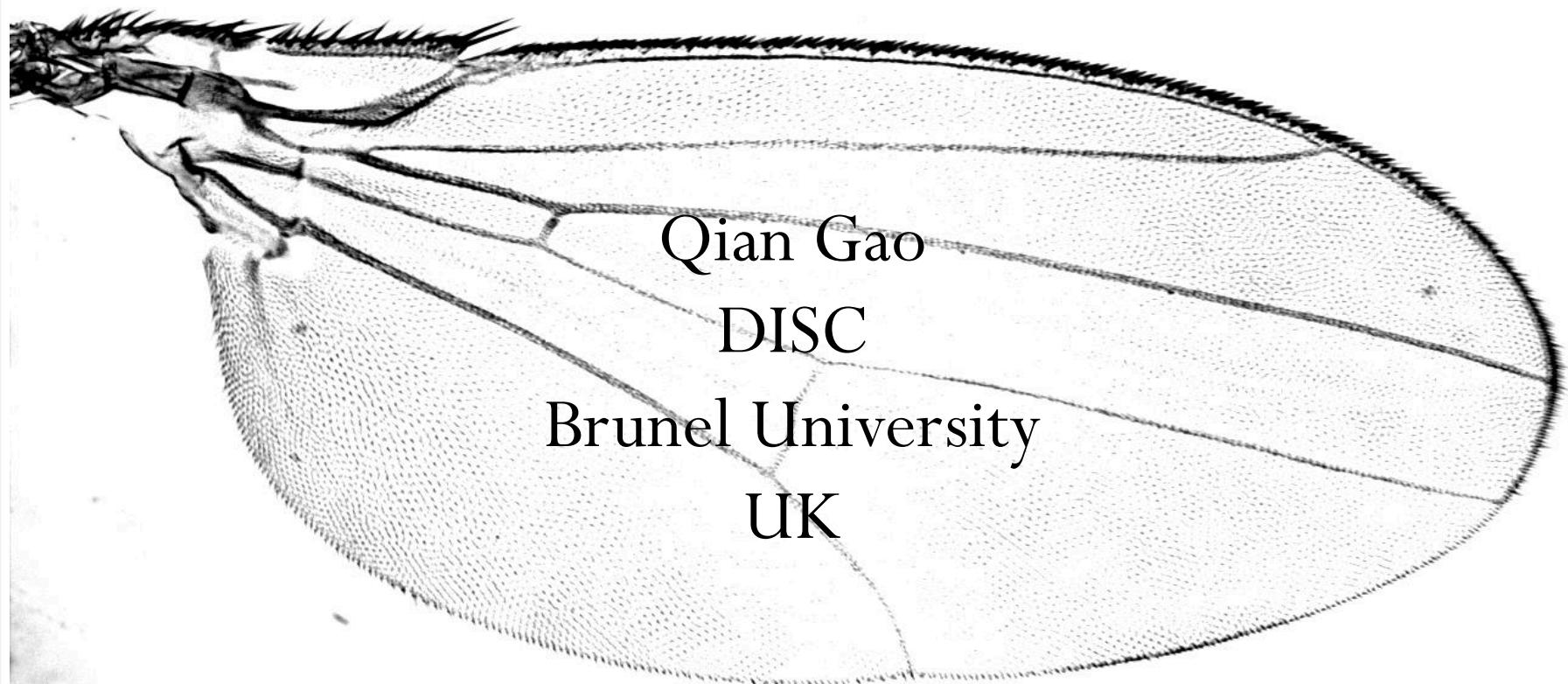
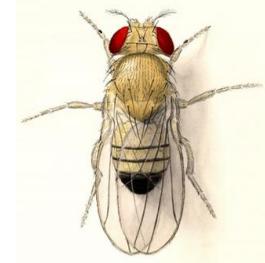


Investigating Planar Cell Polarity by Modelling



Overview

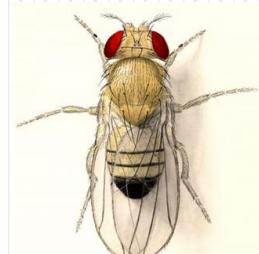


1. Planar cell polarity
2. Modelling approach
3. Some progress in modelling

Planar Cell Polarity Signalling

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Department of Biosciences
Brunel University
UK

ICSB Tutorial, Heidelberg 2011



Aims and Objectives

- The role of *Drosophila Melanogaster* as a system model
- To outline Planar Cell Polarity (PCP)

Drosophila melanogaster

- History of research into genetics and development
- Many genetic and cellular technologies developed, genomes sequenced
- 61% of human disease genes have fly homologues (similar)

The procedures

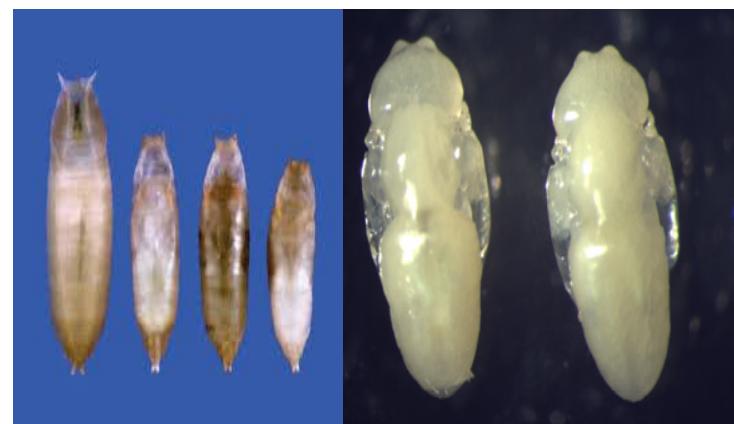
What is great about flies:

- **Short life cycle** - 10 days in a single vial
- **Cheap** - small, simple food source
- **'Small' genome** (100 megabases)
 - Around 14,000 genes
 - Only 3 major chromosomes
- Good vertebrate model
 - 75% human genes found in *Drosophila*

Dissecting fly wings to analyse wing hair patterns: knock-out and knock-in genes



Immuno-fluorescence imaging: imaging of a developing wing

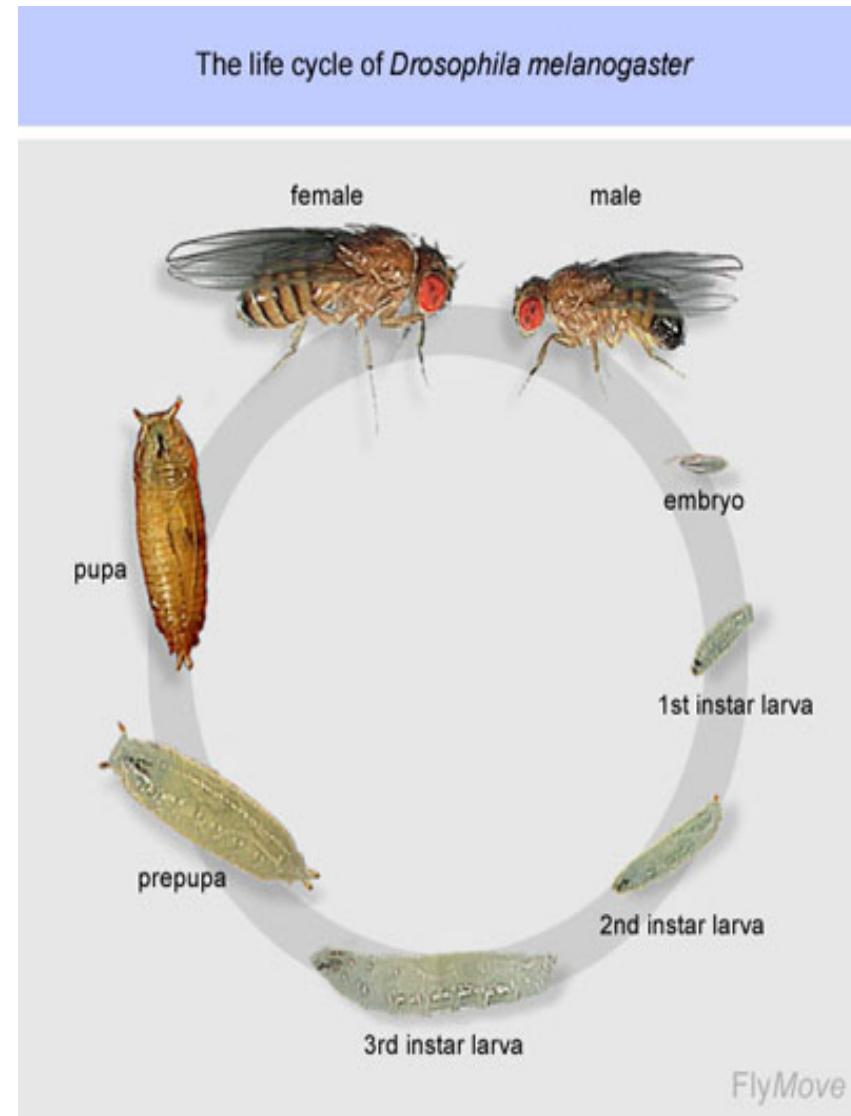


Disadvantages

- Cannot be frozen/stored - **require continuous culture**
- **No targeted mutagenesis** - e.g. homologous recombination
- **Able to fly!**

