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Characterization of Murine Astrocyte Cultures in 3D Hydrogel Matrices

An Approach to Nuclei Segmentation and Assisted Cell Classification

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Thesis to obtain the Master of Science Degree in

Biomedical Engineering

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March 2025

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Motivation

Astrocytes in Neuroscience Research

- Most abundant glial cell type in the mammalian CNS
- Very heterogeneous cell population (Bocchi et al., 2025)
- Potential for therapeutic applications via direct neural reprogramming (Pereira et al., 2024)

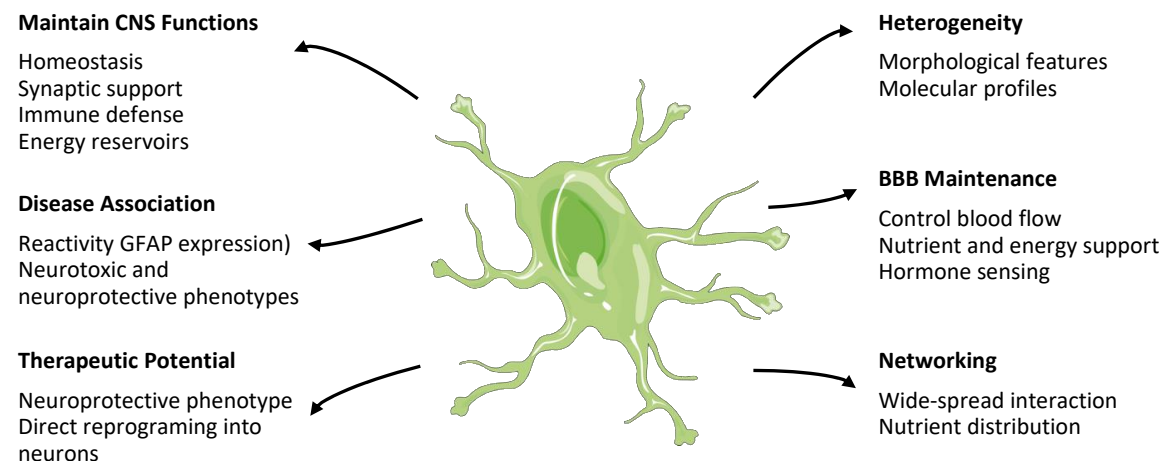


Figure 1.1. Astrocytes main characteristics. Adapted from Garland et al. (2022).

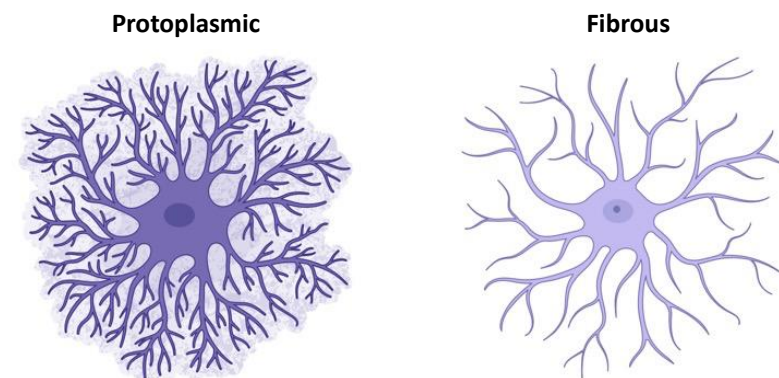


Figure 1.2. Astrocytes morphological heterogeneity. Created in BioRender.

Motivation

Optimize Hydrogel Models

- Limitations of current 2D models to mimic brain structure and physicochemical properties
- Potential of resembling brain physicochemical properties and microenvironment

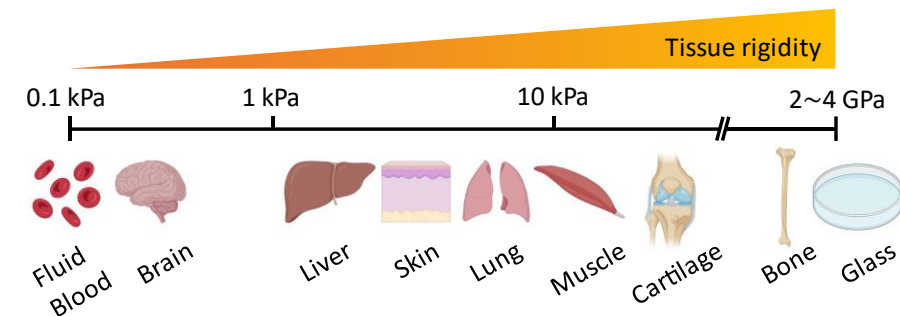


Figure 1.3. Rigidity scale of different tissues and glass (adapted from Park et al. 2022).

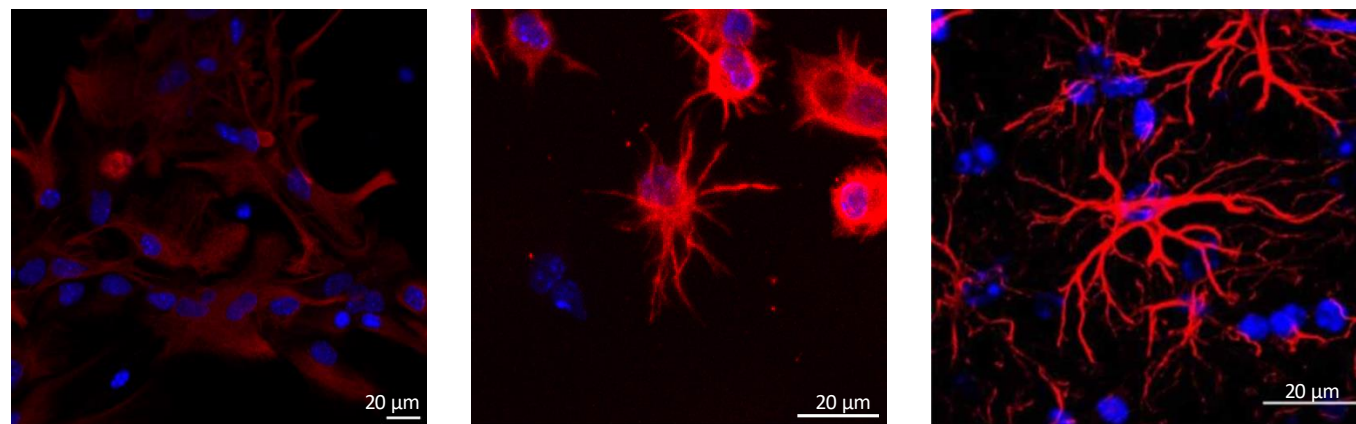


Figure 1.4. Astrocytes morphology grown in different environments. (A) and (B) 7 DIV cultures (from this Master's thesis), and (C) Ex vivo morphology from Matias et al. (2019).

