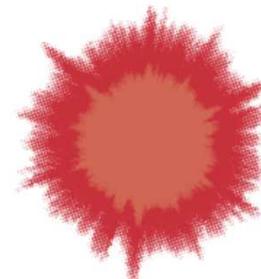


EMT during cancer initiation and progression

Lab: prof. Geert Berx

Dr. Bram De Craene

**American Biotechnology and Life-Science Trade Mission
October 21 2104**



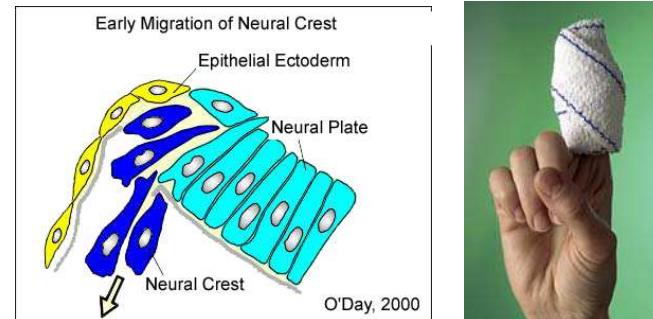
IRC
A VIB-UGENT DEPARTMENT

Epithelial to Mesenchymal Transition (EMT)

EMT is a morphogenetic process in which cells loose their epithelial characteristics and gain mesenchymal properties

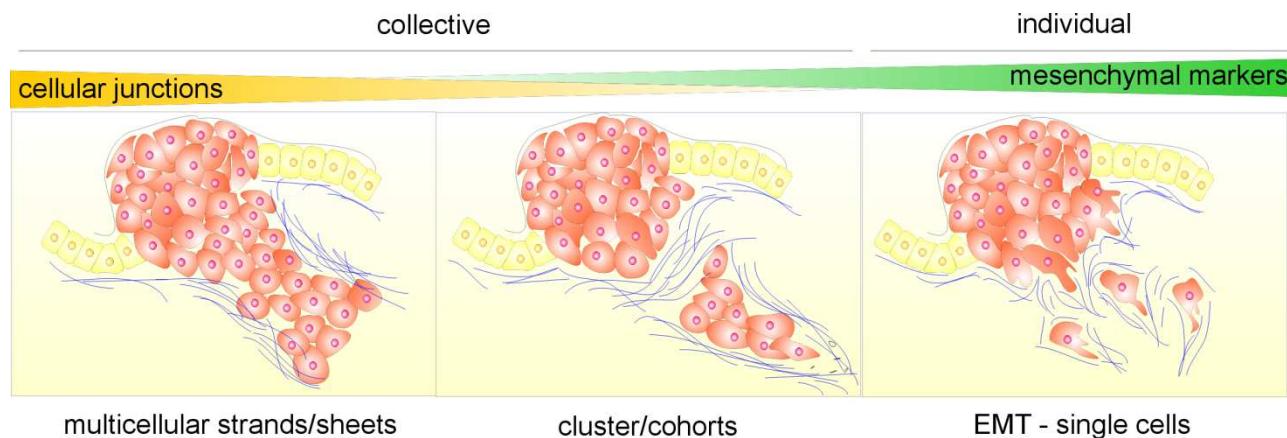
Normal regulation during development/adulthood

- Neural crest
- Mesoderm
- Palate
- Cartilage
- Wound healing

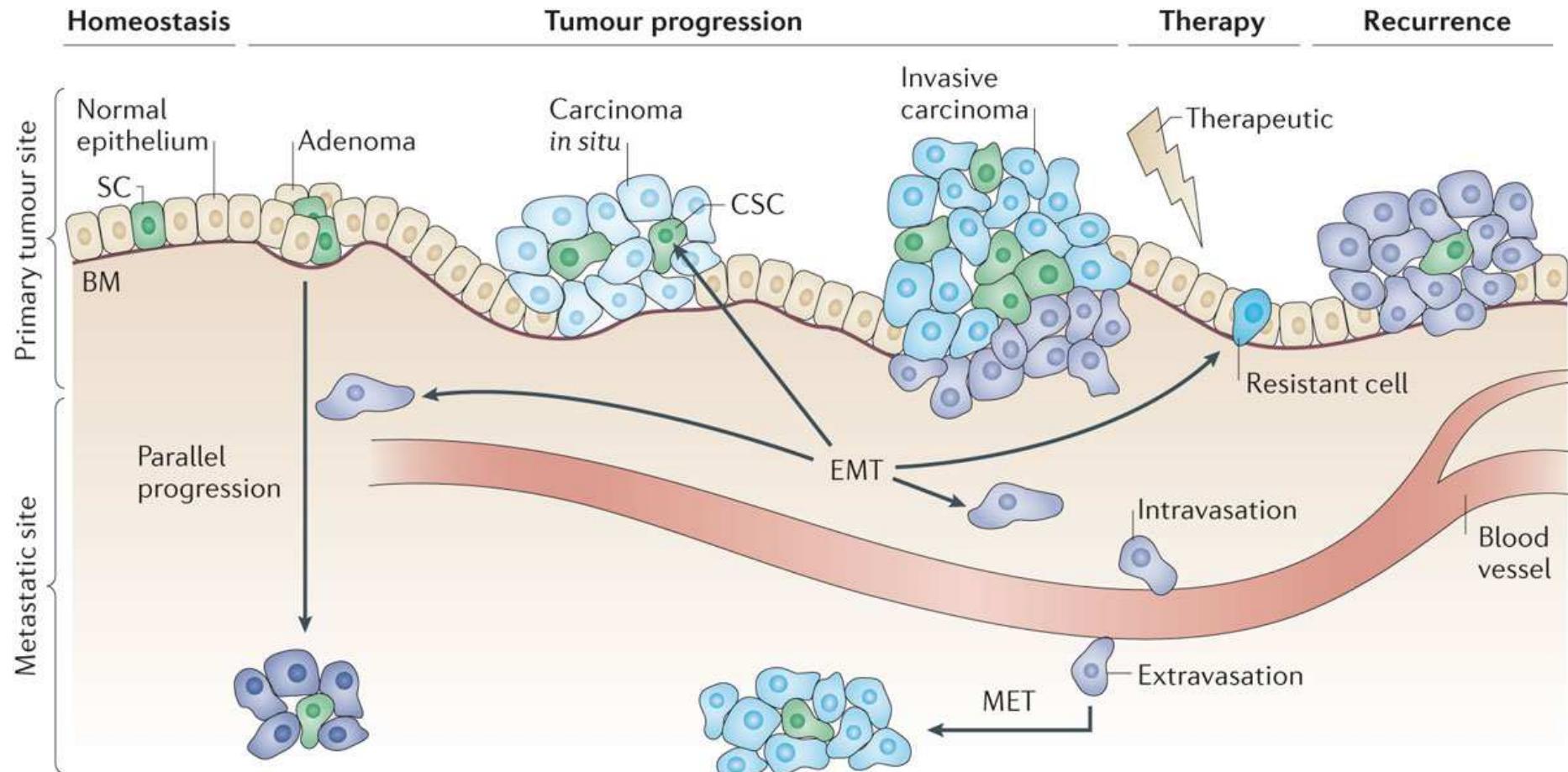


Aberrant regulation late in development or adult:

- Tumour malignancy, invasive tumour cells
- Fibrosis
- Drug resistance
- Leukemias