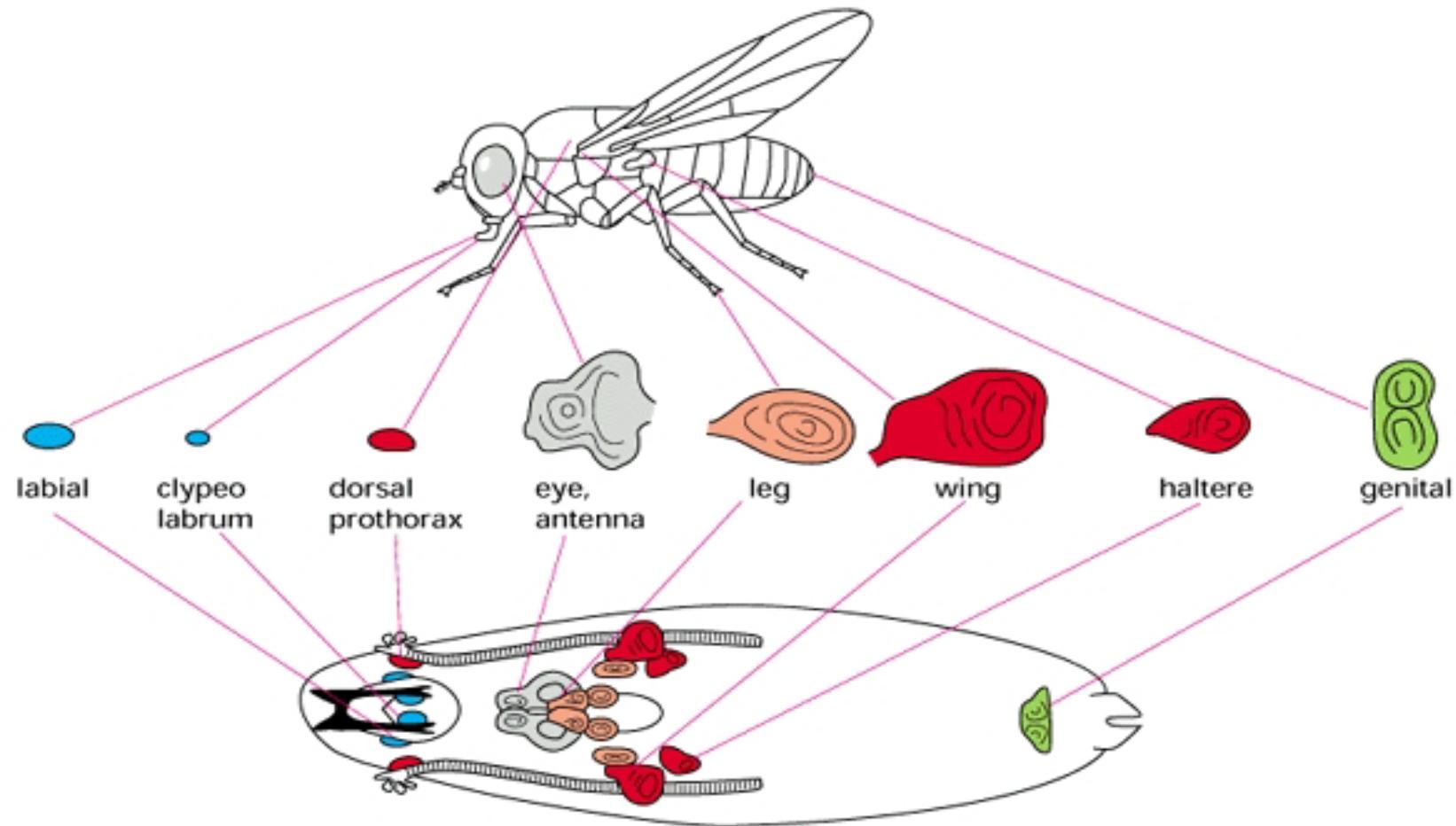
Drosophila melanogaster: life cycle

- **Embryo stage:** 24 hours
- **Larva stage:** 4 days
(Pupariation)
- **Pupa stage:** 4 days
- **Adult stage**
- Life cycle: ~9-10 days



