

Role of extracellular matrix in age-related declines of muscle regeneration

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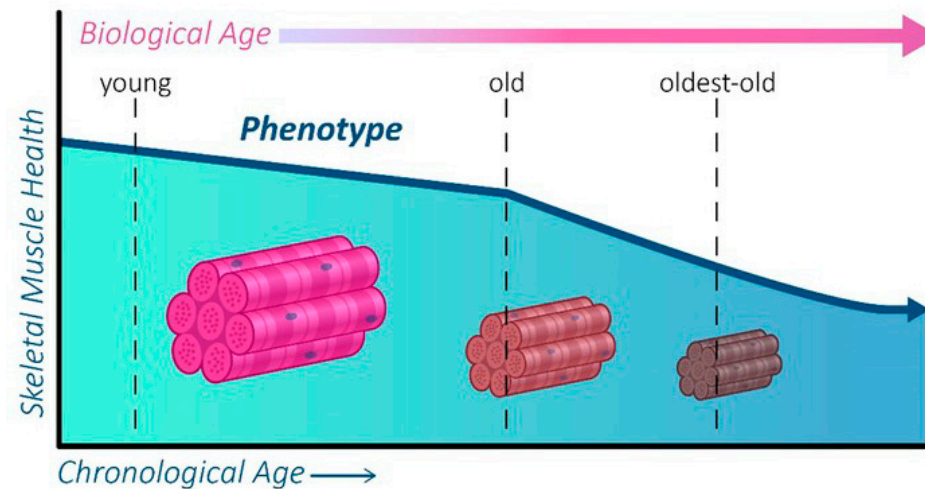
Biochemistry of Aging

- The hallmarks of aging are widely known.
- However, how this takes place on a chemical level is only partially understood.
- We can use biochemistry and bioinformatics to develop a mechanistic understanding of aging.



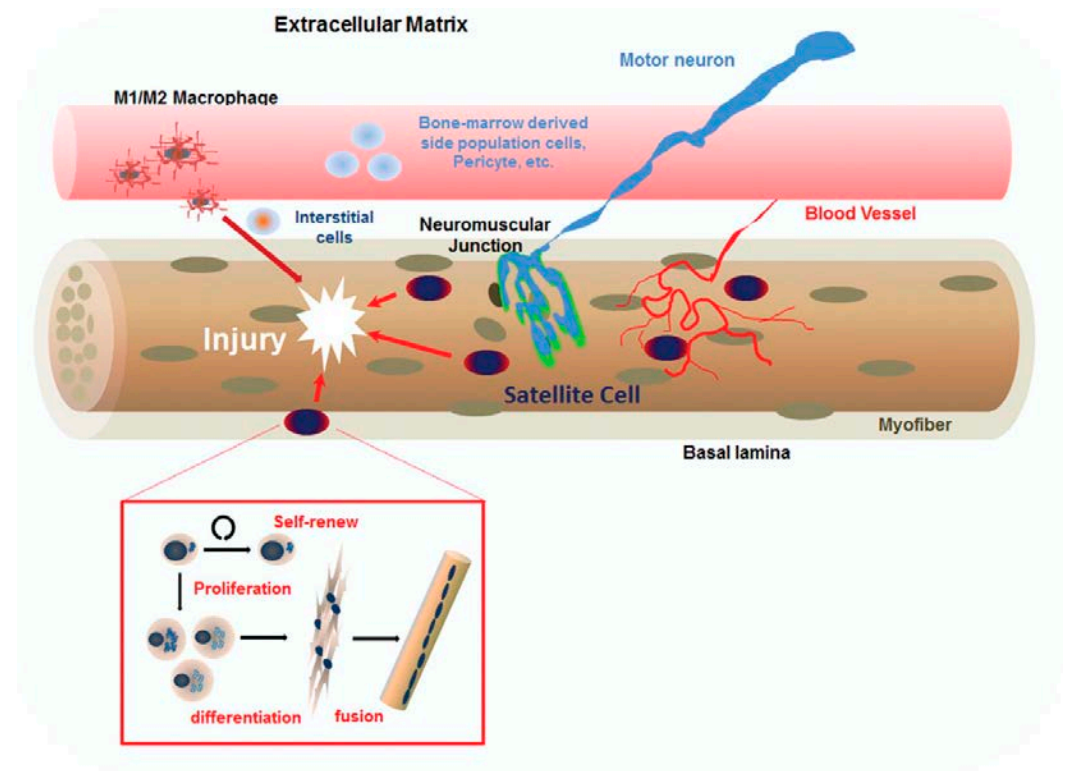
Aging effects on Skeletal Muscles Regeneration

