



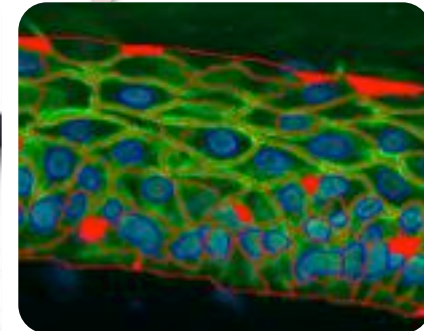
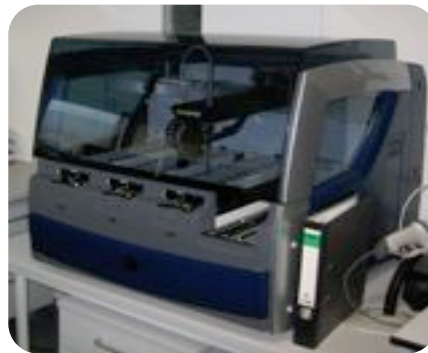
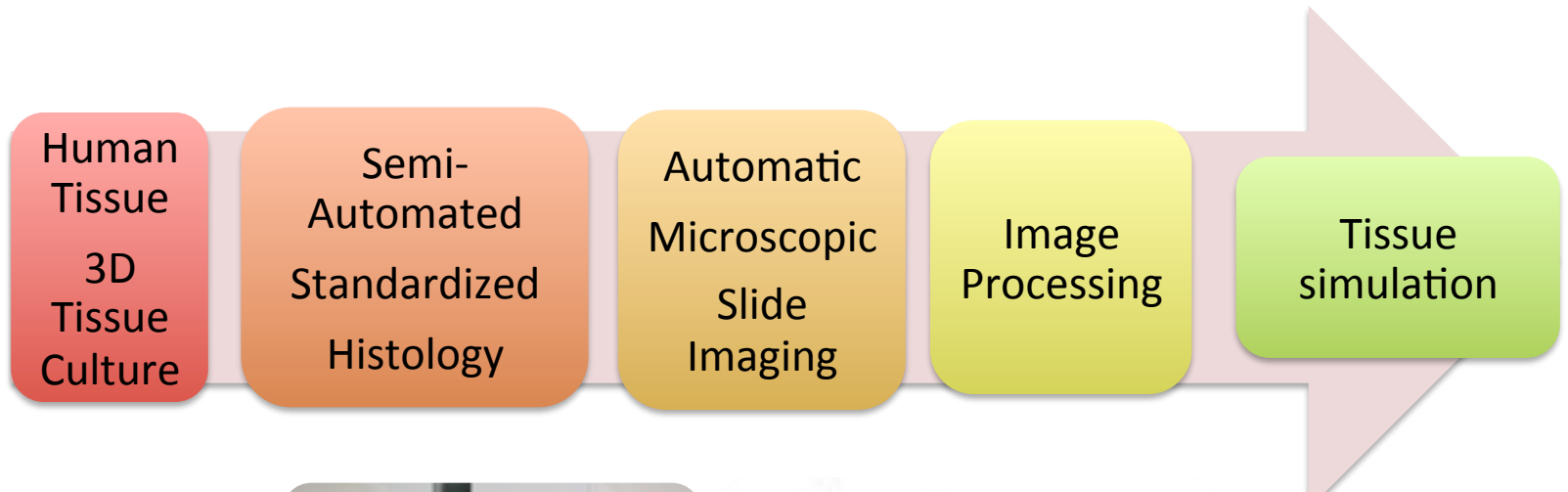
# The mechanism of wound shielding & more using WSI

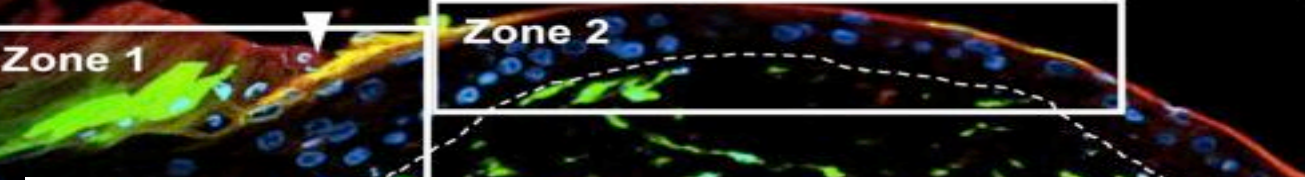
Niels Grabe

*National Center for Tumor Diseases Heidelberg (NCT) /  
Institute of Pathology, University Hospital Heidelberg*

# Specialized Pipeline:

## From Quantitative Tissue Analysis to Tissue Simulation





**JCB**

# Wound healing revised: A novel reepithelialization mechanism revealed by in vitro and in silico models

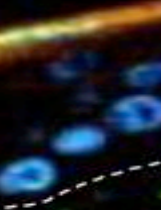
Kai Safferling,<sup>1,2</sup> Thomas Sütterlin,<sup>1,2</sup> Kathi Westphal,<sup>1,2</sup> Claudia Ernst,<sup>1,2</sup> Kai Breuhahn,<sup>3</sup> Merlin James,<sup>1,2</sup> Dirk Jäger,<sup>1,2</sup> Niels Halama,<sup>1,2</sup> and Niels Grabe<sup>1,2</sup>

<sup>1</sup>Hamamatsu Tissue Imaging and Analysis Center, BIOQUANT, and <sup>2</sup>Department of Medical Oncology, National Center for Tumor Diseases, University of Heidelberg, 69117 Heidelberg, Germany

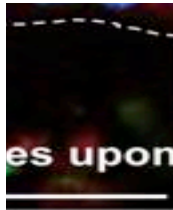
<sup>3</sup>Institute of Pathology, University Hospital of Heidelberg, 69120 Heidelberg, Germany



New CM



Zone 2



60h after green CMFDA at 0h



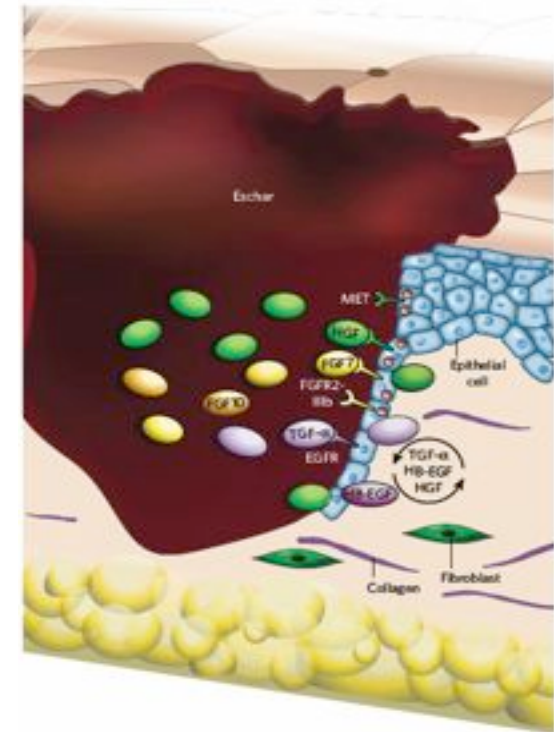
# Understanding Wound Healing is Fundamental in Skin Research

- Uncovers fundamentals of skin homeostasis
- Studies differentiation and migration
- Reveals cross-talk epidermis-dermis
- Route to cancer invasion

**Table 1 | Soluble mediators of re-epithelialization**

Ligand	Receptor	Type of receptor	Signaling proteins	Role in re-epithelialization	References
HGF	MET	Receptor tyrosine kinase	Unknown, possibly ERK1 and ERK2, AKT, GAB1, PAK1 and/or PAK2	Stimulation of keratinocyte migration and probably proliferation	43
FGF7, FGF10 and FGF22	FGFR2-IIIb, possibly FGFR1-IIIb	Receptor tyrosine kinase	Unknown, possibly ERK1, ERK2, AKT and/or STAT3	Stimulation of keratinocyte proliferation and migration	44-46
Heparin-binding EGF and other EGF-family members	EGFR (also known as ERBB1), possibly ERBB2, ERBB3 and/or ERBB4	Receptor tyrosine kinase	Unknown, possibly ERK1 and ERK2, AKT and/or STAT3	Stimulation of keratinocyte proliferation and migration	30, 47
TGF- $\beta$	TGF- $\beta$ receptor I and TGF- $\beta$ receptor II	Receptor serine/threonine kinase	SMAD3 and others, including SMAD2 and MAPK	Inhibition of keratinocyte proliferation and survival	30, 51, 52
Acetylcholine	M3 receptor	G-protein-coupled receptor	Ca <sup>2+</sup> -dependent guanylyl cyclase, cyclic GMP and PKG, leading to inhibition of RHO	Inhibition of keratinocyte migration	54
	M4 receptor	G-protein-coupled receptor	Adenylyl cyclase, cyclic AMP and PKA, leading to activation of RHO	Stimulation of keratinocyte migration	54
Catecholamines, including adrenaline	$\beta_2$ -Adrenoceptor	G-protein-coupled receptor	Activation of phosphatase PP2A, resulting in dephosphorylation and inhibition of ERK1 and ERK2	Inhibition of keratinocyte migration	55
Polysaturated fatty acids	PPAR- $\alpha$ and PPAR- $\beta$ *	Nuclear receptor	Direct activation of target genes by binding to the promoter/enhancer of these genes	Stimulation of keratinocyte migration and survival	56-58

EGF, epidermal growth factor; EGFR, EGF receptor; ERK, extracellular-signal-regulated kinase; FGF, fibroblast growth factor; FGFR1-IIIb, IIIb isoform of FGF receptor 1; GAB1, growth-factor-receptor-bound protein 2 (GRB2-associated binding protein); HGF, hepatocyte growth factor; M3, muscarinic receptor subtype 3; PAK, p21-activated kinase; PKA, cyclic-AMP-dependent protein kinase; PKG, cyclic-GMP-dependent protein kinase; PPAR, peroxisome-proliferator-activated receptor; SMAD3, SMAD-family member 3; STAT3, signal transducer and activator of transcription 3; TGF- $\beta$ , transforming growth factor- $\beta$ . \*PPAR- $\beta$  ligands might be fatty acids.

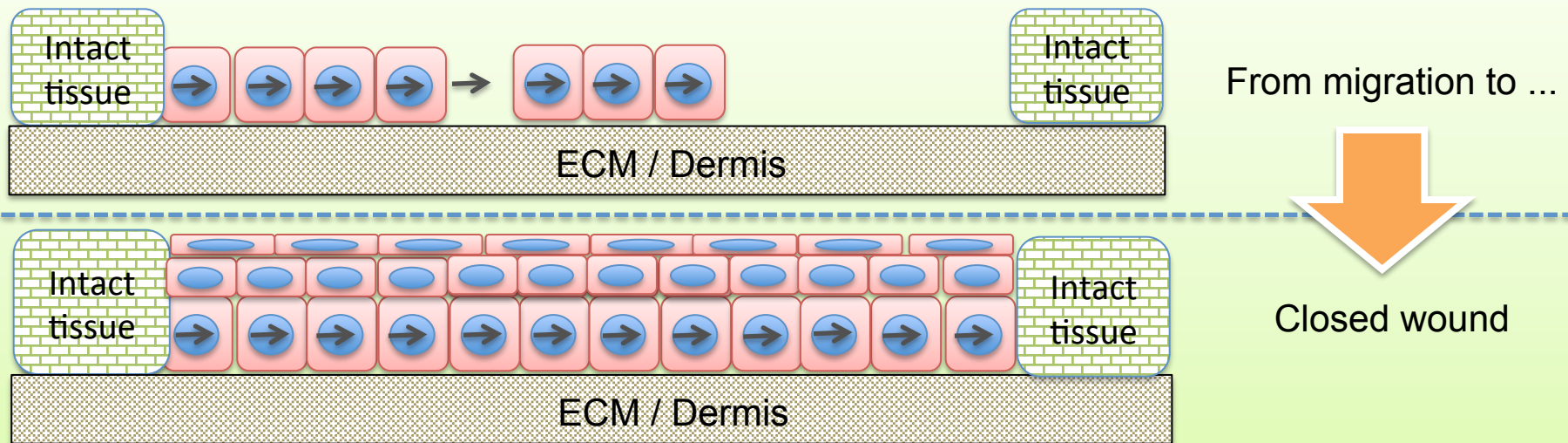


# Wound healing as a higher level process?

Unsolved since 40 years: Krawczyk WS (1971) A pattern of epidermal cell migration during wound healing. *The Journal of Cell Biology* 49:247–263.

