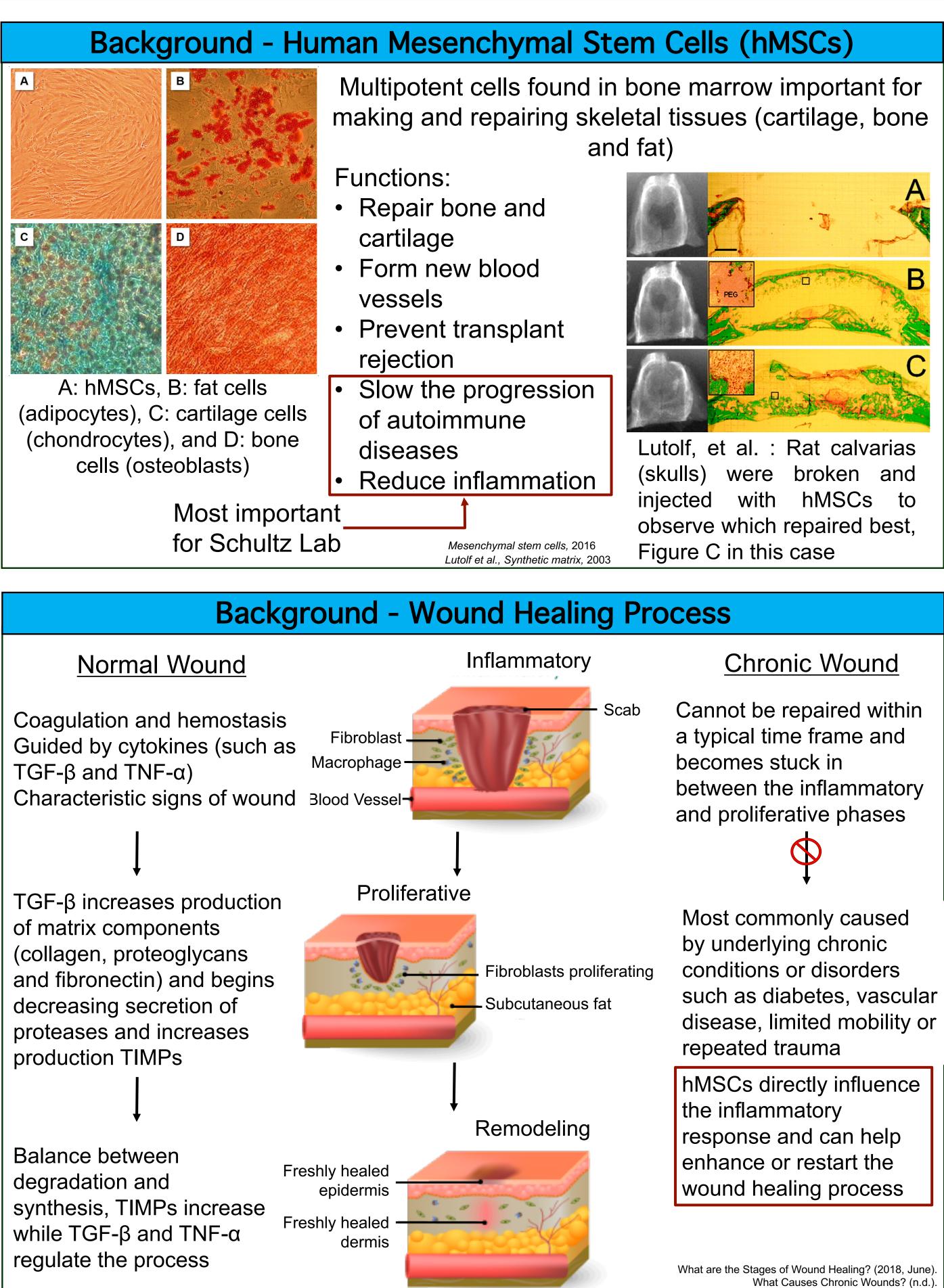
Interaction of Human Mesenchymal stem cells with synthetic scaffolds in the presence of cytokines Hannah E. Knudsen, Jenna A. Catalano, John A. McGlynn, and Kelly M. Schultz Department of Chemical and Biomolecular Engineering, Lehigh University, Bethlehem PA, 18015