A. Monolayer PDL

B. Hydrogel substrate

C. Murine brain

Motivation

Automate Biological Image Analysis

Challenges:

- Biological image annotation is very time consuming
- 3D stack annotation is challenging

Opportunities:

- Potential to automate annotation process
- Advances in computer vision technologies allow accurate identification of objects
- High-throughput characterization of cultures

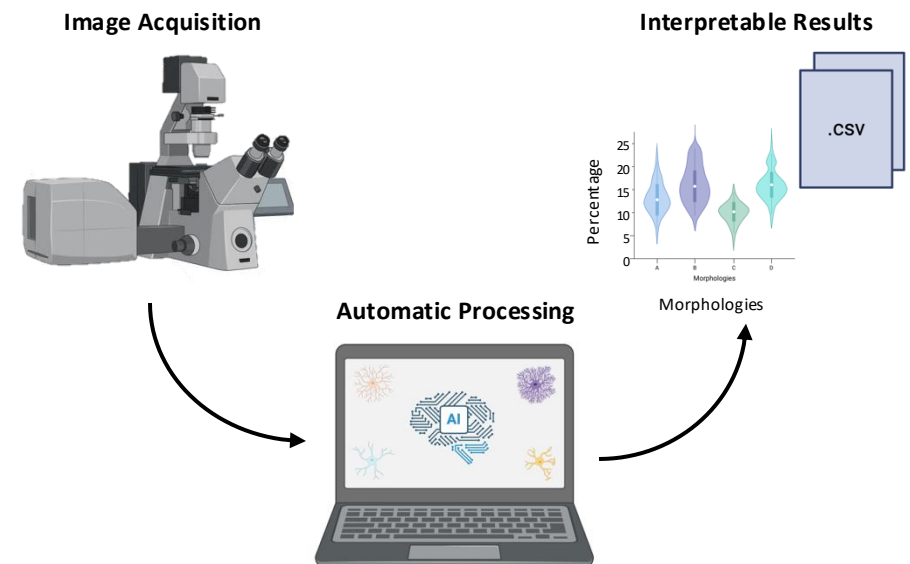


Figure 1.5. Pipeline overview for automatic image analysis.

Research Objectives

Computational Work

- Train nuclei segmentation models for hydrogel cultures
- Co-develop GUI for z-stack image annotation

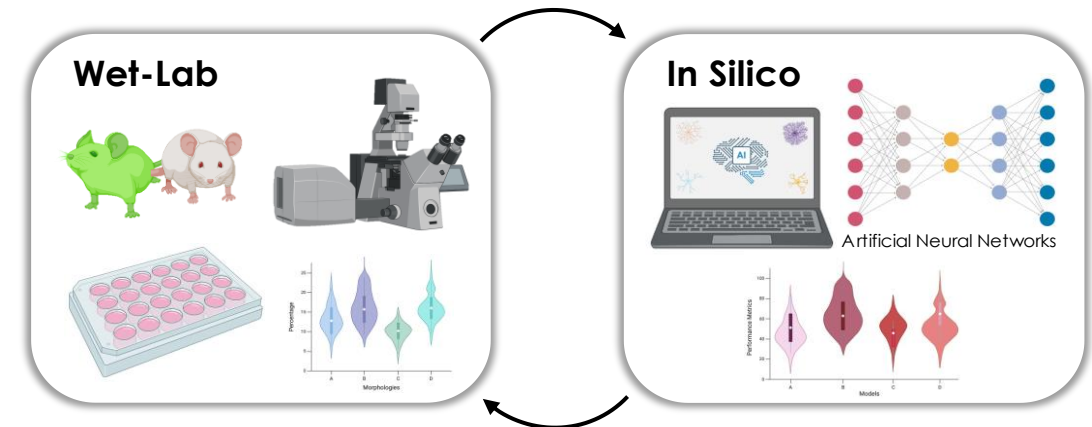


Figure 1.6. Illustrative scheme of project tasks.

Experimental Work

- Prepare murine gray matter-derived astrocyte cultures in hydrogels
- Compare astrocyte 3D **morphology** in different hydrogel substrates (MVG, VLVG, Fibrin, FibrinVLVG)
- Analyse of **expression astrocyte markers** in different hydrogels and PDL cultures (SOX9, GFAP)
- Study the **effect of CaCl₂ concentration** used for hydrogel gelling on cell viability, morphology and astrocyte marker expression

02 Computational Work

Machine Learning for Biomedical Data Analysis

Biomedical Data Analysis

- Drug discovery
- Precision healthcare
- Protein structure prediction
- Omics/Multi-omics
- Risk prediction prognosis
- Clinical decision support
- Therapeutic assistance
- Medical and **biological image analysis**

Biological Imaging

- Observe biological processes and the cellular and molecular levels
- High volumes of data
- Absence of standardized protocols
- High variability of experimental conditions

Need of Automation

- ❖ High-throughput data analysis
- ❖ Enhance reproducibility of data analysis

Vs.