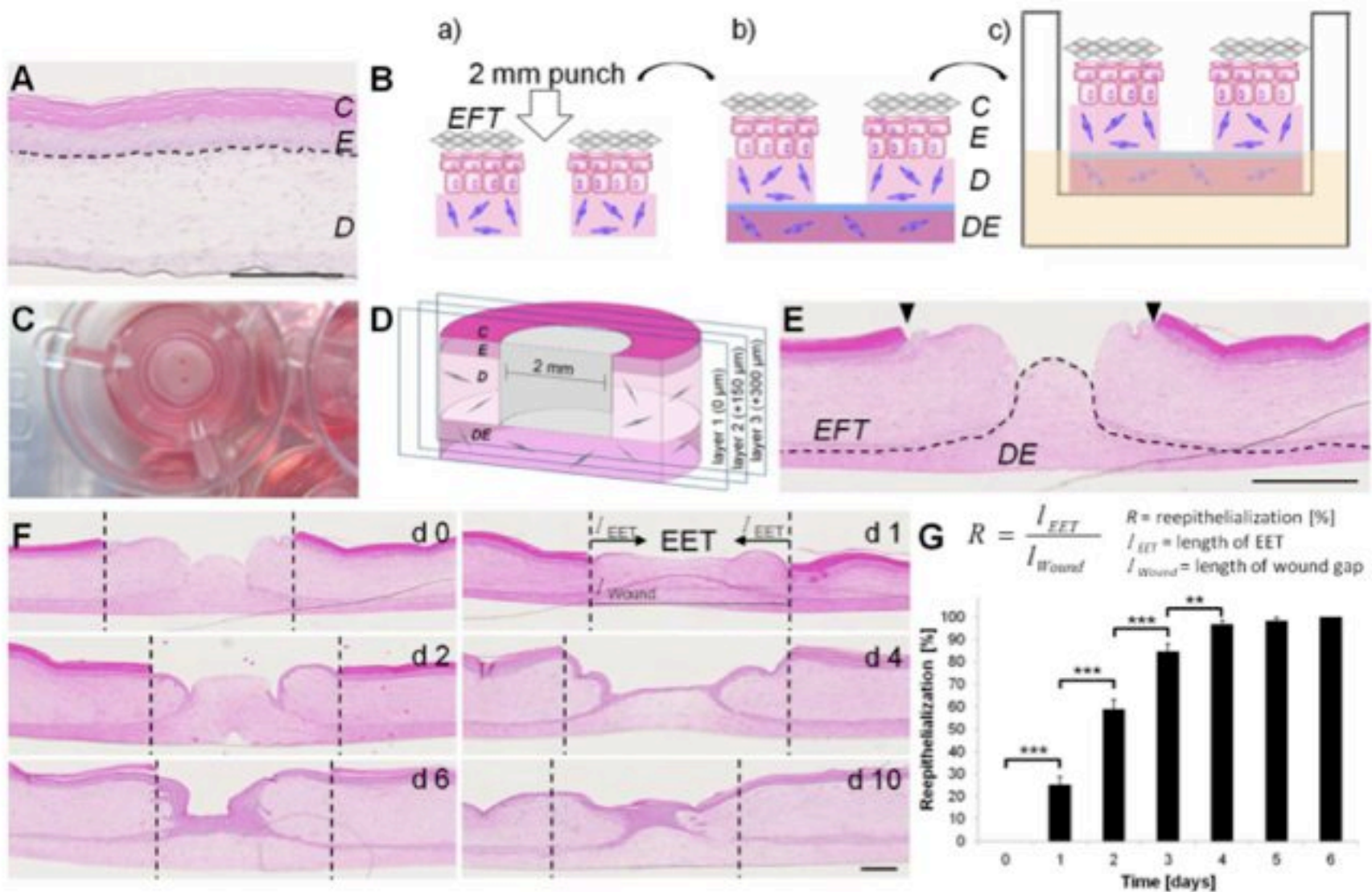
*Main questions:*

1. *What is the role of 2D/3D migration?*
2. *By which mechanism 3D epithelium is built from 2D migration?*
3. *What is the role of the surrounding tissue?*
4. *How is this higher-level process orchestrated?*