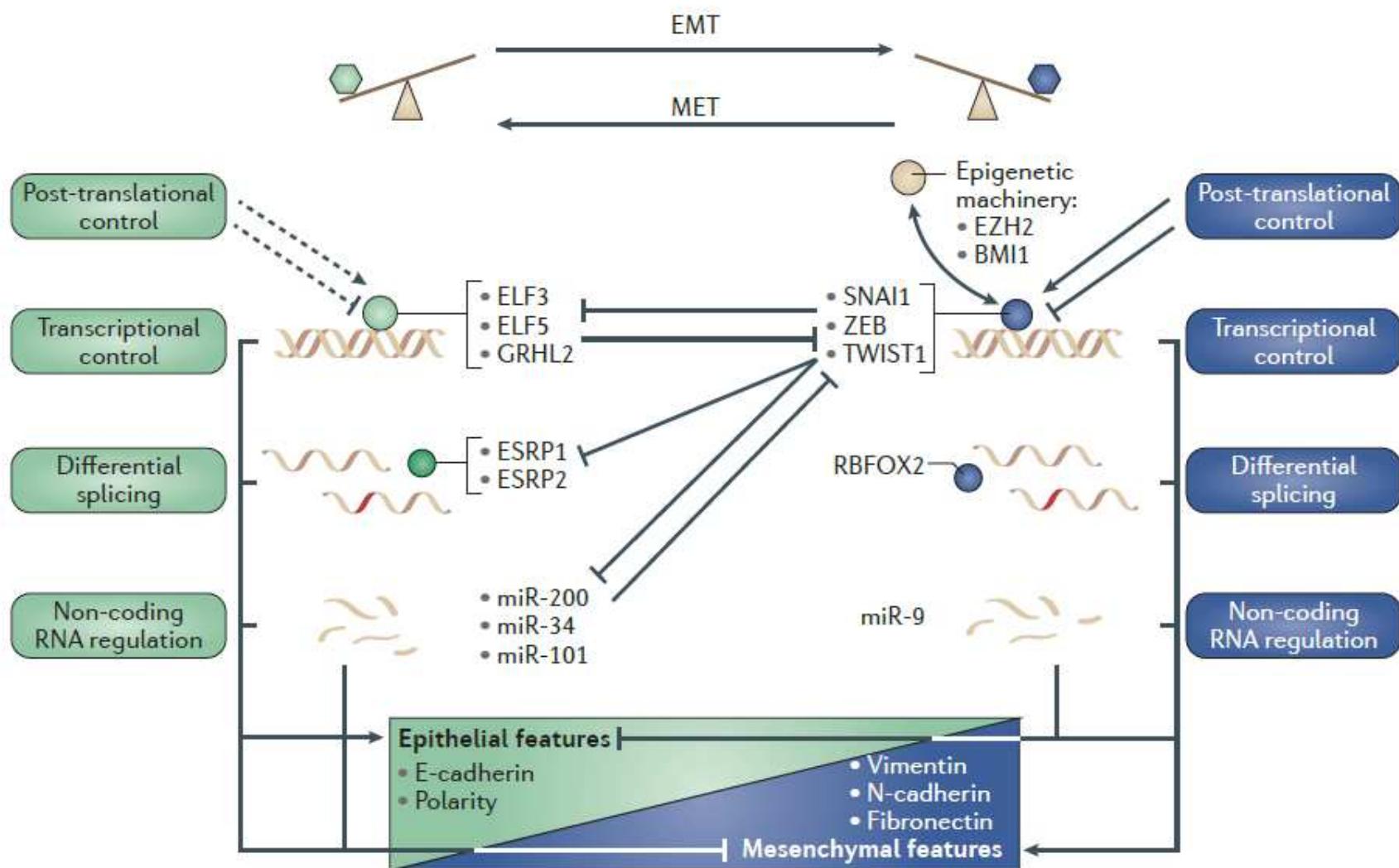


Role of EMT during cancer progression



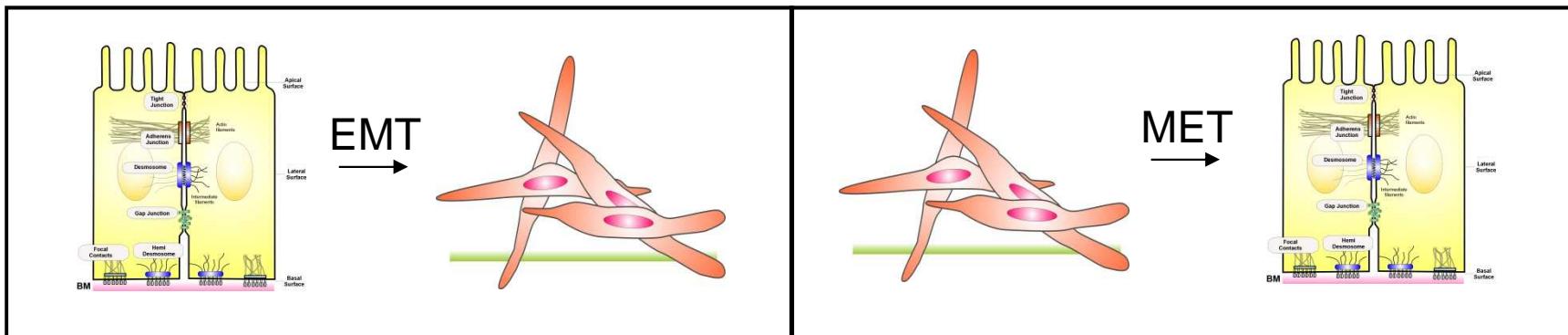
De Craene and Berx; 2013 Nature Reviews | Cancer

