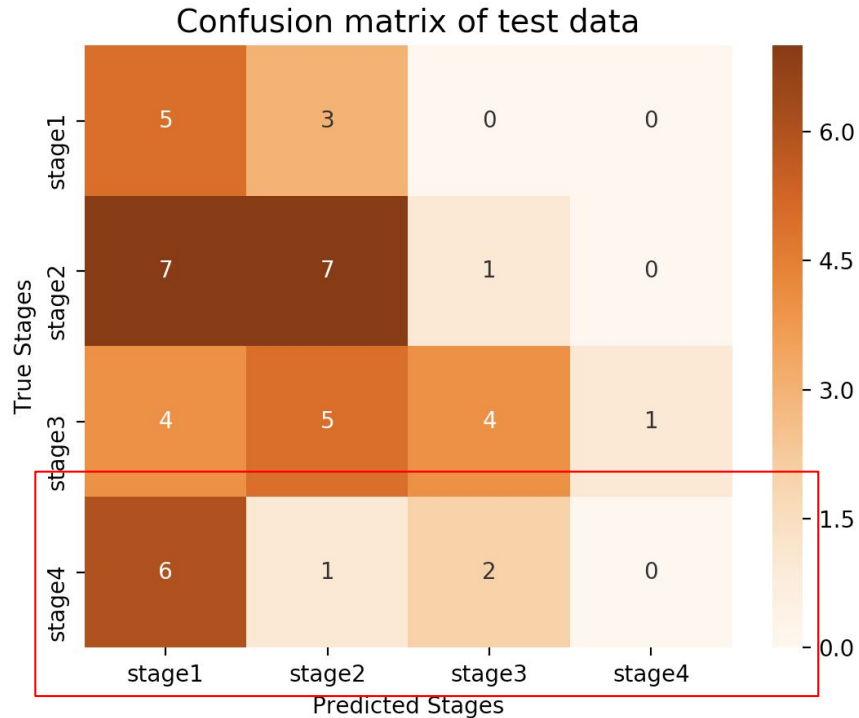
Hold-out data performance

❏ **46 Patients** Prediction accuracy: 34.8% and 47.0%



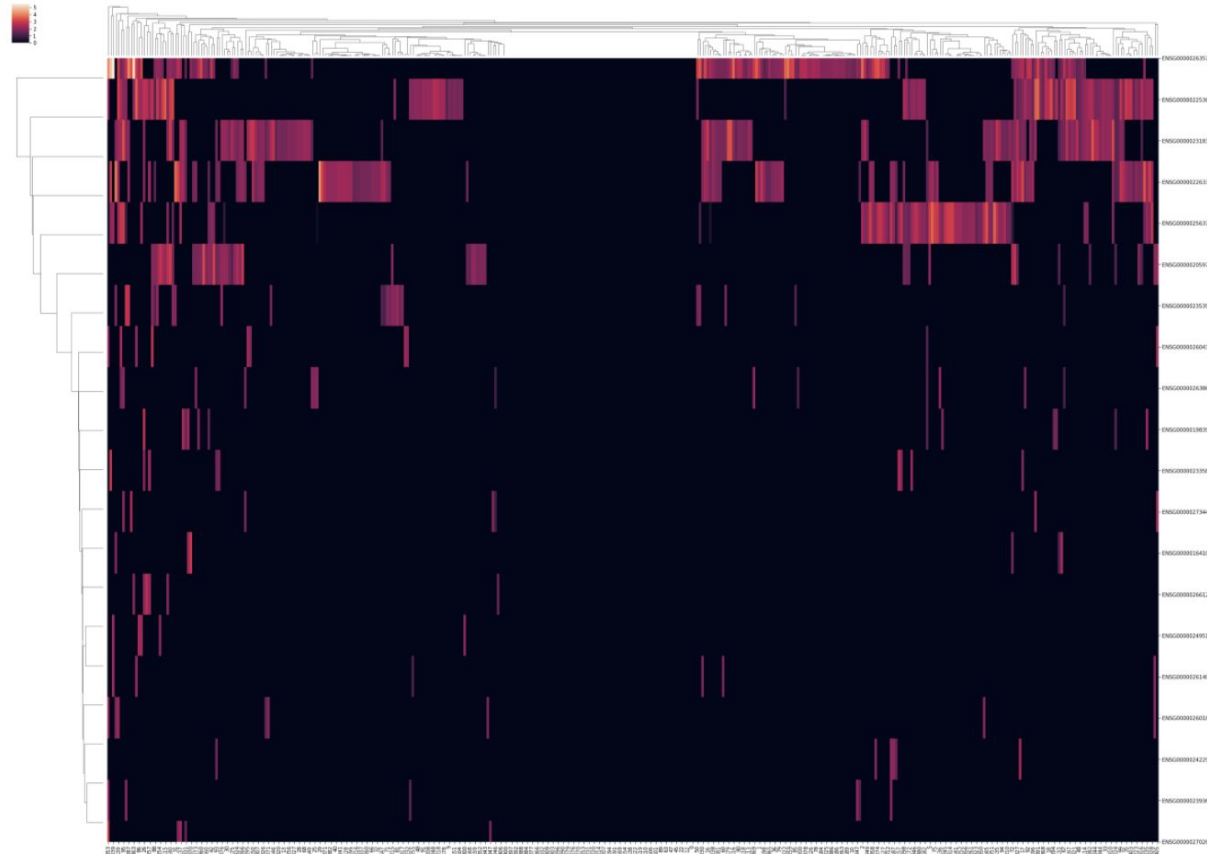
Hold-out data performance

❏ **46 Patients** Prediction accuracy: 34.8% and 47.0%



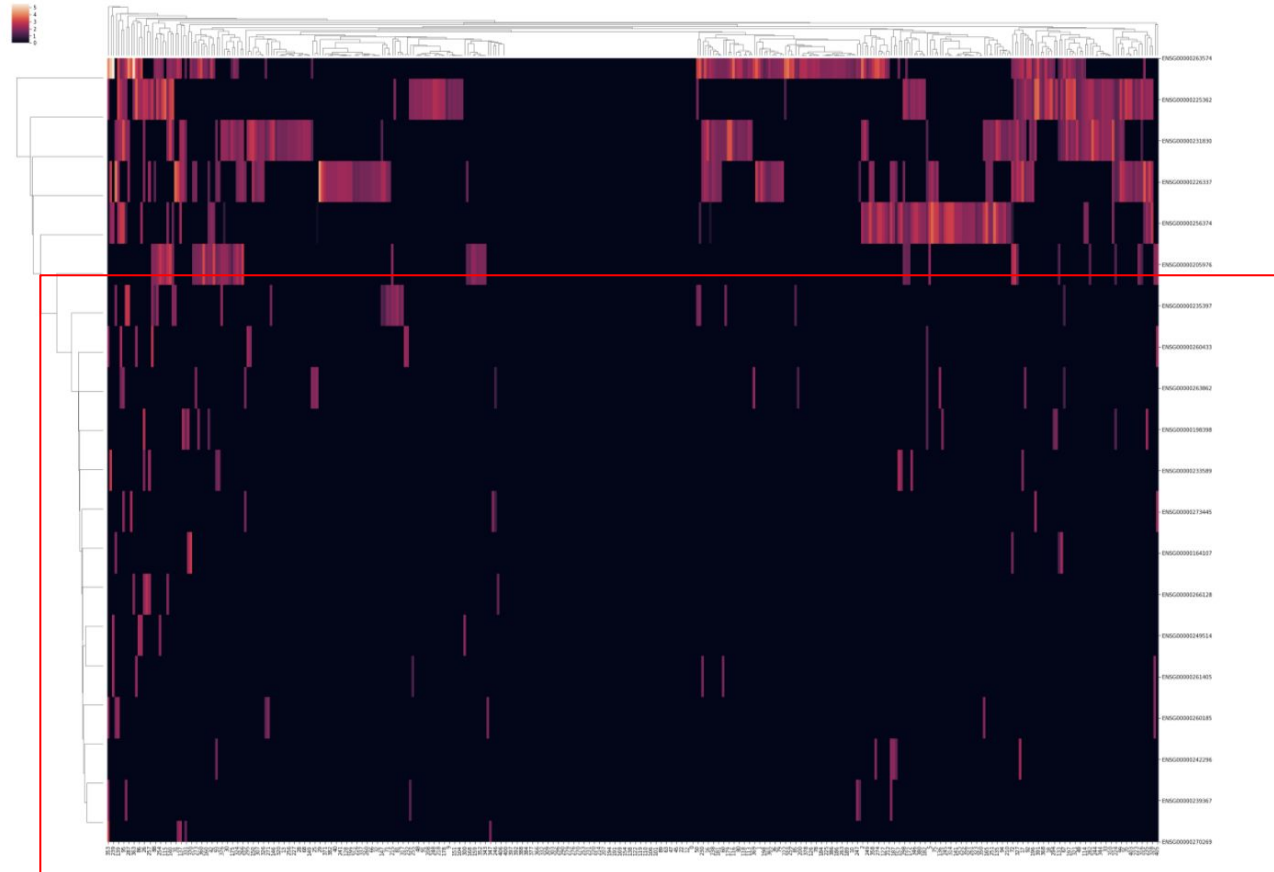
Cluster heatmap of 20 important genes

□ Stage 4