Hurdles of Automation

- ❖ Heterogeneity of datasets
- ❖ High data variability

Deep Learning for Biological Image Analysis

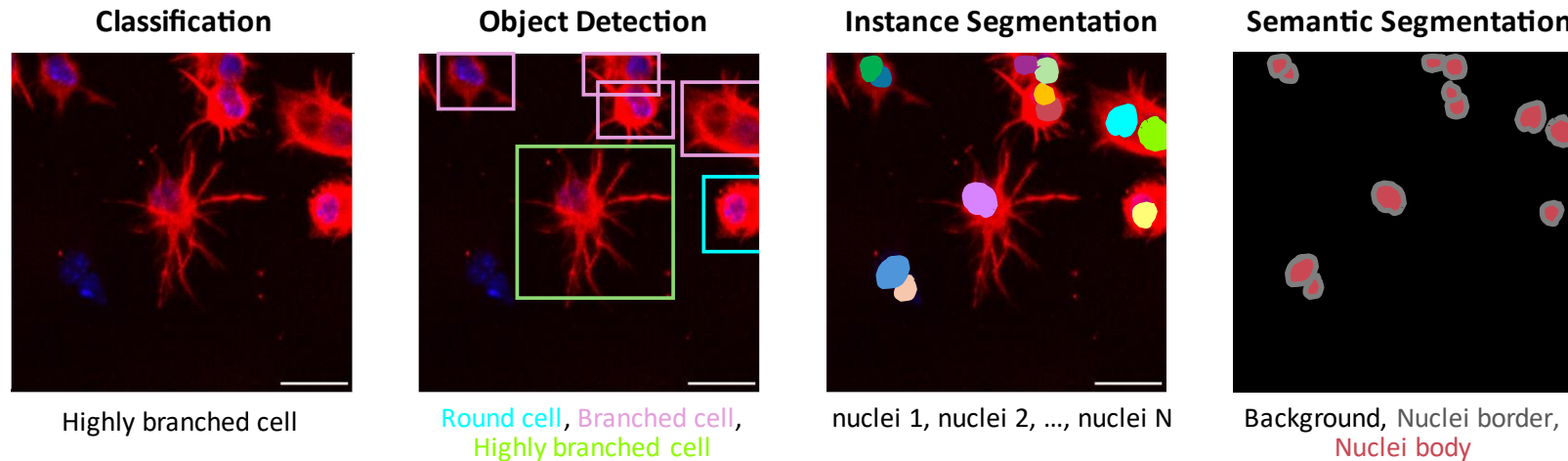


Figure 2.3. Outlook of computer vision tasks in the context of astrocyte culture characterization.

Convolutional Neural Networks (CNNs)

- Sparse interactions
- Parameter sharing
- Equivariance to translation

Open-source Architectures for Segmentation

- U-Net, nnU-Net
- Mask R-CNN
- Stardist 2D, Stardist 3D
- Cellpose and Cellstitch

3D Nuclei Segmentation

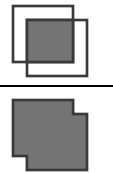
Training and Testing Dataset

- 16 z-stack images
- Murine-derived astrocyte cultures in 3D hydrogel matrices
- 200.63 x 200.63 x 200 μm^3
- DAPI and GFAP channel

Architectures

- nnU-Net
- Stardist 3D
- Cellstitch

Evaluation Metrics

- $IoU = \frac{\text{Area of overlap}}{\text{Area of union}} = \frac{A \cap B}{A \cup B}$ 
- $precision_{\tau} = \frac{TP}{TP+FP}$
- $recall_{\tau} = \frac{TP}{TP+FN}$
- $mAP_{\tau} = \frac{TP}{TP+FP+FN}$
- $F1\ score_{\tau} = \frac{2 * precision * recall}{precision + recall}$
- $PR = \frac{\# predictions}{\# GT predictions} = \frac{TP+FP}{TP+FP+FN}$
- $AR = \frac{\# aggregated predictions}{TP+FP+FN}$

Results: 3D Nuclei Segmentation

nnU-Net Outperforms Stardist 3D and Cellstitch

- nnU-Net closer to optimal values in all assessed metrics
- Stardist 3D **underestimates** the number of nuclei
- Cellstitch **overestimates** the number of nuclei

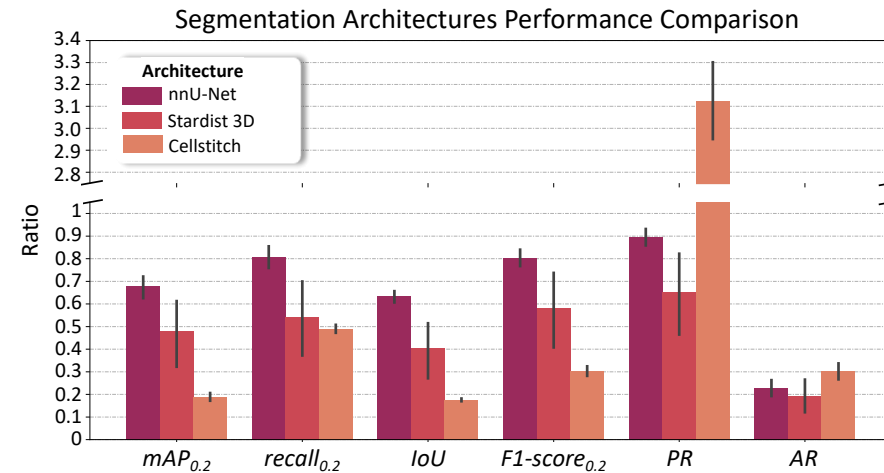


Figure 2.5. Performance of nnU-Net, Stardist 3D and Cellstitch. Error bar: 95% confidence interval.

Results: 3D Nuclei Segmentation

Improving nnU-Net Performance

- Transform the training dataset
- FOV range relates to nuclei relative size in the images

Table 2.3. Depiction of datasets generated for nnU-Net model enhancement.

Features	dataset 1	dataset 2	dataset 3	dataset 4	dataset 5
Images	preprocessed	raw	raw	raw + augmented	raw + augmented
Size	16	16	16	120	150
Selected Channels	DAPI + GFAP	DAPI + GFAP	DAPI	DAPI	DAPI
Augmentation	–	–	–	rotations, vertical and horizontal flips	rotations, vertical and horizontal flips, zoom
Digital Size (512,512,101,Z)	Z = 2	Z = 2	Z = 1	Z = 1	Z = 1
Training FOV Range (μm^2)	200.63	200.63	200.63	200.63	100.31 – 802.50
nnU-Net model	<i>model 1</i>	<i>model 2</i>	<i>model 3</i>	<i>model 4</i>	<i>model 5</i>