Regulatory levels of EMT

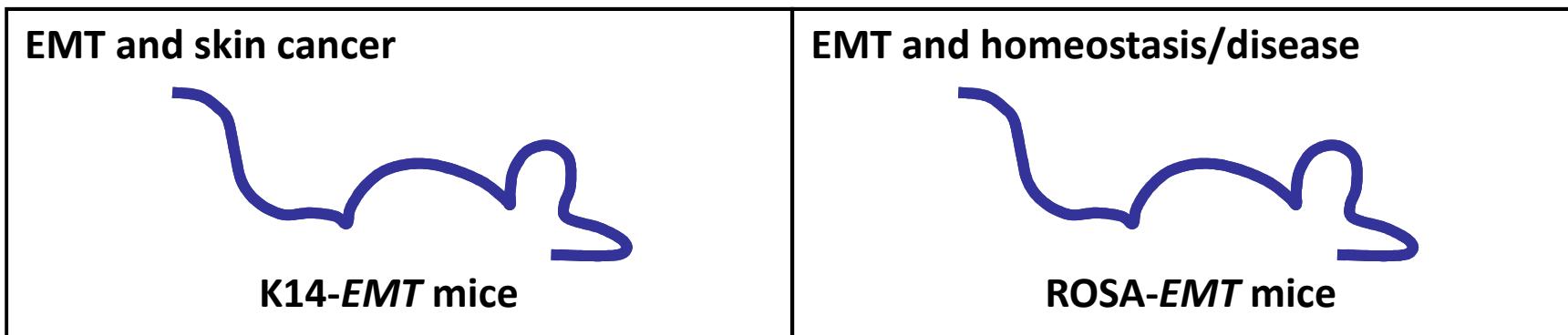


Study of molecular mechanisms of EMT driven cancer progression

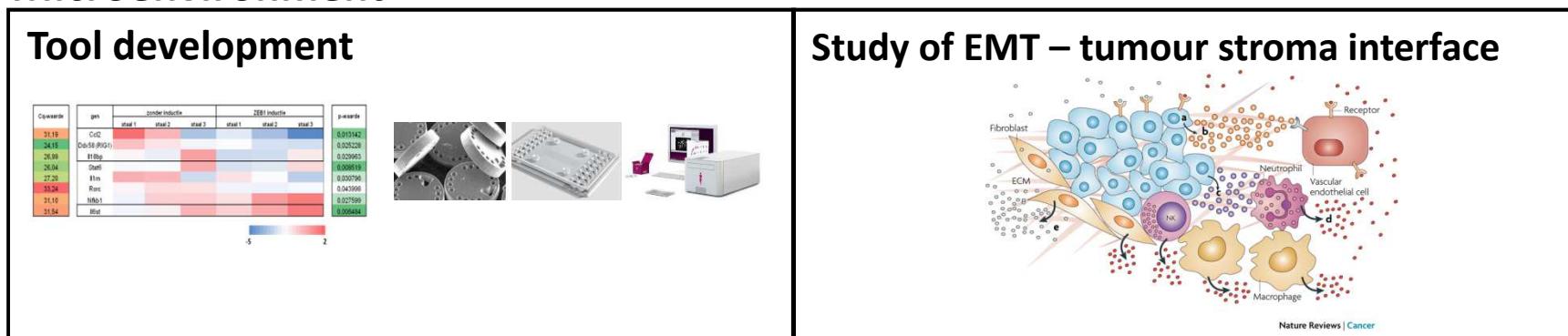
In vitro



In vivo

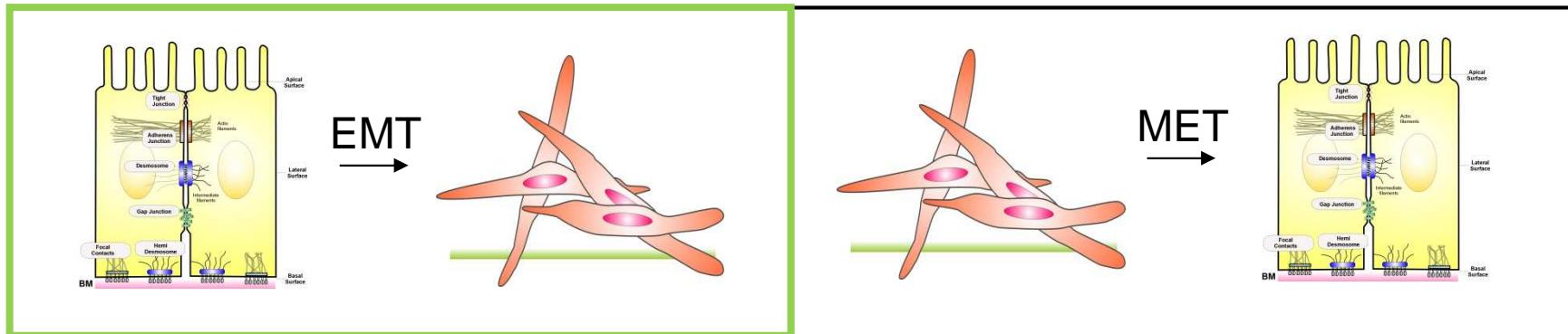


Microenvironment

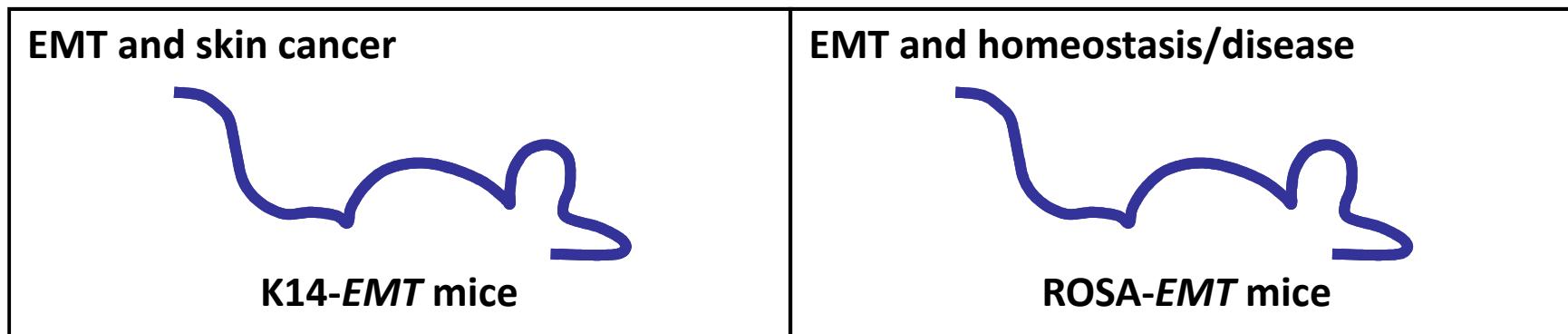


Study of molecular mechanisms of EMT driven cancer progression

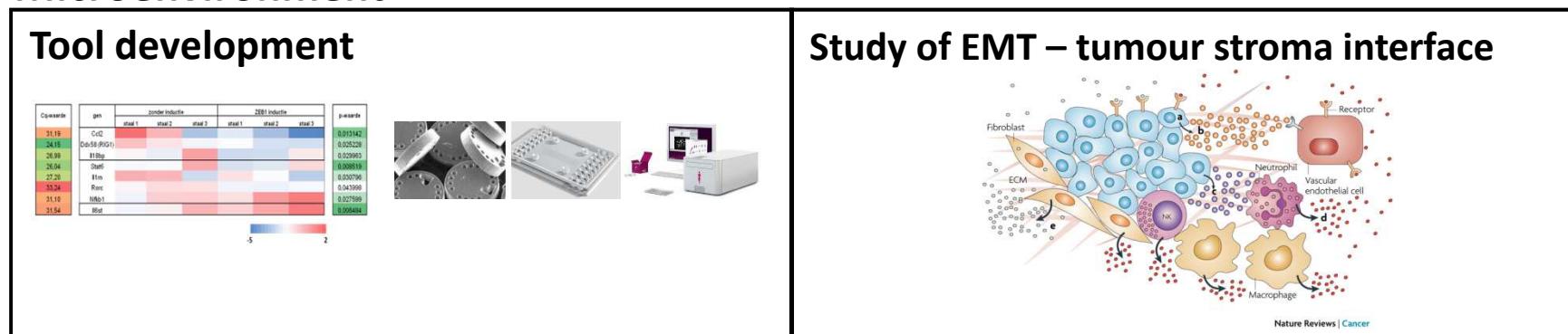
In vitro



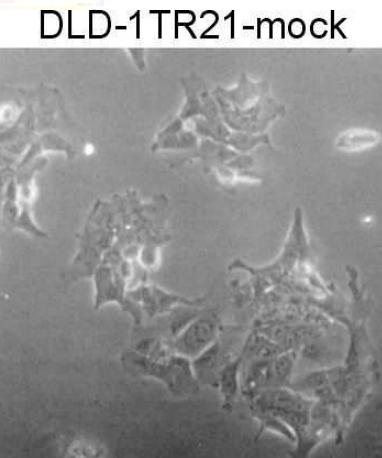
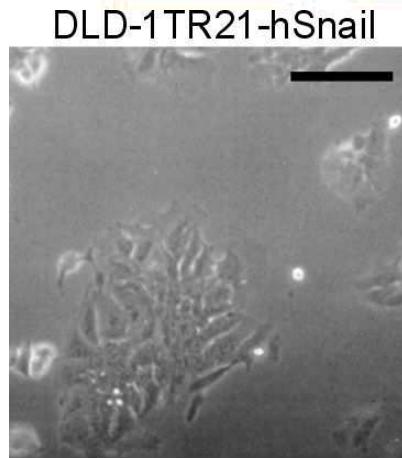
In vivo



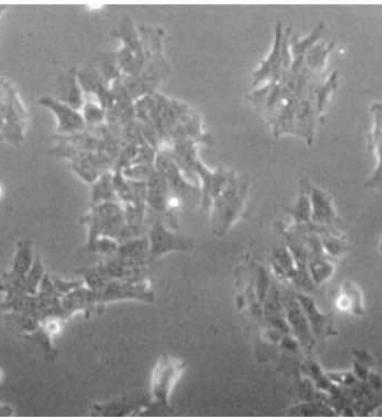
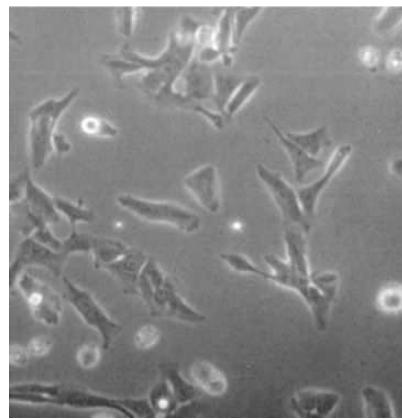
Microenvironment



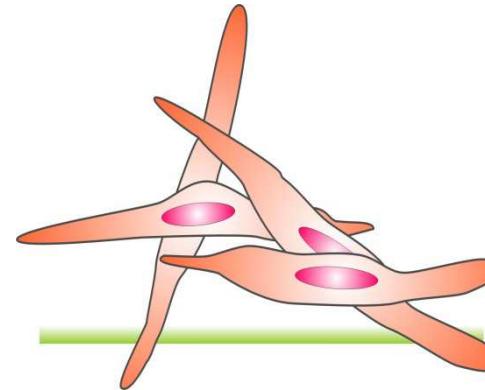
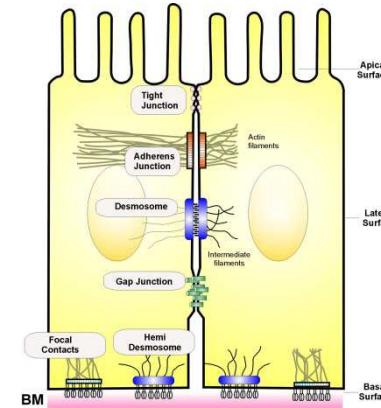
In vitro: Colorectal cancer cells with inducible expression of hSnail or hSlug