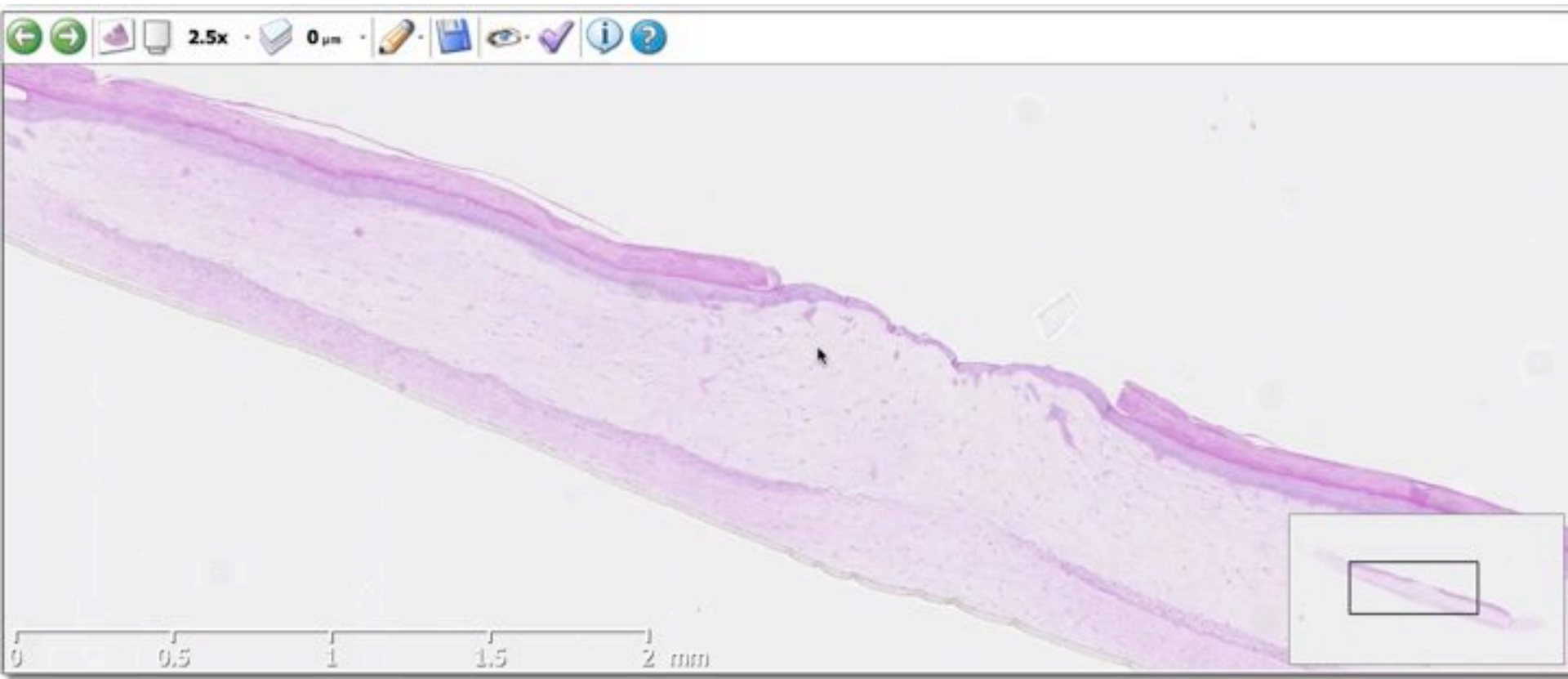




# 3D Punch Wound Model



# 3D Culture after 4 days of wound healing



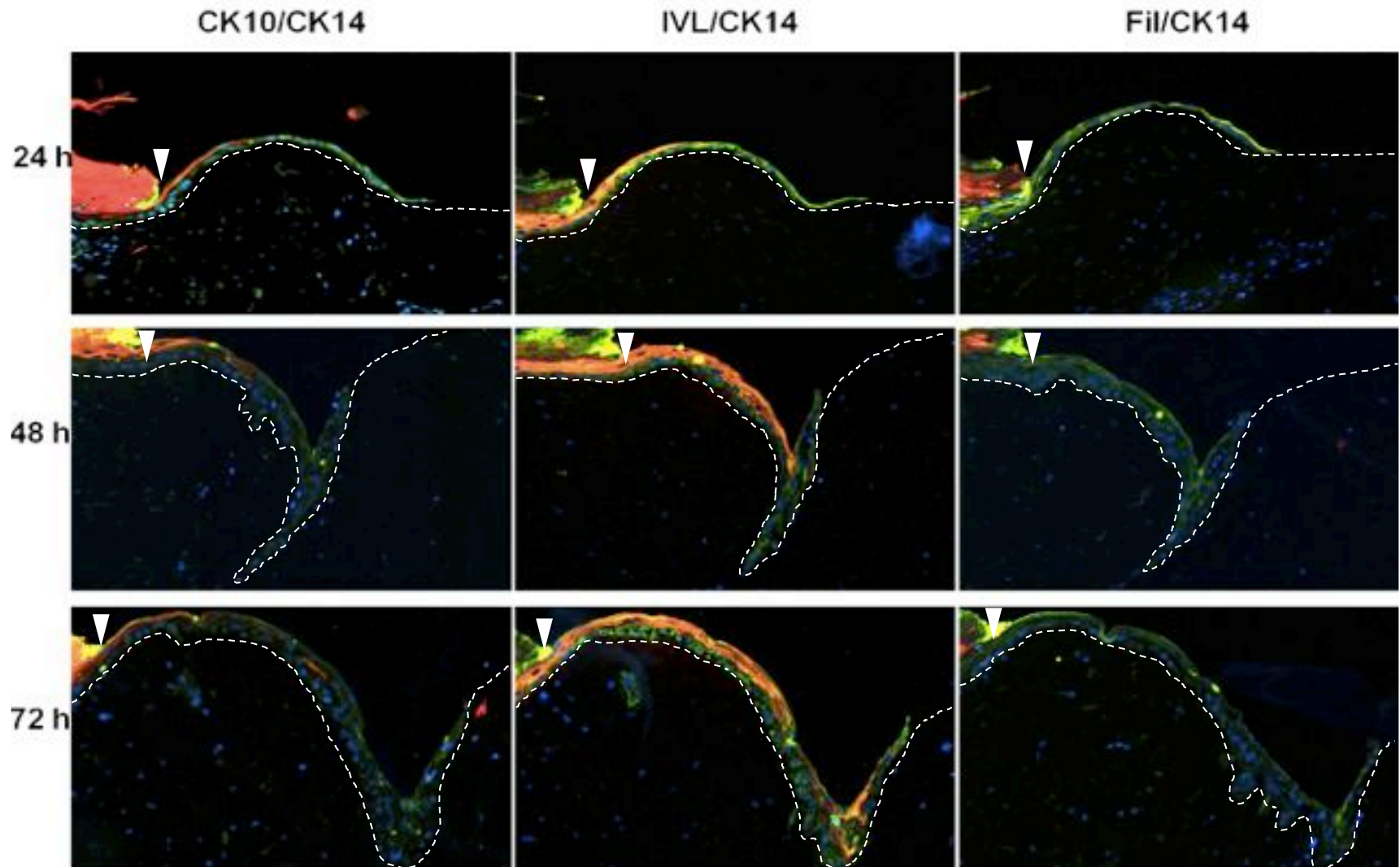


0 2.5 5 7.5 10 mm

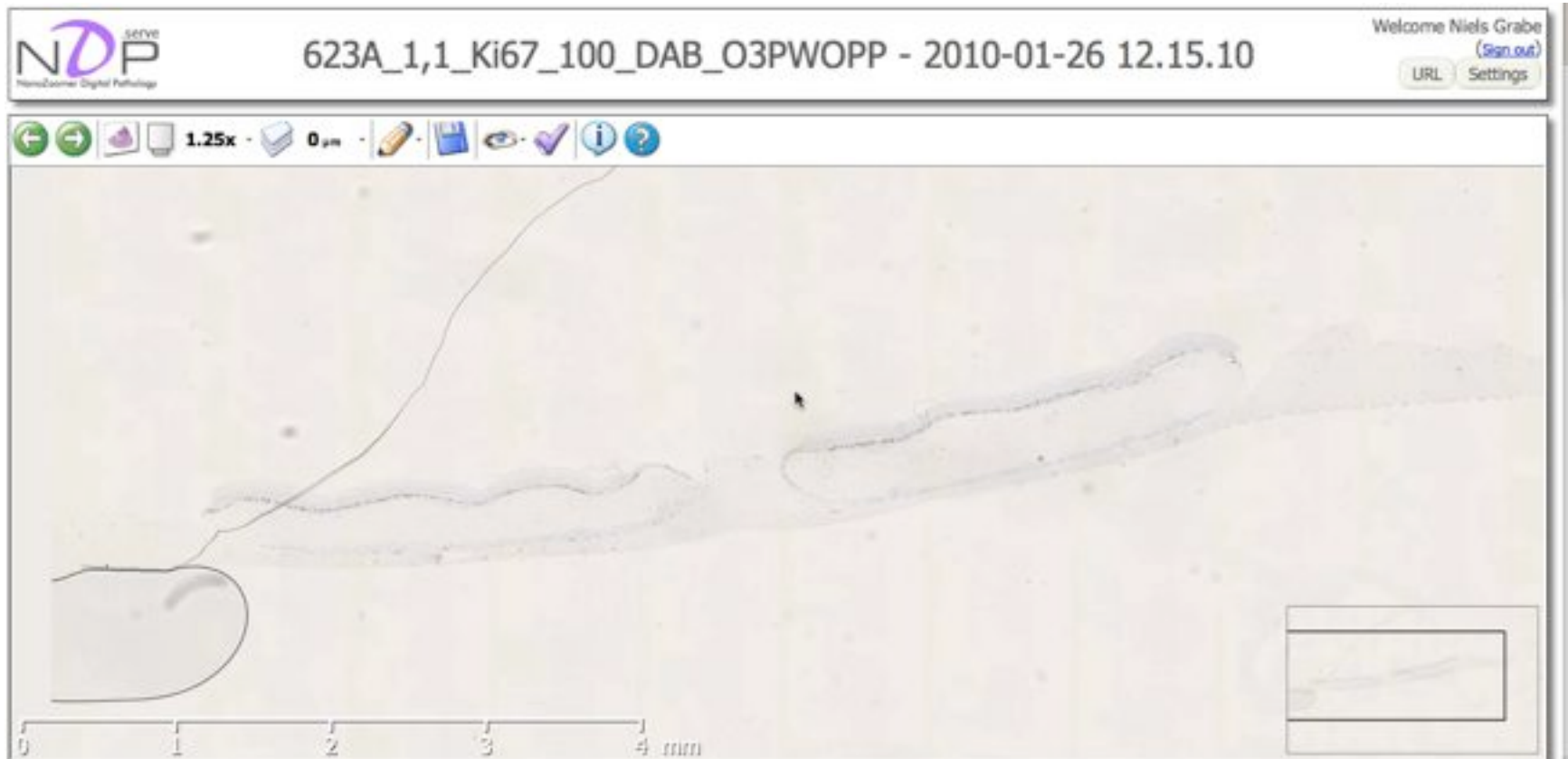




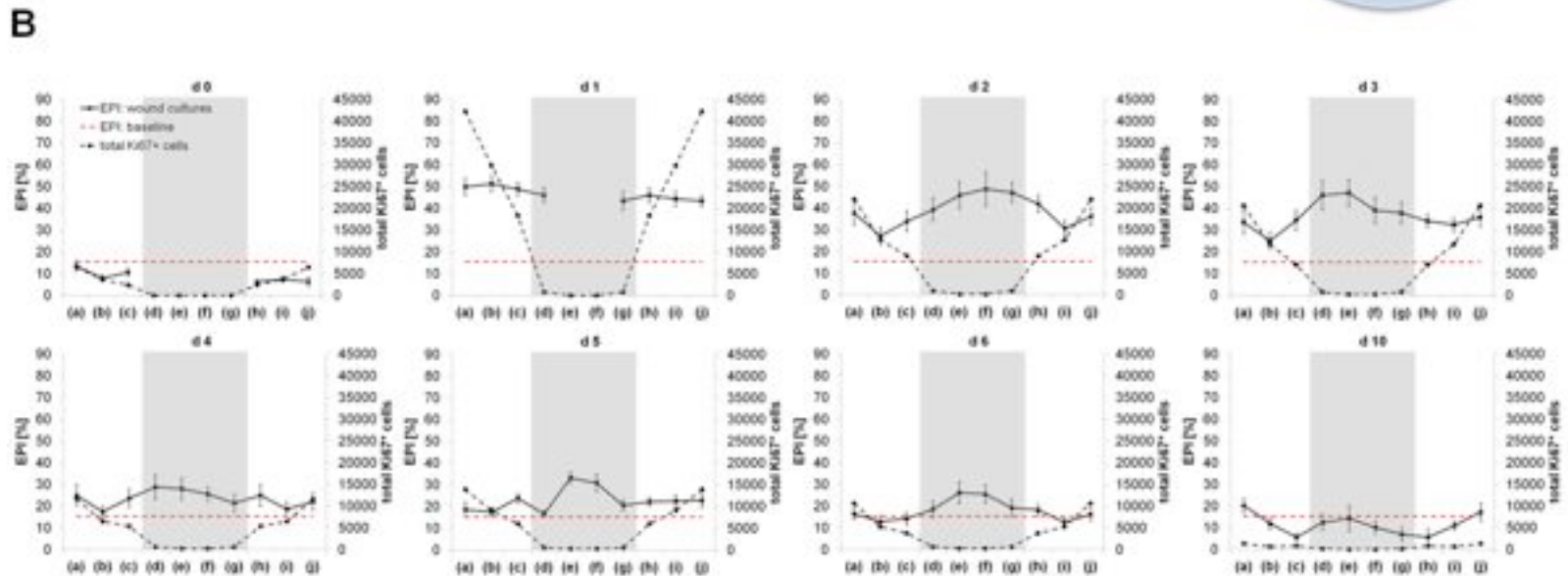
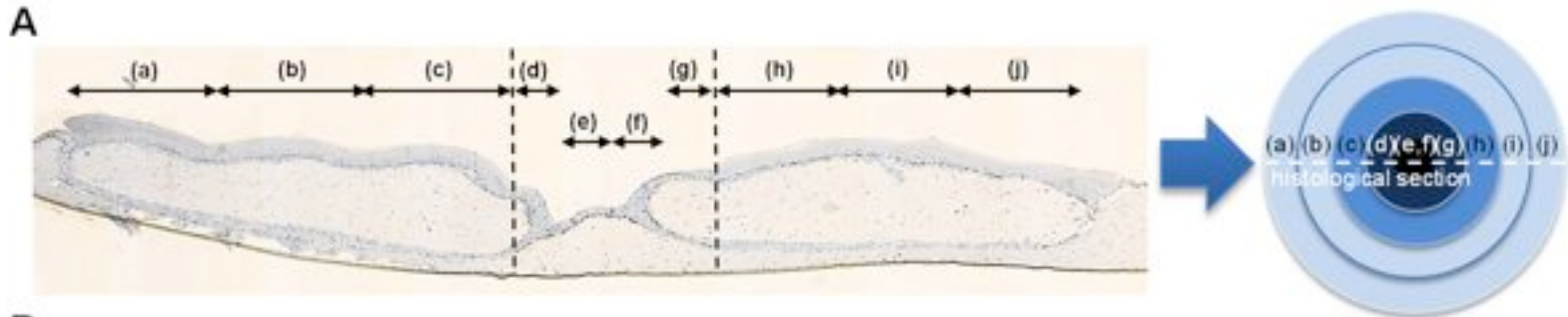
# Wound closure happens by a continuously extending triangular (3D) tongue



# Ki-67 Proliferation in Extending Epidermal Tongue

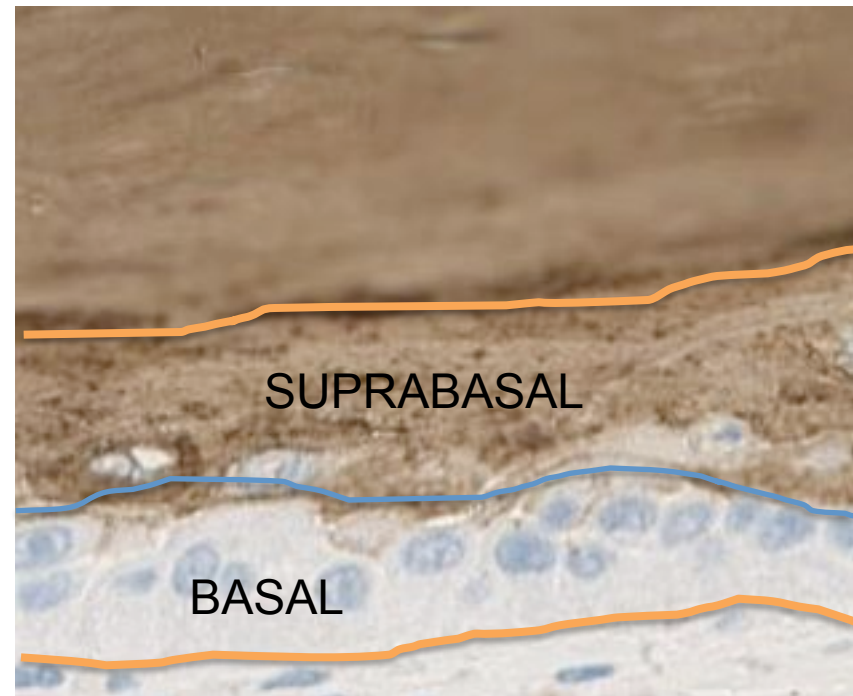
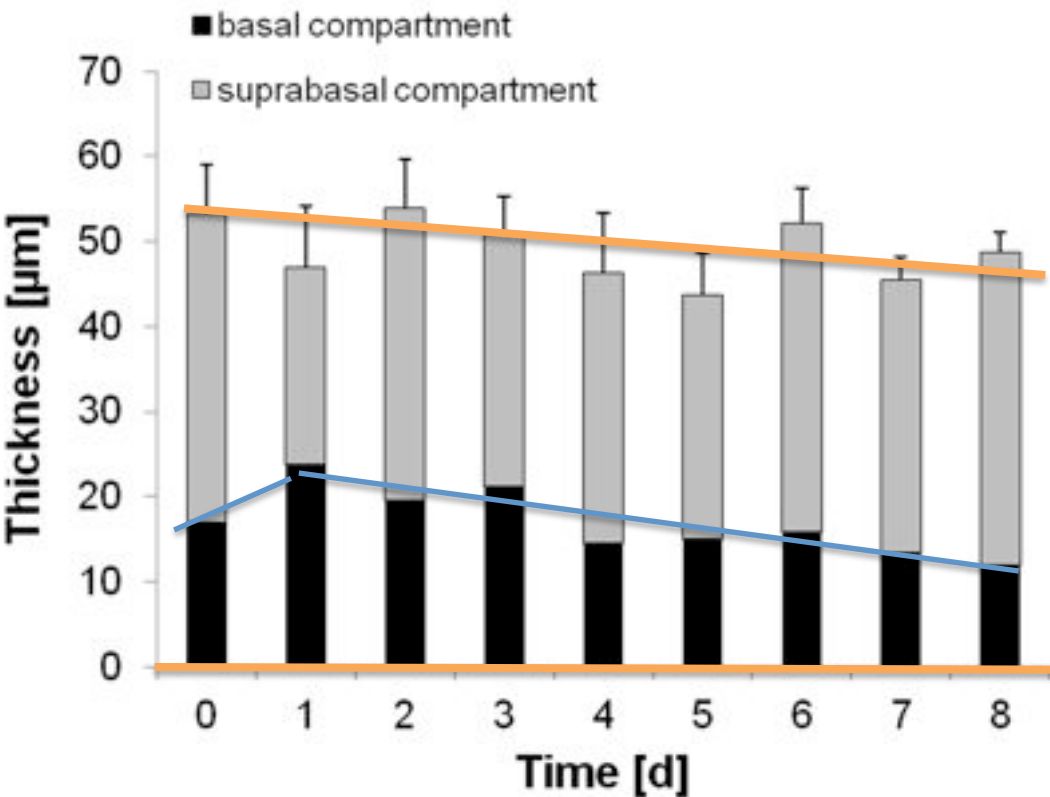


# Spatio-temporal profile of proliferation (image processed Ki67<sup>+</sup>)

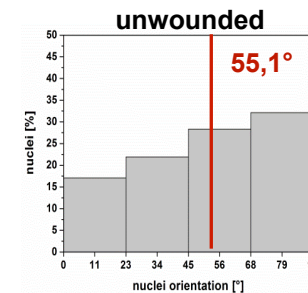
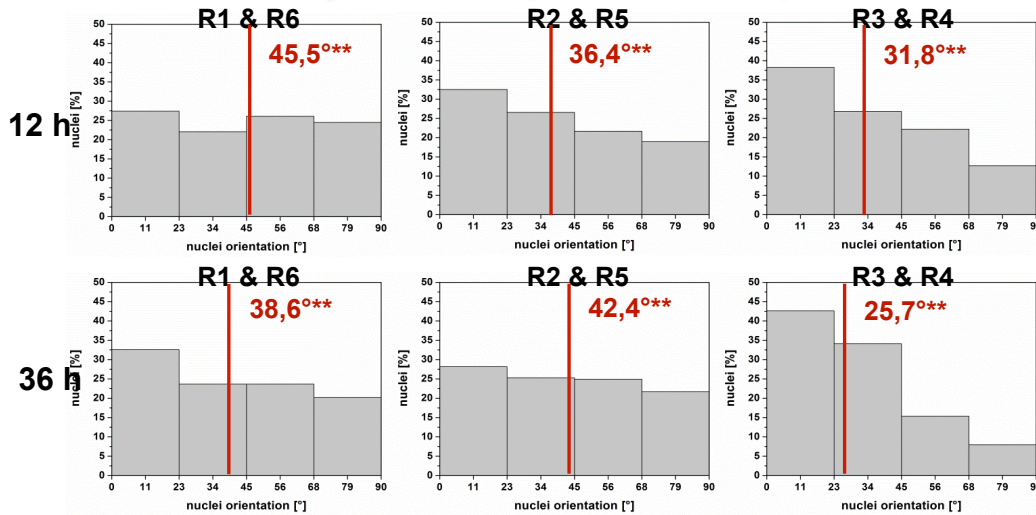
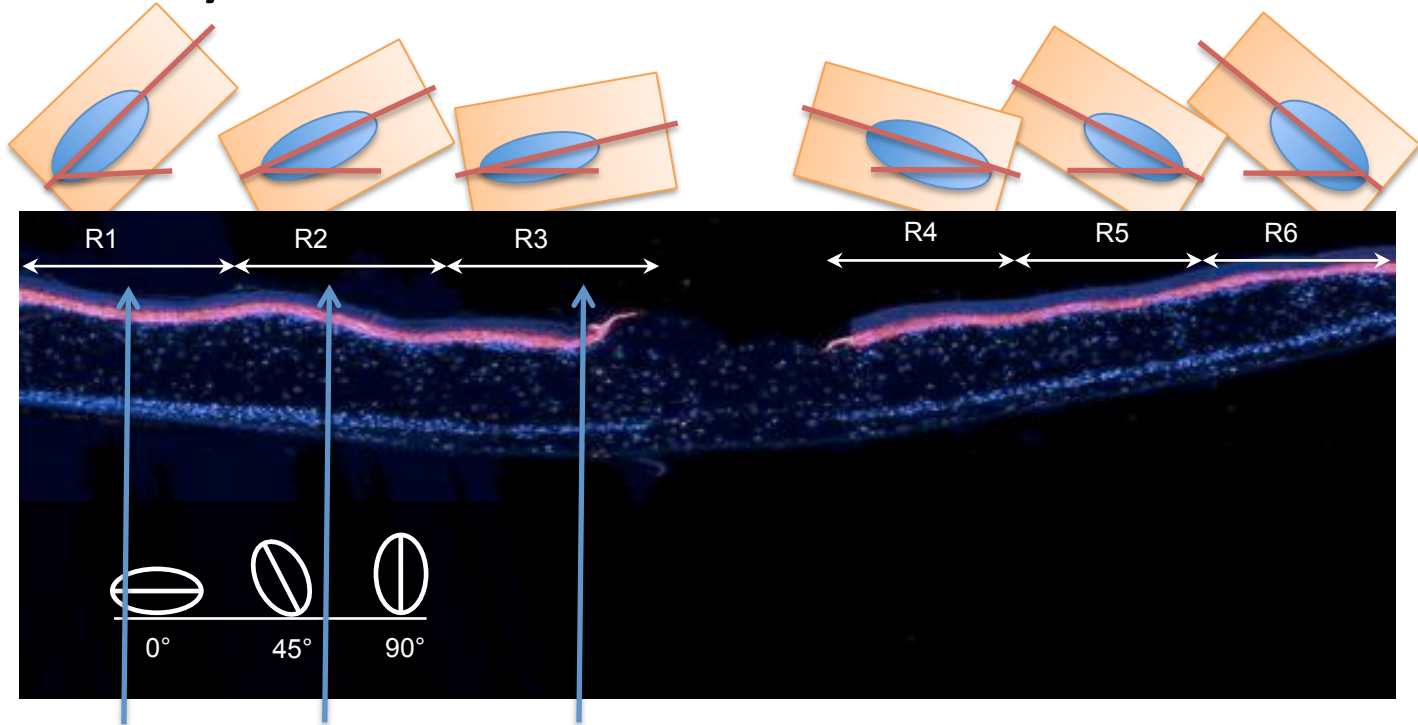


The grey area = the actual wound bed => does factually NOT contribute new cells but the surrounding tissue !!

