

# Exploring the Membrane Lipid Environment for Glucose Transporter 1

Lauren Strohbehn, Dr. Larry Louters, Dr. Brendan Looyenga  
Calvin College

## Background

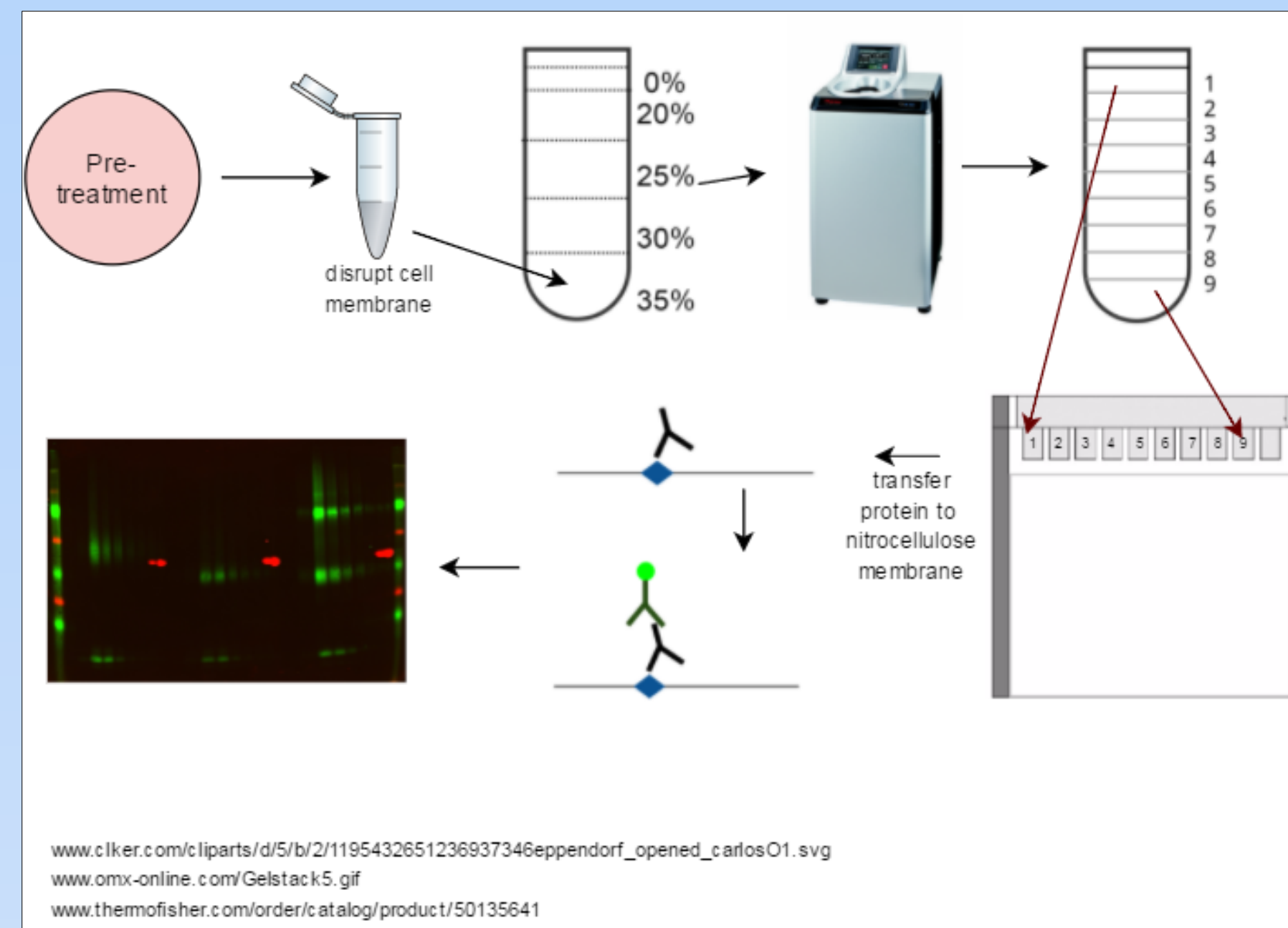
- Glucose is a preferred cellular metabolite and it requires a membrane transport protein, such as GLUT1, to enter cells.
- Abnormal glucose regulation is linked to serious diseases, including cancer, diabetes, and Alzheimer's disease. A deeper understanding of this protein may be important for developing new therapeutic strategies.
- The activity of membrane proteins are influenced by their lipid environment. This study focuses on understanding the association of GLUT1 with compact membrane microdomains called lipid rafts.
- Previous work suggests that lipid rafts play only a subtle role in the regulation of GLUT1 activity. In addition, the GLUT1-containing lipid rafts in L929/EGFP mouse fibroblast cells appeared to be an unusual type of raft.
- The purpose of my research this summer was to investigate the nature of this raft-like structure, and what targets GLUT1 to these domains.