Abstract

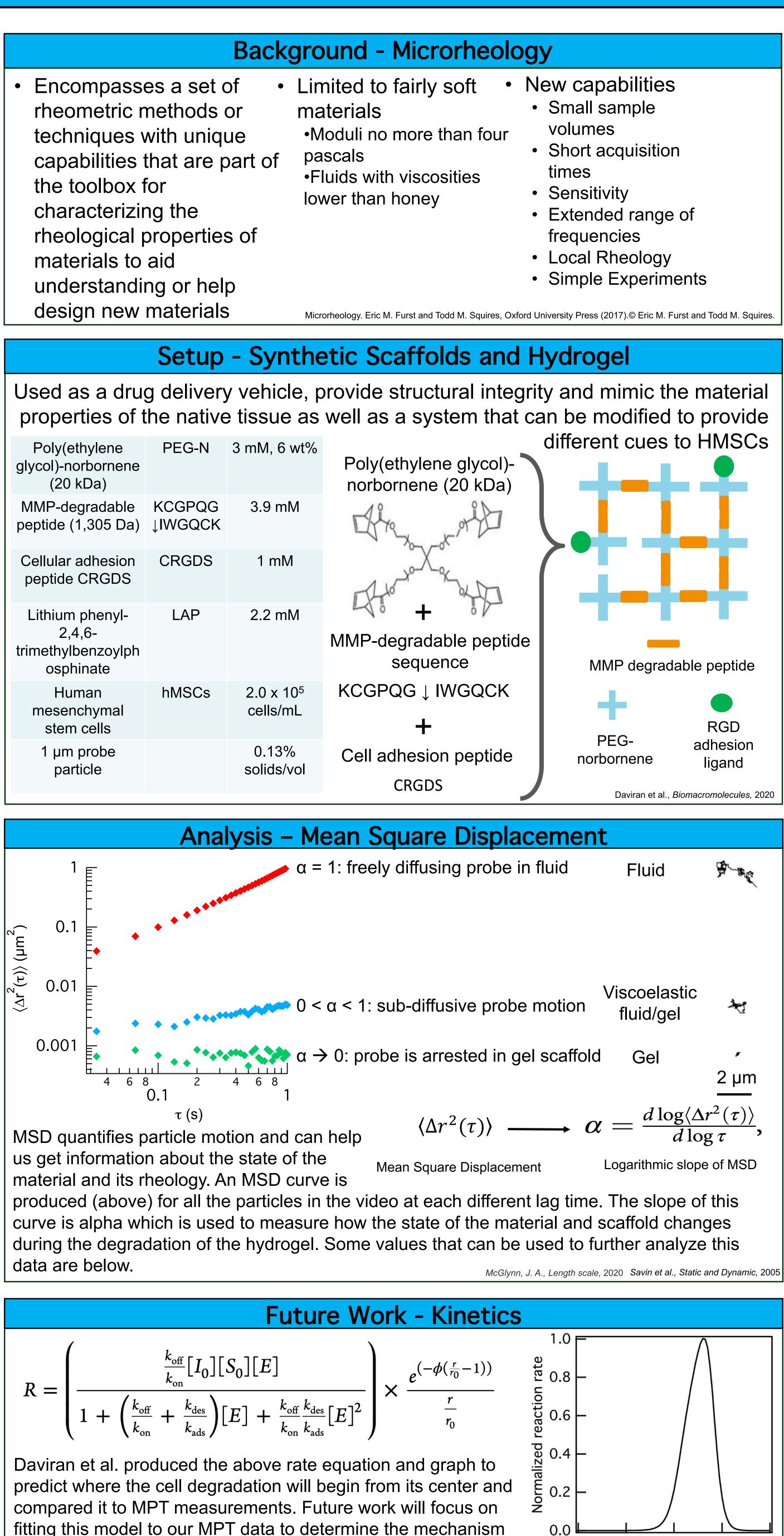
A normal wound heals by progressing through three phases: inflammatory, proliferative and remodeling. From previous research, we know human mesenchymal stem cells (hMSCs) are key to progressing healing through all stages and can prevent and help resolve chronic wounds, which are stuck in the inflammatory phase. hMSCs directly influence the body's inflammatory response by decreasing the amount of pro-inflammatory cytokines and increasing the number of anti-inflammatory cytokines. Cytokines are proteins that act as signaling molecules to evoke biological processes, and include tumor necrosis factor alpha (TNF- α) and transforming growth factor beta (TGF- β). When secreted by a wound, these cytokines can elicit a response that regulates inflammation at the site by increasing cell-secretion of matrix metalloproteinases (MMPs) or tissue inhibitor of metalloproteinases (TIMPs). MMPs degrade the extracellular matrix (ECM) while TIMPs inhibit MMP degradation. In the Schultz Lab, we are characterizing implantable cell-laden materials that can direct cells to a wound using natural cues. We are characterizing poly(ethylene glycol)-peptide hydrogels that mimic the *in vivo* environment inside the body with multiple particle tracking microrheology (MPT). MPT measures the change in matrix degradation and cell motility of the cells after the addition of cytokines in the incubation environment. We are also developing a kinetic model for TGF- β using Michaelis-Menten enzymatic inhibition kinetics which will describe the mechanism of gel degradation. The results from MPT and the kinetic model can then be compared to determine the mechanism of matrix degradation and future scaffold design that will guide cell behavior during wound healing.