- Loss of mass and function known as sarcopenia.
- Impaired ability to regenerate after injury due to deficiency in the number and functionality of stem cells.
- Satellite cell function is controlled by both intrinsic and extrinsic regulatory cues, whose integration determines the success of muscle regenerative responses.



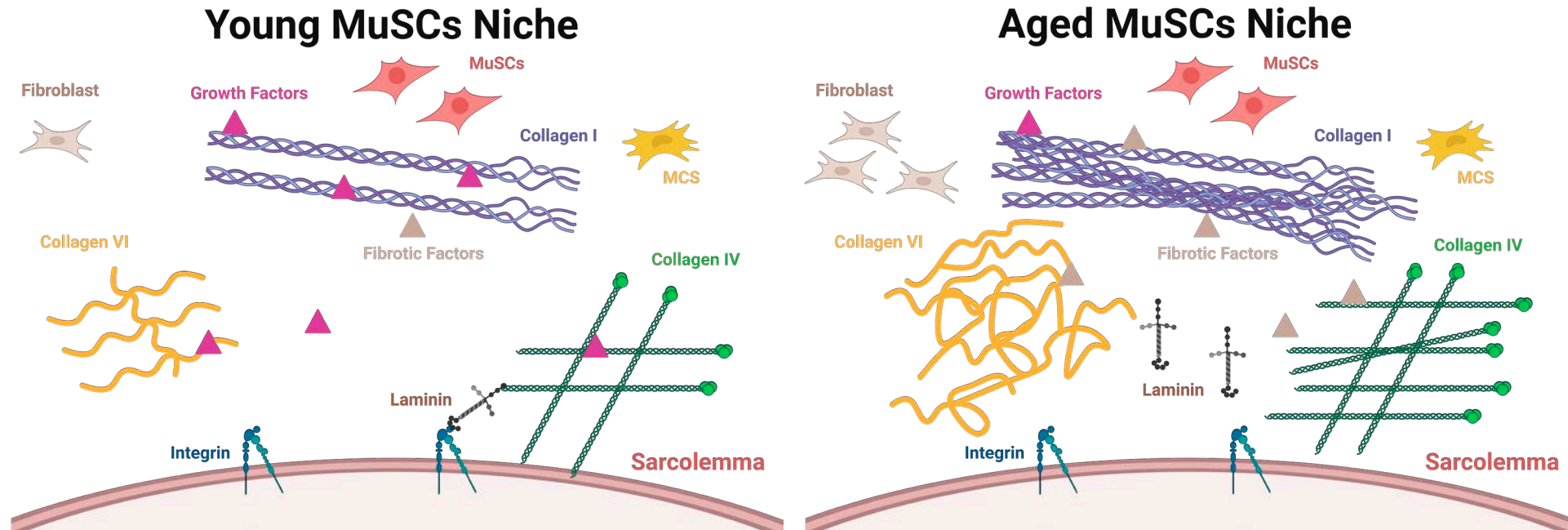
Intrinsic and Extrinsic Regulation of Muscle stem cells (MuSCs)

- The intrinsic factors include damage of key cellular macromolecules, such as proteins, lipids, and nucleic acids.
- The extrinsic factors affecting MuSCs describe MuSCs niche which includes:
 - The extracellular matrix (ECM)
 - The muscle fiber itself.
 - Motor neurons.
 - Inflammatory cytokine and cells.
 - Fibro/adipogenic precursor cells, and other nonhematopoietic cell types.