Enhanced nnU-Net Models Achieve Outstanding Prediction Ratios

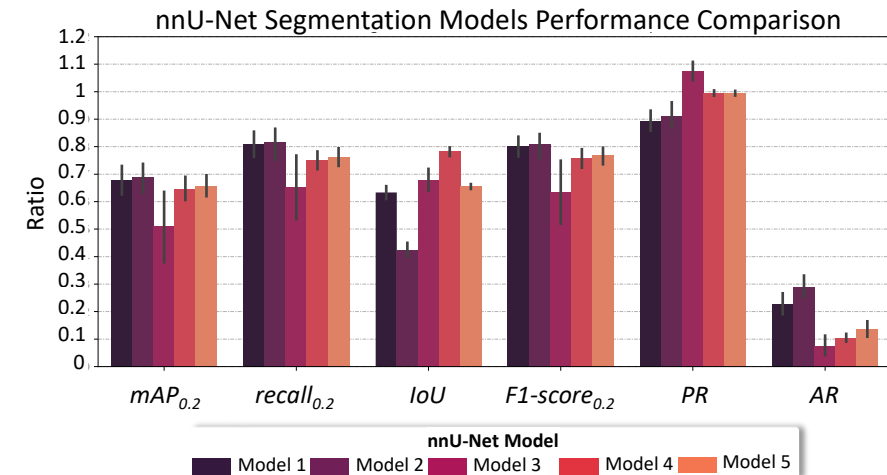


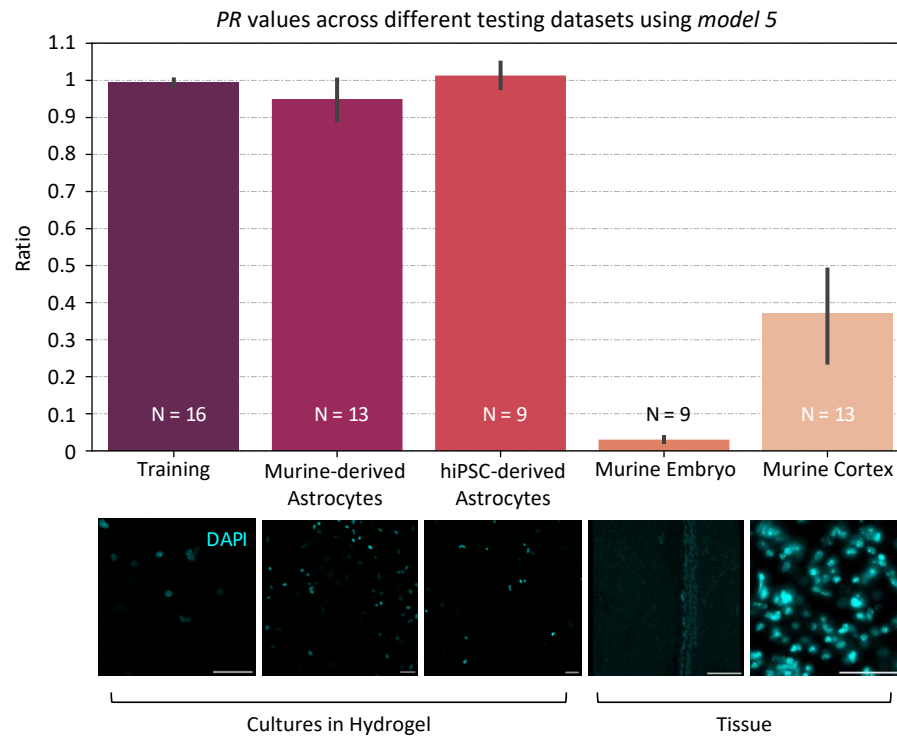
Figure 2.6. Performance of nnU-Net enhanced models. Error bar: 95% confidence interval.

Table 2.4. Scoring of trained nnU-Net models.

Model	$recall_{0.2}$	$precision_{0.2}$	$mAP_{0.2}$	IoU	PR	AR	Score
<i>model 1</i>	2	2	2	4	5	4	20
<i>model 2</i>	1	1	1	5	4	5	17
<i>model 3</i>	5	5	5	2	3	1	21
<i>model 4</i>	4	4	4	1	2	2	17
<i>model 5</i>	3	3	3	3	1	3	16

Results: 3D Nuclei Segmentation

nnU-Net model successfully generalizes to unseen hydrogel cultures datasets



$$PR = \frac{\# \text{ predictions}}{\# \text{ GT predictions}} = \frac{TP + FP}{TP + FP + FN}$$

Figure 2.8. nnU-Net model 5 performance. **(A)** PR across the different testing datasets using model 5 as predictor. Error bar: 95% confidence interval. **(B)** Maximum intensity projections of images of each dataset. Scale bar = 50 μm . Z-stack depth = 12 μm .

GUI for Annotation of Z-stack Images

Image Visualization:

- Z-stack images: multilayer and multichannel
- Automatically adjusts channels to image loaded

Automatic Detection of Cells:

- Nuclei segmentation model to draw boxes around cell nuclei
- Additional facility to manually add missed instances
- Visualization of individual cell instances

Cell Classification:

- Semi-automatic prediction of markers expression
- Classification of cell morphology

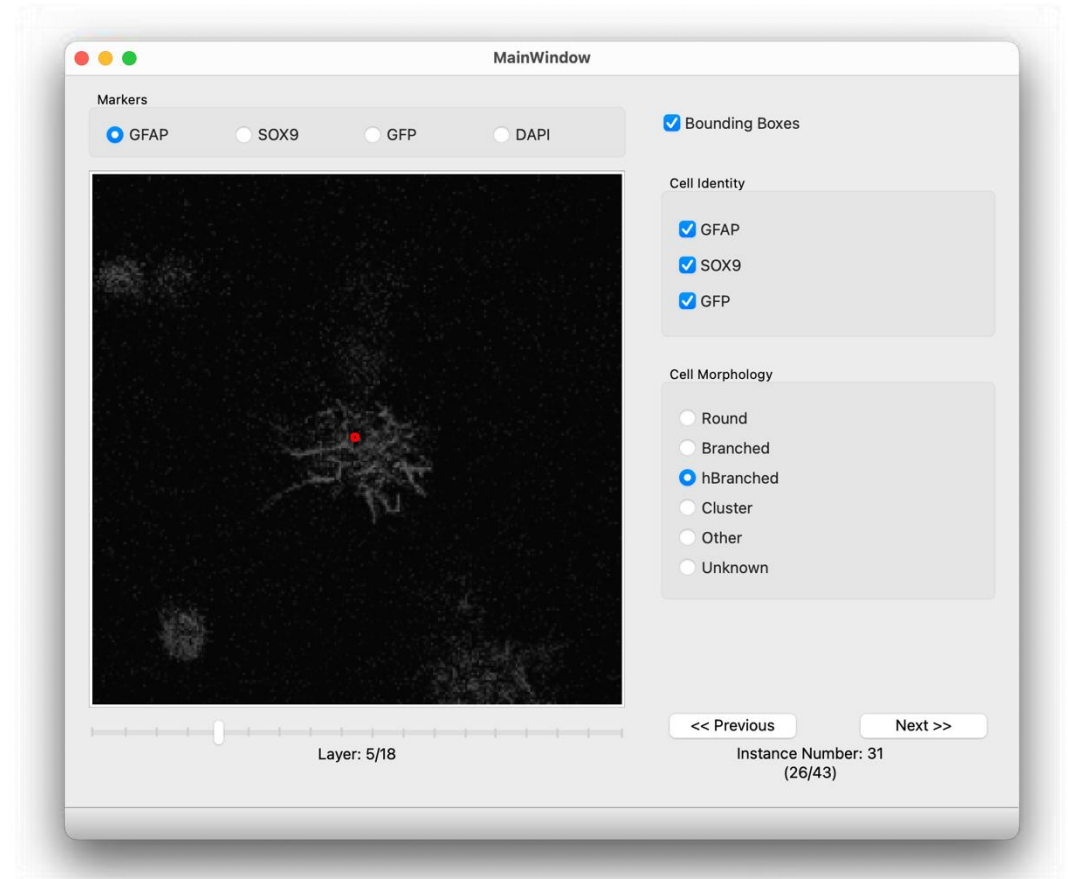


Figure 2.11. GUI cell annotation window.