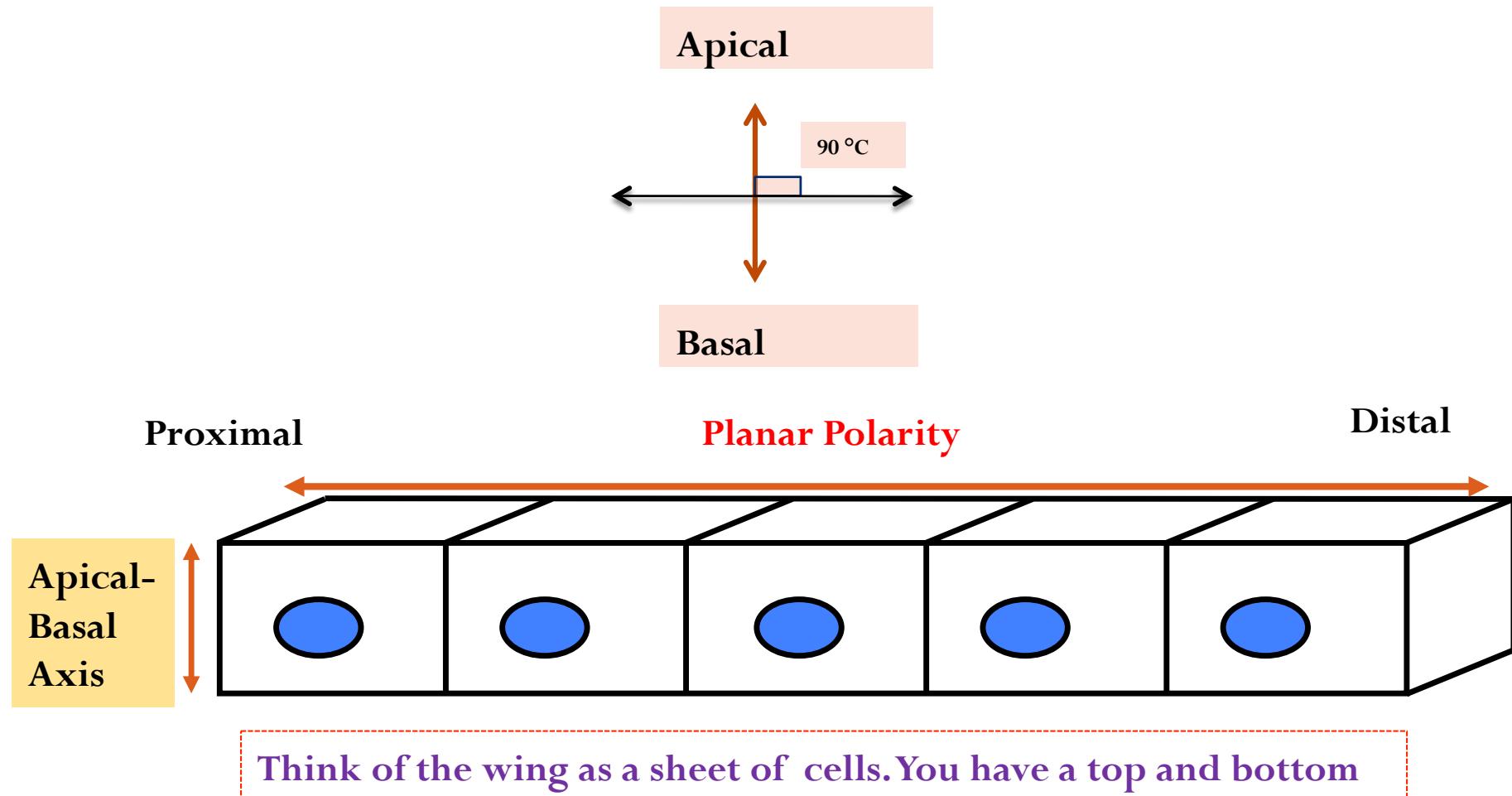
Drosophila Development

- Patterning: groups of cell aggregate and later become organs

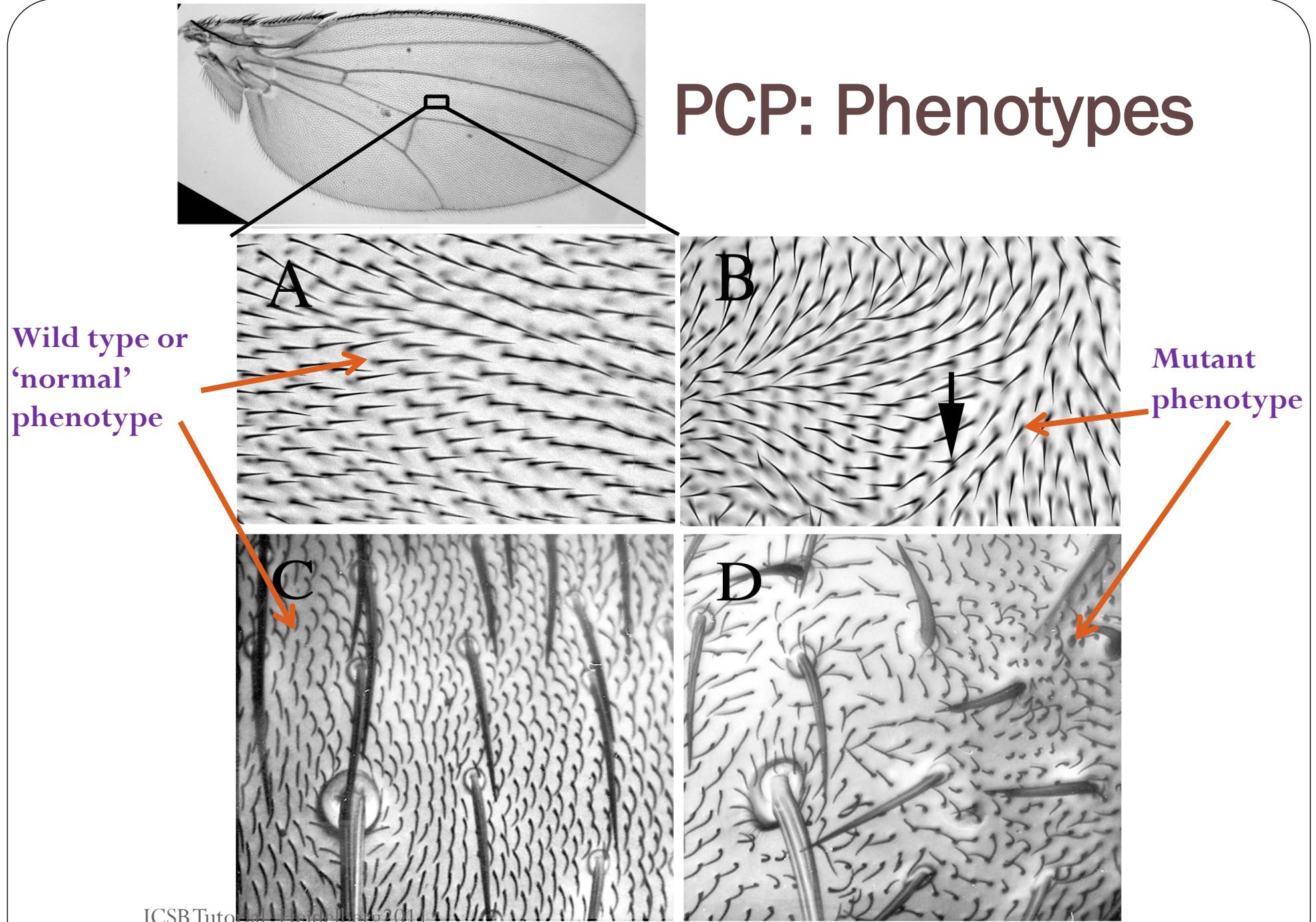


Planar Cell Polarity

- Mechanism through which a number of tissue types determine the polarity of their cells perpendicular to their cellular apical-basal axis.
 - in mammals: ear (sensory hair cell), eye (Equipotent R3/R4 cells) epithelia

