

Selected recent publications

Poulton et al. (2017) Journal of Cell Biology, 216, 1255-1265.
 Choi, W. et al. (2016) Journal of Cell Biology 213, 243-260
 Pronobis et al. (2015) eLife 4, e08022
 Poulton et al., (2014). Developmental Cell 30, 731-745
 Bilancia et al., (2014) Developmental Cell, 28, 394-408

The Peifer lab: Choosing cell fate and assembling the body plan: how it goes right in embryos and wrong in cancer



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Our goal: Define how cells choose fate and assemble tissues and organs, and determine how this goes wrong in cancer

How did my body get built?

We combine two major model systems:

Drosophila embryonic development

Normal and tumor-derived cultured mammalian cells

In our own lab and via collaboration we take a very multidisciplinary approach

Modern genetic tools including RNAi and CRISPR

Cutting edge cell biology and microscopy

Computational image analysis

Biochemical approaches including mass spec

How do cells remodel their cytoskeletons and cell-cell junctions so that they can change shape and move during morphogenesis without disrupting epithelial integrity and barrier function?

Cadherin-catenin complexes mediate cell adhesion and link junctions to the cytoskeleton

We're also using shRNA knockdown and SIM superresolution microscopy to define how mammalian cells link junctions and the cytoskeleton to maintain tissue integrity

Loss of Cdc42 in vivo disrupts critical cell shape changes like apical constriction during mesoderm invagination

Combining live cell imaging and computational tools allow us to put numbers on dynamic cytoskeletal events revealing new insights

How do cells establish and maintain apical-basal polarity and how does loss of polarity contribute to oncogenesis?

Apical-basal polarity is critical to assemble and maintain the function of all epithelial tissues

The early Drosophila embryo is the premier system to study establishment of apical-basal polarity

We are defining the network of proteins that act in polarity establishment

We are continuing to expand this network and define the mechanisms by which proteins act

How do cells regulate their actin cytoskeletons during cell migration and shape change and how do oncogenic mutations alter this?

Our challenge: to determine how cells use this complex actin regulatory toolkit during normal development and how it goes wrong in disease

We use Drosophila and cultured cells to explore this at ALL levels of organization

Molecular machines Cells Whole animals

We also explore how the oncogenic kinase Abl regulates the cytoskeleton in both epithelial cells and the nervous system

Genetics and live imaging provide insights into how cells make protrusions and thus migrate in a dish and in vivo

Lamellipodia Filopodia

How do tumor suppressors like APC and Axin tightly regulate cell-cell Wnt signaling to allow proper cell fate choices and avoid oncogenic transformation?

The tumor suppressors APC and Axin, mutated in most colorectal cancers, are negative regulators of the Wnt pathway

We're using human colorectal cancer cells to define the workings of this megadalton Multiprotein machine

Combining SIM Superresolution microscopy with FRAP provided the first look inside this machine revealing a role for APC in stabilizing Axin polymerization

Functional tests in flies and cultured cells revealed a key role for phosphorylation by GSK3

Current Funding

NIH 1 R35 GM118096 (MIRA Award)
 Regulating cell fate and shaping the body plan during morphogenesis and their alteration during oncogenesis July 2016-June 2021

Peifer Lab Trainees

18 Ph.D. Students and 20 postdocs have or are training in our lab. Alumni have gone on to success in diverse careers in science. Examples include:

Tony Harris, Professor, U Toronto
 Don Fox, Assistant Professor at Duke
 Nasser Rusan, Investigator & Lab Head, NHLBI
 Catarina Homem, PI, CEDOC, Lisbon
 Karen Hales, Professor, Davidson College
 Ed Rogers, Senior Scientist, Janelia
 Joe Louriero, Senior Investigator I, Novartis Institutes for Biomedical Research
 Lizz Grevengoed, Physician Assistant with Western Oncology and Hematology

Full list at: <http://peiferlab.web.unc.edu/alumni/>

How do cells maintain genome stability in the face of mutations affecting proteins that help build the mitotic spindle or regulate key cell-cycle checkpoints?

Centrioles were thought to be the key organelle that nucleates the mitotic spindle

We found loss of centrosomes slowed but did not prevent Spindle assembly, but many cells lost genome stability and underwent apoptosis

Mutations affecting human Centrosomal proteins are the primary genetic cause of microcephaly

Centrosome plus SAC loss leads to massive aneuploidy and polyploidy, a hallmark of most advanced-stage cancer cells

Genotype	# Normal	# Aneuploid	# Polyploid	% Abnormal
wt	140	4	0	2.8%
mad2	108	7	1	6.9%
sas-4	129	15	0	10.4%
mad2; sas-4	12	34	14	80.0%

While fly brains can tolerate Centrosome loss, Combined loss of centrosomes plus the spindle assembly checkpoint (SAC) leads to microcephaly