-Dox



+Dox

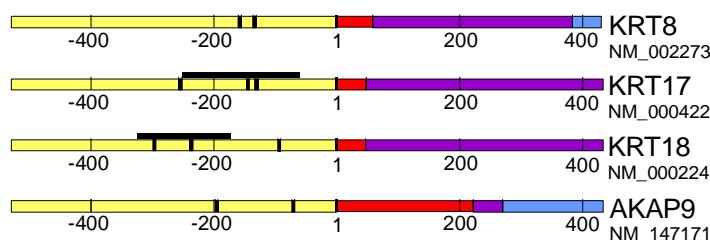


De Craene *et al.*, 2005

Identification of hSnail target genes by ChIP analysis

Epithelial differentiation

Cytoskeletal organization

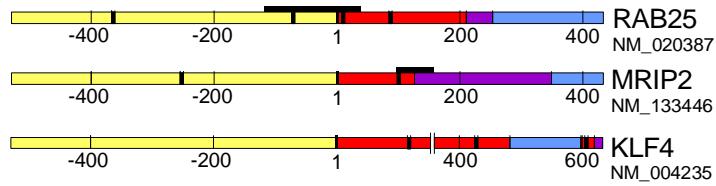


Cell adhesion

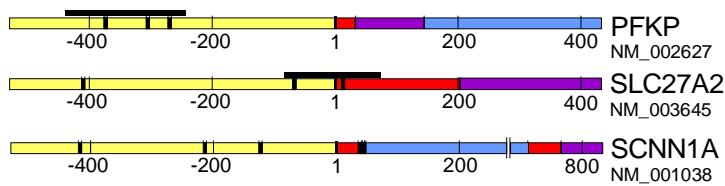


- promoter region
- 5' UTR
- ORF

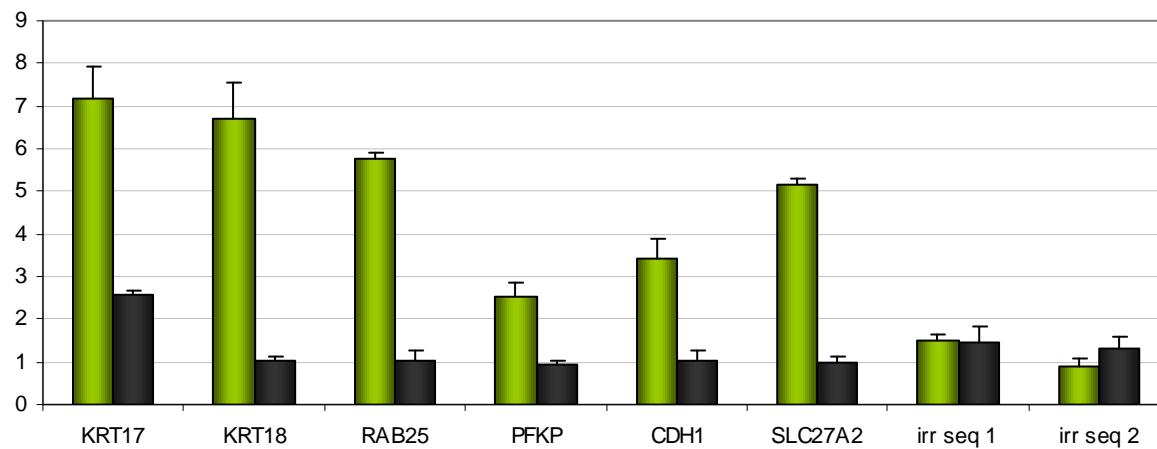
Signaling



Metabolism



- E-box CACCTG or CAGGTG
- amplicon in ChIP analysis



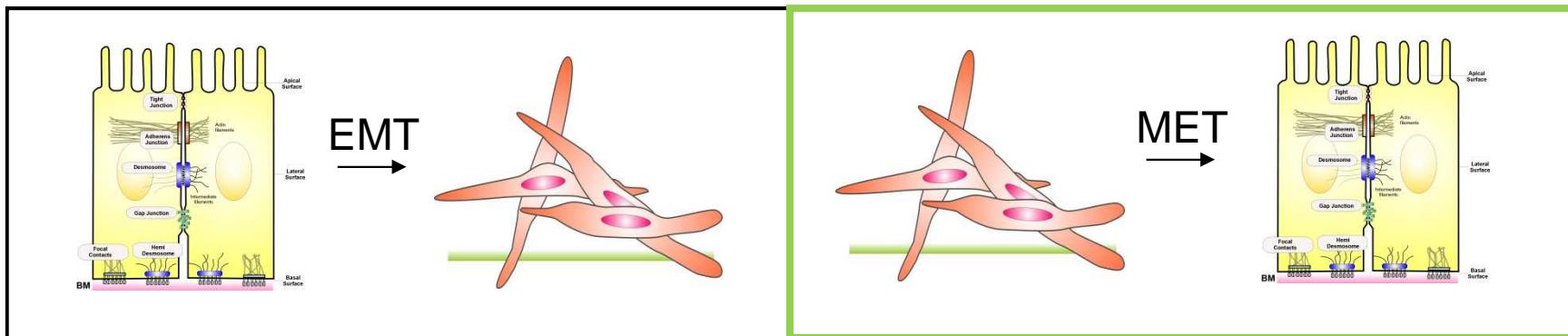
■ DLD-1TR21-EGFP-hSnail

■ DLD-1TR21-EGFP

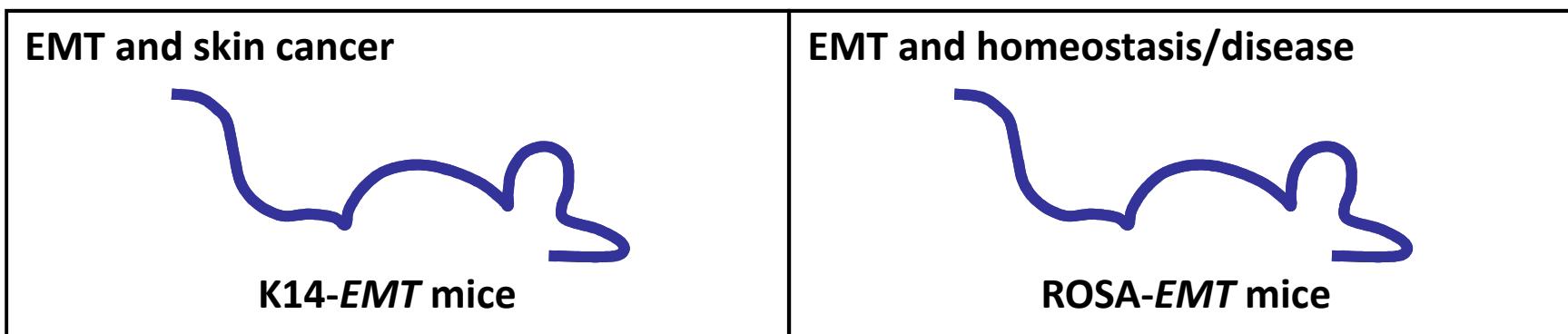
De Craene et al., 2005

Study of molecular mechanisms of EMT driven cancer progression

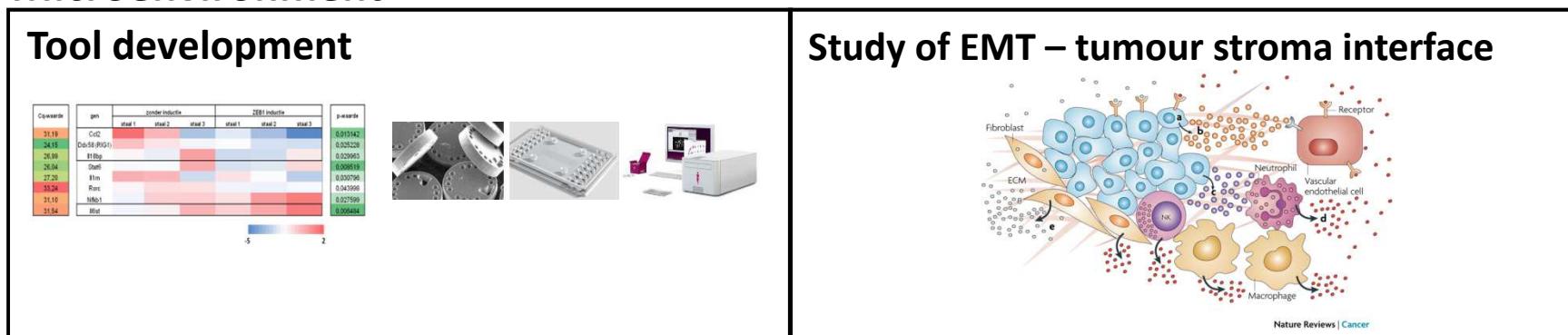
In vitro



In vivo



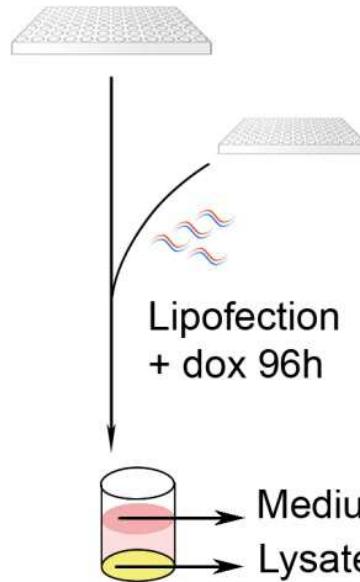
Microenvironment



Construction of *in vitro* screening system for functional EMT interactors

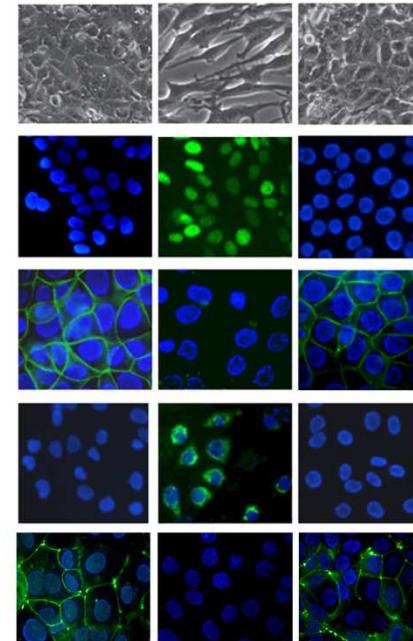
EMT sensor based on luciferase reporter gene driven by the vimentin promoter

Seeded cells



siRNA library

- dox + dox + dox
+ siZEB1



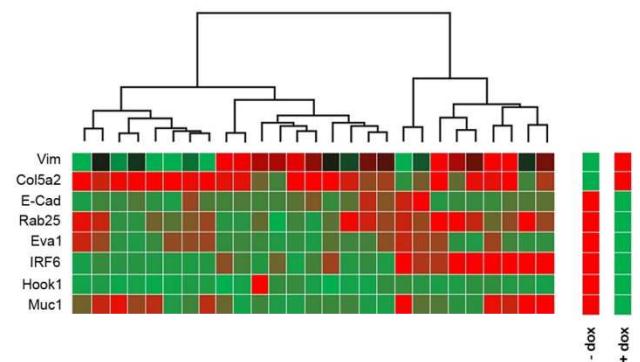
ZEB1 (myc)

E-cad

Vim

Occ

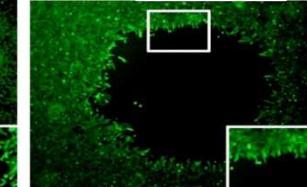
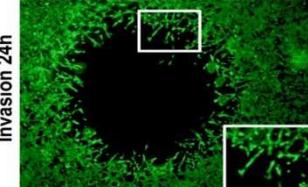
qPCR analysis



migration assay

shcon

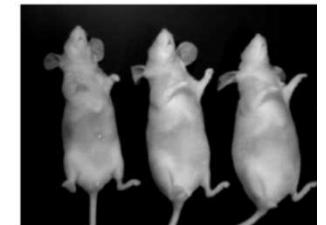
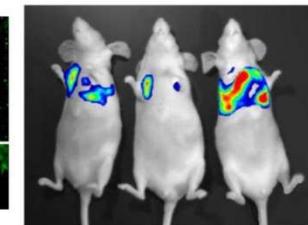
shTG