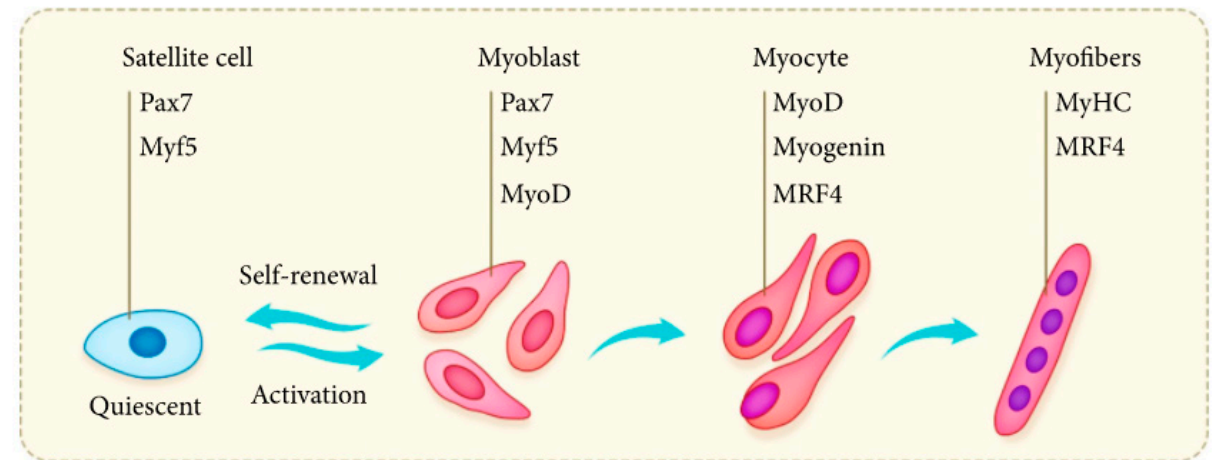
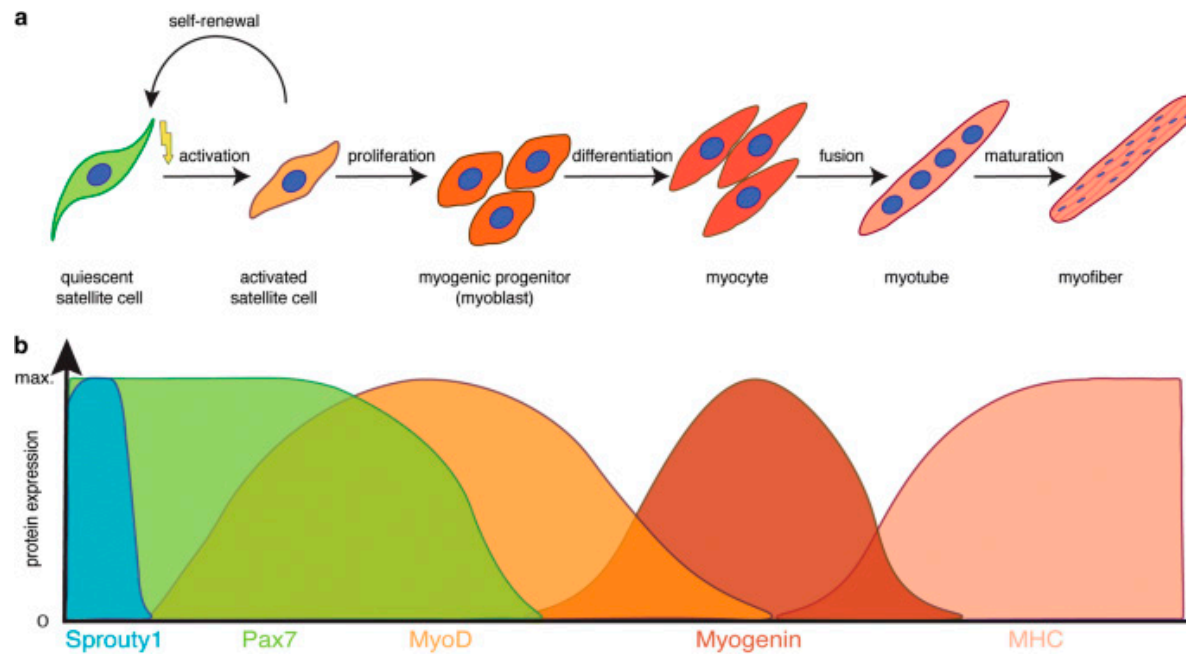
Aging Effects on Muscle Stem Cells (MuSCs) Niche

- Aging causes increased extracellular matrix (ECM) stiffness which hinders skeletal muscle contractibility and affects the regeneration of the neighboring muscle stem cells (MuSCs) post injury.
- These effects are reversible if the MuSCs were transferred to a soft environment, but the molecular mechanisms of this remain poorly understood.