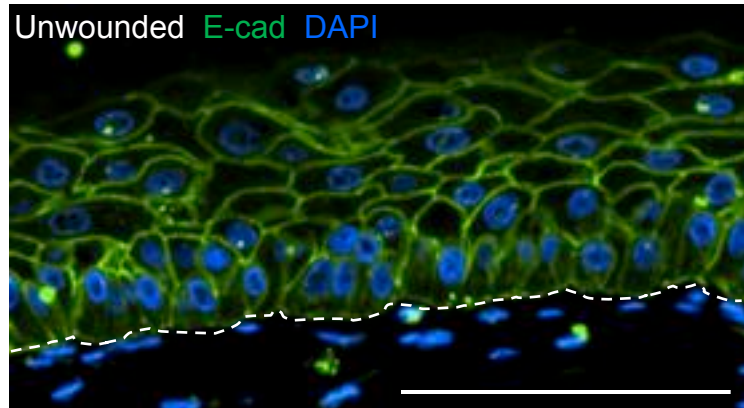
No great change in thickness of skin layers:  
=> where do the newly produced cells go to?



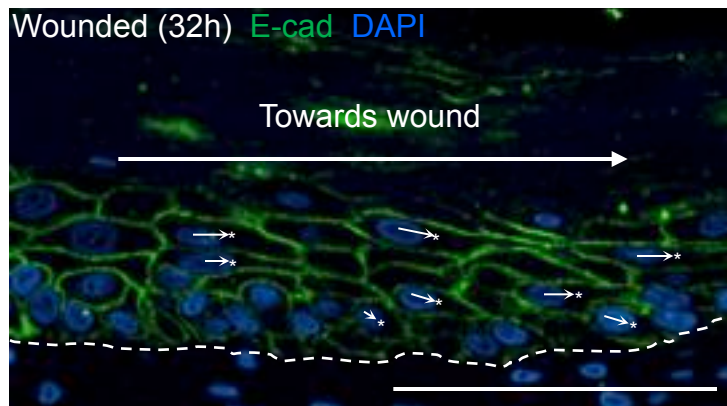
# Increasing collective cell rotation in the basal layer of the intact tissue after wounding



# Keratinocytes of the intact tissue surrounding show cell elongation and nuclear displacement

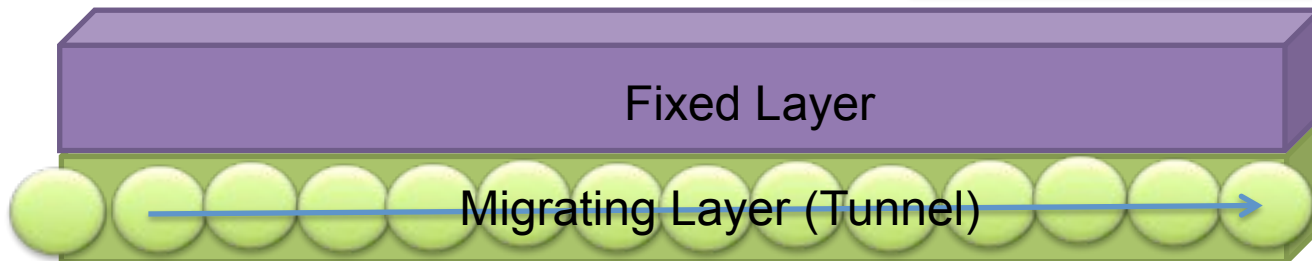
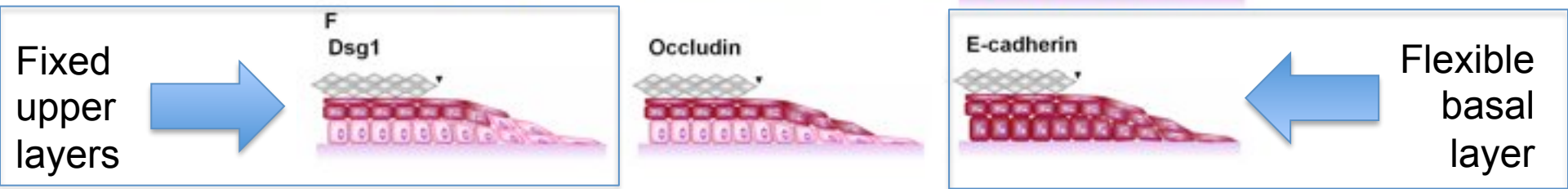
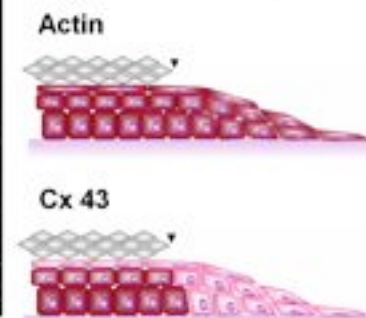
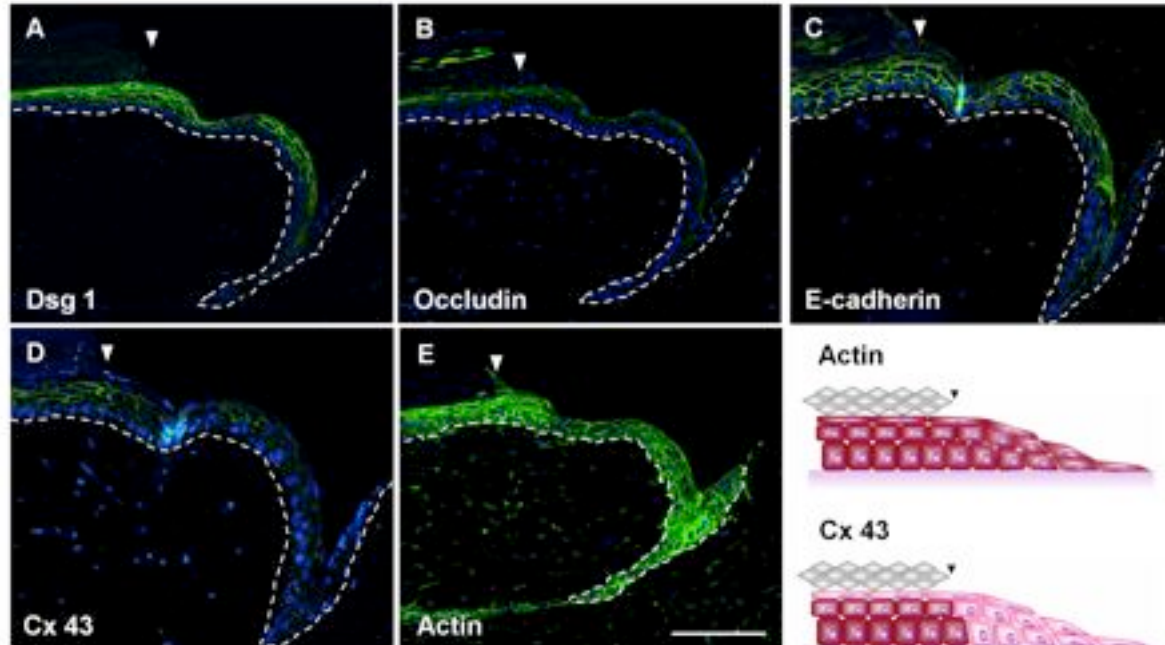


Control



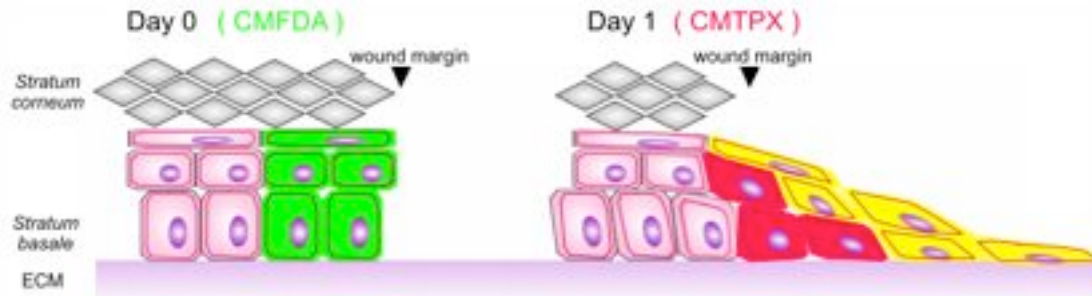
- Cell elongation
- Nuclear displacement

# Tunelling of collective migration in intact tissue and full neoepidermis

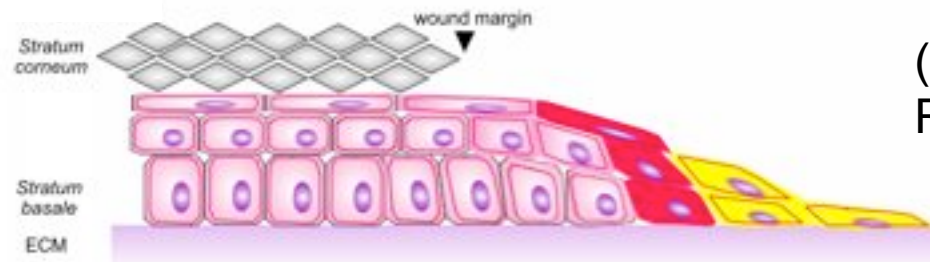


# Current models of tongue extension:

Hypothetical double labeling experiment:

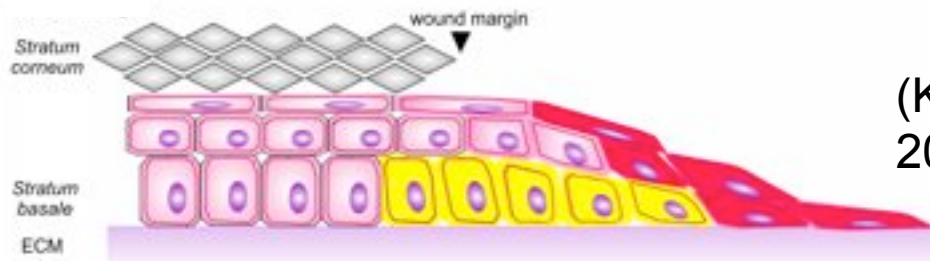


Tractor-tread theory



(Woodley DT, 1996; Radice, 1980)