Maxson et al.. Concise review: role of mesenchymal stem cells in wound repair, 2012

	Ba	ckg	round - N		
rheometric techniques capabilities the toolbox characteriz rheological materials te	zing the I properties of o aid ding or help		Limited to fa materials •Moduli no m pascals •Fluids with v lower than he		
Setup - Synthetic Sca					

Poly(ethylene glycol)-norbornene (20 kDa)	PEG-N	3 mM, 6 wt%	Poly(ethy norborne
MMP-degradable peptide (1,305 Da)	KCGPQG ↓IWGQCK	3.9 mM	They a
Cellular adhesion peptide CRGDS	CRGDS	1 mM	The to to
Lithium phenyl- 2,4,6- trimethylbenzoylph osphinate	LAP	2.2 mM	MMP-degra sec
Human mesenchymal stem cells	hMSCs	2.0 x 10 ⁵ cells/mL	KCGPQG
1 µm probe particle		0.13% solids/vol	Cell adhe CR



	Future We	ork - Ki
R =	$\left(\frac{\frac{k_{\text{off}}}{k_{\text{on}}}[I_0][S_0][E]}{1 + \left(\frac{k_{\text{off}}}{k_{\text{on}}} + \frac{k_{\text{des}}}{k_{\text{ads}}}\right)[E] + \frac{k_{\text{off}}}{k_{\text{on}}}\frac{k_{\text{des}}}{k_{\text{ads}}}[E]^2}\right)$	$\times \frac{e^{(-\phi(\frac{r}{r_0})}}{\frac{r}{r_0}}$

of matrix degradation.

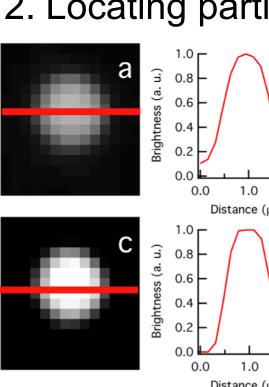
 r/r_{o}

Method - Multiple Particle Tracking Microrheology (MPT)

. Filter noise in images

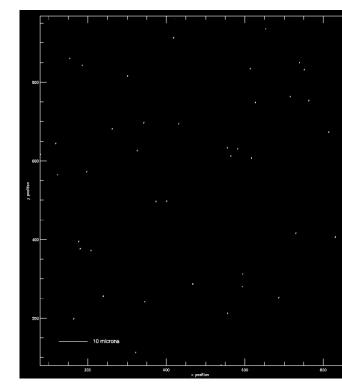
Neighboring pixels will be averaged together and background gradients are removed to achieve uniform illumination which will remove potential bright spots that would have previously been $A_{\xi_n}(x, y) =$ identified as a particle.