## Objectives

- Determine the composition of the raft-like domains in L929 cells
- Determine what post-translational modifications might alter GLUT1 membrane location
- Confirm that my procedures also isolate traditional lipid rafts in other cell lines

## Methods

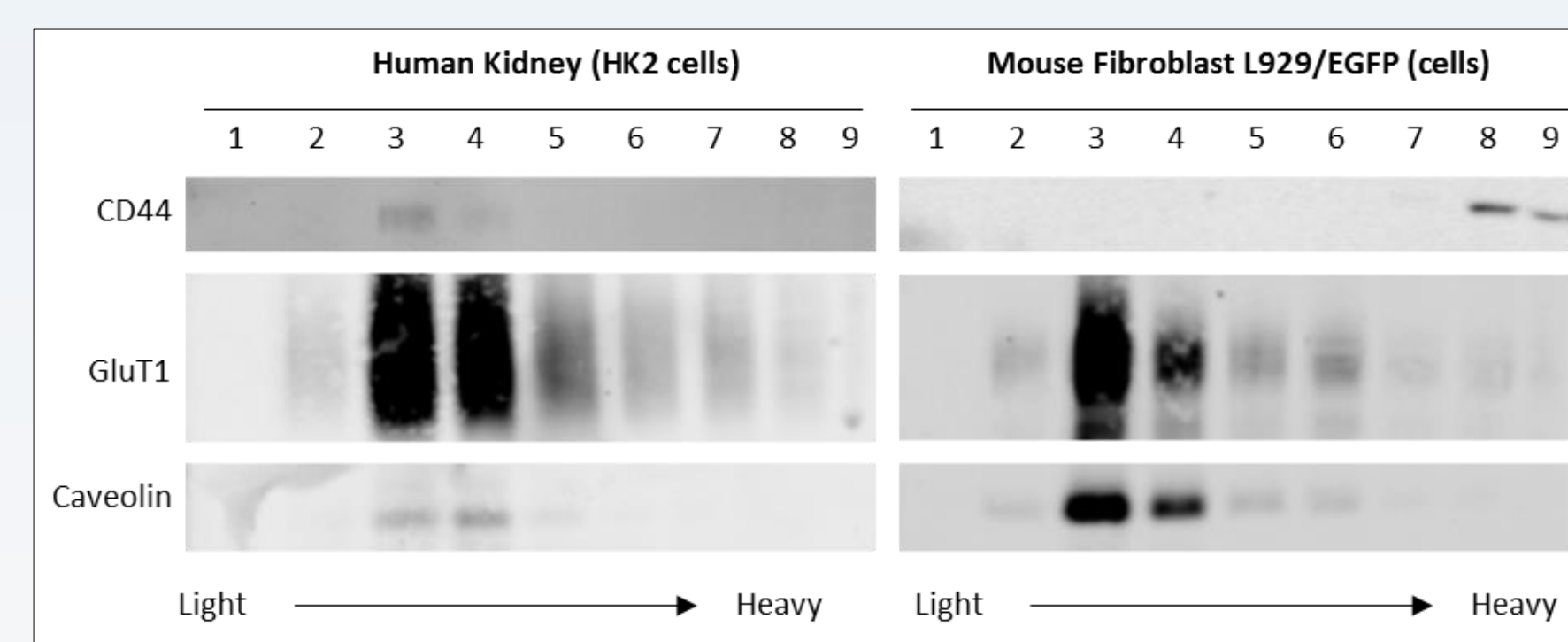
### Lipid raft isolation technique



- Cells are pre-treated
- Cells are collected and lysed
- Membrane is broken apart and loaded in an ultracentrifuge tube that has denser liquid towards the bottom
- Ultracentrifuge is spun; lipid rafts float towards the top due to the lipid that remains associated with the protein
- Western blot is performed to determine relative GLUT1 amount in each successive density fraction
- GLUT1 distribution in treatment and control is compared