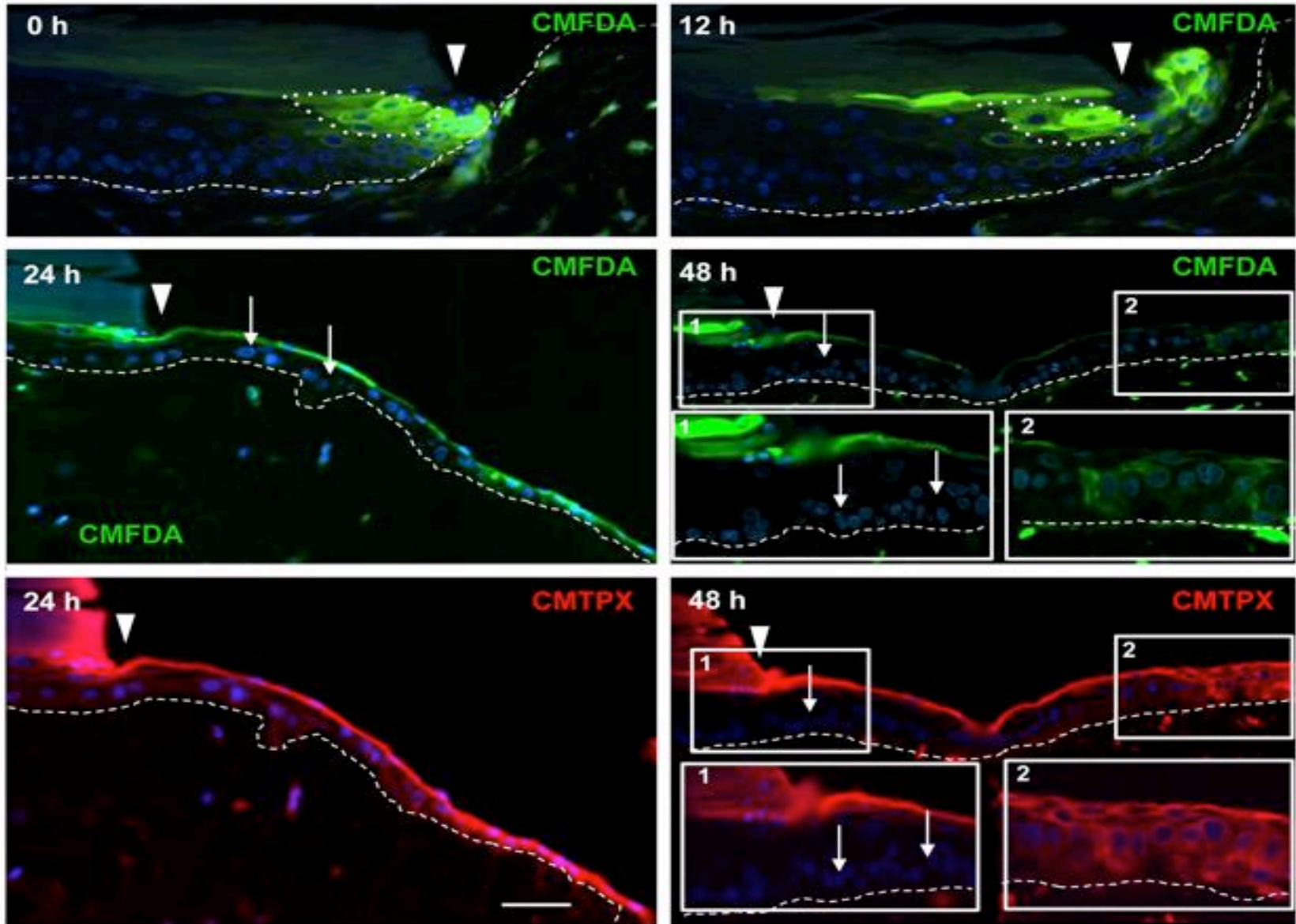
Leap frog theory



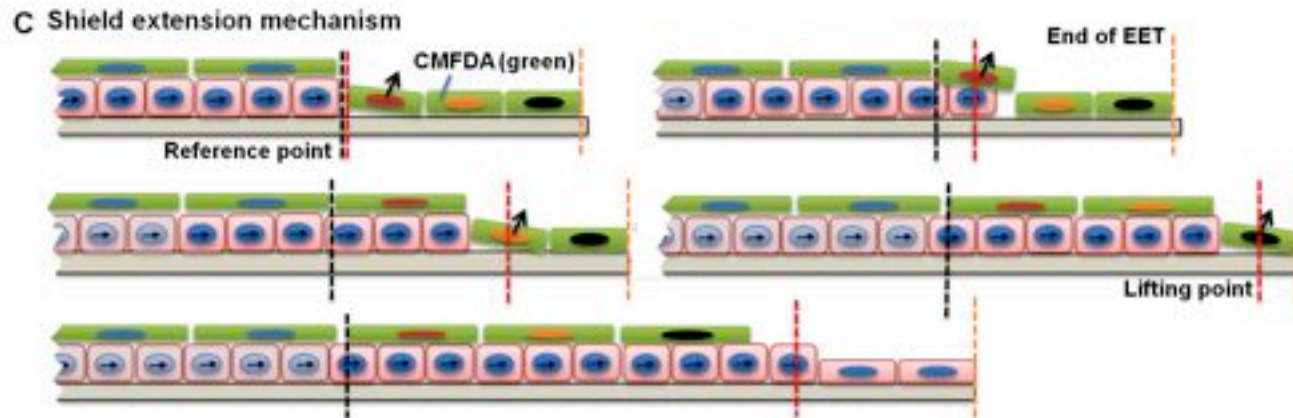
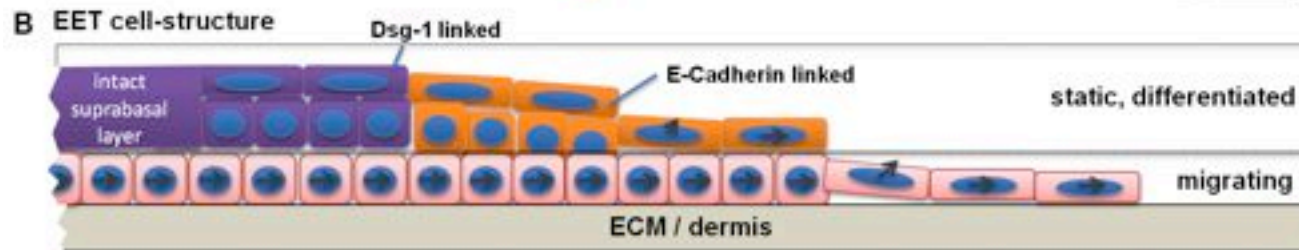
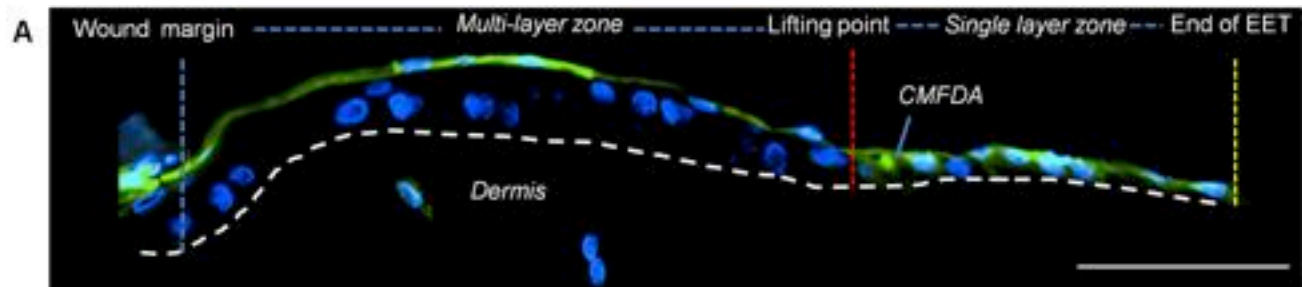
(Krawczyk, 1971; Usui et al., 2005; Paladini et al., 1996)



# Double Labelling Experiment in 3D in vitro culture: Green: @ 0h, Red: @ 24h

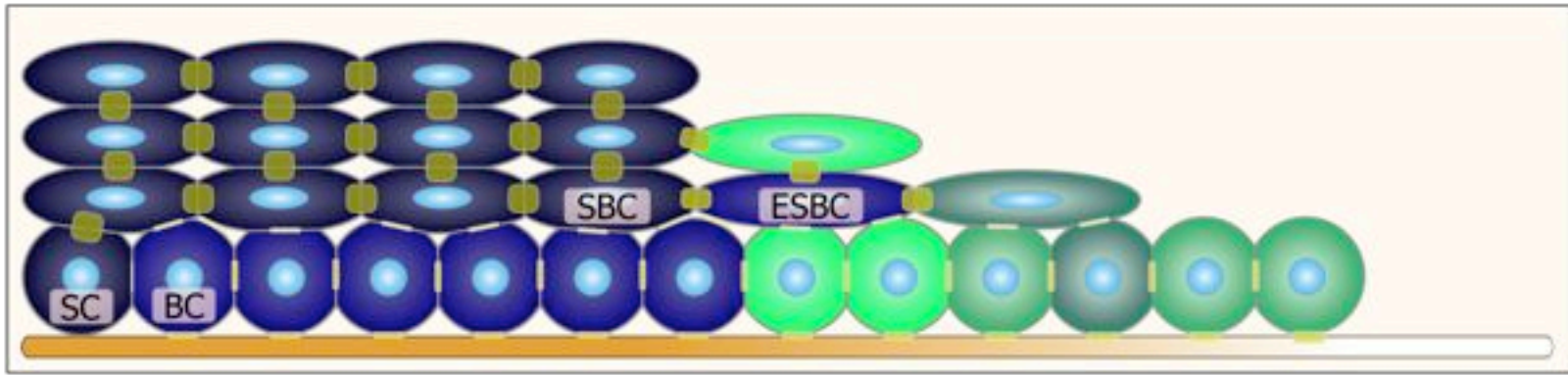


# Novel shield extension mechanism creates the 3D neoepidermal structure



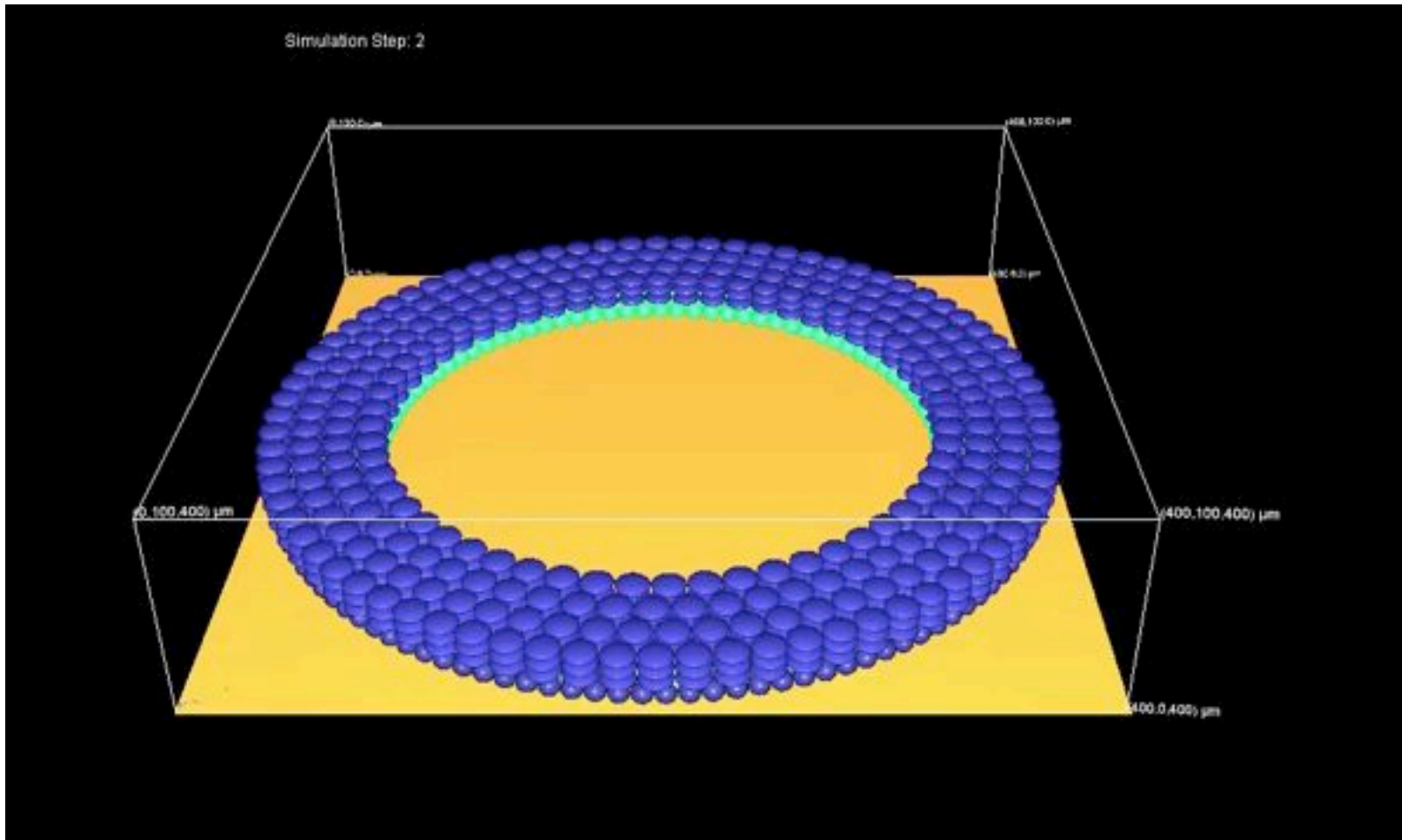


# Modeling adhesions

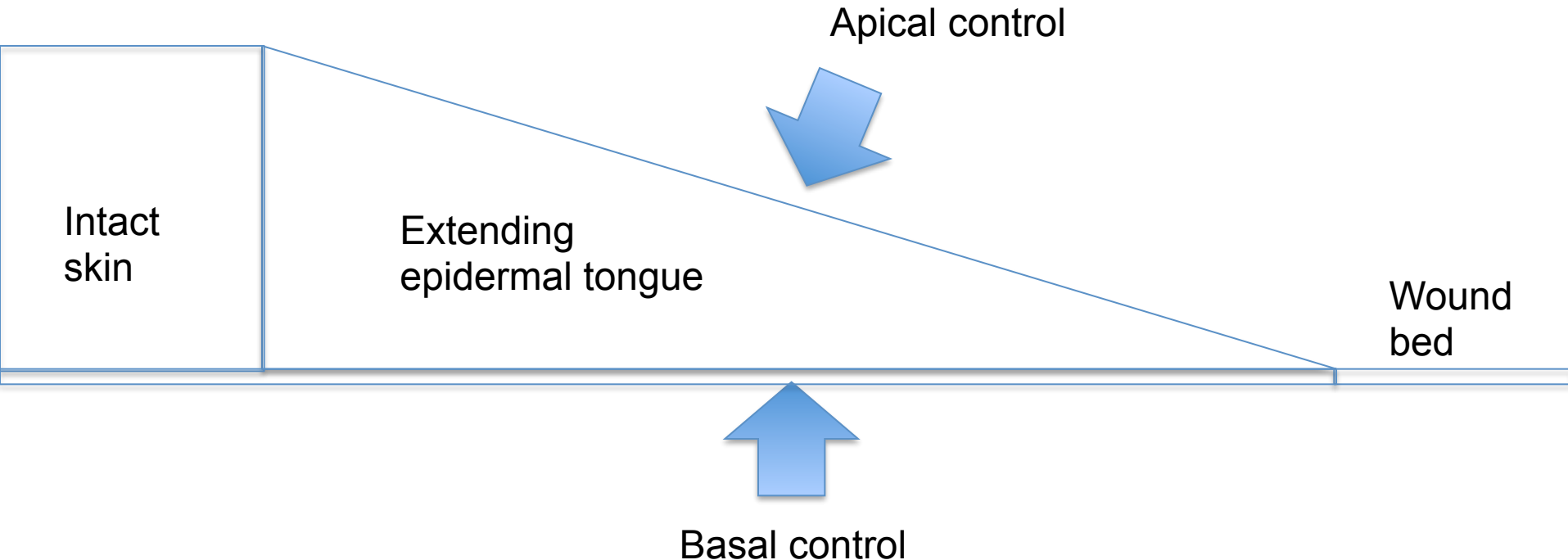


adhesion	Stem Cell (SC)	Basal Cell (BC)	Suprabasal Cell (SBC)	Early Suprabasal Cell (ESBC)	Basalmembrane (BM)
SC	fixed pos.	1% —	90% ■	----	fixed pos.
BC	1% —	21% —	5% —	10% —	20% - 45% —
SBC	90% ■	5% —	90% ■	30% - 70% ■	----
ESBC	----	10% —	30% - 70% ■	40% - 80% ■	----
BM	fixed pos.	20% - 45% —	----	----	----