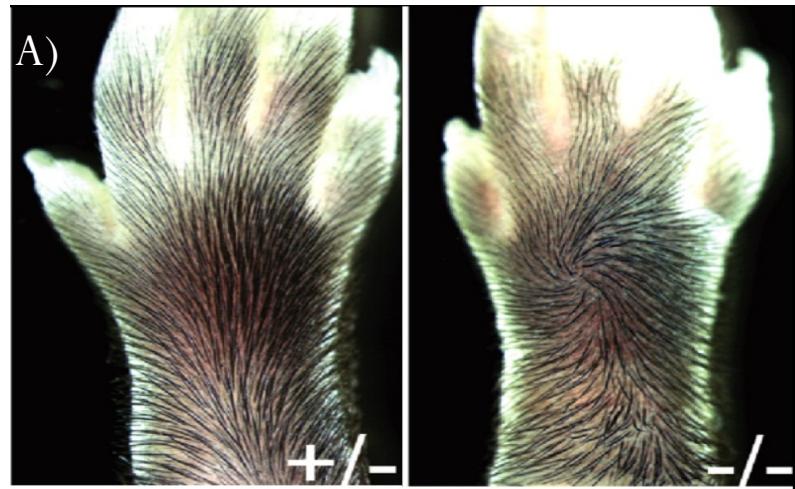


PCP: Phenotypes

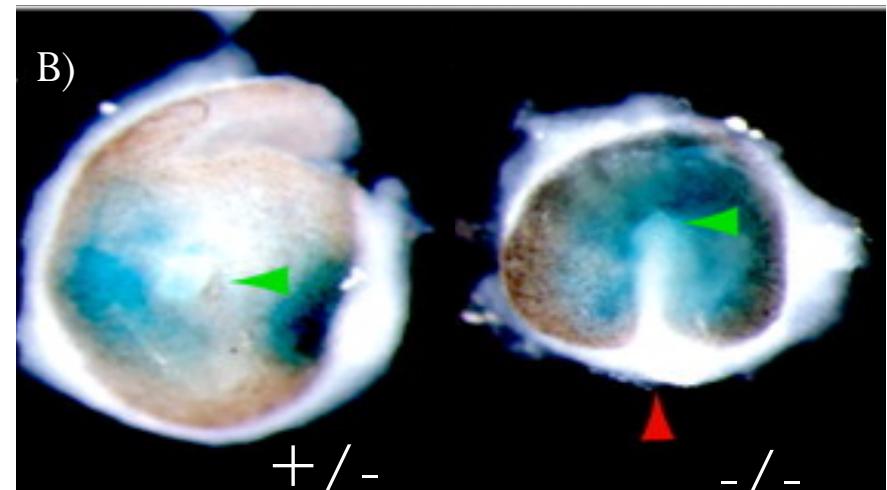


PCP: Vertebrate Phenotypes

Fz6 Knockout: Hair Polarity



Fz5 Knockout: Coloboma

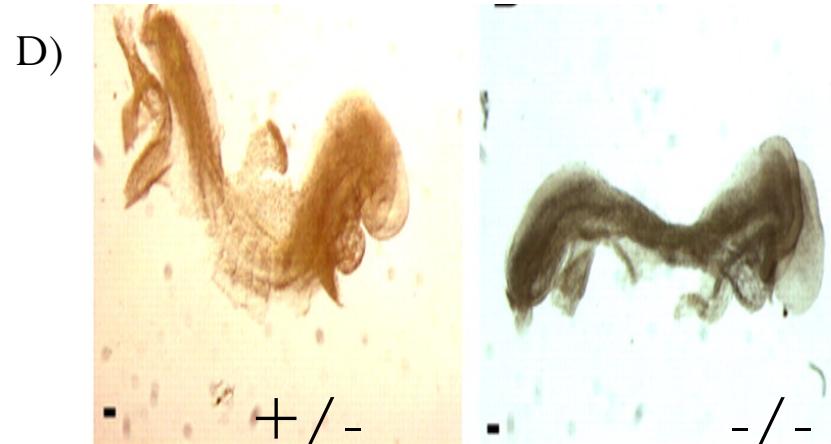


Dact1 Knockout: Urogenital Defects

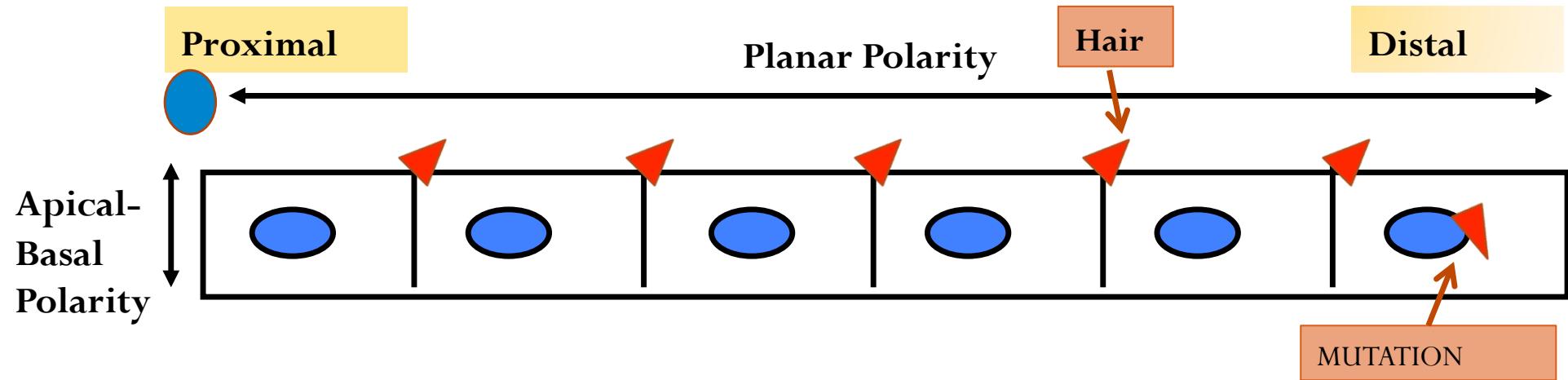


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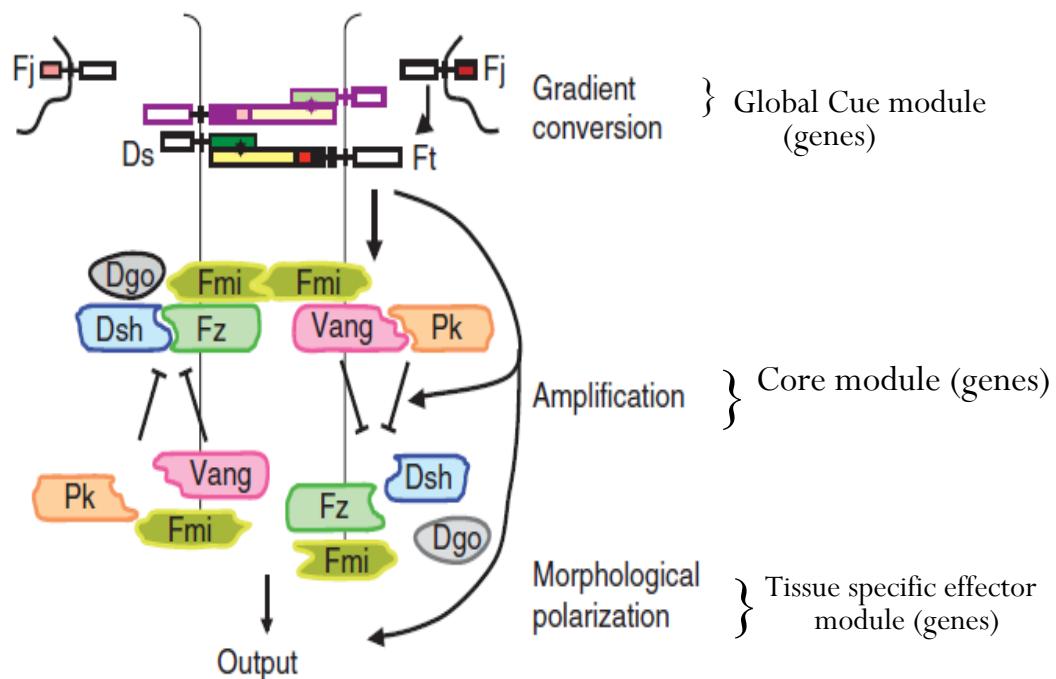
Vangl2 Knockout: Impairs Cranial Neurulation



Planar Cell Polarity: in the fly wing

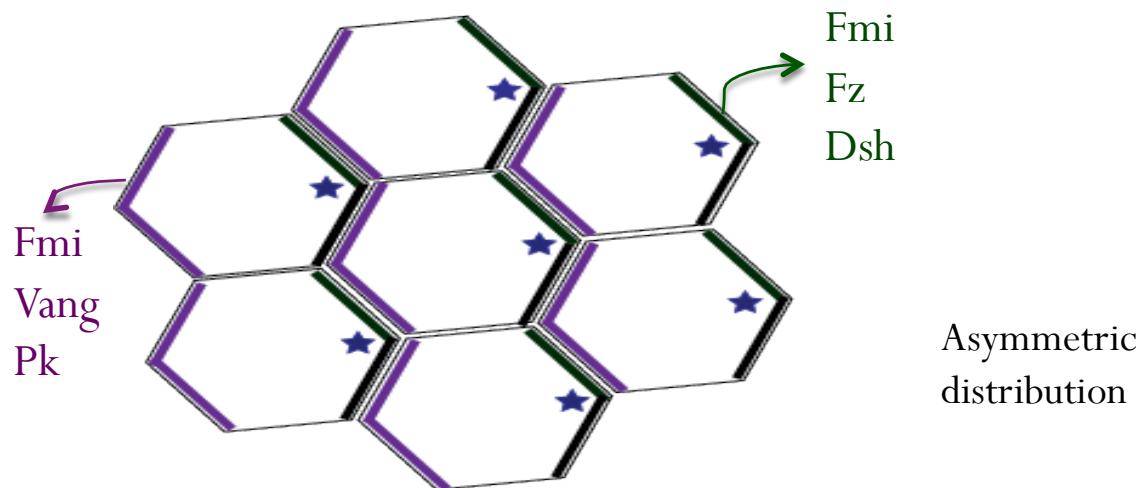
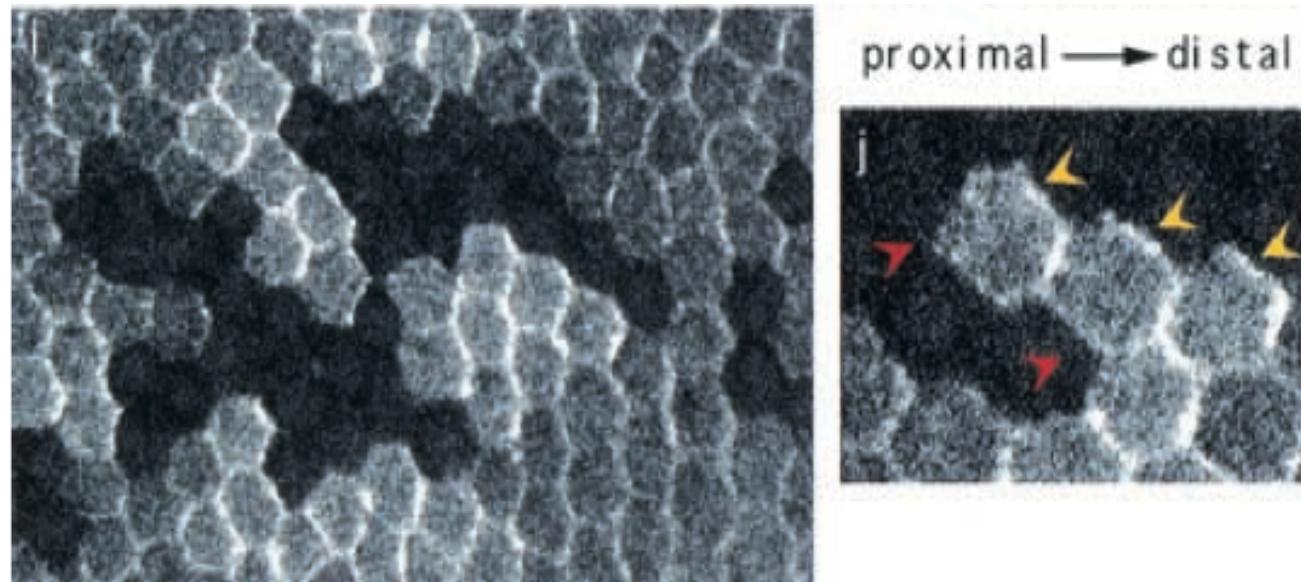