Differentiation of Muscle Stem Cells in Healthy Young Mice

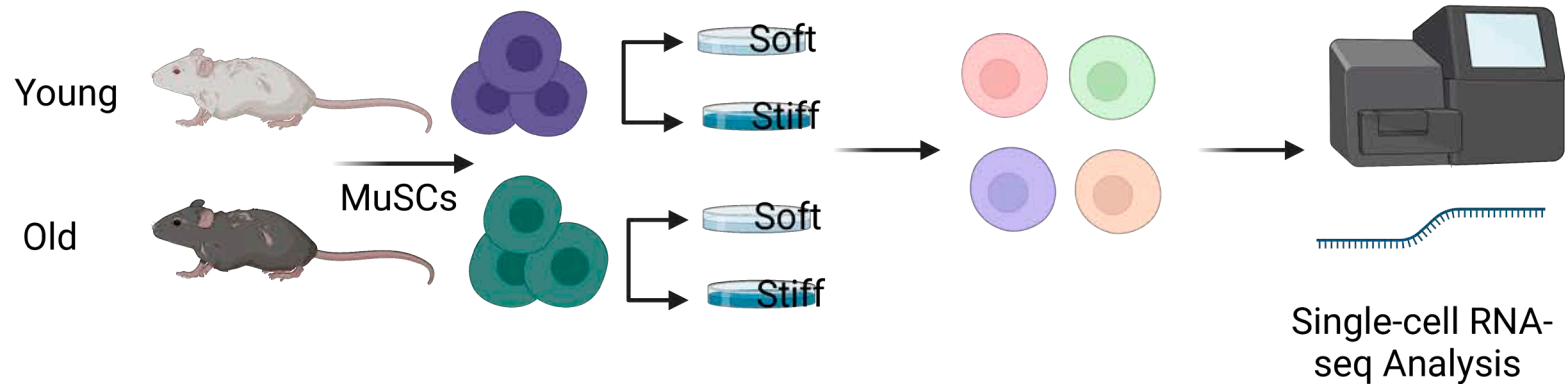
- Stages of differentiation is dictated by time-dependent gene expression patterns.
- What happens to aged MuSCs in stiff environments?