# Total multi-cellular 3D wound closure model



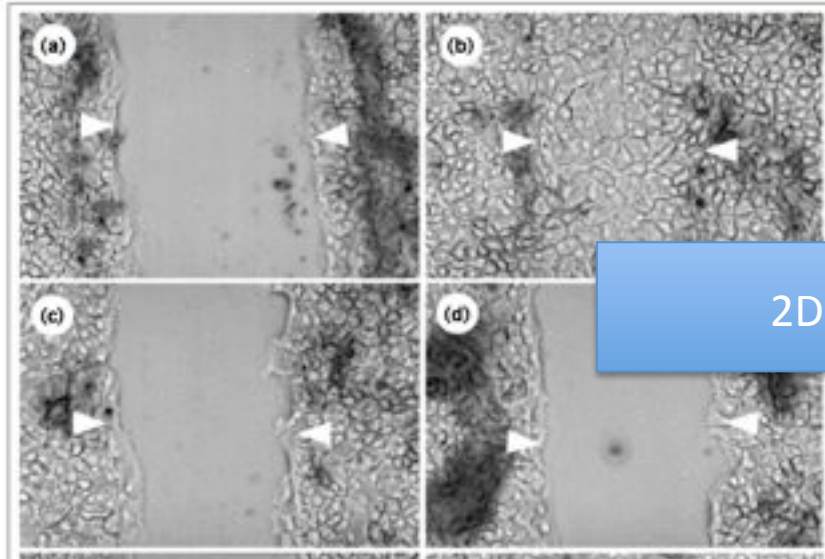
# Modeling resulted in two potential control mechanisms of epidermal shield extension



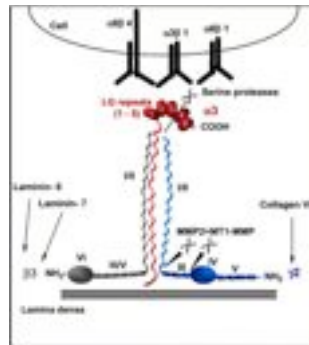
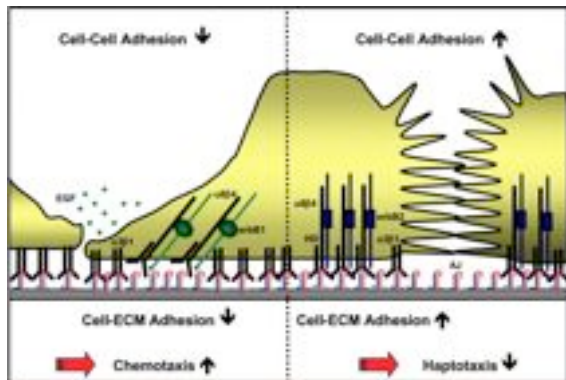
Two possible models of shield extension mechanism:

- Lifting occurs by slow moving leaders and faster followers by ECM deposition of leaders
- Shield extension occurs by adhesion specific to the upper layer

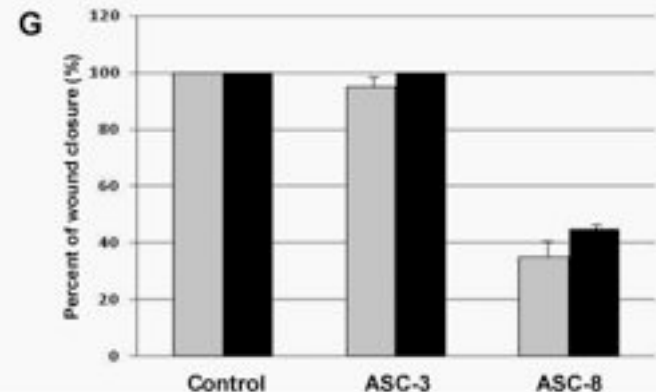
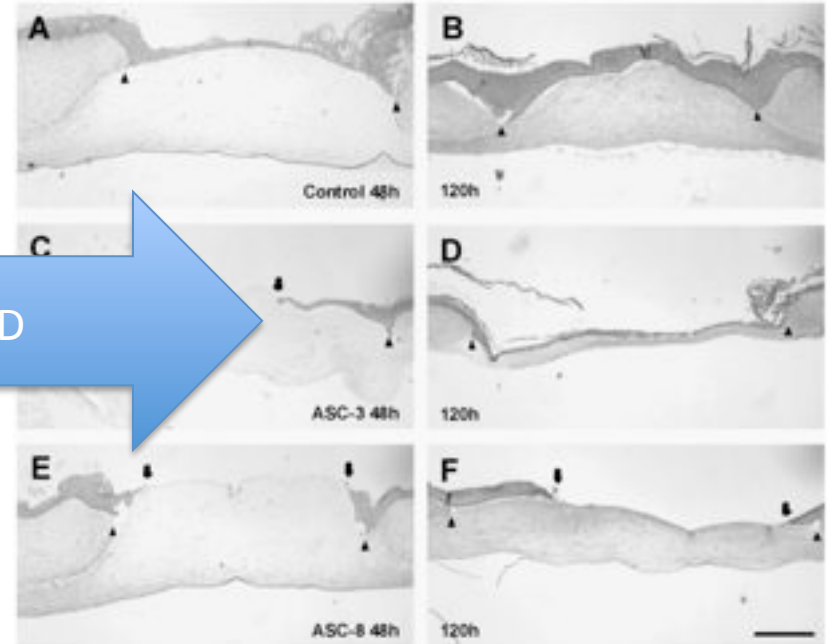
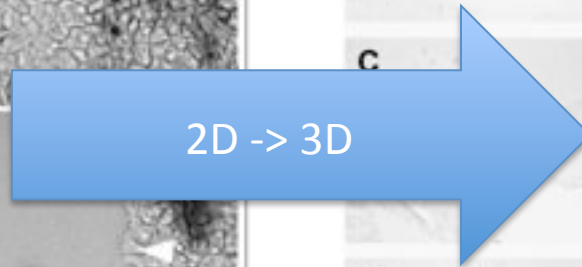
# Blocking of laminin-5 delays wound healing but does not perturb epithelial tongue formation



Laminin-5 inhibits migration in scratch assay:  
 Nguyen, Curr. Op. Cell Biol. 2000

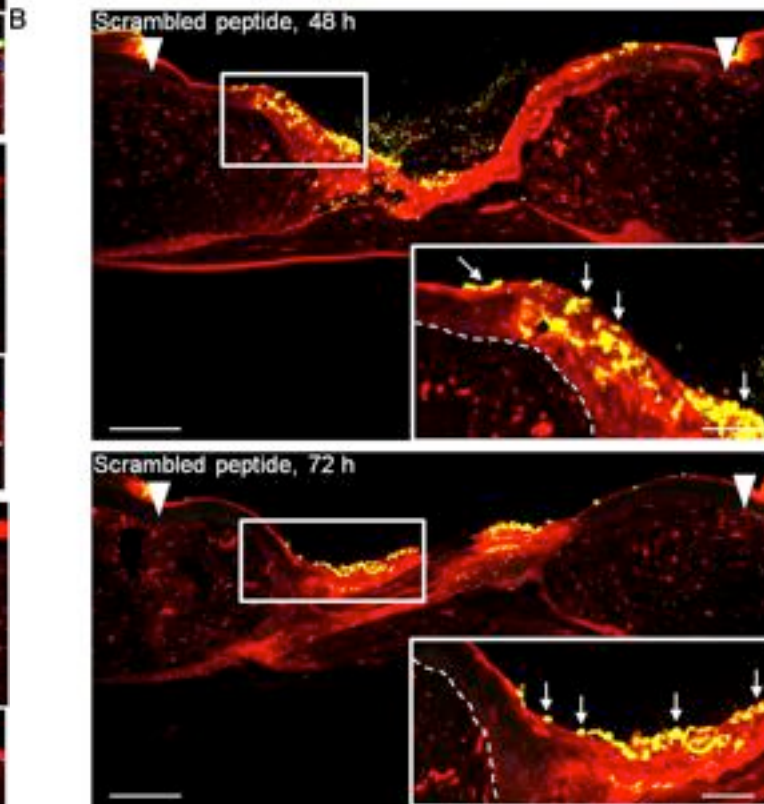
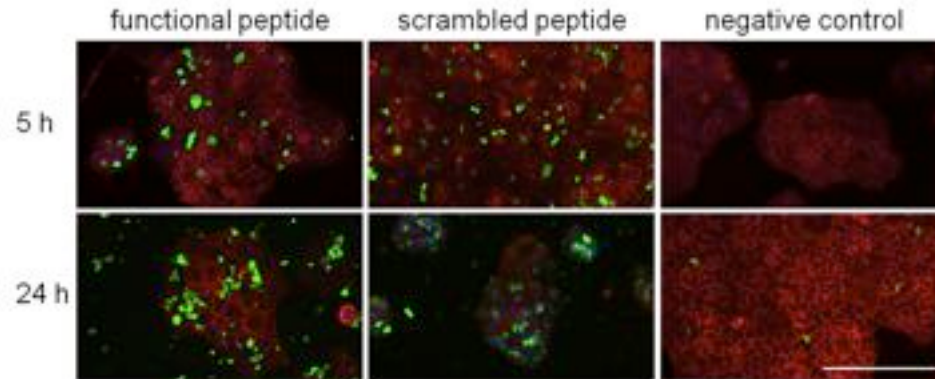
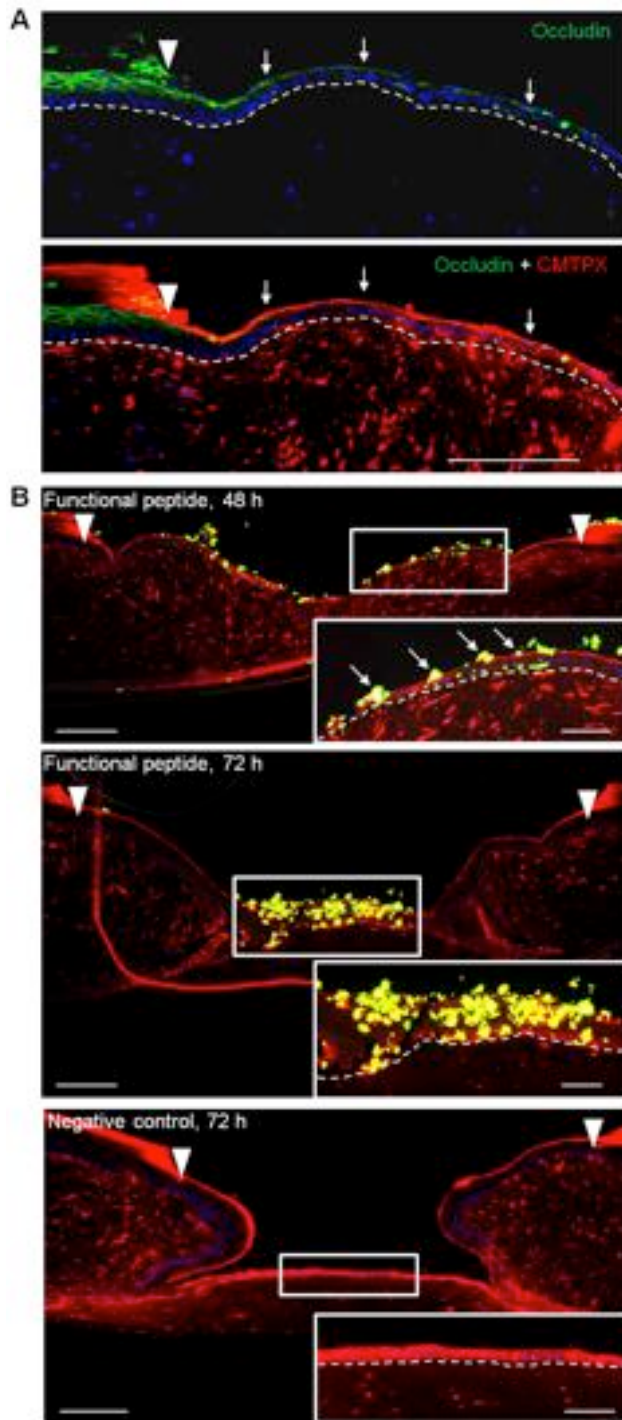


Hinterman and Quaranta, Matrix Biology  
 Volume 23, Issue 2, May 2004, P. 75–85

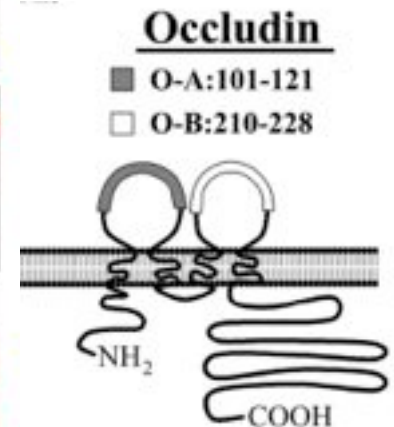


Integrin-Blocking Antibodies Delay Keratinocyte Re-Epithelialization in a Human Three-Dimensional Wound Healing Model, Garlick Group, PLoS One. 2010; 5(5): e10528.

