# Documentation of the mechanistic, single-cell model

This supplementary material provides a summary of published evidence in support of the nodes and their relationships at the core of our model (Figure 3).

## Input nodes

These nodes embody external influences and have no associated logical function.

## Dpp

Decapentaplegic is a BMP ligand (a TGF- $\beta$  superfamily member), secreted from the stretched FCs that overlay the nurse cells, and from stage 10 onwards in the centripetally migrating follicle cells [1]. It signals to the follicular epithelium through receptors Wit [30] and Tkv.

At stage 10, Dpp/BMP signaling activity can be detected in a narrow band along the anterior border of the follicular epithelium, slightly larger at the dorsal side [2,3].

### Grk

Gurken is a TGF- $\alpha$  like protein secreted by the oocyte. It accumulates around the oocyte nucleus in the dorsal-anterior corner from stage 8 [4,5] and signals to the overlying follicular epithelium until the formation of the vitelline membrane at stage 10B [6].

### Br\_adj

This node denotes Br presence in neighboring cells (only those directly adjacent).

### Aos\_ext

This node denotes Aos presence in neighboring cells.

### Rho\_ext

This node denotes Rho presence in neighboring cells.

#### earlyBMP

*dpp* expression is first detectable at the end of stage 8 in approximately 20 to 30 somatic follicle cells at the anterior tip of the egg chamber [1]. However, there is evidence that the BMP pathway is active in early (<stage 6) egg chambers [7].

#### earlyEGF

Gurken is a TGF- $\alpha$  like protein secreted by the oocyte. In early stages, it is located in the posterior pole [4,5], where it activates EGF signaling in the overlying follicle cells.

## Core nodes

The behavior of these "internal nodes" depends on other nodes (described in the bullet lists) following rules given in Figure 3.

#### Aos

*argos* expression begins around stage 10A in the dorsal centripetal follicle cells, before extending to the dorsal midline. The patch splits around stage 11-12 into a pattern reminiscent of the late pattern of *rho* [6,8-10]. It is targeted by

- Pnt, which induces *aos* in the neuroectoderm [11].
  Overexpression of *pntP1* may reduce expression of EGF signaling targets *rhomboid* and *69D* (unpublished data mentioned by [12]).
  Pnt may induce *aos* in the FCE (unpublished data mentioned by [6]).
- dpERK: in addition to the *argos* expression pattern, we know that in intervein cells in the wing imaginal disc, *aos* transcription is repressed by Cic in a cell autonomous way; Cic itself is targeted for degradation through the EGF signaling pathway [3,13–15].
- Mid: when EGFR is ectopically activated in the epithelium, *aos* transcription is restricted to the anterior side [8], which suggests repression by Mid.

## Br

Broad-Complex (Br-C), is a transcription factor. It is a key marker of the roof region, expressed in two large patches of cells on each side of the dorsal midline [2,7,16–21]. *broad* mRNA is initially present at low levels everywhere, then repressed in an antero-dorsal T-shaped domain, before increasing in the roof domains and finally decreasing everywhere else.

br expression is controlled by two promoters [22]; only the BrL promoter is considered here.

- Mirr is required both to upregulate Br-C in the normal appendage primordium, and to downregulate it in the entire dorsal epithelium [20]. Fuchs et al. showed that Mirr activates the BrL promoter (late, Roof-specific Br pattern) and represses the BrE promoter (Early, ubiquitous Br pattern) [22].
- Pnt represses BrL (promoter controlling *broad* late expression) [22].

## dpERK

Activation of the EGF pathway may be visualized directly through MAPK or ERK phosphorylation (indicated with dpERK), starting from a midline-centered pattern, which expands to encompass a larger part of the dorsal-anterior domain, before regressing to a spectacle-like pattern and finally to the L-shape pattern characteristic of the floor domain [6,9,10,21]. It is targeted by

- Gkr activates EGFR [4,8].
- Argos inhibits EGFR signaling by sequestering Spitz [23]. Secreted Spitz (after being cleaved by Rhomboid) activates EGFR [24–26].
- X, corresponding to a hypothetical enhancement of EGFR activity via a juxtacrine signal from the roof.

## Mid

Midline expression marks the posterior domain of the follicular epithelium [27].

- earlyEGF: early, posterior Grk signalling activates Mid expression [27].
- earlyBMP: there is indirect evidence that BMP signaling affects the posterior boundary of the appendage primordia [8,18,28]. Considering the contrasting results of BMP pathway disruption on *br* expression [3,28–30], we postulate that this effect has to occur early, and propose it functions through repression of *mid*.

## Mirr

The transcription factor Mirror is activated in a large dorsal-anterior patch of the epithelium by low levels of EGFR activity [28,31,32].

- dpERK; *mirr* is induced in a large dorsal-anterior patch of the epithelium by low levels of EGF activity through Cic [14,28,31,32].
- Mid represses *mirr* and its target *br* [27].

## Pnt

The gene encoding the transcription factor Pointed is expressed along the dorsal midline at stage 10A, and in the roof region at stages 10B-11 [3,12].

• dpERK: *pnt* is induced by high levels of EGF signaling activity [2,11,12].

## Rho

*rho* expression pattern seems to correlate with EGF signaling activity as it evolves from the whole dorsal-anterior region in early stages to the characteristic one-cell wide floor region, adjacent to the roof, in late stages [4,8,33,34]. Rhomboid, a protease, cleaves Spitz, which is then secreted [25,26].

- Mirr: ectopic expression of *mirr* induces *rho* ectopic expression [31,35]. There is evidence that some anterior clue is required to induce *rhomboid* [8,18,29], and *mirror* could relay that signal to Rho.
- Br, which is required cell-autonomously to repress *rho* expression. Broad binding sites are predicted in the promoter region of *rho* [36].
- dpERK: *rho* transcription is repressed by the transcription factor CF2 [34]; CF2 is downregulated by the EGFR pathway [37].
- Dpp signaling represses *br* in the anterior follicle cells, via pMAD; strong *dpp* overexpression represses Br entirely [2,3].

## Integration nodes

#### A

Combines the values of Aos and Aos ext.

Aos and Aos\_ext: Argos is secreted [38]. The *argos* gene encodes a diffusible factor that regulates cell fate decisions in the Drosophila eye. Rule for the epithelial model: level 1: at least 5 cells among the proper cell and 126 neighbors (direct and or at a distance up to 6 cells) express Aos.

## S

Combines the values of Rho and Rho\_ext, to account for Spitz signaling.

• Rho and Rho\_ext: Rho activates Spitz, which is then secreted [25,26]. Rules for the epithelial model:

level 2: autocrine (Rho at level 2) or at least 6 cells over six (direct) neighbors level 1: autocrine (Rho at level 1) or between 2 and 5 cells among the six (direct) neighbors.

## X

Hypothetical mechanism. EGFR activity seems to be enhanced in cells in contact with the roof region ("spectacle-shaped" pattern in [9,10,21]). Moreover, Rho is expressed along the high-Notch side of a high/low Notch border; ectopic *rho* expression may be induced by ectopic high/low Notch barriers generated by Notch-clones [36].

• Br\_adj and Br. Br is taken as a marker of the roof. As such, it activates X in the neighboring cells, while inhibiting it in the roof itself.

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