- Physical interactions/signalling relationships

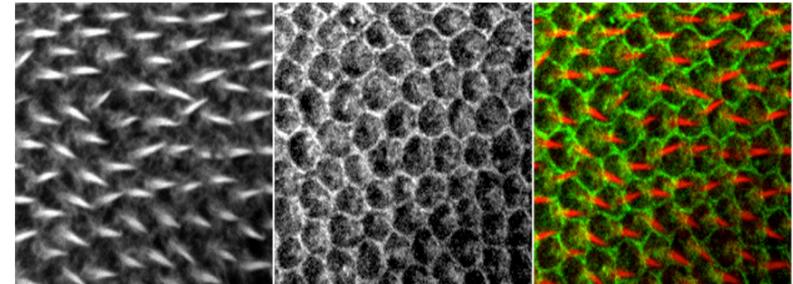
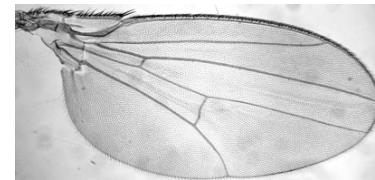
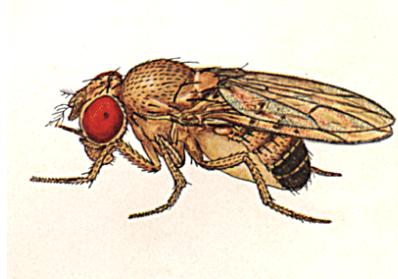


Core : Asymmetric Protein Localisation

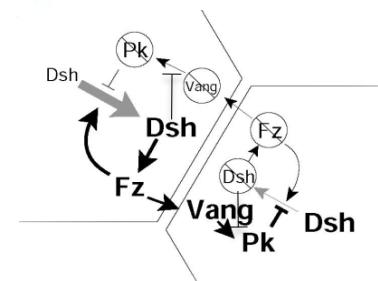
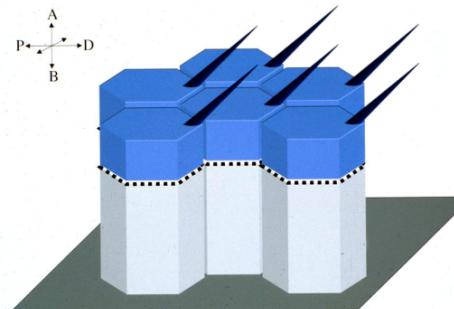
GFP- GREEN
FLUORESCENCE
PROTEIN



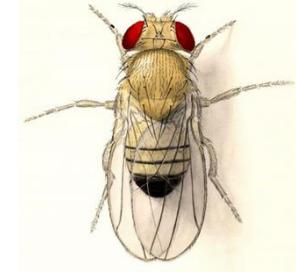
Multiscale from signalling to organs



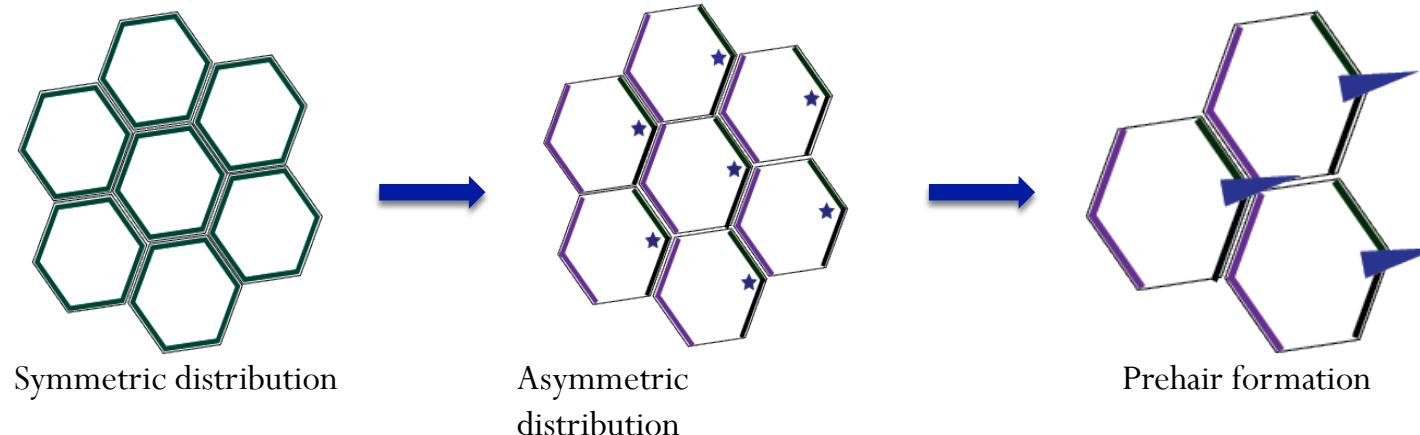
- Drosophila wing hairs point distally virtually error free.
- Hexagonally packed, planar (30,000)
- Planar cell polarity: the polarization of a field of cells within the plane of a cell sheet.
- Human pathology:
 - ✓ Cochlear hair cells
 - ✓ Spina bifida
 - ✓ Oncogenic Wnt pathway



Biological Model



- **PCP proteins** involved in the intercellular signalling: Frizzled (Fz), Dishevelled (Dsh), Prickle (Pk), Flamingo (Fmi) and Van-Gogh (Vang) .
- **A core machinery** mediates a competition between the proximal and distal proteins: Frizzled (Fz), Dishevelled (Dsh), Prickle (Pk) and Van-Gogh (Vang). Flamingo (Fmi) localises at both distal and proximal edges.
- **Feedback loops**: cells tend to align cell polarity as asymmetric distribution.

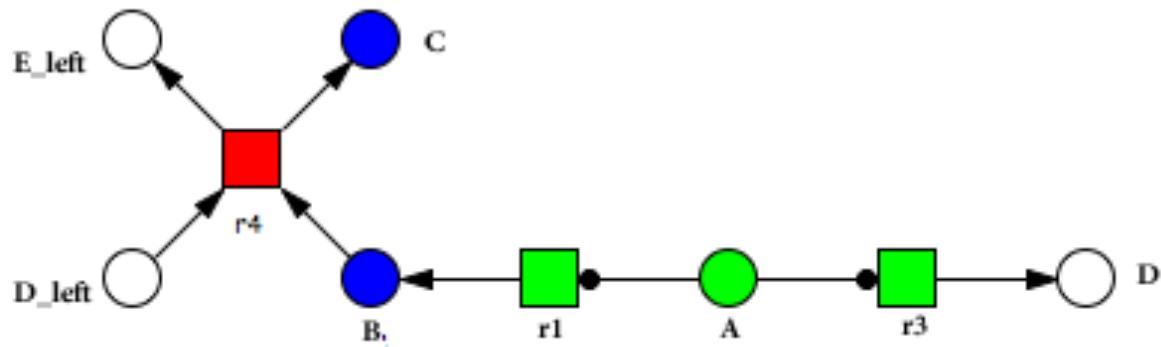


Coloured Petri Nets (CPN)

- Tokens distinguished via their colours.
 - Each place gets a colour set, specifying the kind of tokens which can reside on the place.
 - Each transition gets a guard, specifying which coloured tokens are required for firing.
 - Each arc gets an arc expression specifying the kind of tokens flowing through it
-
- Allows for the discrimination of species (molecules, metabolites, proteins, secondary substances, genes, etc.).
 - Colours can be used to distinguish between sub-populations of a species in different locations (cytosol, nucleus and so on).