Experimental Design for Harvesting MuSCs

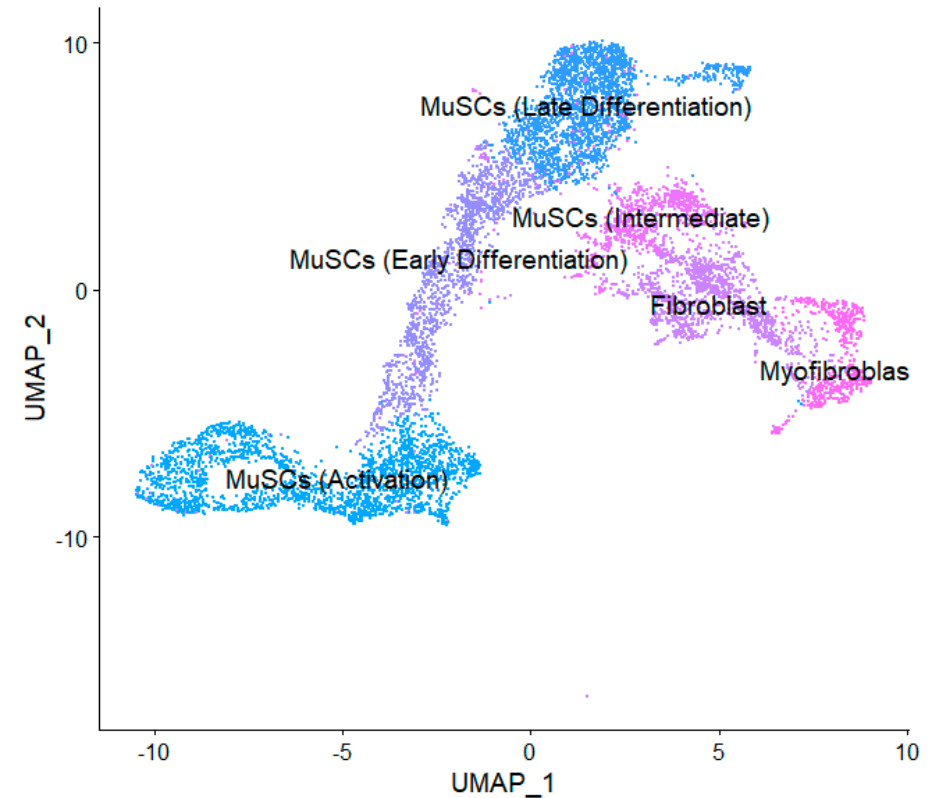
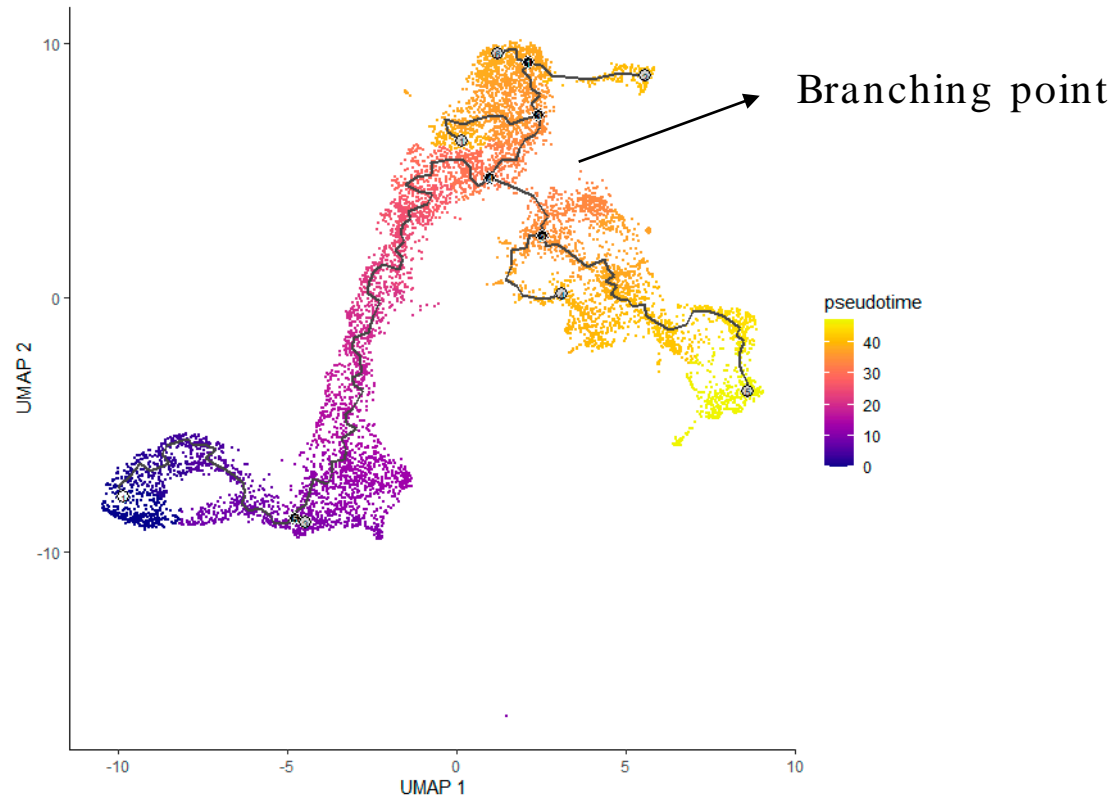
MuSCs derived from both aged and young mice were planted on fabricated silicone-based organic polymers (PDMS) substrates to mimic the stiffness of young (elastic modulus (E): 12 kPa) or aged muscle (E: 29 kPa).

Followingly, cells were recovered for scRNA-seq.