03 Experimental Work

Hydrogels

Alginate Hydrogels – MVG and VLVG

- Derived from brown algae – sodium alginate
- Polysaccharide-based hydrogel
- RGD motifs for cell adhesion

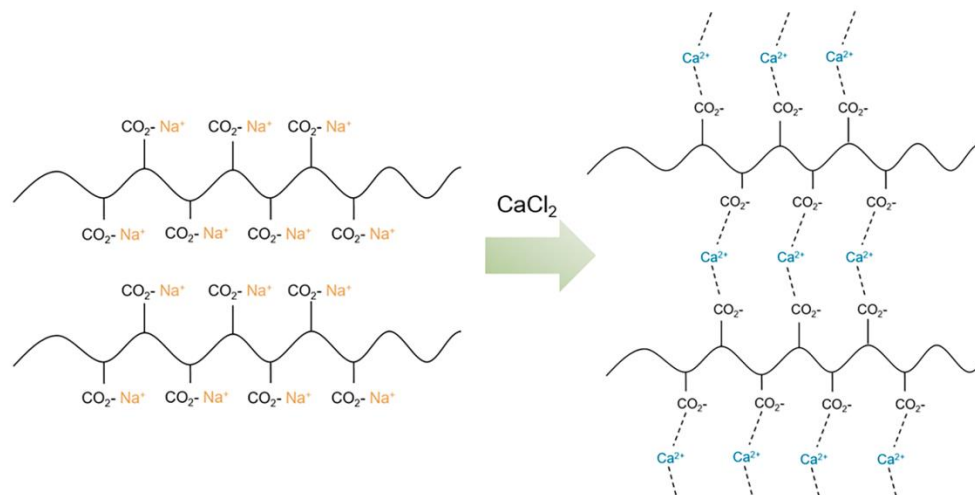


Figure 3.1. Sodium alginate crosslinking.

Fibrin Hydrogels

- Can be found in the ECM
- Fibrinogen is upregulated in response to lesion
- Might mimic closer brain injury than healthy conditions – fibrin is involved in coagulation cascade

Table 3.1. Summary of the hydrogels and crosslinking method used.

Abbreviation	Material	Crosslinking
MVG	Sodium alginate	100mM CaCl ₂
VLVG	Sodium alginate	175mM CaCl ₂
Fibrin	Fibrin + Thrombin	–
FibrinVLVG	Composite (VLVG alginate + fibrin, 1:1)	5, 50, 100, 175mM CaCl ₂ *

* Concentration dependent on experiment

MVG = Medium viscosity guluronate (222.000 g/mol; guluronic acid (GA) > 60%; 0.4% DOS RGD peptide)

VLVG = Very low viscosity guluronate (32.000 g/mol, GA > 60%, 0.376% DOS RGD peptide)

Astrocyte Markers

GFAP

- Glial Fibrillary Acidic Protein
- Cytoskeletal astrocyte marker
- Upregulated in reactive astrocytes and pathological conditions (Miller, 2018)
- Particularly prominent in fibrous astrocytes (Jurga, 2021)

SOX9

- SRY (sex determining region Y)-box transcription factor 9
- Nuclear transcription factor
- Pan-astrocyte marker, generally markers astrocytes in the CNS (Jurga, 2021) – labels almost all astrocytes

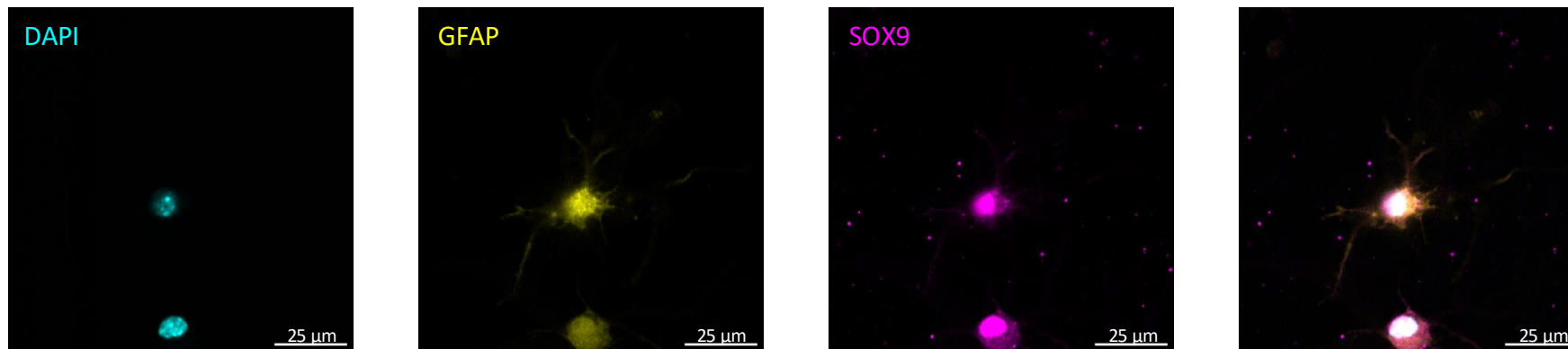


Figure 3.2. Astrocyte markers.

Validation of Alginate and Fibrin-Alginate Hydrogels

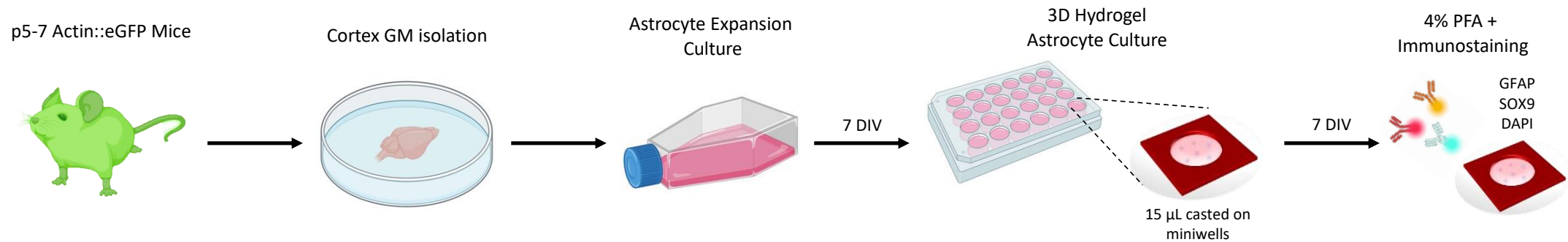


Figure 3.3. Procedure for the study of morphology and astrocyte marker expression in alginate and fibrin-alginate hydrogels.