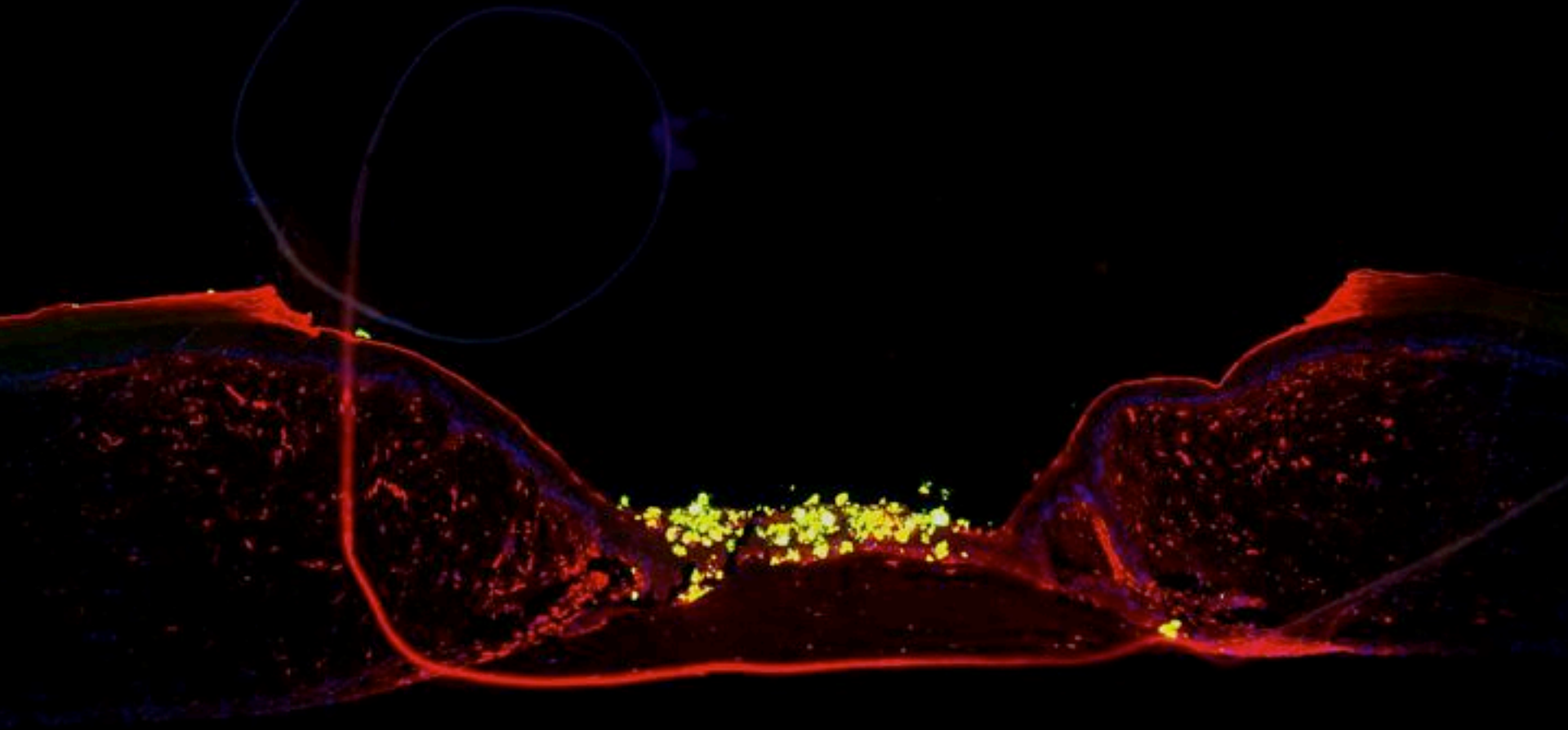
# Perturbing Occludin



**Peptide O-B 210-228:**  
 Biotin-SQIYALCNQ (bpa)  
 YTPAATGLYVD-NH2

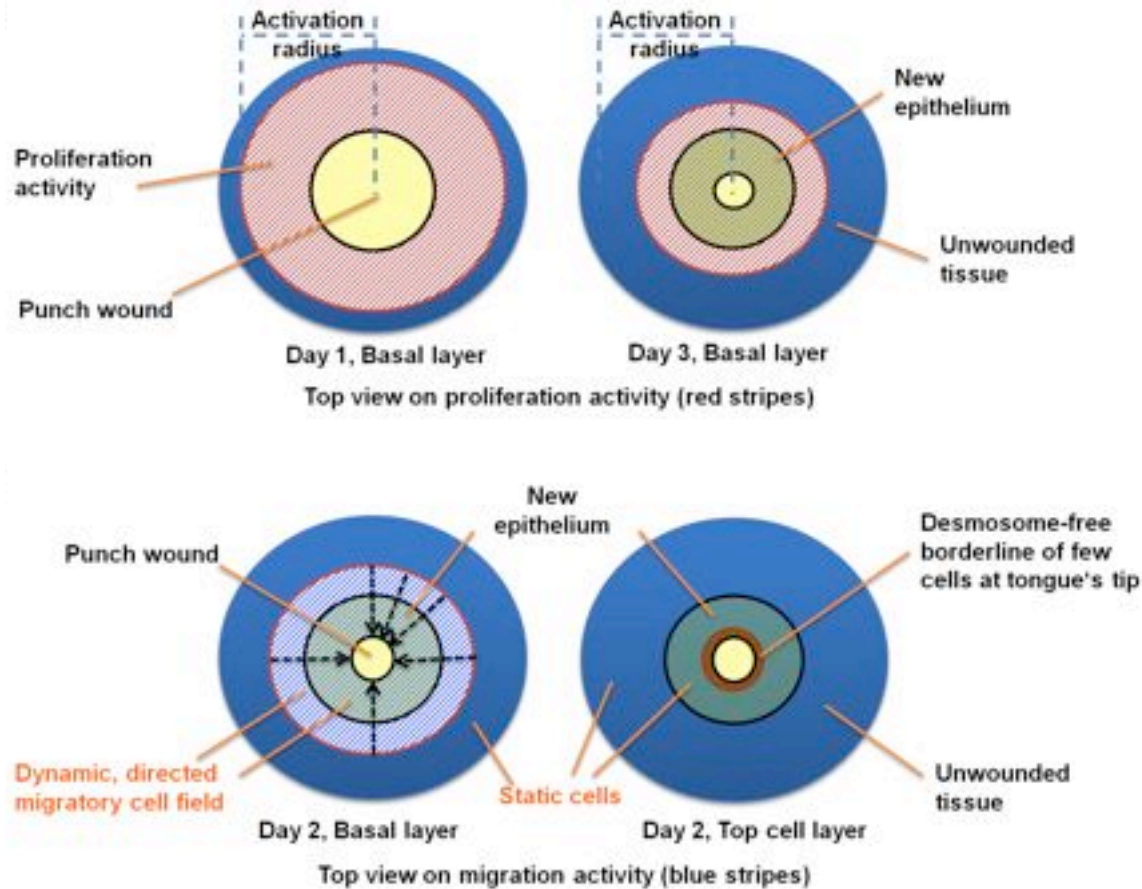


# Mechanism validation by tight-junction blockade

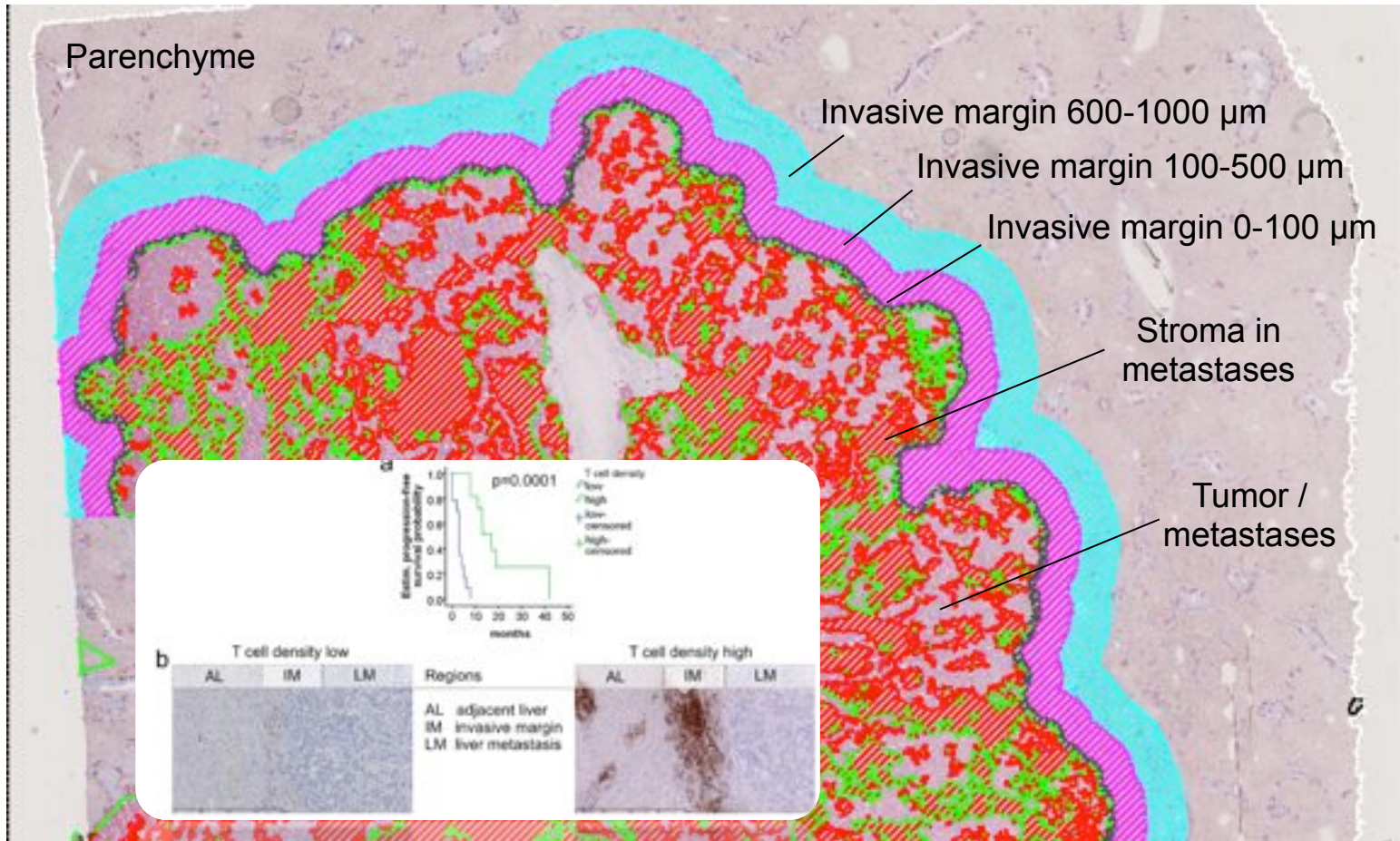




# From shield extension to total wound closure

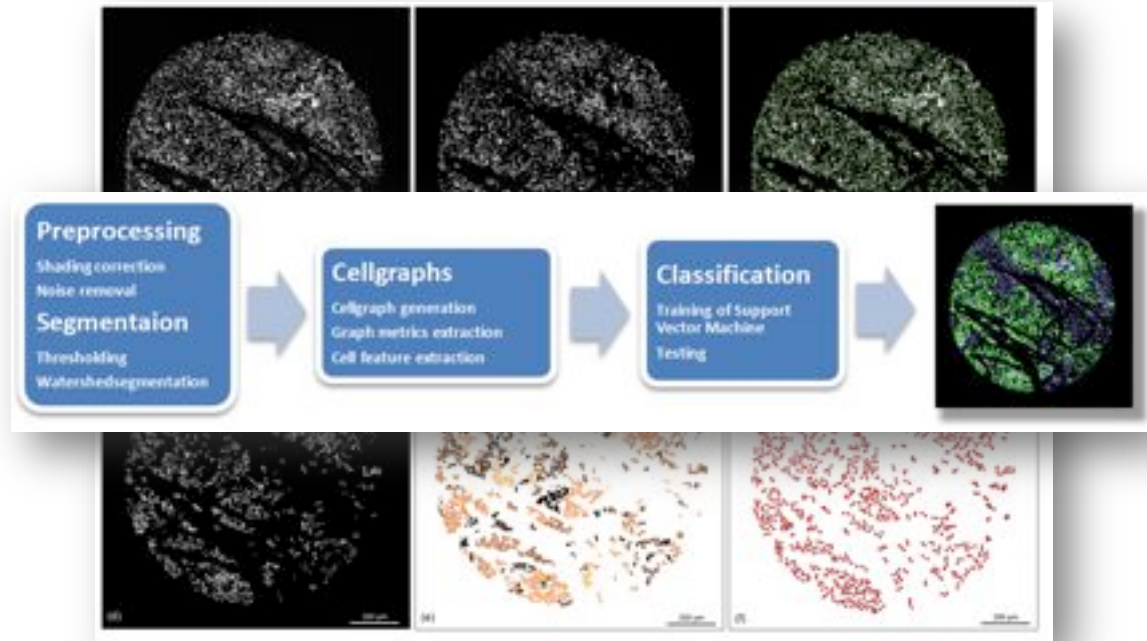
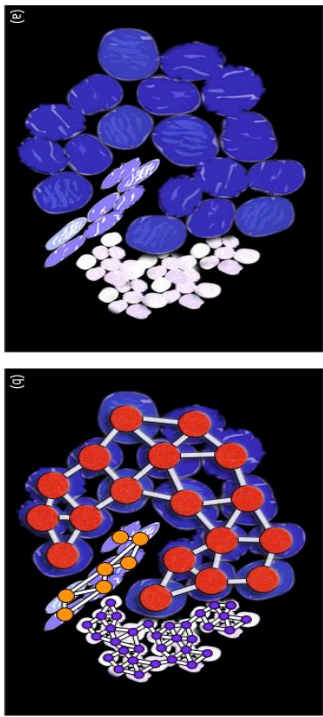