Lineage Branching Observed in Trajectory Analysis

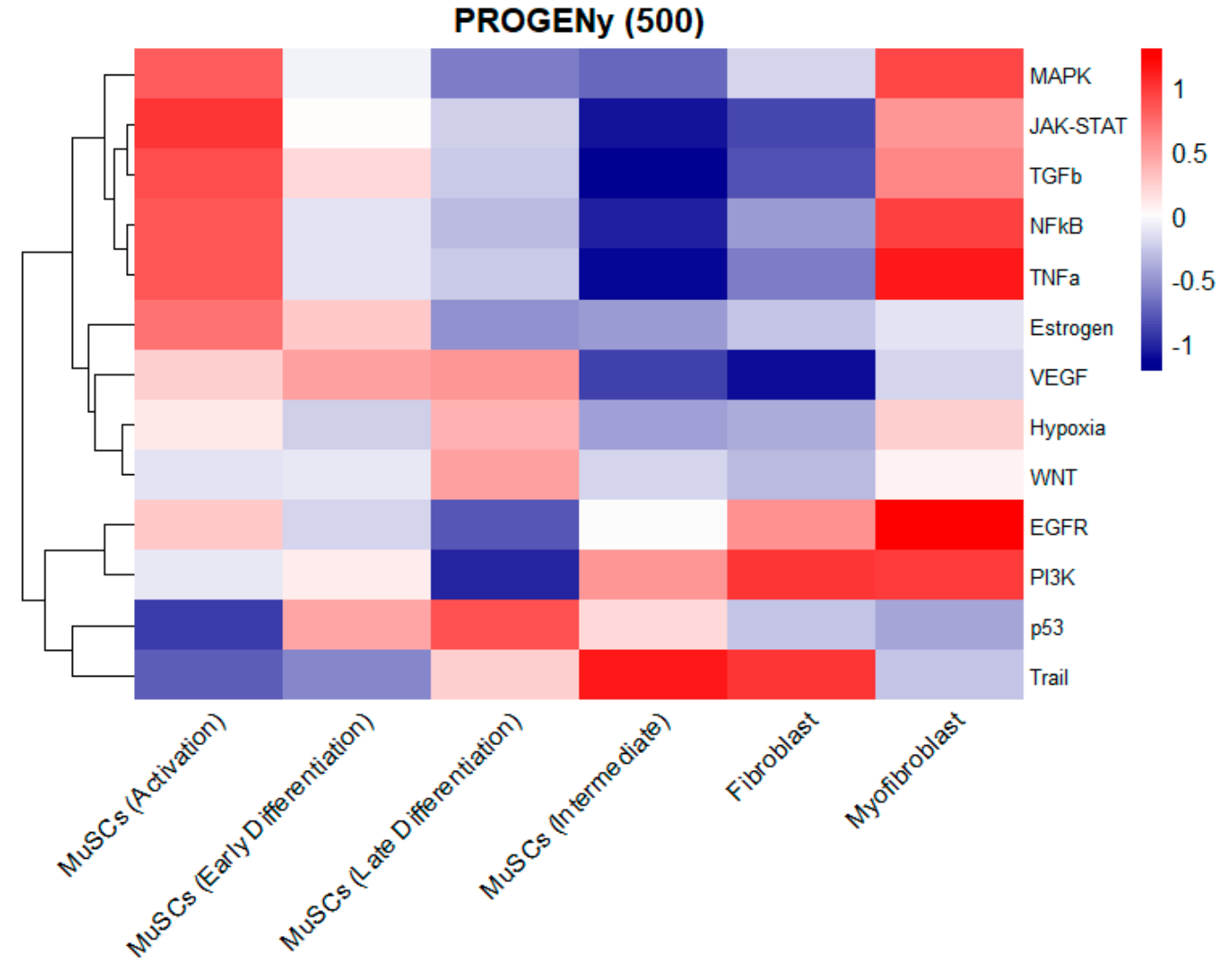
- The Monocle3 R package was used for trajectory analysis.
- Cell subpopulations are organized according to pseudo-time to infer developmental transitions.