Experimental conditions:

- 0.5% MVG – 100 mM CaCl_2
 - 0.5% VLVG – 175 mM CaCl_2
 - Fibrin + 0.5% VLVG (FibrinVLVG) – 175 mM CaCl_2
 - PDL (2D control)
- } 10^6 cells/mL
 → 0.25×10^6 cells/mL

Experiment:

- 3 biological replicates
- 2-4 technical replicates each

Morphology Categorization

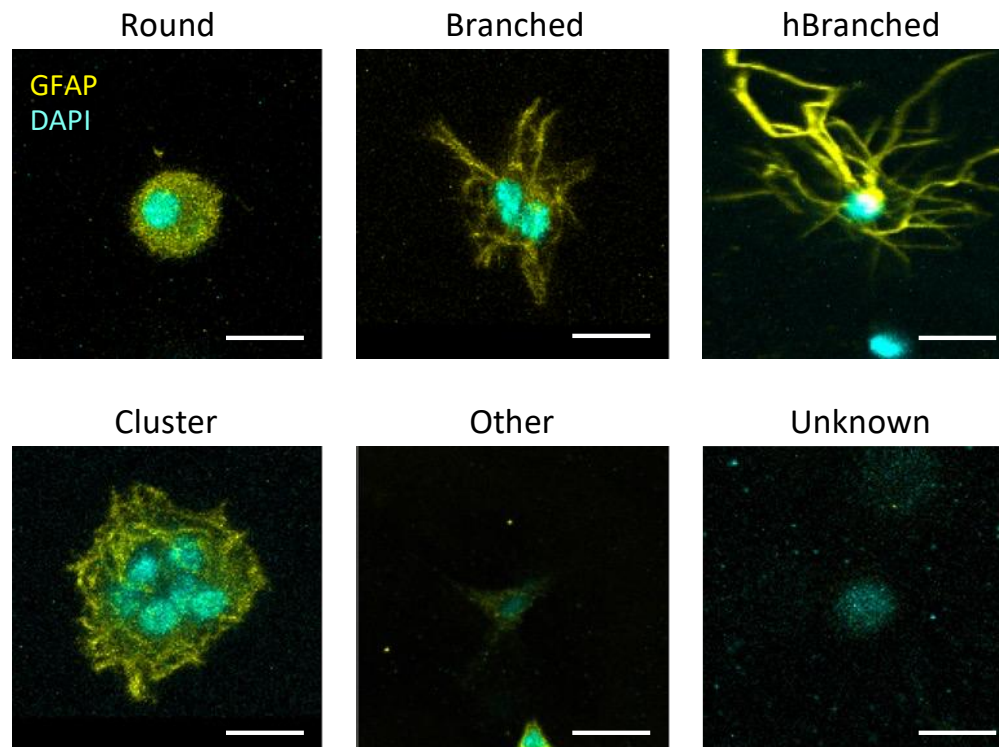


Figure 3.4. Representative cells of each morphology class.

Round cells: No branching

Branched cells: Some branching – cells with small branches and/or only 1 or 2 large branches, or branches that expand concentrically rather than radially

hBranched cells: Highly branched – cells with at least 3 large branches expanding radially

Clusters: cell agglomerates with at least 3 nuclei and no specific morphology

Other: Cells which morphology does not fit in any of the descriptions above

Unknown: Cells that are not stained with any cytoplasmic marker – impossible to categorize morphology

Result: FibrinVLVG hydrogels support **in vivo-like morphologies** and **increased GFAP⁺ reactive astrocytes**

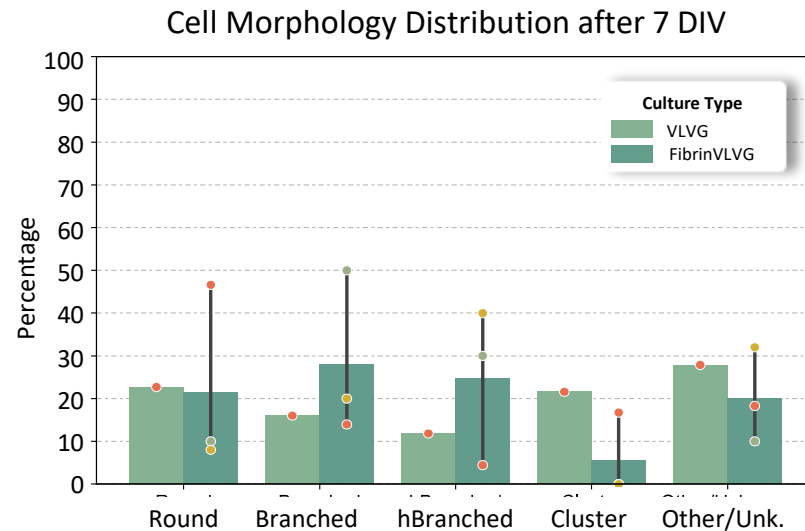


Figure 3.5. Distribution of cell morphology in VLVG and FibrinVLVG after 7 DIV.

- FibrinVLVG have **less clusters** than VLVG hydrogels (22% versus 6%)
- FibrinVLVG have **more branched cells** (53% versus 38%)

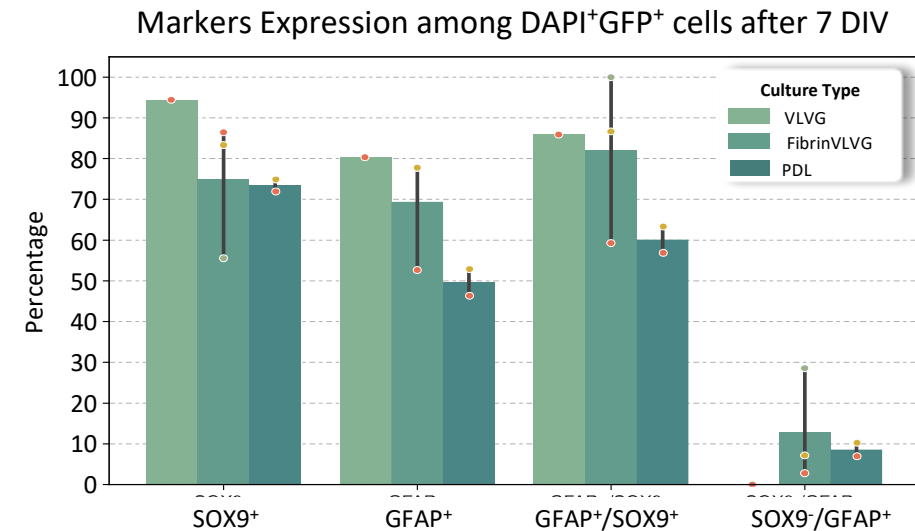


Figure 3.6. Distribution of astrocyte marker expression in VLVG and FibrinVLVG after 7 DIV.