experimental metastasis

shcon

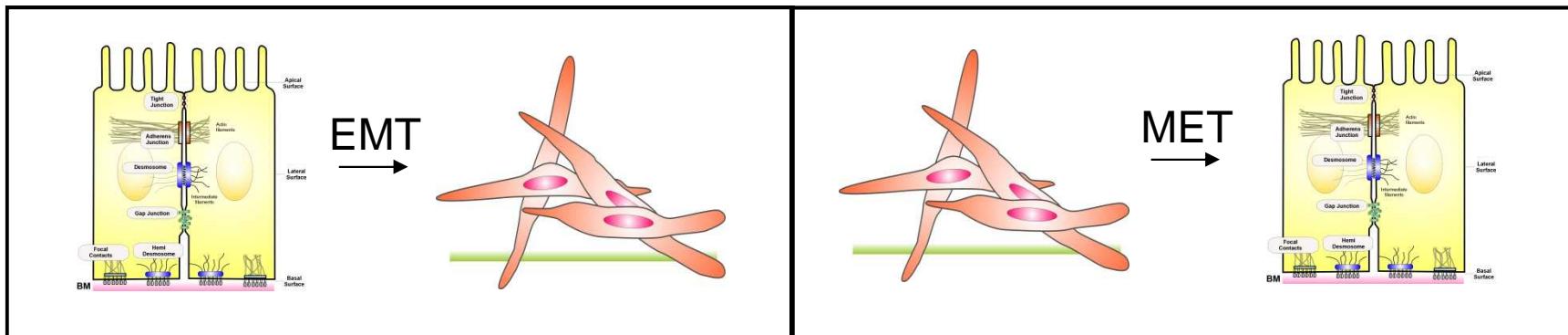
shTG



Gheldof et al., in preparation

Study of molecular mechanisms of EMT driven cancer progression

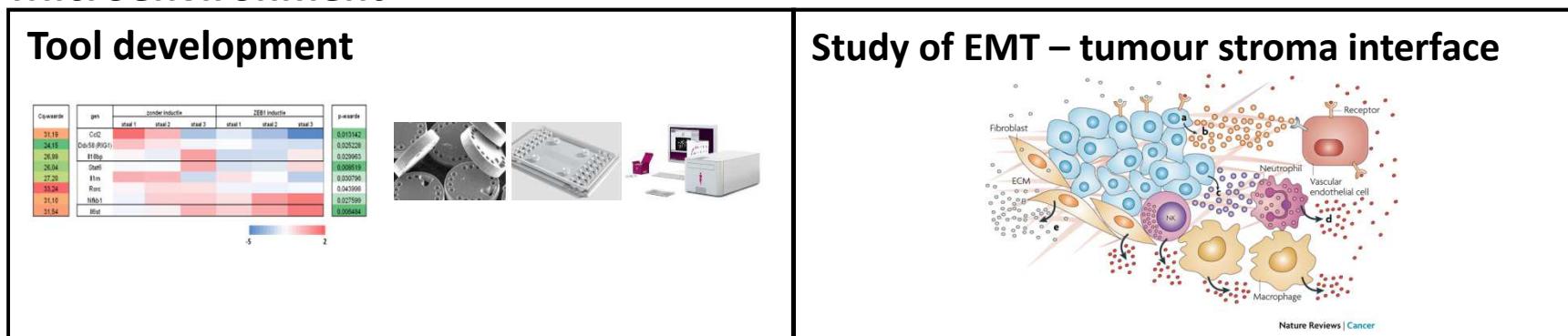
In vitro



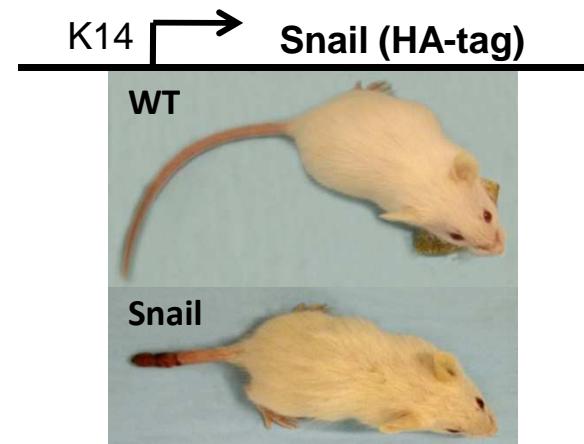
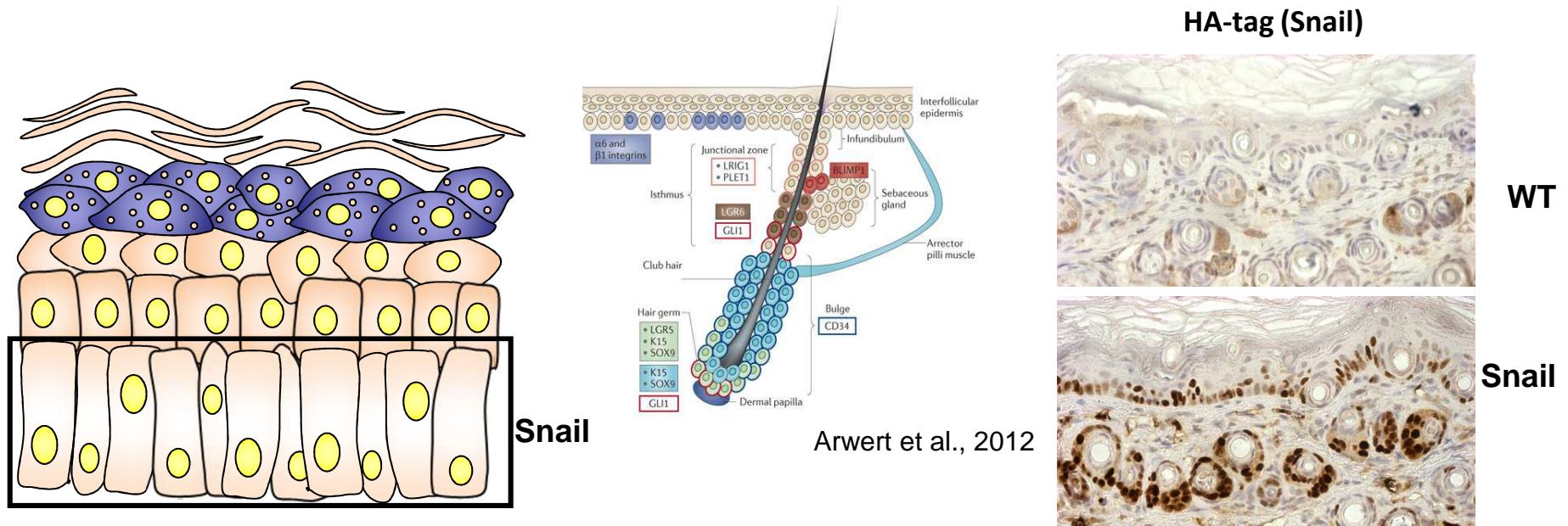
In vivo



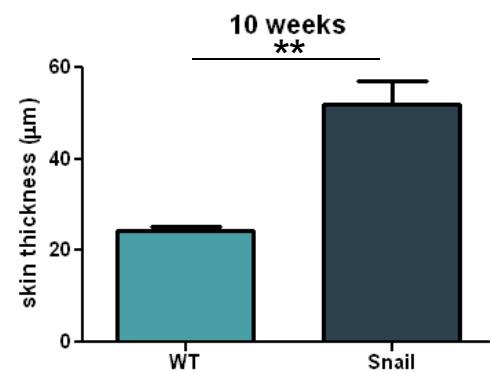
Microenvironment



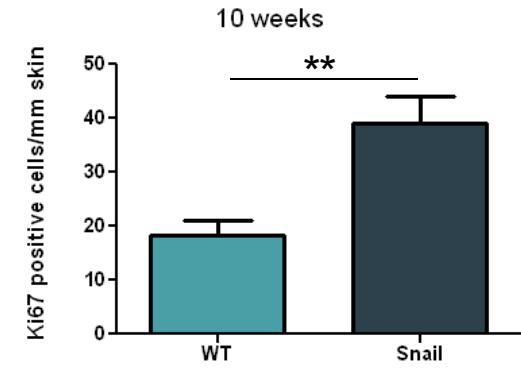
Mouse model: K14-driven overexpression of Snail: K14-Snail mice



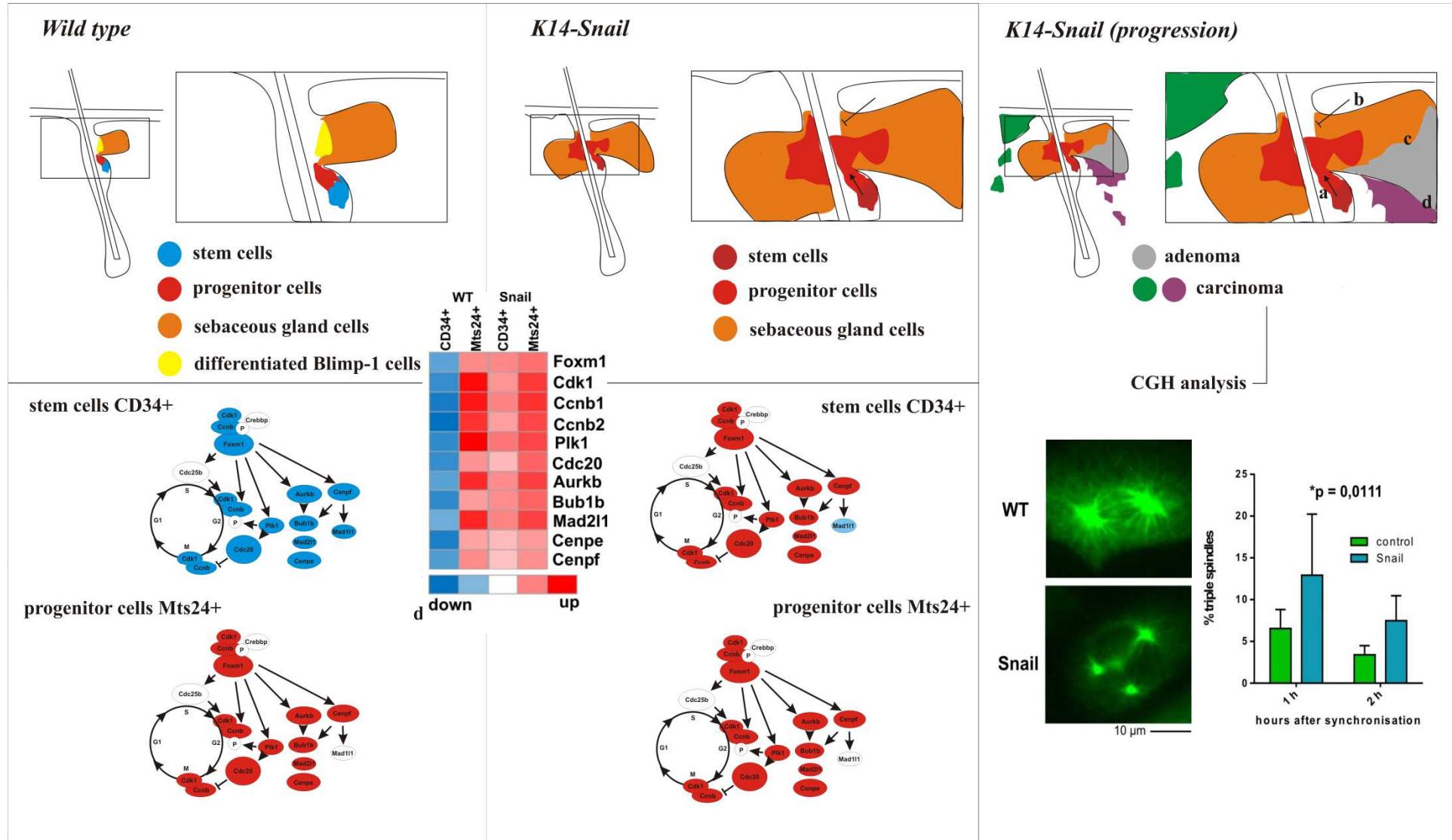
Increased epidermal thickness
in K14-Snail mice



Increased amount of Ki67 pos. cells
in the epidermis of Snail mice



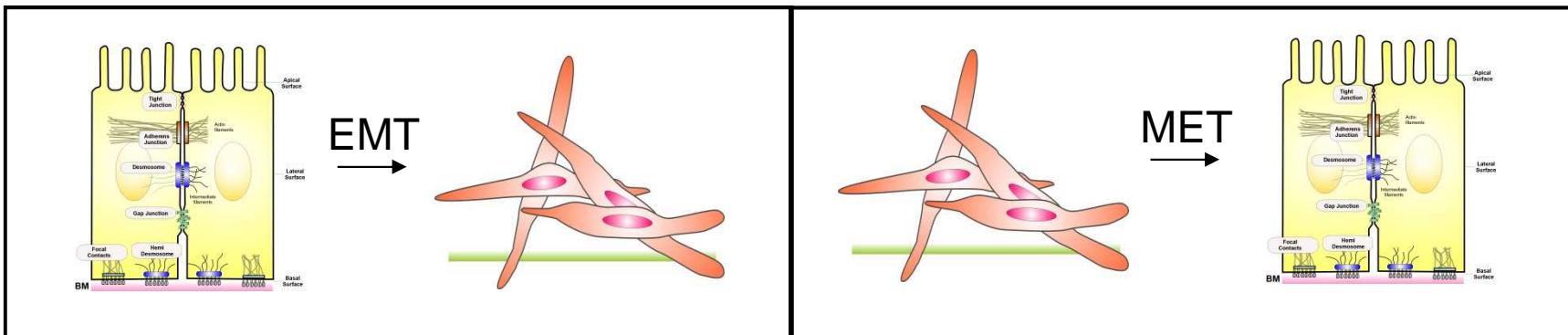
Snail induces FoxM1, a central regulator of the mitotic program and chromosome stability