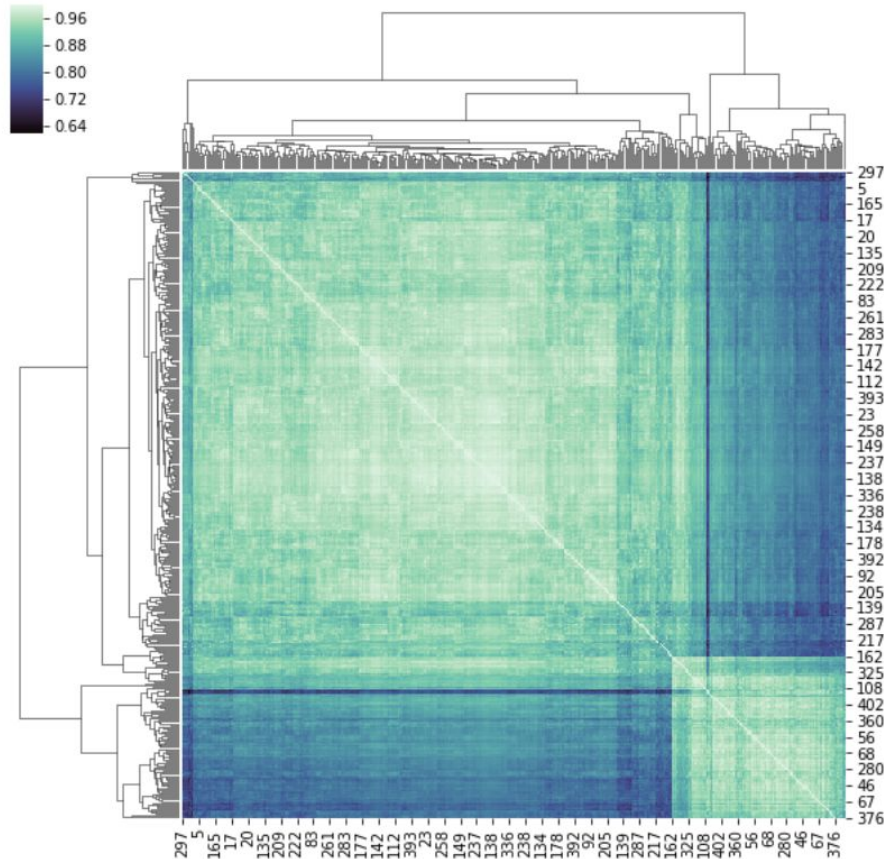


Cluster heatmap of 20 important genes

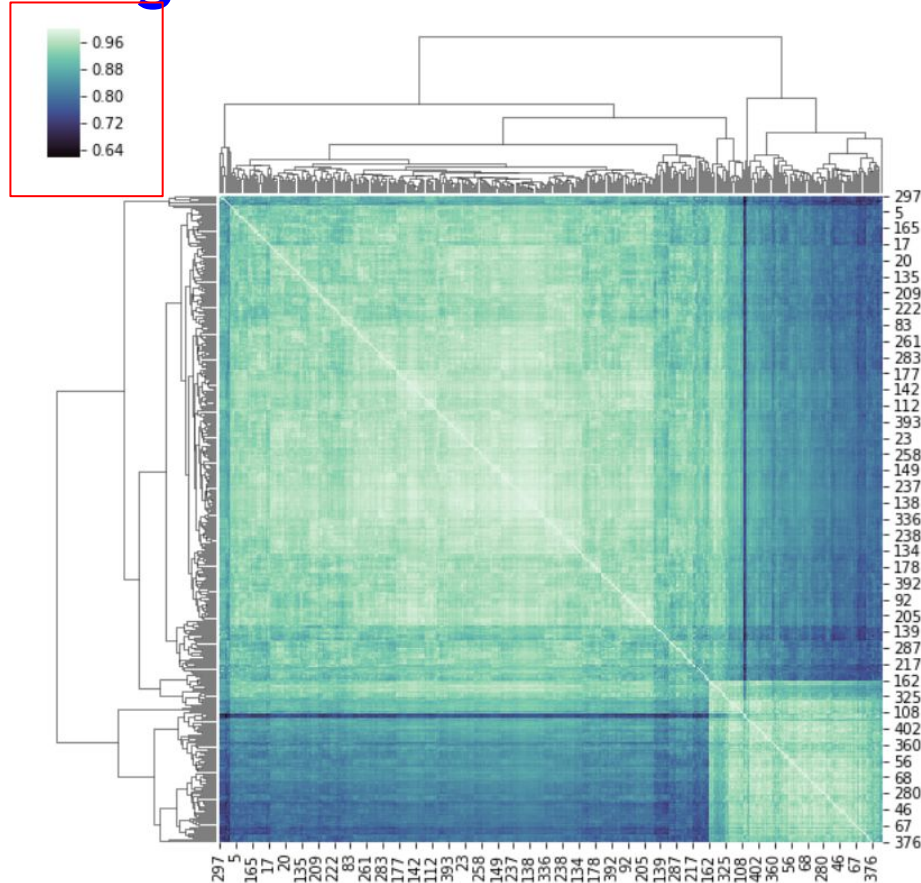
□ Stage 4



Stage 4 Correlation Matrix important genes



Stage 4 Correlation Matrix important genes



Objective 2

Explore genomic pathways
and their association with
AMD stages

Objective 3

Explore pathway networks