- Astrocyte proportions, **SOX9⁺ (%) exceed 70%**
- Hydrogels show **higher GFAP⁺/SOX9⁺ reactive profiles** (*t*-test, *p* = 0.15)

Effects of Different CaCl_2 Crosslinking Concentrations

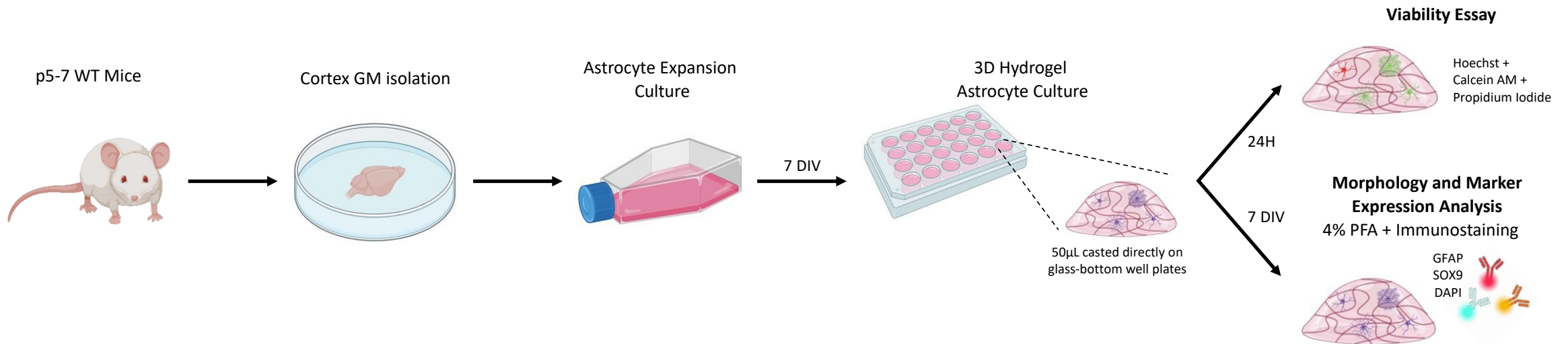


Figure 3.7. Procedure to study the effects of CaCl_2 concentrations used for hydrogel crosslinking.

Experimental conditions:

- FibrinVLVG crosslinked with 5mM, 50mM, 100mM and 175mM CaCl_2
 - Fibrin (0mM CaCl_2)
 - PDL
- } 10^6 cells/mL
- 0.25×10^6 cells/mL

Independent experiments:

1. Viability essay:
 - 3 biological replicates
 - 2-4 technical replicates each
2. Morphology and marker expression analysis
 - 5 biological replicates for each condition
 - 2 technical replicates

Result: Different CaCl_2 concentrations **do not impact** cell viability on FibrinVLVG hydrogels

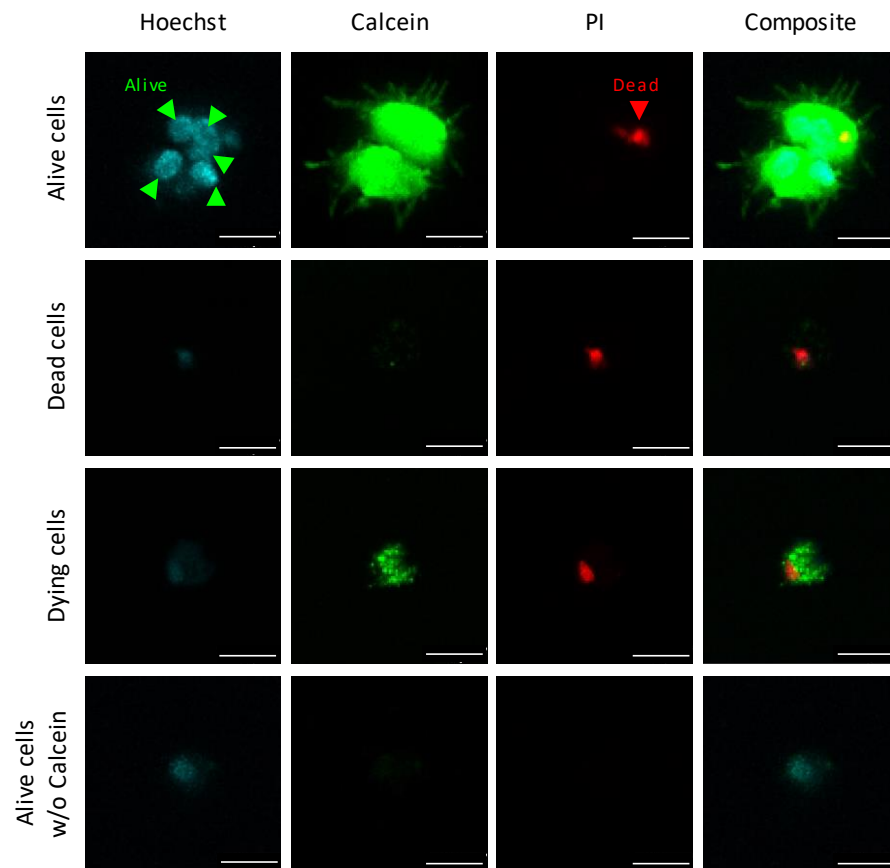


Figure 3.8. Representative cells of each viability class.

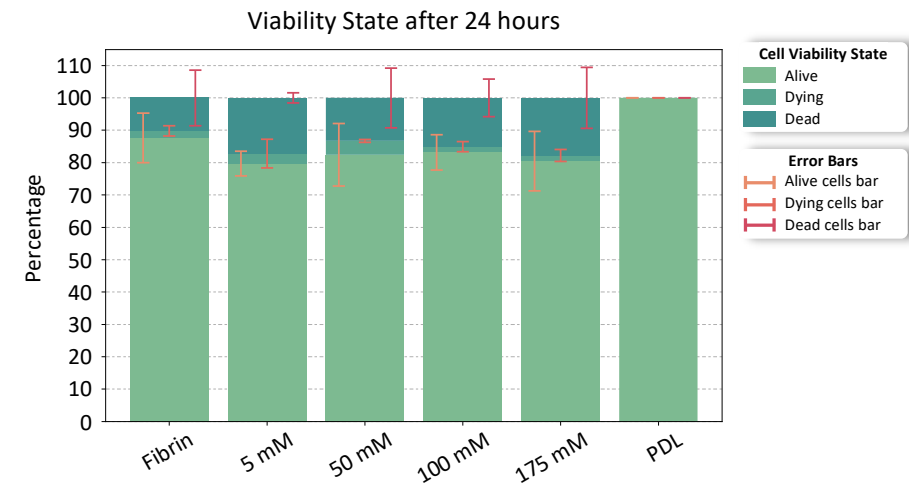


Figure 3.9. Viability assay analysis on VLVG, FibrinVLVG hydrogels and monolayer cultures.