## Results

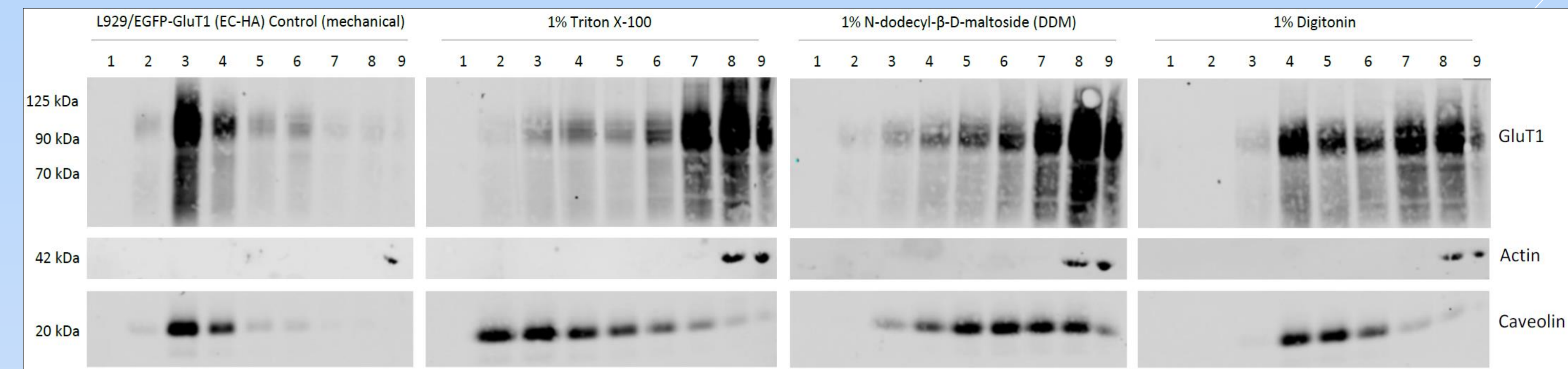
### L929/EGFP mouse fibroblast cells do not contain common raft protein markers in the lipid raft fraction



- Both GLUT1 and caveolin (a protein found in a type of low density domain) are in the raft fractions
- CD44, a common raft marker, is still found in raft fractions in HK2 cells (as expected) but pellets down in L929/EGFP cells

### Other isolation methods don't capture GLUT1 in low density fractions

Mild detergents are a common way of isolating lipid rafts. However, only mechanically disrupting the cell membrane isolates GLUT1 in low density fractions, suggesting that the low density microdomains in L929/EGFP cells are of unique composition.



### Attempts to cause GLUT1 shift in order to investigate composition of domains, or determine what targets GLUT1 to these domains

Drug/treatment	Effect of drug/treatment	Implications	GLUT1 shift?
<b>Triton X-100, DDM, Digitonin</b>	Detergents; permeabilize cell membrane	Is GLUT1 in traditionally isolated lipid rafts?	<b>yes</b>
<b>Myriocin</b>	Inhibits sphingolipid synthesis	Is sphingolipid organization an essential component of L929 lipid rafts?	<b>slight</b>
<b>Sphingomyelinase</b>	Catalyzes breakdown of a type of sphingolipid	Is sphingolipid organization an essential component of L929 lipid rafts?	<b>slight</b>
<b>M-β-CD</b>	Removes cholesterol	Cholesterol helps organize traditional lipid rafts. Are L929 rafts cholesterol enriched?	<b>no</b>
<b>Nocodazole</b>	Inhibits microtubule formation	Is microtubule structure essential?	<b>no</b>
<b>Latrunculin A</b>	Disrupts actin cytoskeleton formation	Are GLUT1 low density microdomains tethered to the cytoskeleton?	<b>no</b>
<b>2-Bromopalmitate</b>	Inhibits palmitoylation	Does palmitoylation target GLUT1 to low density microdomains?	<b>no</b>
<b>Kifunensine</b>	Inhibits glycosylation	Does glycosylation target GLUT1 to low density microdomains?	<b>no</b>

## Conclusions

- GLUT1 translocation to lipid rafts is not the primary mechanism of activation
- L929 cells contain a unique type of raft-like structure. Sphingolipids (but not cholesterol or actin) seem to be important in organizing these domains.
- Neither palmitoylation nor glycosylation targets GLUT1 to L929 "lipid rafts"

## Acknowledgments

- Dr. Larry Louters
- Dr. Brendan Looyenga and lab
- Past and present Louters lab members
- Rich Huisman, Scott Prentice, and Lori Keen

## Future Directions

- Explore more ways to disrupt these microdomains in order to determine their composition
- Inhibit other post-translational modifications of GLUT1 to try and determine why GLUT1 is in these raft-like structures
- Conduct these experiments on other cell lines



Funding Source:  
National Institutes of Health  
Ken and Marcia Wierda



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