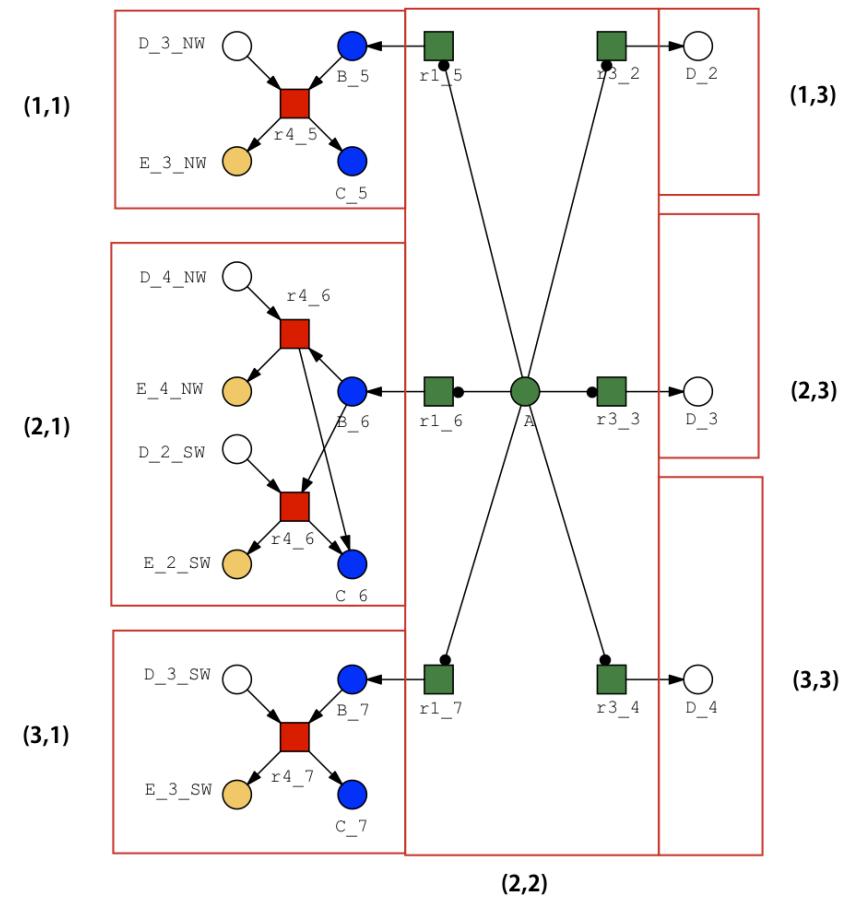
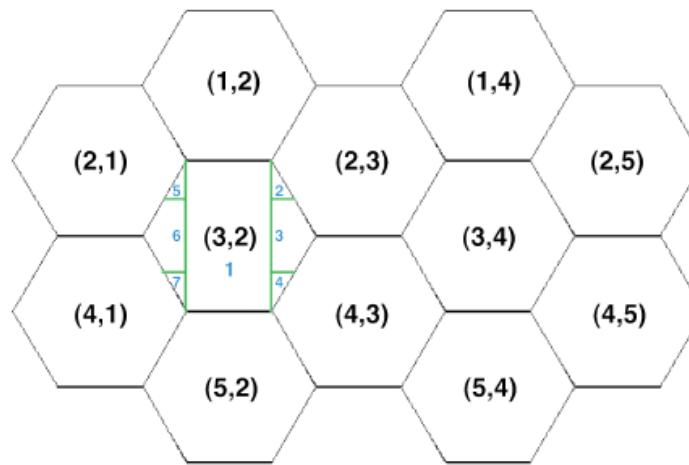
Single cell

Abstract level



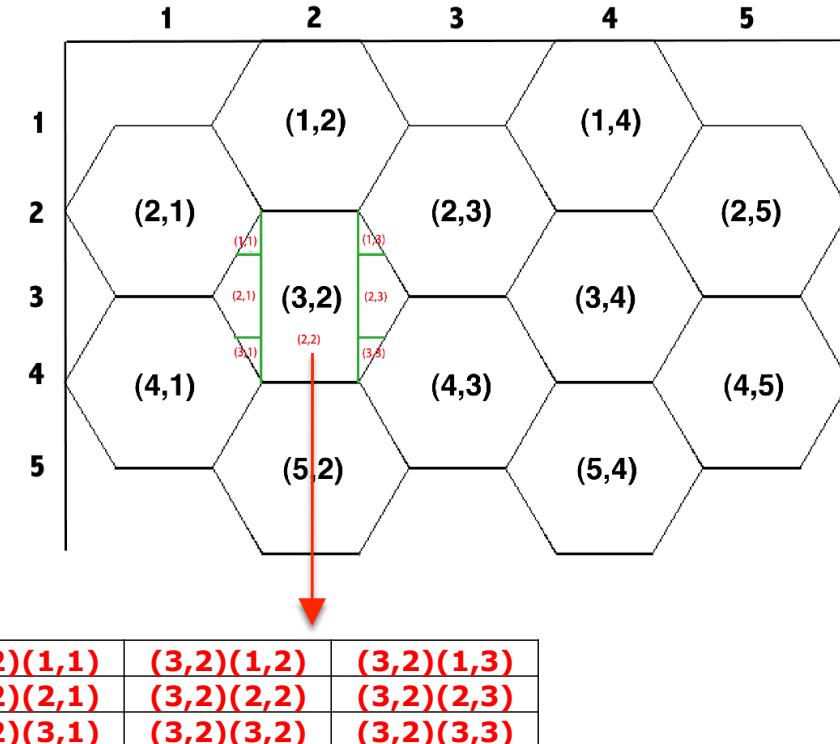
- Four spatial regions as labelled (*Labelled colours are **not** CPN colour sets*)
- D_{left} & E_{left} : two molecular species (places) from the left-hand side neighbouring cell(s)

Petri net model for a single cell



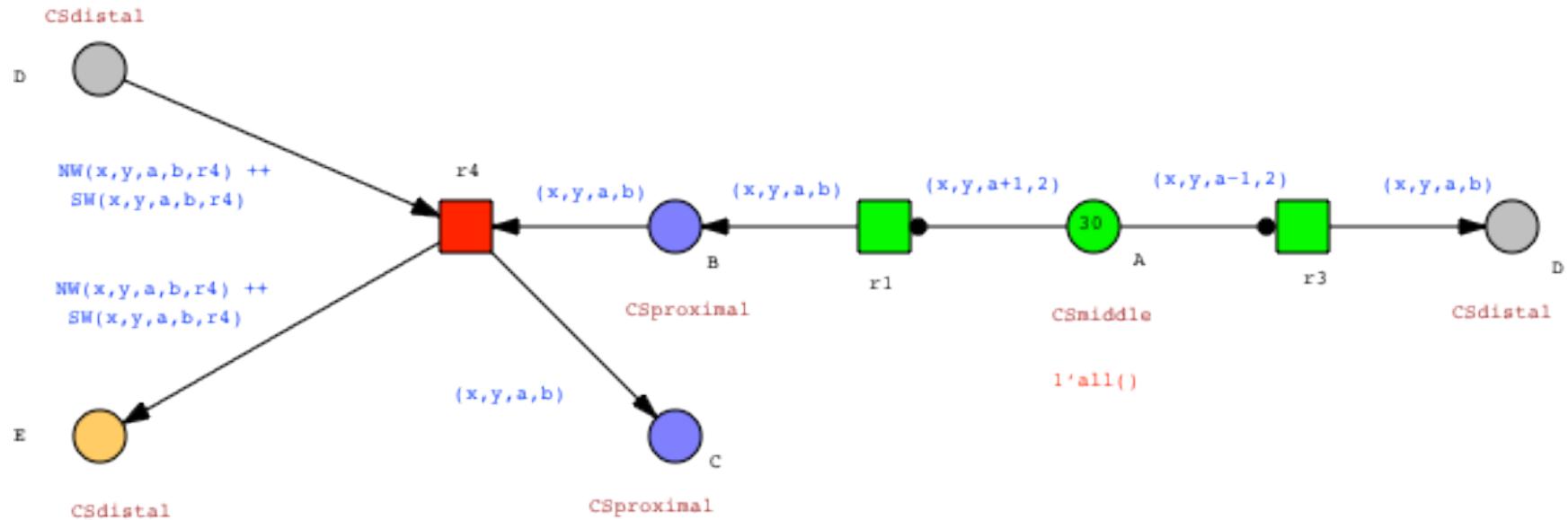
Hierarchical organisation

- Hierarchically coloured



Colourset = $\{\dots, \{((3,2)(1,1)), ((3,2)(1,2)), ((3,2)(1,3)), \dots, ((3,2)(3,3))\}, \dots\}$

CPN model of cells with 7 compartments in a 2-D lattice



- 4 spatial regions: communication, proximal, transport and distal.
- Seven virtual compartments $((1, 1), (2, 1), \dots, (3, 3))$.
- Each place or transition belongs to a specific compartment.
- NW and SW denote two left neighbours of the current cell.