De Craene et al., 2014

Study of molecular mechanisms of EMT driven cancer progression

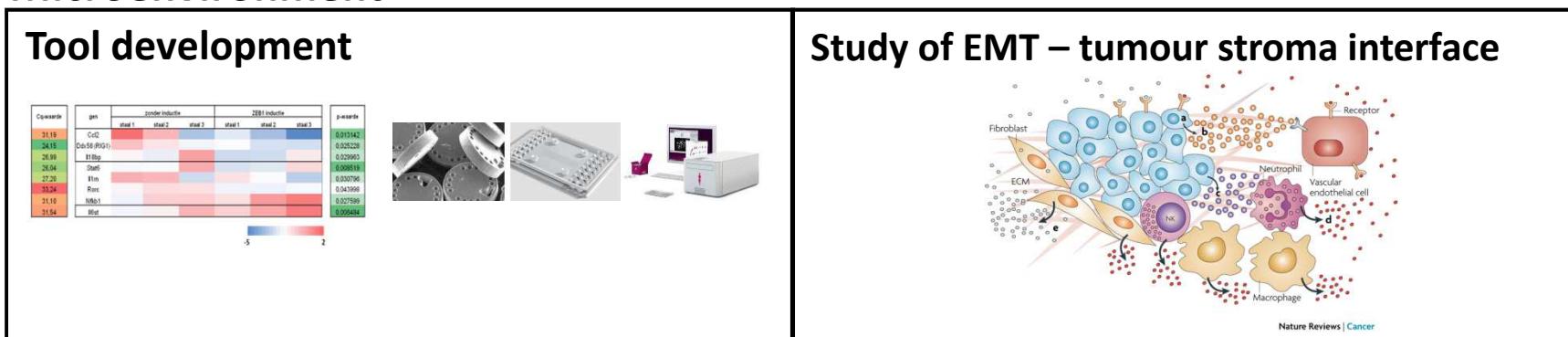
In vitro



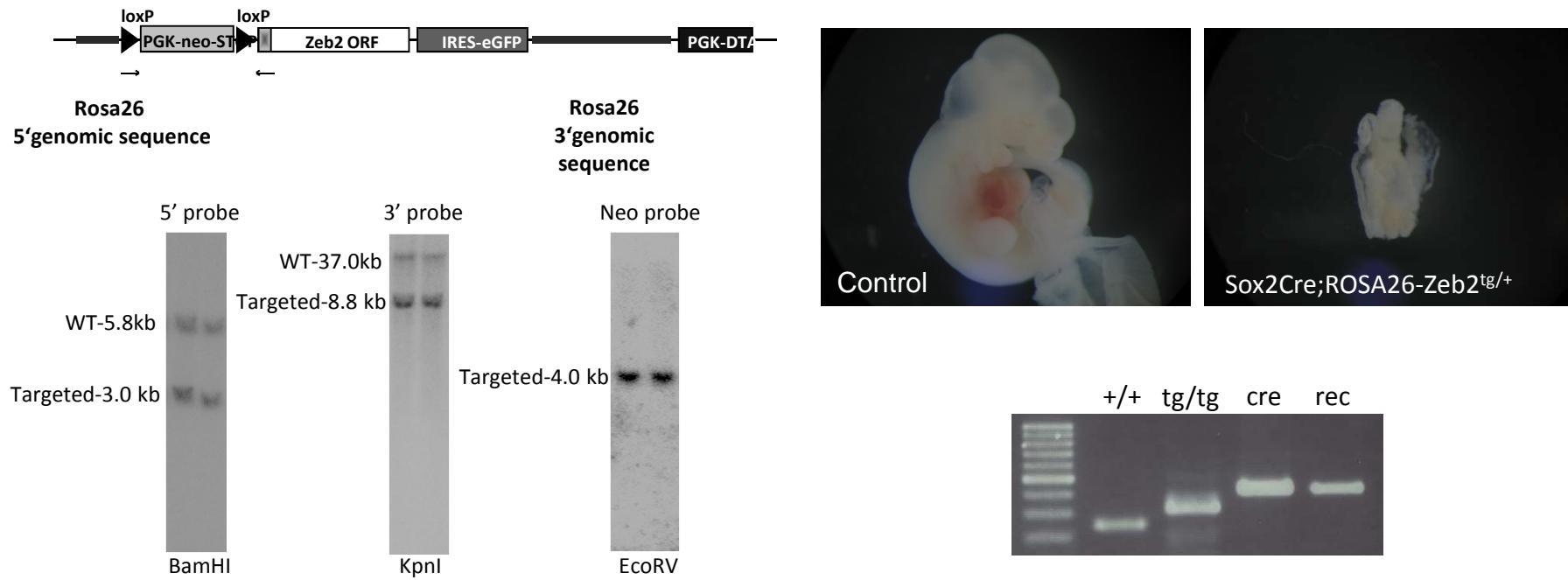
In vivo



Microenvironment

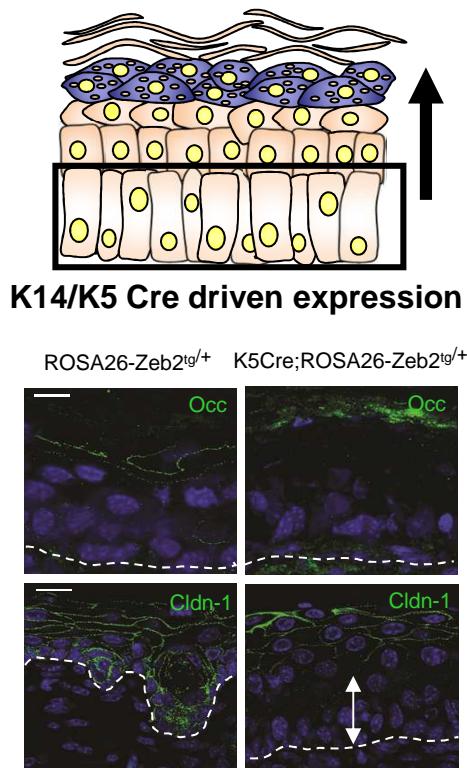


Generation of a conditional transgenic mouse model for Zeb2

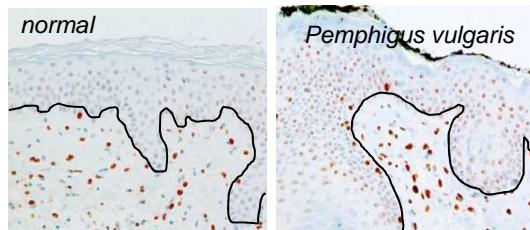


Crucial role of the EMT transcription factor Zeb2 in homeostasis and disease

skin/epidermis

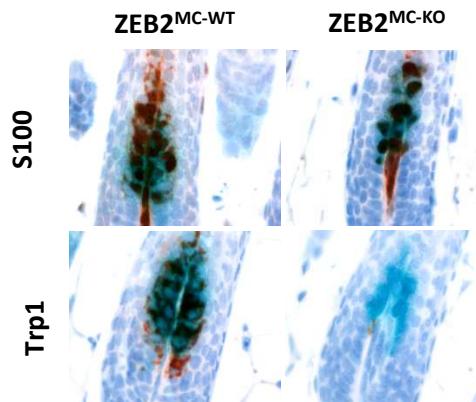


Expression in human Pemphigus vulgaris

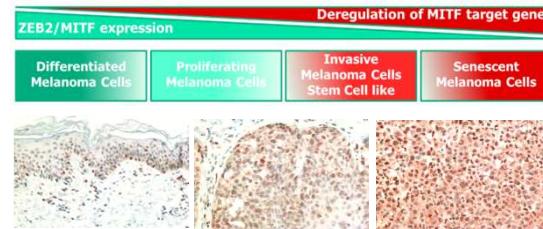


Tatari et al., 2014

melanocyt

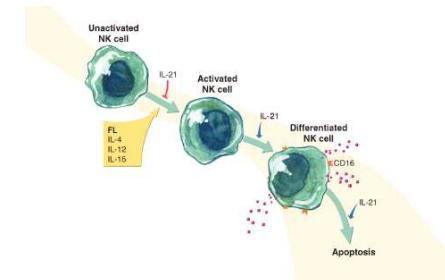


Role in melanoma

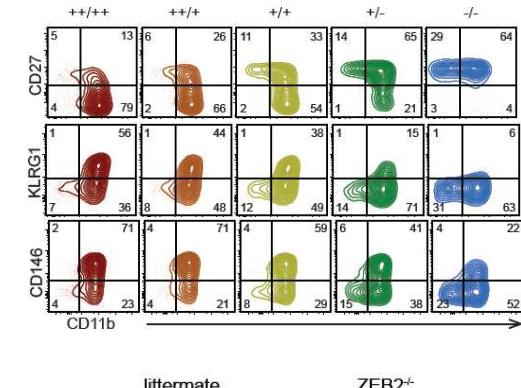