- **No correlation** between calcium levels and proportion of live or dead cells (Pearson and Spearman tests)
- Hydrogel cultures **might have increased cell stress** comparing to monolayer cultures

Result: Different CaCl_2 concentrations **impact** cell morphology and astrocyte marker expression

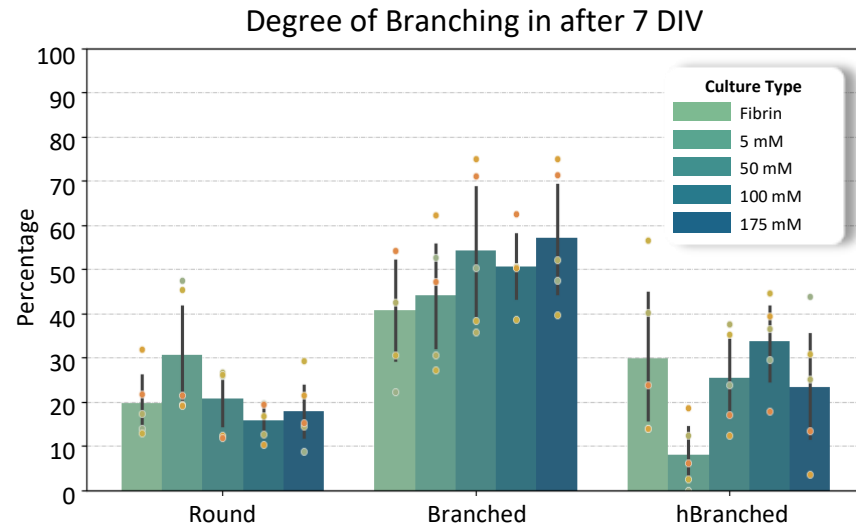


Figure 3.10. Effects of CaCl_2 concentration on cell morphology after 7 DIV.

- **Higher Ca^{2+} concentrations** show tendency to exhibit **more branched cells** ($r = 0.32$, $p = 0.12$)
- **Possibly Ca^{2+} increases stiffness** of the hydrogels allowing more branched structures

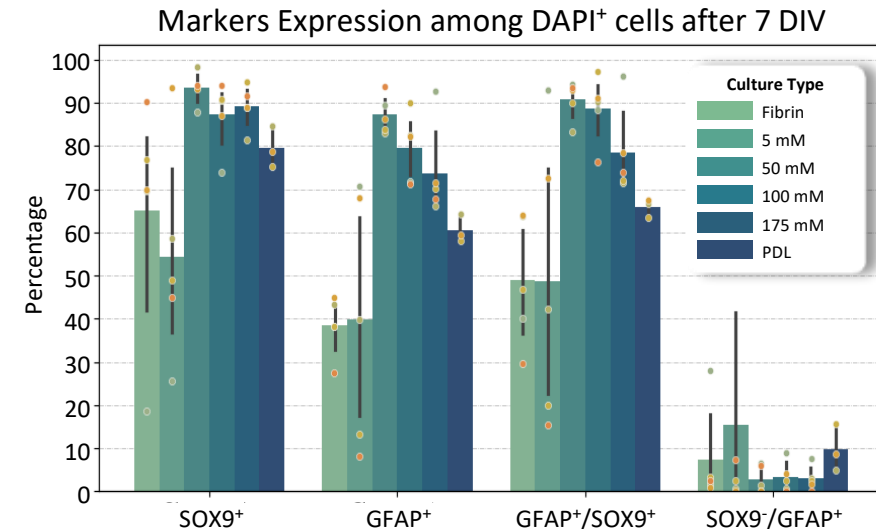


Figure 3.11. Effects of CaCl_2 concentration on astrocyte marker expression after 7 DIV.

- Fibrin and 5mM show high variation and low reliability (excluded from evaluation)
- **Higher Ca^{2+} concentration** used for gelation generally leads to **less GFAP⁺ expression** ($r = -0.6$, $p = 0.02$)
- Hydrogels show **higher GFAP⁺/SOX9⁺ reactive profiles** (t -test, $p = 0.02$)

04 Conclusion

Conclusions



Experimental validation of hydrogel matrices

- ❖ FibrinVLVG hydrogels (50 – 175 mM CaCl_2) stable for 7 DIV cultures
- ❖ Elevated CaCl_2 concentrations did not decrease cell viability
- ❖ CaCl_2 concentration **tendency to increase cell branching** (no significant effect)
- ❖ Higher CaCl_2 concentrations produced **less GFAP⁺ reactive** profiles



Computational solutions to nuclei segmentation and assisted cell classification

- ❖ High performing nuclei segmentation model – error rate <5% on hydrogel images
- ❖ GUI for z-stack image annotation targeting cell nuclei

Limitations and Future Work

EXPERIMENTAL WORK

Limited understanding of hydrogel properties

- ❖ Interactions scaffold-substrate
- ❖ Hydrogel physical and chemical properties

Limited understanding of astrocyte cultures

- ❖ Phenotypic studies (e.g. RNA-seq, gene expression)
- ❖ Cell function (e.g. motility, calcium imaging)
- ❖ Increase biological replicates

COMPUTATIONAL WORK

Scarcity of annotated data and Limited generalizability

- ❖ Fine-tuning pretrained models with annotated data
- ❖ Increase dataset size and diversity

Limited GUI versatility

- ❖ Real time cell position prediction
- ❖ Configurable classification labels

Limited cell characterization

- ❖ Analysis of nuclei properties
- ❖ Automatic morphology classification

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Thank you!

Acknowledgements:

Supervisors:

Prof. Rui Henriques
Dr. Thomas Distler

Examination Committee:

Prof. João Sanches
Prof. Rui Henriques
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Götz Lab:

Dr. Magdalena Götz
Dr. Giacomo Masserdotti
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Daniela Cimino
Maroussia Hennes

Marr Lab:

Dr. Amirhossein Kardoost
Dr. Carsten Marr

Maria Beatriz Sereno Fernandes

March 2025