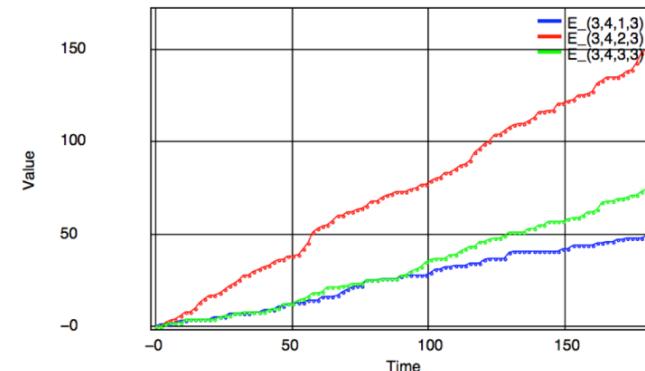
Declarations for the CPN model

Declaration

```
Constant M = int with 5 ;
Constant N = int with 5;
Constant C = int with 3;
Constant R = int with 3;
colourset Row = int with 1 - M;
colourset Column = int with 1 - N;
colourset ComR = int with 1 - R;
colourset ComC = int with 1 - C;
colourset CSr4 = enum with c5,c6_1,c6_1,c7;
colourset CS1 = product with Row × Column;
colourset CS2 = CS1 with x%2 = 1&y%2 = 0|x%2 = 0&y%2 = 1;
colourset CS = product with Row × Column × ComR × ComC;
colourset CS4 = CS3 with x%2 = 1&y%2 = 0|x%2 = 0&y%2 = 1;
colourset CSdistal = CS4 with b = 3;
colourset CSproximal = CS4 with b = 1;
colourset CSmiddle = CS4 with b = 2;
Variable x : Row;
Variable y : Column;
Variable a : ComR;
Variable b : ComC;
Variable r4 : CSr4;
Function CSproximal NW(Row x,Column y,ComR a,ComC b);
Function CSproximal SW(Row x,Column y,ComR a,ComC b);
```

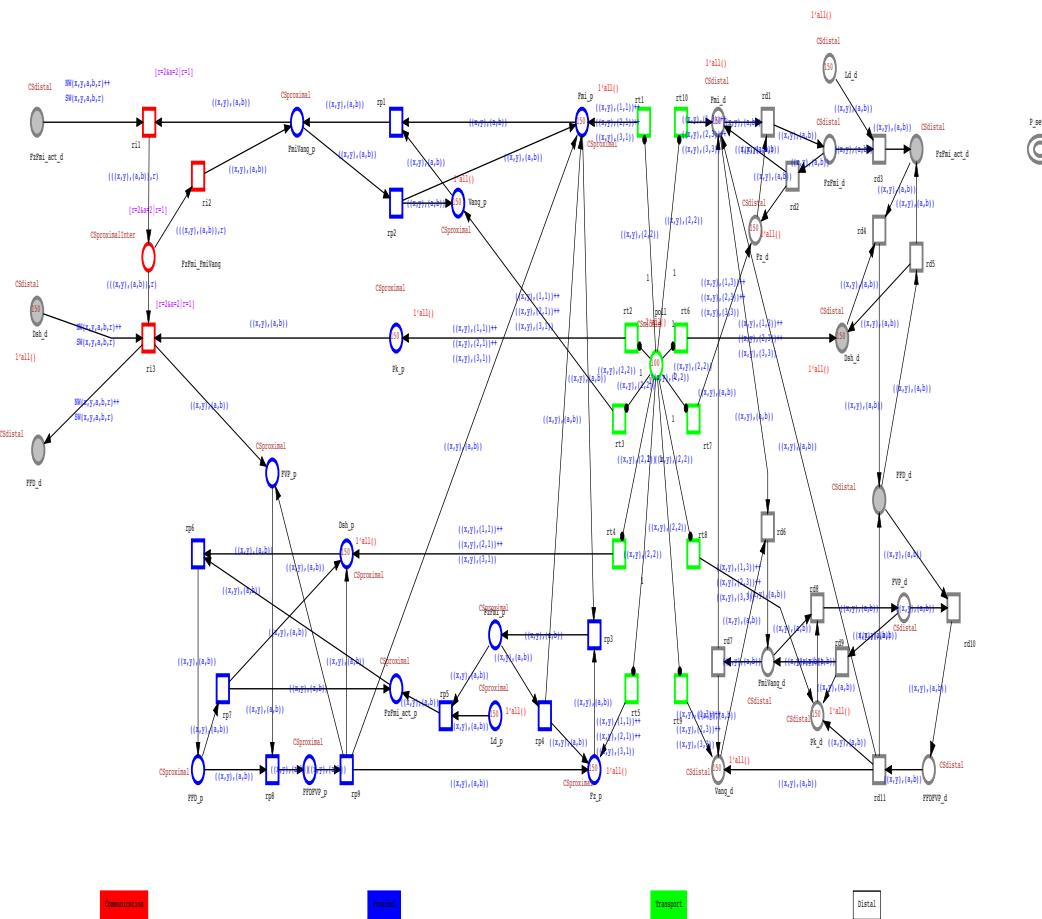
Analysis Results

Stochastic simulation results



		0		0		
	72	(1,2)	97	69	(1,4)	77
(2,1)	135		49	(2,3)	148	45
	46		58		45	50
	(3,2)	168		(3,4)	170	
	46		48		39	0
(4,1)	167		42	(4,3)	165	43
	76	(5,2)	88		68	(5,4)
			0			84
						0

CPN model of PCP signalling



Declarations for
the coloured
Petri net model

Type	Declaration
con	$M = \text{int with } 15;$
con	$N = \text{int with } 15;$
con	$R = \text{int with } 3;$
con	$C = \text{int with } 3;$
cs	$\text{Row} = \text{int with } 1 - M;$
cs	$\text{Column} = \text{int with } 1 - N;$
cs	$CS1 = \text{product with } \text{Row} \times \text{Column};$
cs	$CS_Cell = CS1 \text{ with}$
	$x \% 2 = 1 \& y \% 2 = 0 x \% 2 = 0 \& y \% 2 = 1;$
cs	$ComR = \text{int with } 1 - R;$
cs	$ComC = \text{int with } 1 - C;$
cs	$CS_ComP = \text{product with } ComR \times ComC;$
cs	$CS2 = \text{product with } CS_Cell \times CS_ComP;$
cs	$CSdistal = CS2 \text{ with } b = 3;$
cs	$CSproximal = CS2 \text{ with } b = 1;$
cs	$CSmiddle = CS2 \text{ with } a = 2 \& b = 2;$
cs	$CSInter = \text{int with } 1 - 2;$
cs	$CS3 = \text{product with } CSproximal \times CSInter;$
cs	$CSproximalInter = CS3 \text{ with } r = 2 \& a = 2 r = 1;$
var	$x : \text{Row};$
var	$y : \text{Column};$
var	$a : ComR;$
var	$b : ComC;$
var	$r : CSInter;$
fun	$CSproximal\ NW$ $(Row\ x, Column\ y, ComR\ a, ComC\ b, CSInter\ r)$ $\{[(!(x=1 y=1)) \& (r=1 \& a=1 \& b=1 r=2 \& a=2 \& b=1)]$ $((x-1,y-1),(a+1,b+2));\}$
fun	$CSproximal\ SW$ $(Row\ x, Column\ y, ComR\ a, ComC\ b, CSInter\ r)$ $\{[(!(x=M y=1)) \& (r=2 \& a=2 \& b=1 r=1 \& a=3 \& b=1)]$ $((x+1,y-1),(a-1,b+2));\}$
fun	$\text{bool}\ MutReg(Row\ x, Column\ y)$ $\{x>=4 \& x<=8 \& y>=3 \& y<=7;\}$

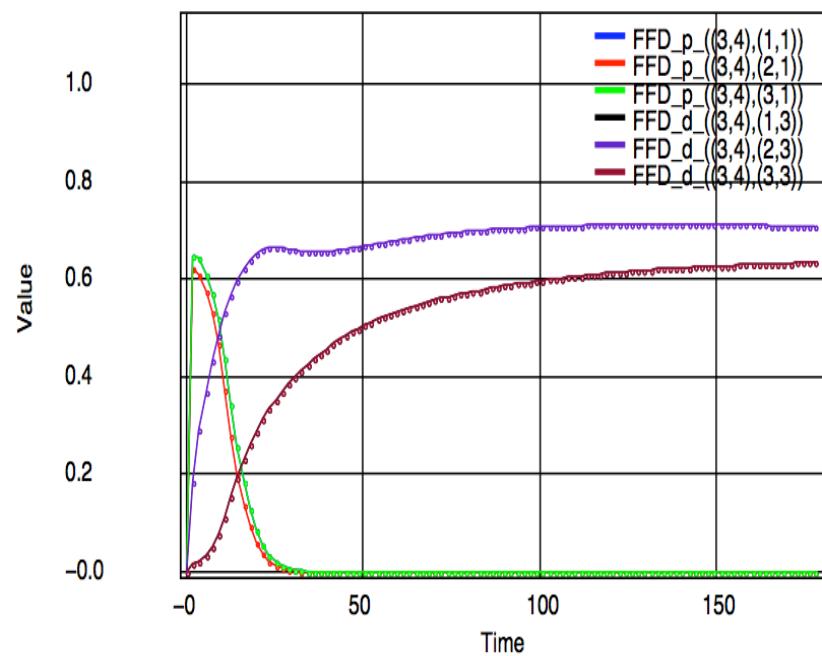
Some Statistics

Grid($M \times N$)	Size			Time (seconds)			
	Cells	Places	Transitions	Unfolding	Unfolding/Cells	Simulation	Simulation/Cells
5 × 5	12	924	984	0.665	0.0546	1.192	0.0993
10 × 10	50	3,850	4,100	2.528	0.0506	5.152	0.1030
15 × 15	112	8,624	9,184	6.108	0.0545	11.952	0.1067
20 × 20	200	15,400	16,400	11.705	0.0585	25.272	0.1264
50 × 50	1,250	96,250	102,500	139.594	0.1117	220.735	0.1766