What is a pathway?

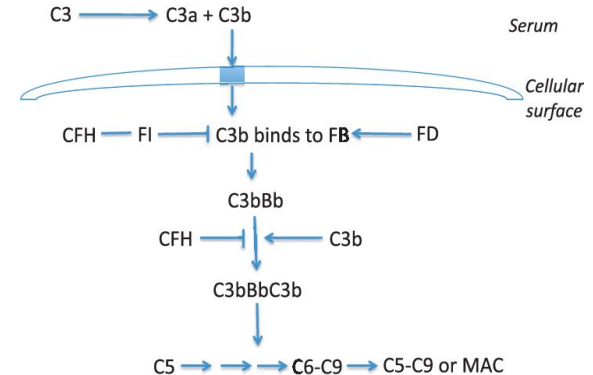
A pathway is a linked **series of actions** among genes in a cell that **lead to** a certain product or a **change**.

Genes affected in disease

CFH
C3
C5
.
.
C9



Complement pathway



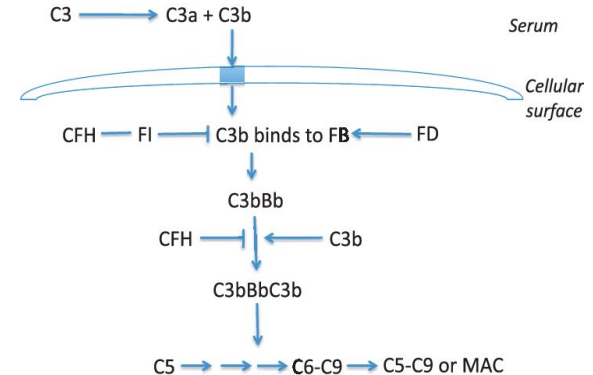
What is a pathway?

A pathway is a linked **series of actions** among genes in a cell that **lead to** a certain product or a **change**.

Genes affected in disease

CFH
C3
C5
.
.
C9

Complement pathway



What is a pathway?

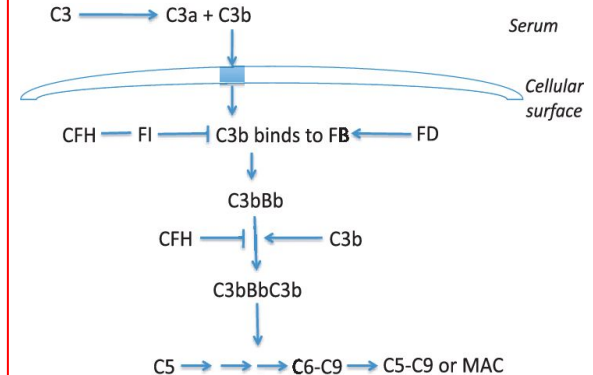
A pathway is a linked **series of actions** among genes in a cell that **lead to** a certain product or a **change**.