TRAIL Activation Trend in the Fibrogenic population

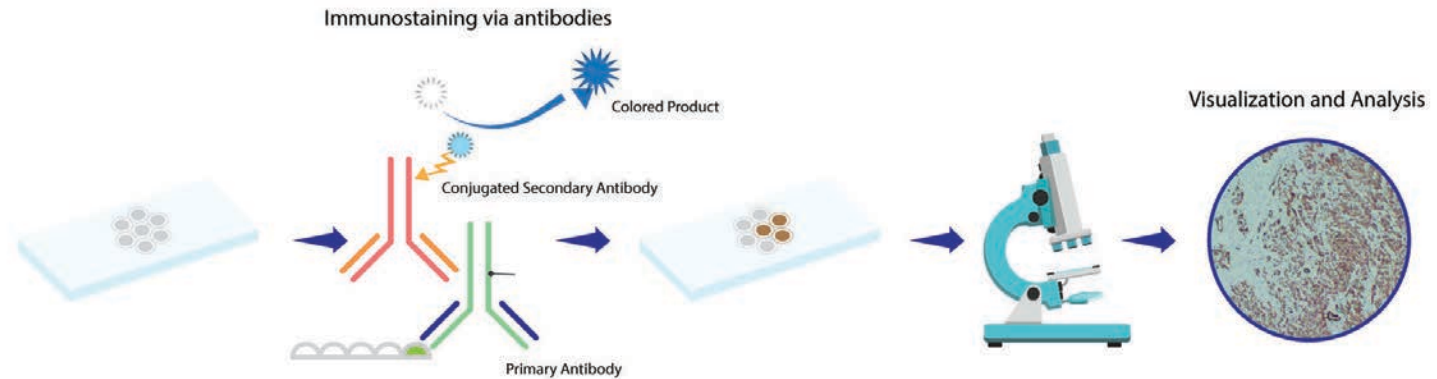
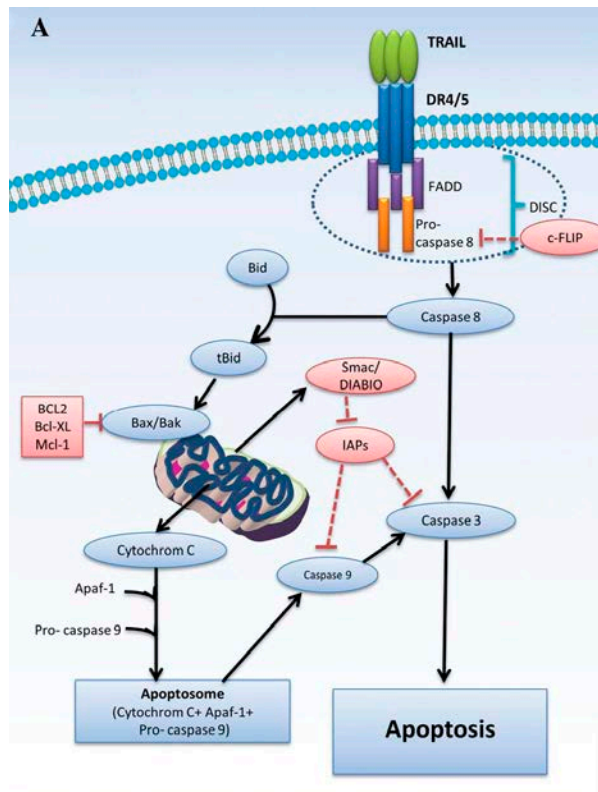
Three pathways were activated in the fibroblastic population:

1. TRAIL was following a trend along the trajectory as it was highly activated in fibroblasts followed by the MuSCs (Intermediate), and slightly activated in the MuSCs (late differentiation).
2. the Phosphoinositide 3-kinases (PI3K) pathway that inhibits apoptosis by activating Akt.
3. Epidermal Growth Factor Receptor (EGFR) that mediates apoptosis through Stat3.