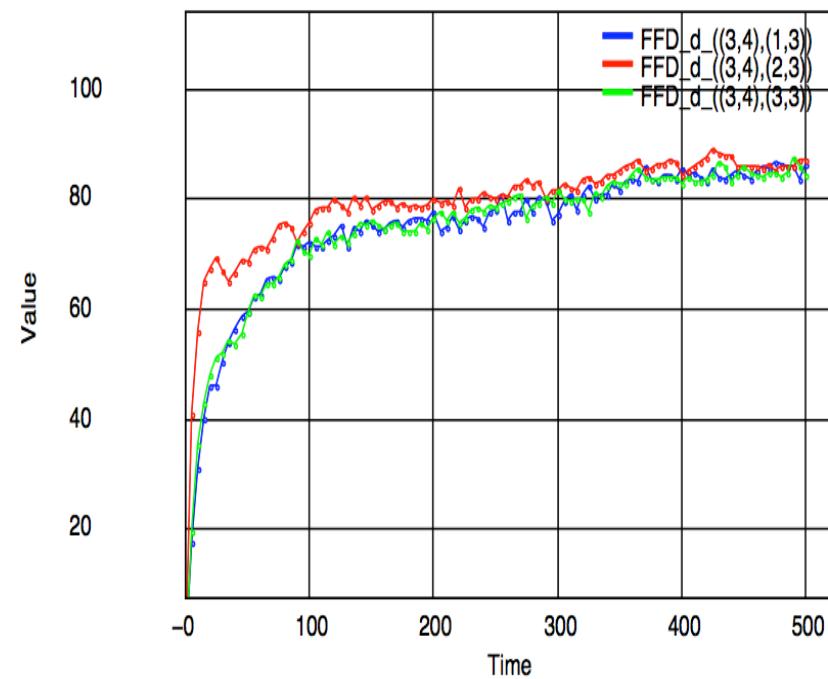
^{a)} done on PC Windows Vista (TM) business Intel(R) Xeon(R), CPN 2.83GHz, Memory(RAM) 4.00 GB;

Simulations

Continuous simulation

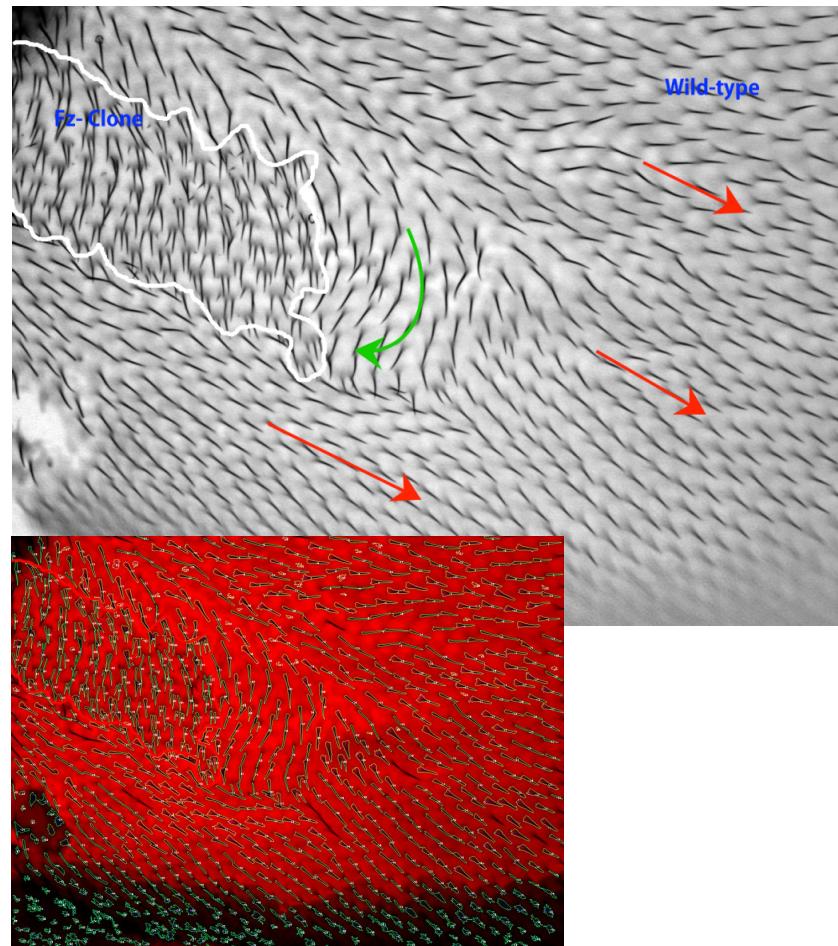


Stochastic simulation (average of 10 runs)

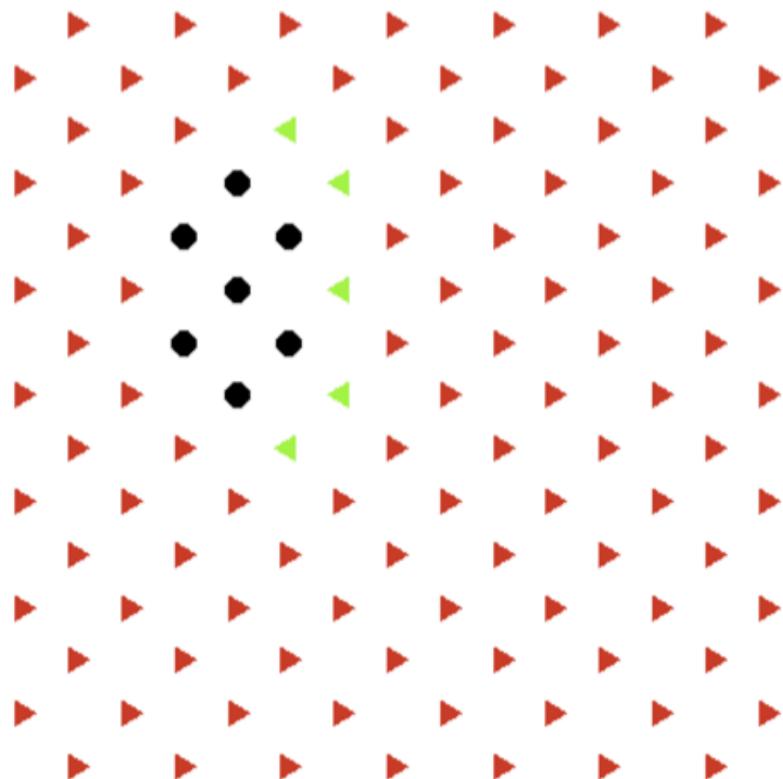


FFD accumulates at the distal edge of the cell rather than the proximal edge at the end of signalling.

Experimental vs In-silico



FFD at distal vs FFD at proximal over Tissue



ICSB Tutorial, Heidelberg2011

- Q. Gao, F. Liu, D. Gilbert, M. Heiner & D. Tree. A Multiscale Approach to Modelling Planar Cell Polarity in Drosophila Wing Using Hierarchically Coloured Petri Nets. **CMSB 2011**, Paris, France.

Downloads

- Snoopy for standard PN & CPN:

[http://www-dssz.informatik.tu-cottbus.de/DSSZ/Software/
Snoopy](http://www-dssz.informatik.tu-cottbus.de/DSSZ/Software/Snoopy)

- CPN models for PCP:

<http://people.brunel.ac.uk/~cspgqqg>

Acknowledgements

Brunel:

David Gilbert



David Tree



Cottbus:

Monika Heiner



Fei Liu



The End...

Thank you