Genes affected in disease

CFH
C3
C5
.
.
C9



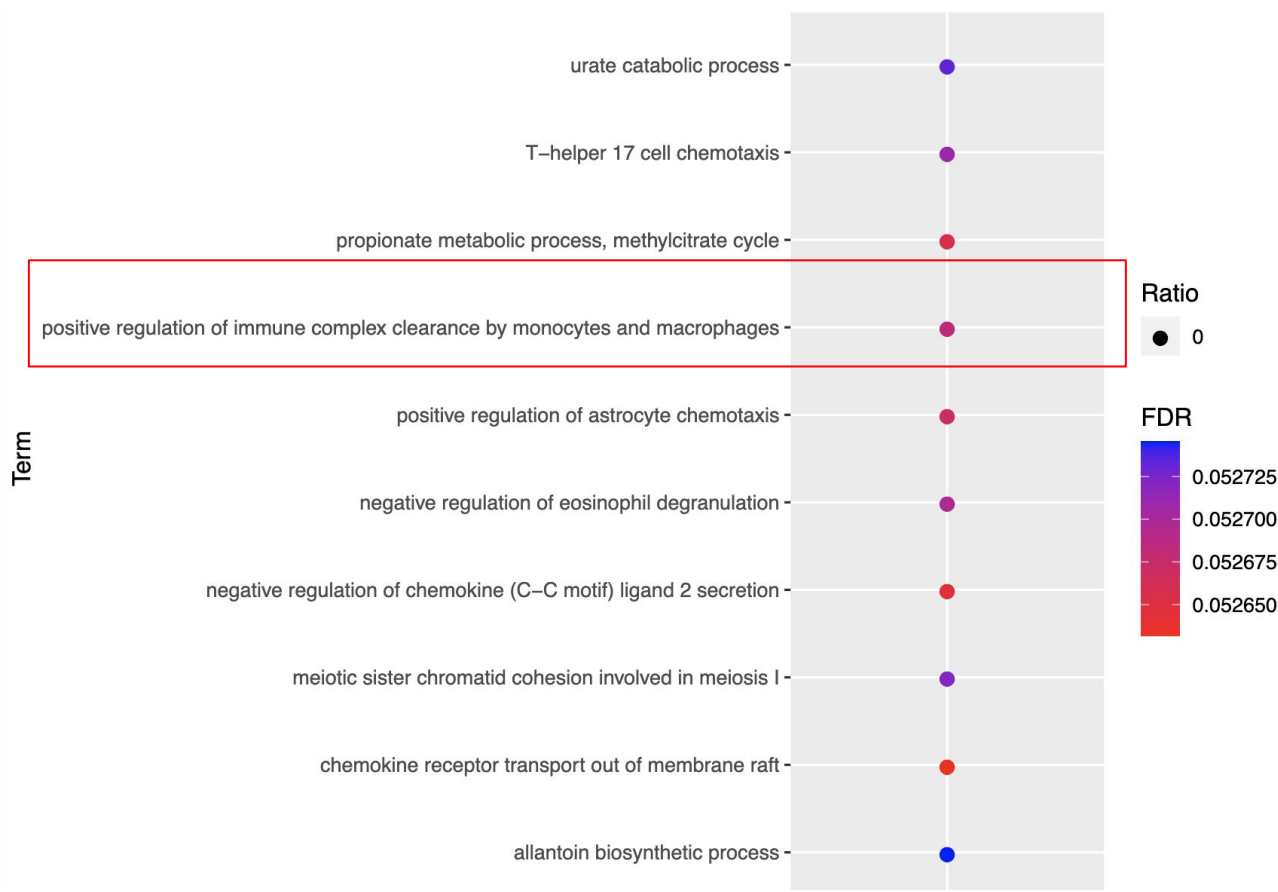
Complement pathway



GO Enrichment Analysis



GO Enrichment Analysis



Next step: Pathway Networks

- Network comprises **multiple pathways** that are already known **to exist and interact** with each other
- **Holistic view** of pathways with other experimental datasets

Thank You!