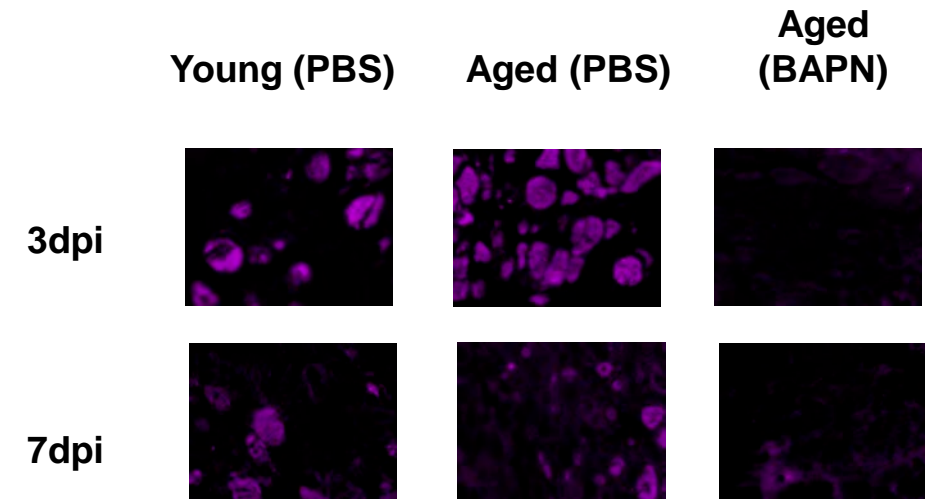
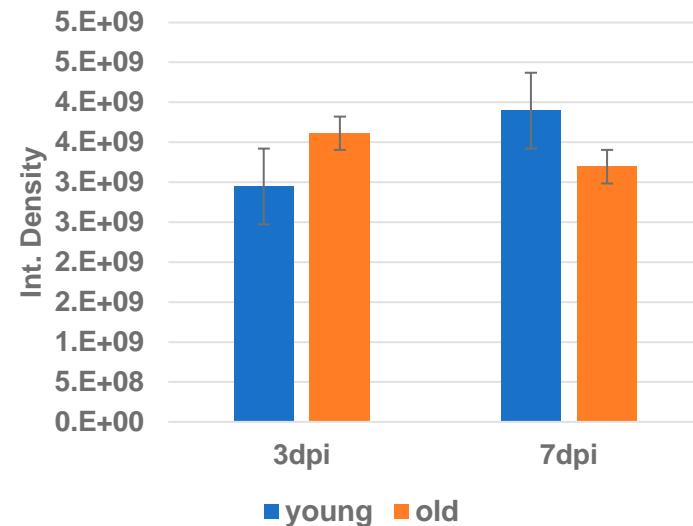
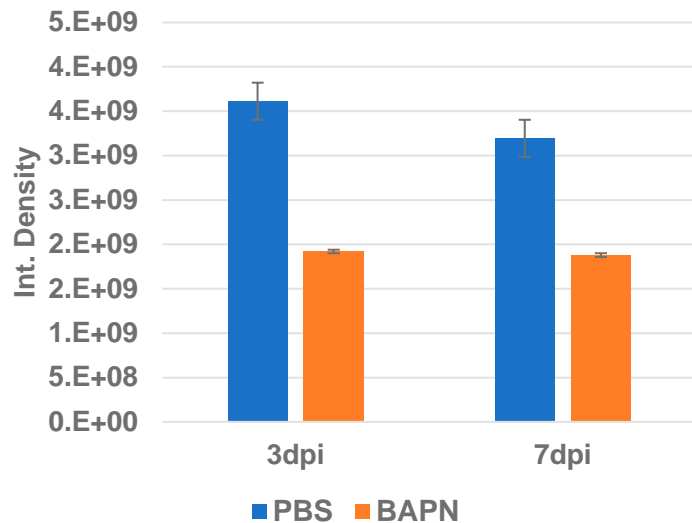
In-vivo Validation with TRAIL Immunostaining

- Caspase-3, a TRAIL downstream protein associated with cell apoptosis was stained and quantified.



In-vivo Validation with TRAIL Immunostaining

- Whole tissues were harvested from young and aged animals.
- β -aminopropionitrile (BAPN), a collagen anti-crosslinking agent used to reverse stiffness, was applied to aged tissues mimicking the soft tissues environment and then those tissues were also stained for caspase-3.
- Caspase-3 was found to be downregulated in aged animals treated with BAPN, suggesting a connection between aging, fibrogenic conversion, matrix stiffness, and apoptosis



Conclusions and Acknowledgments

Single-cell RNA sequencing data were analyzed from muscle-derived stem cells

A novel mechanism for fibrogenic conversion was identified that involved the inflammatory mediator TRAIL

This mechanism was validated using immunohistochemical staining of muscle tissue sections

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