Denecker et al., 2014, Vandamme et al., in prep

immune cells



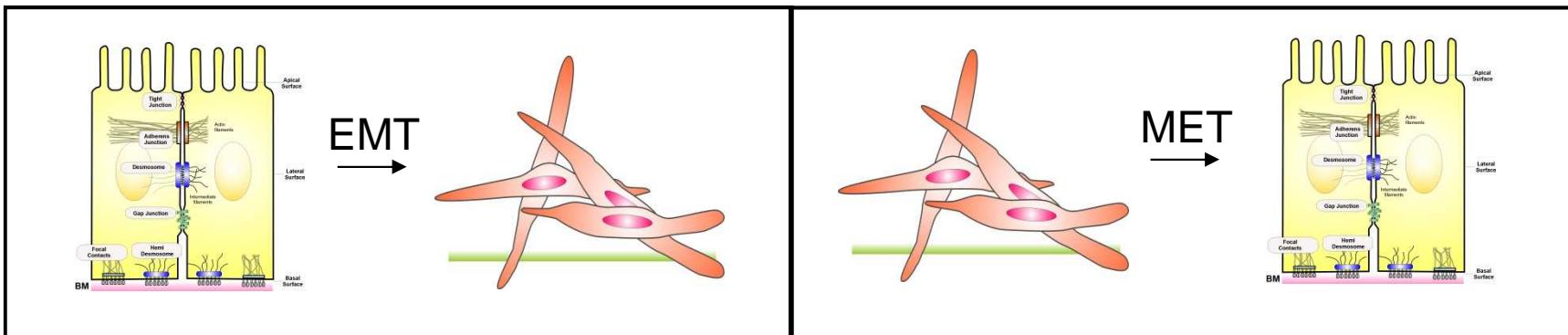
Cre driven expression or deletion



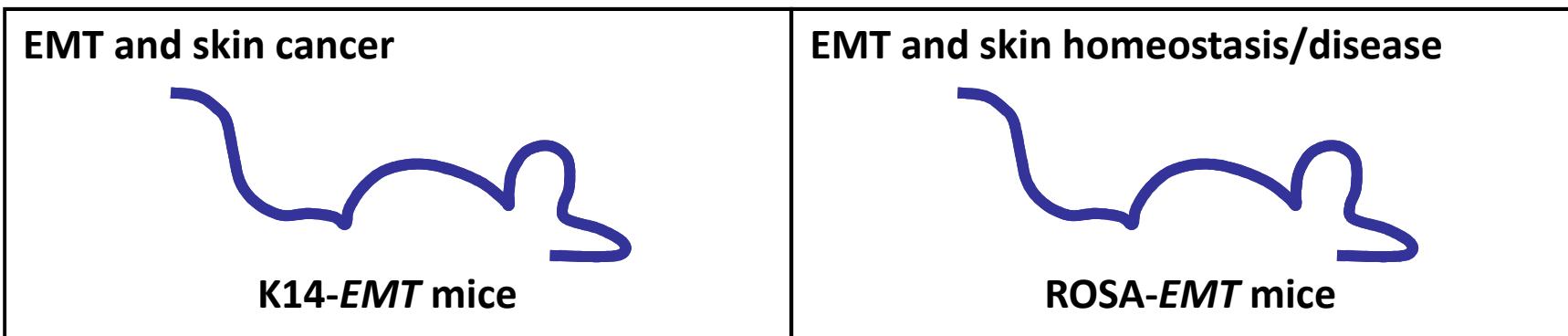
Van Helden et al., submitted

Study of molecular mechanisms of EMT driven cancer progression

In vitro

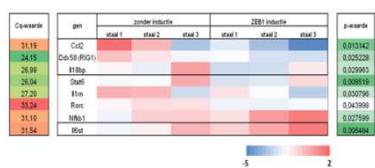


In vivo

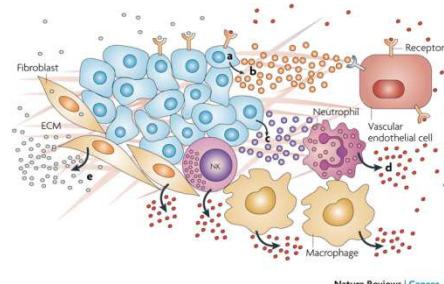


Microenvironment

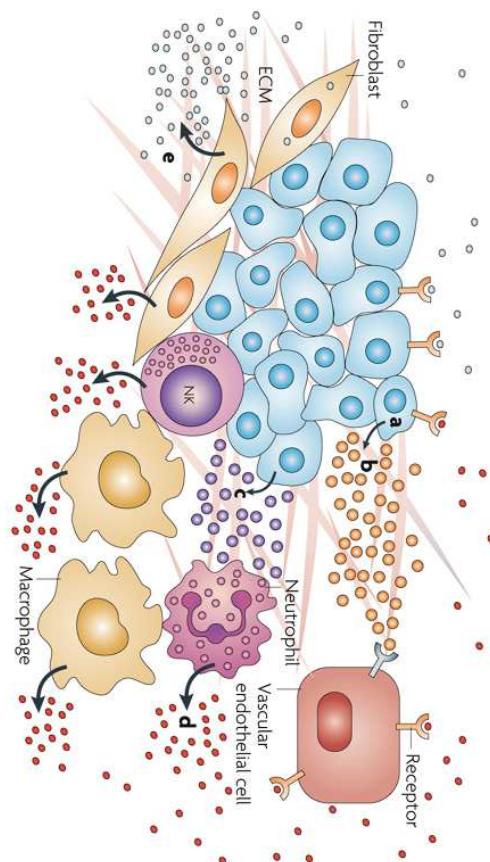
Tool development



Study of EMT – tumour stroma interface



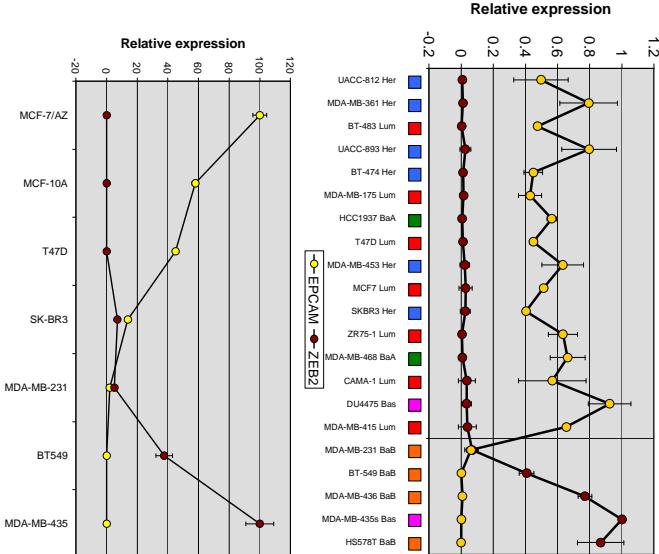
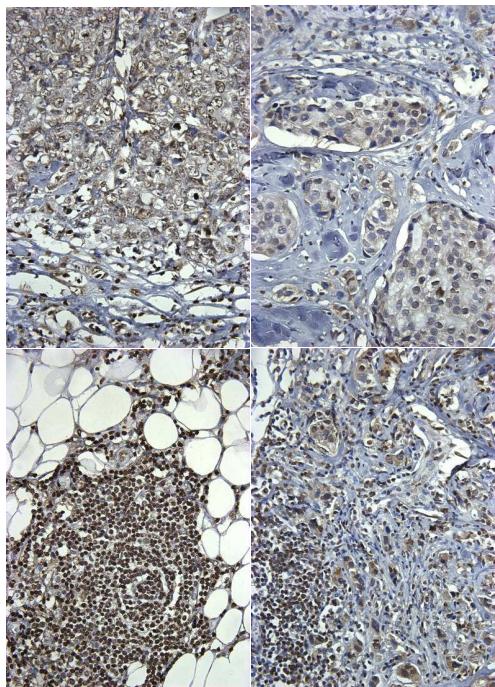
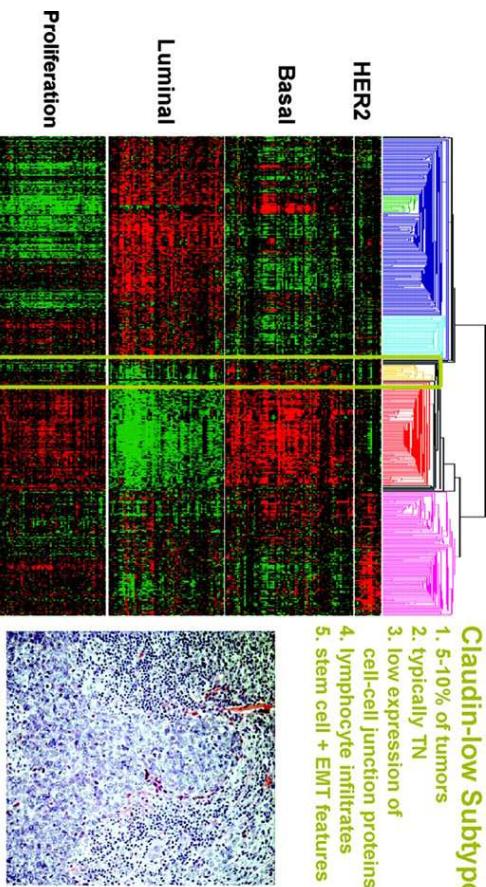
Tumour, microenvironment & tumour associated cells Contribution of Zeb2 in tumour cells?