# Standardized Cancer Immune Cell Profiling





# Cell Networks for Tissue Segmentation



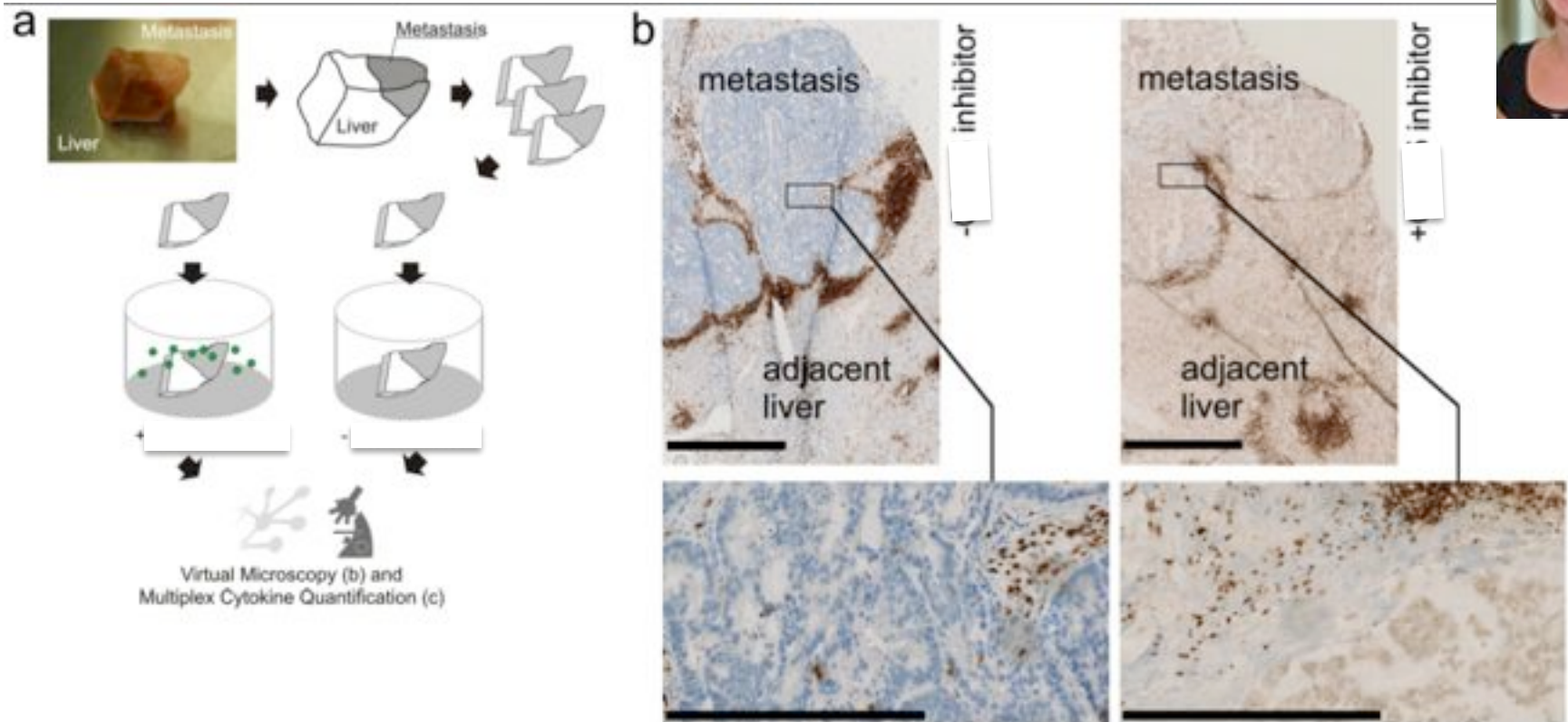
	Training set	Test set					Average 1-5
		1	2	3	4	5	
Overall	88.47(±06.68)	87.65(±08.19)	90.30(±06.44)	88.68(±07.19)	88.76(±06.98)	88.59(±09.83)	88.80(±07.73)
Tumor	89.26(±10.20)	87.56(±13.29)	87.83(±12.47)	88.00(±17.64)	88.98(±10.01)	87.71(±14.13)	88.02(±13.51)
Stroma	85.14(±10.95)	81.19(±11.62)	91.45(±06.21)	82.97(±15.12)	80.02(±12.35)	86.90(±13.69)	84.67(±11.80)

Lahrman B, Halama S, Sinn HP, Schirmacher P, Jaeger D, Grabe N.

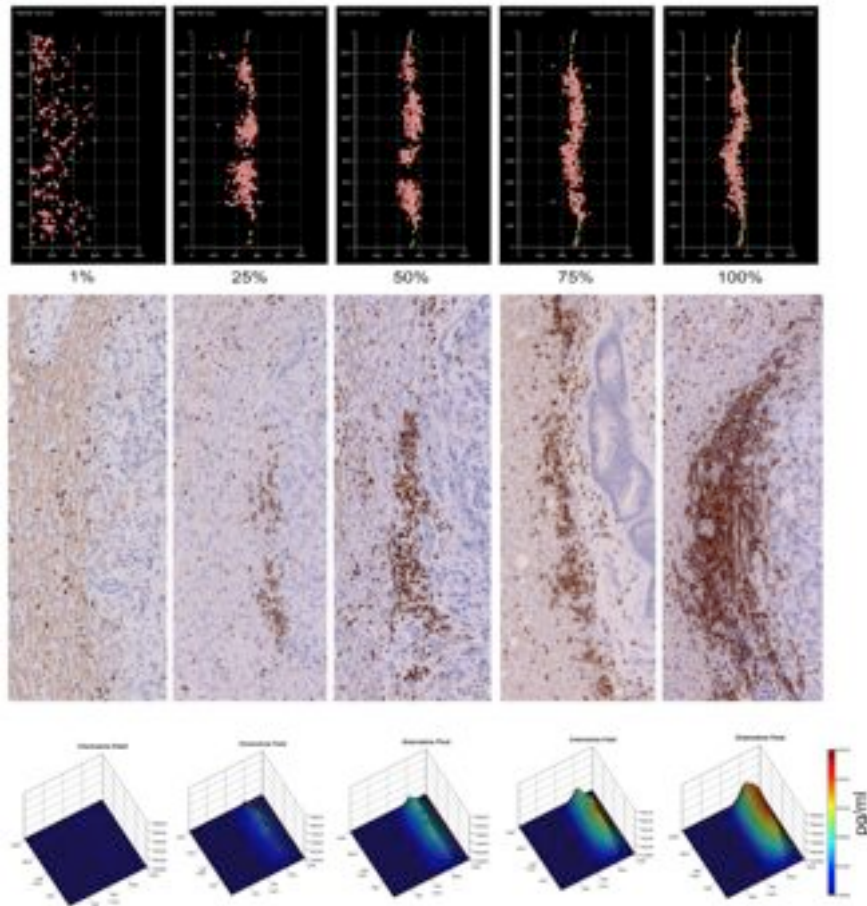
**Automatic Tumor-Stroma Separation in Fluorescence TMAs Enables the Quantitative High-throughput Analysis of Multiple Cancer Biomarkers**

*PLoS ONE*. December 2011;Vol 6(12):e28048

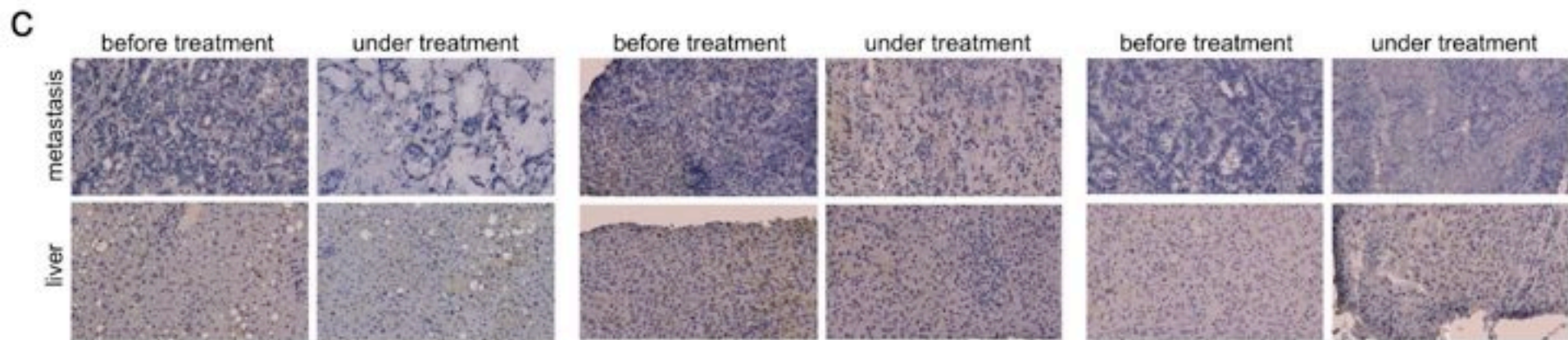
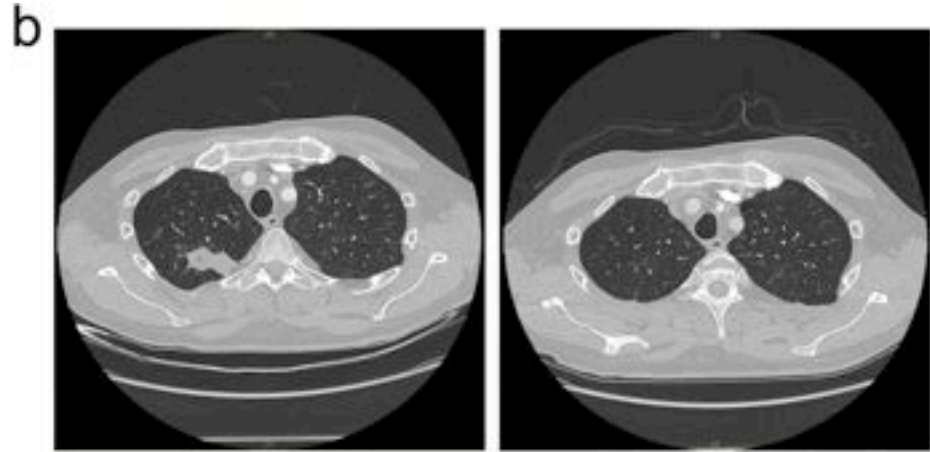
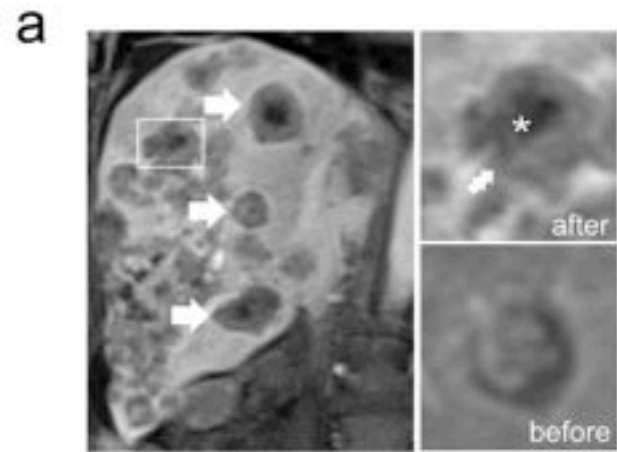
# Cancer Modeling



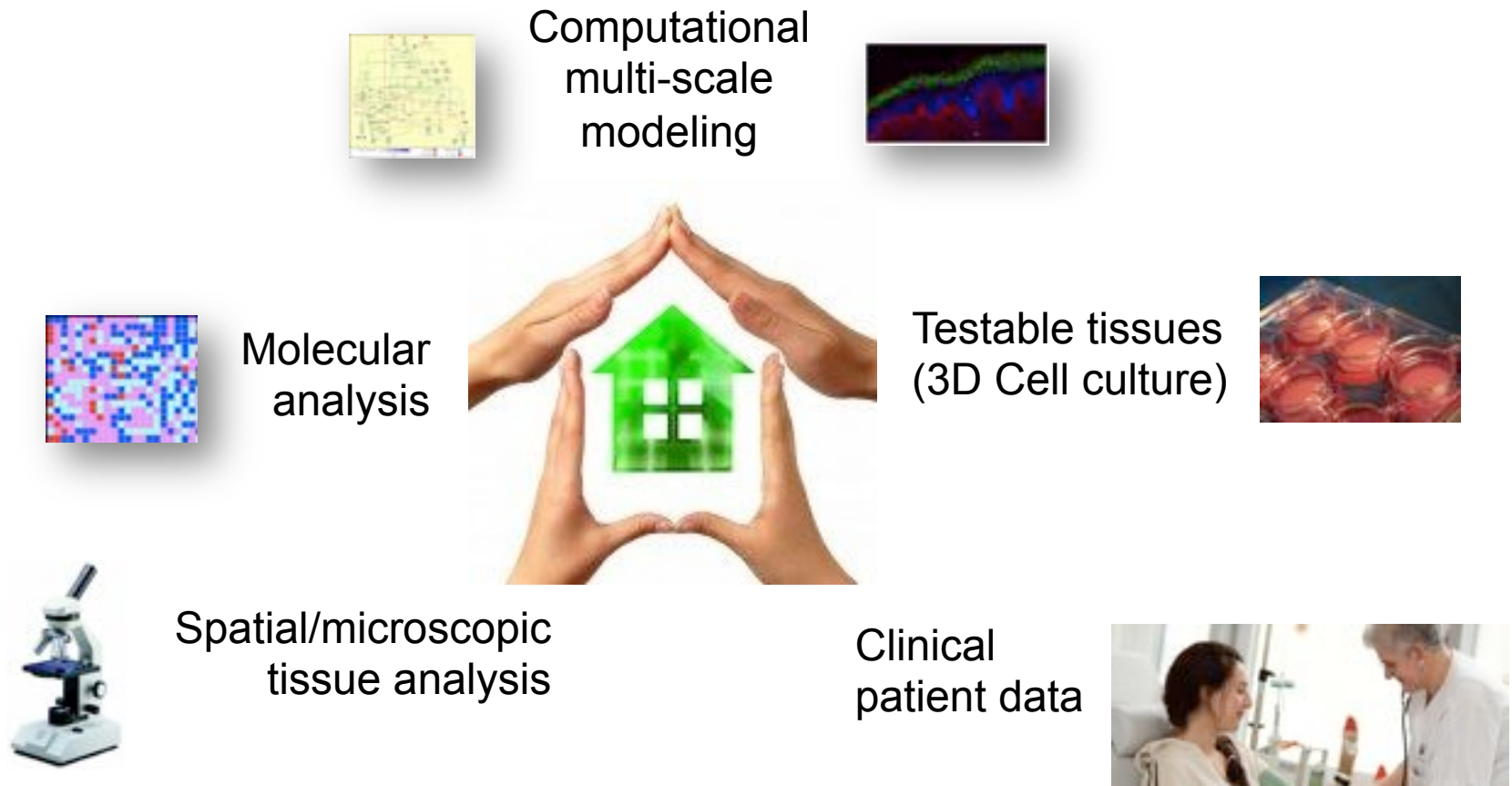
# Computational Simulation of Immune cell Profile



# Patient Metastatic Response



# „House“ of Medical Systems Biology



What is Medical Systems Biology? Integration of these levels in a way closer than ever before driven by technology to generate new points of intervention.