2. Locating particles in images



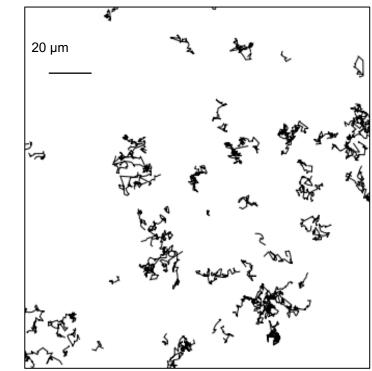
Particles are identified based on $\begin{pmatrix} \boldsymbol{\varepsilon}_x \\ \boldsymbol{\varepsilon}_y \end{pmatrix} = \frac{1}{m_0} \sum_{i^2+j^2 \leq w^2} \binom{i}{j} A(x+i,y+j),$ brightness and distance between others. Only the top 30-40% of particles are selected to avoid darker Offset Distance pixels being selected just because $R_g^2 = \frac{1}{m_a} \sum_{i=1}^{n} (i^2 + j^2)A(x + i, y + j).$ they are bright. Eccentricity and radius of gyration are calculated to Squared Radius of Gyration exclude particles that may not be the correct size or not particularly circular.

3. Refining particle positions



Positions connected to form trajectories using a probability distribution function that describes Brownian motion. To limit the particles that swap positions, the particle separation distance should be greater than the distance the particle moves between each frame. Particles that disappear and reappear between frames are taken into account by specifying the max amount of frames in a row that a particle will be considered different if it reappears. Trajectories of particles can also be rejected if the particle does not appear for enough frames in a row.

4. Linking positions into trajectories



The algorithm creates a text file with the particles x and y coordinates, brightness, radius of gyration squared, eccentricity, frame number and particle ID.

. Able to understand how cells degrade our material and

- change the way the cells move
- 2. Able establish design rules so hMSC scaffolds can last longer in the body and correctly deliver cells to a wound

This research is important in the field of biomaterials to evolve biological systems by mimicking the body's environment and characterizing the necessary design rules. hMSCs play an important role in each phase of the wound healing process and implantable synthetic hMSCs would prevent wounds from becoming chronic.

ouis and Cona, What are Mesenchymal Stem Cells (MSCs)?, 2020

Professor Kelly Schultz, Professor Angela Brown, John McGlynn, Schultz Lab Graduate Students, David and Lorraine Freed Research Symposium

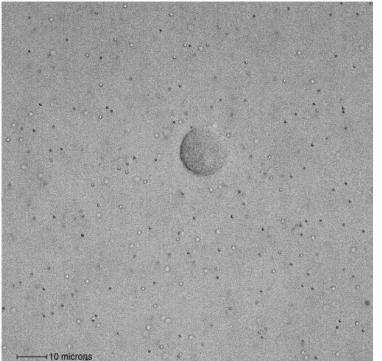
Image: Sector of the sector

MPT uses Brownian motion to extract bulk rheological properties

 $A_w(x, y) = \frac{1}{(2w+1)^2} \sum_{i,j=-w}^{w} A(x+i, y+j).$ Boxcar Average particle separation distance > W > particle radius $\sum_{i,j=-w}^{w} A(x+i, y+j) \exp\left(-\frac{i^2+j^2}{4x^2}\right)$ $\left[\sum_{i=-w}^{w} \exp\left(-i^2/4\xi^2\right)\right]$

Gaussian Average

5. Cell Centers



The cell centers from each experiment were measured by tracing the outline of the cell and used to calculate the cell center.

Muscle

Myocardium

Multiple differentiati

Conclusions

Acknowledgments