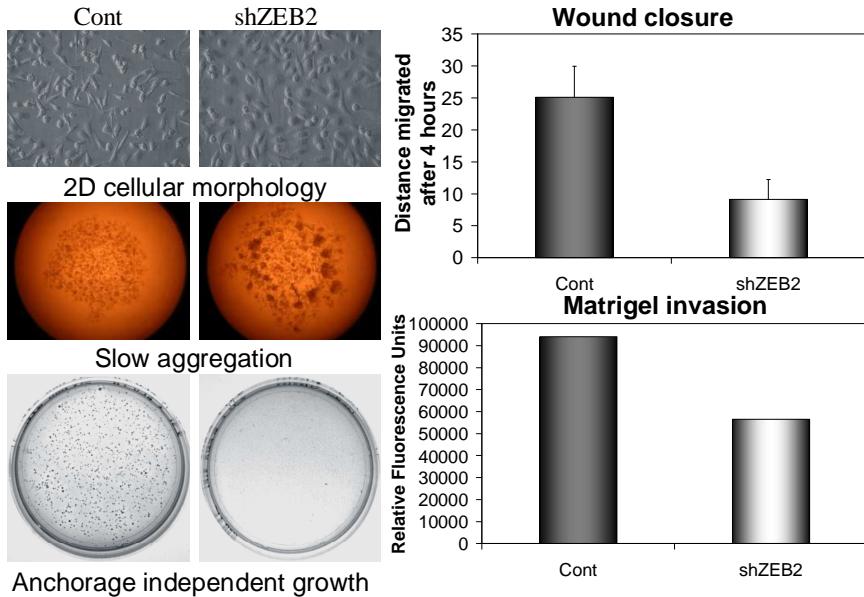
Nature Reviews | Cancer

Claudin-low Subtype

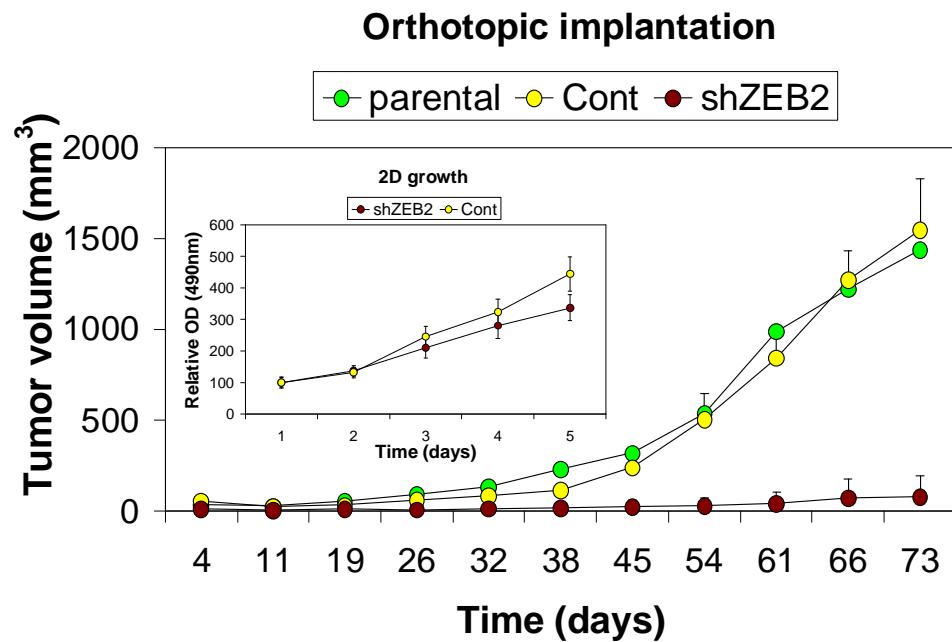
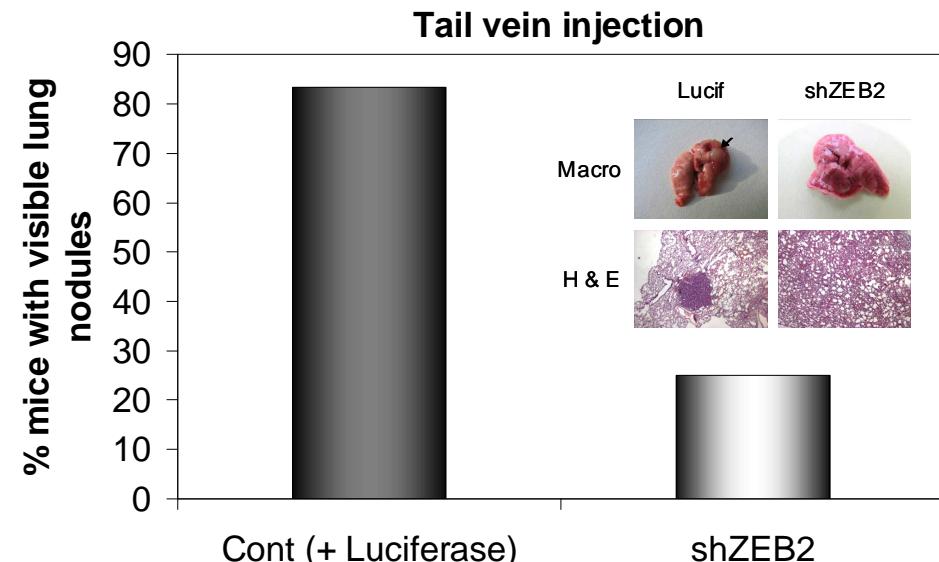
1. 5-10% of tumors
2. typically TN
3. low expression of cell-cell junction proteins
4. lymphocyte infiltrates
5. stem cell + EMT features



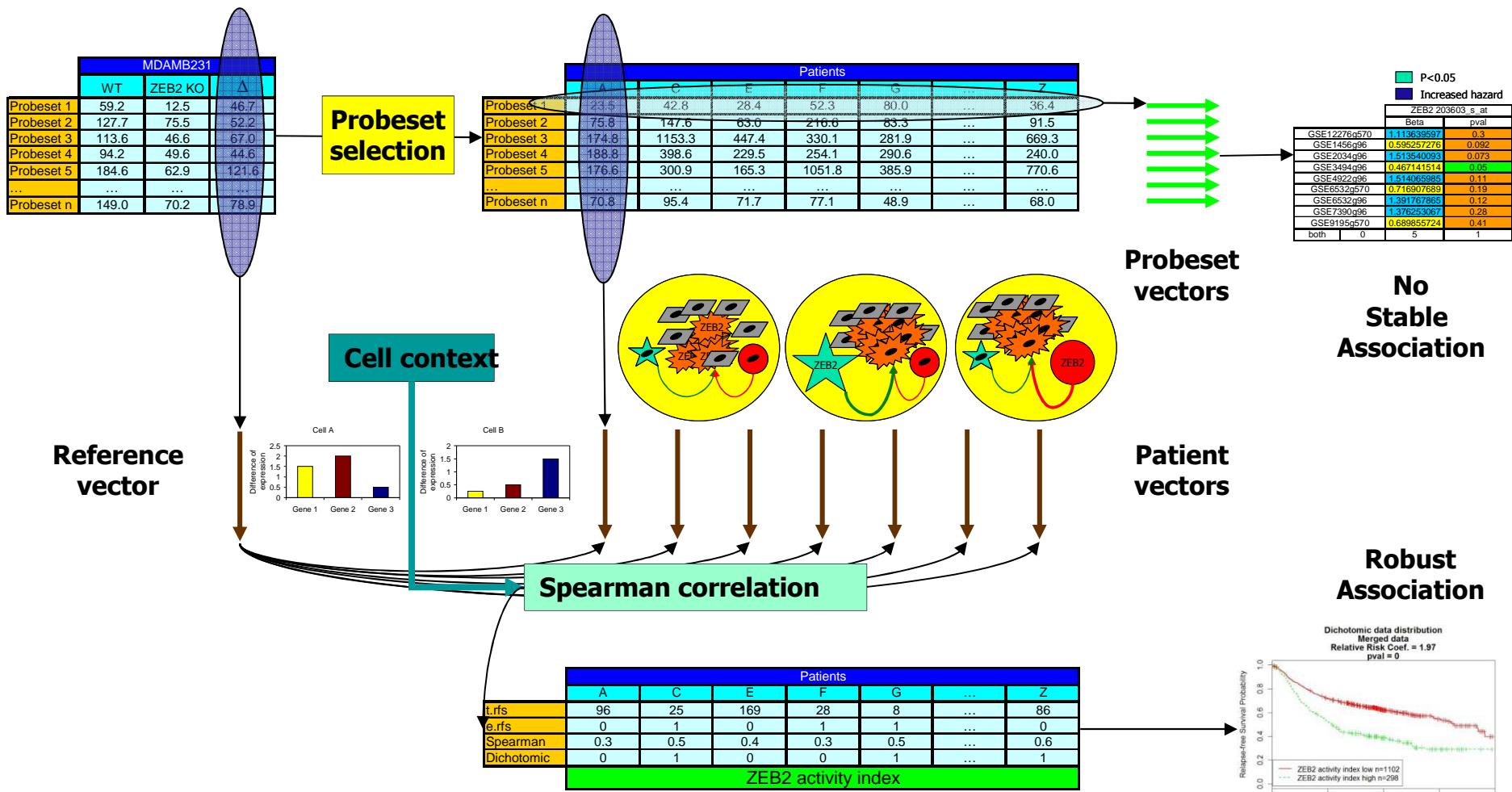
Impact of Zeb2 knock-down in basal breast cancer cells



- **Increased epithelial morphology**
- **Increased aggregation**
- **Reduced 2D and 3D migration**
- **Impaired lung colonization**
- **Minor effect on 2D growth**
- **Impaired anchorage independent growth**
- **Impaired in vivo growth.**



Breast cancer microarray: Zeb2 activity index building



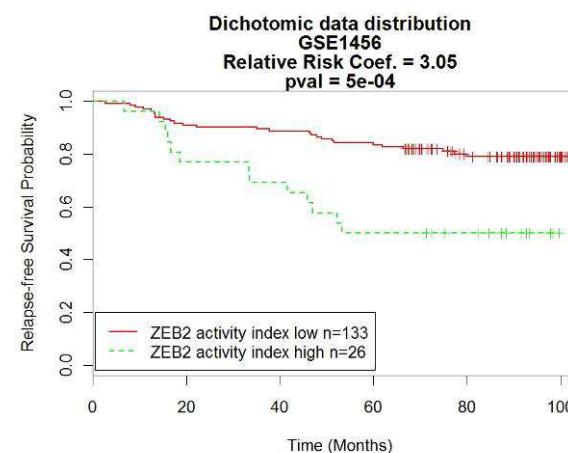
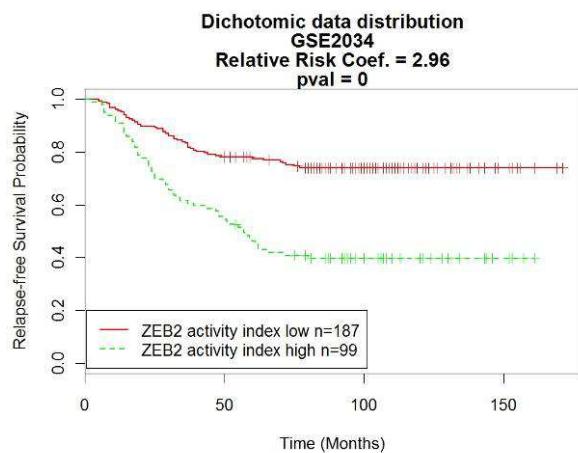
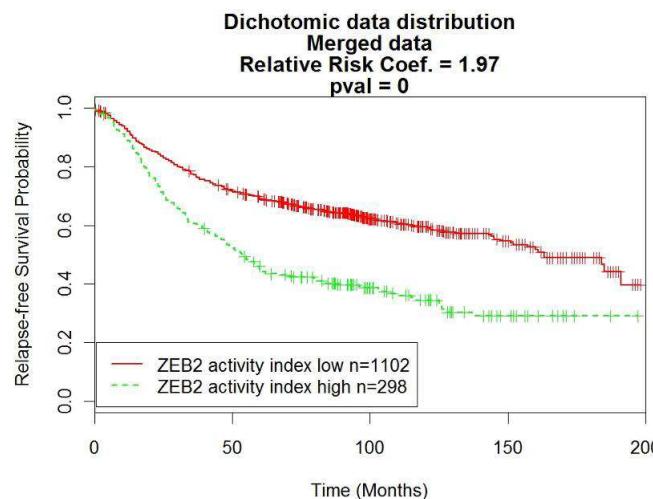
- ZEB2 target genes → ZEB2 activity index.
- Which tumour behaves as if ZEB2 is knocked-down in MDAMB231 cells?

Zeb2 transcription factor activity: survival analysis



Raw ZEB2 activity index

Studies	Pooled data (n=1400)		
Count (tot=9) p<0.05 Rel Risk>1	Rel. Risk	p values	
ZEB2	0	1.002	1.50E-01
Full List n=36	6	24.5	5.30E-11
Selected List n=16	6	5.3	6.90E-12



Raspe et al., submitted

Metagene expression signature for prognosis of breast cancer patients.
Priority date 2010.10.29. Published: WO2012/056047 A1 (05.12.2013)

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and Development
Prof. Dr. Elaine Fuchs

University of Cologne (Germany)
Department of Dermatology
Prof. Dr. Cornelia Mauch

PhD student position available !!!!
Postdoc position available !!!!