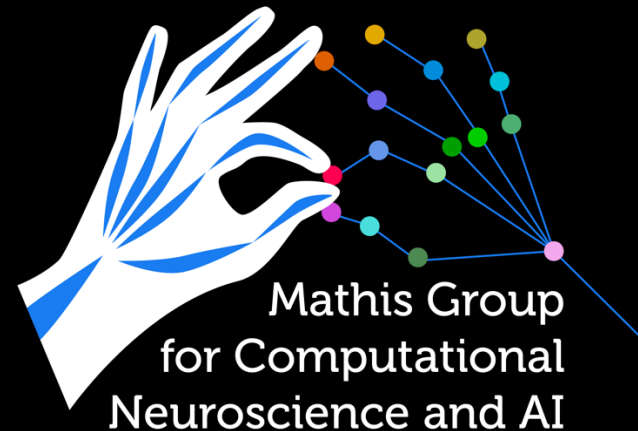
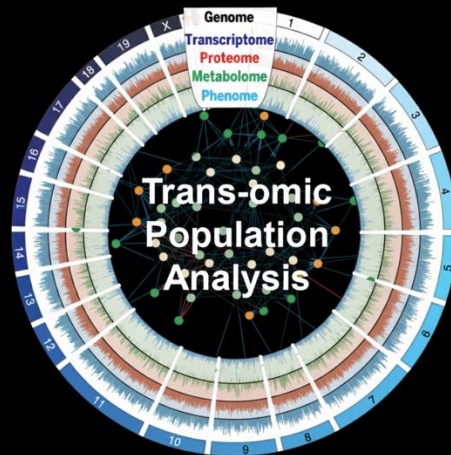


Deep phenotyping via hierarchical learning of mouse movement: Decoding aging through activity

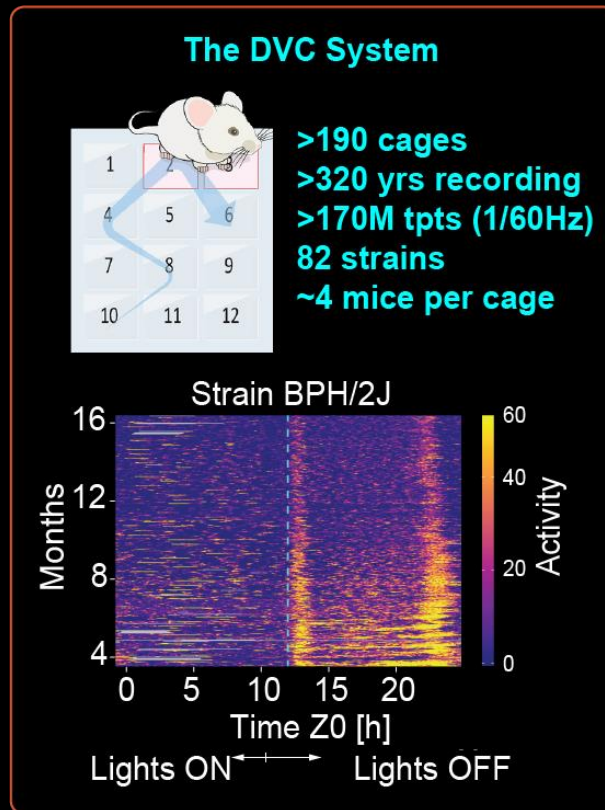
Gaurav Bhole¹, Jon Lecumberi¹, Giacomo von Alvensleben¹, Andy Bonnetto²,
Michal Grudzien², Johan Auwerx¹, Alexander Mathis²

¹Laboratory of Integrative Systems Physiology

²Mathis Group for Computational Neuroscience and AI



The HDP: an integrated platform for Healthspan discovery



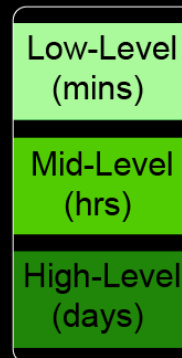
82 inbred strains
>130k samples
>20 tissues
>4k female mice

DVC: 24/7 homecage activity and
environmental monitoring

Classical Statistical
Analysis

Deep Learning - Hierarchical
Masked Autoencoder (hBehaveMAE)

**Unsupervised
Representation
Learning**



Human Presence,
Bedding Change, ...

Day / Night, ...

Aging Patterns,
Strain Identity, ...

Embeddings

**Downstream
Analysis**

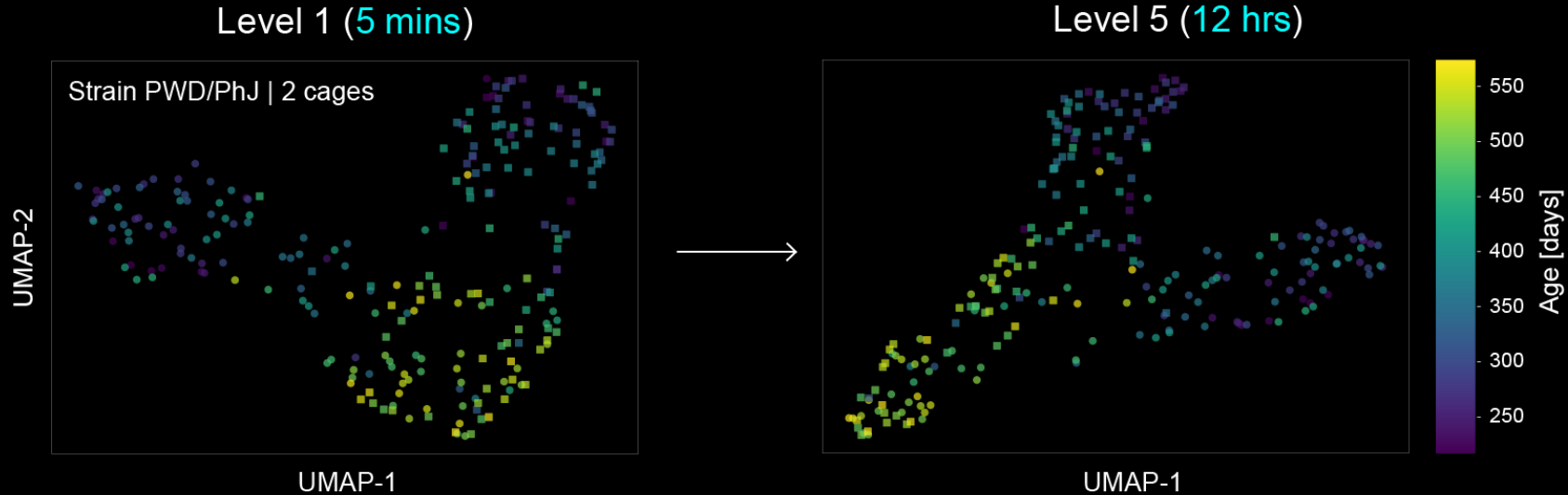
Behavioural
& Aging

.....

Genetic Mapping
of Activity

DVC embeddings recapitulate phenotypic features

UMAPs of unsupervised learnt embeddings show patterns of age separation



Categorical classification tasks

All levels combined

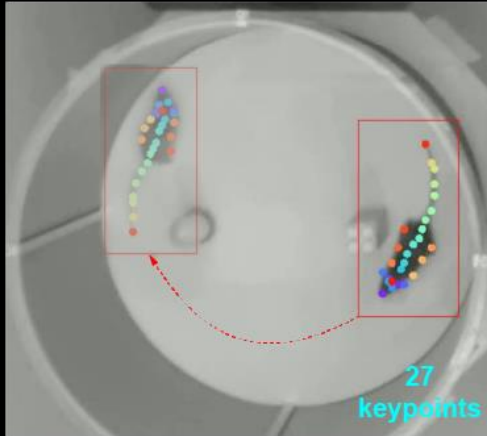
Features	Performance (Metric)
Strain Identity	56.6%* (Accuracy)
Age Prediction	~4 months (RMSE)
Day/Night	In Progress...
...	

*often mispredicted as another genetically similar strain

Future Prospects

Pose estimation

Openfield videos, DeepLabCut

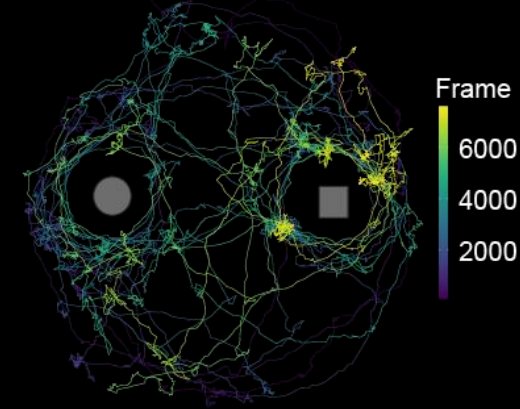


>700 hrs single-animal recordings.
Adult, Old. >3400 videos



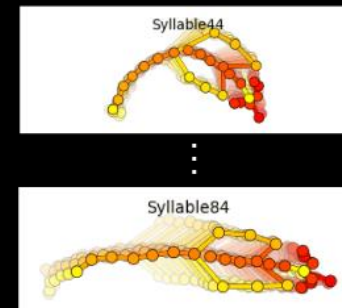
DeepLabCut

Mouse center trajectory



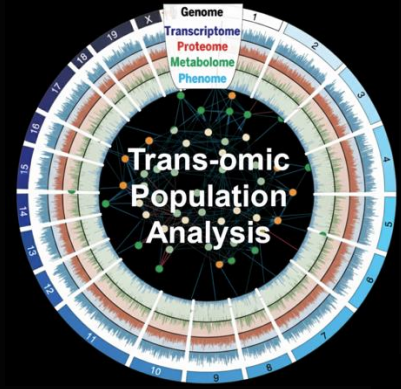
Behavioral syllables extraction

Keypoint-MoSeq

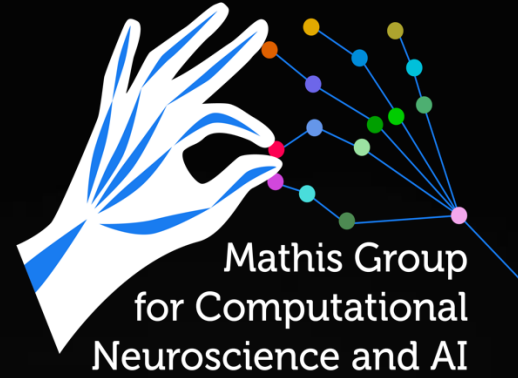


Changes
across strains

Parsing the ageing
and genetics
of mouse behaviour



Credits



Laboratory of Integrative Systems Physiology
Animal Facility Team (UDP)

Mathis Group for Computational
Neuroscience and AI

See you at Poster 5 !!!

