Planar Cell Polarity: A Bridge Too Far?

The mechanisms of planar cell polarity are being revealed by genetic analysis. Recent studies have provided new insights into interactions between three proteins involved in planar cell polarity: Flamingo, Frizzled and Van Gogh.

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We now understand much of how cells know where they are in an embryo, but little of how they know their orientation, anterior from posterior, distal from proximal. Yet we believe that many, perhaps all, epithelial cells are polarised in the plane of the sheet—that they exhibit planar cell polarity, and that this polarity is vital. Planar cell polarity is not used primarily to make structures but more to orient them, making its study conceptually difficult.

But, genetics is the right approach and Drosophila has proved the model of choice—particularly as the genes identified in the fly are conserved in other animals, including vertebrates [1–3]. In the 60s it was argued that pervasive gradients are set up in the main axes of the body; it was suggested that the slope of a gradient could specify the polarity of cells [4,5]. This viewpoint is still very much alive and these gradients are now being identified with the help of genetics.

There is now a resurgence of interest in the mechanisms of planar cell polarity: three new papers [6–8] (one in this issue of Current Biology [7]) report the use of both genetics and molecular techniques to get to one of the two hearts of the matter.

Drosophila cells make oriented structures; examples are hairs and bristles on the wing and abdomen. In the 80s, pioneers such as Adler and Gubb found genes whose mutants altered these polarities [9,10]. Early on frizzled (fz) was identified; and, significantly, it was found that clones of fz– cells repolarised neighbouring wild-type cells so that they point their hairs towards cells with lower Fz activity [9,10]. It helps to think of the fz– cells as sending and the wild-type cells as receiving polarising information [11].

Many different genetic mosaics can be made in Drosophila and, for example, each gene can be tested to see if it is needed in the sending, in the receiving cells or in both. This repolarisation assay has proved an incisive aid in the analysis of planar cell polarity.

The first working models used a small group of genes: prickle (pk), fz, Van Gogh (Vang) (also called strabismus, stbm) and dishevelled (dsh). In the 90s it was found that, just before polarised structures are formed, some of these proteins become localised to one or other ends of the cell [12]. It was suggested that some small initial bias (unknown) is amplified by interactions and feedback between these four proteins to polarise each cell; propagation from cell to cell would be driven by interactions across the cellular space [13]. This model was simulated in a powerful computer [14] and became popular; however, complex computers are no match for simple experiments and the model looked feeble when it was found (in repolarisation assays) that pk and dsh are dispensable in both sending and receiving cells and so, for this central process, could be ignored [11,15–17].

The model suffered further blows when we found that a cell completely lacking fz could be repolarised [11] and that protein localisation itself appeared to be dispensable for repolarisation [11,16].

Flamingo (fmi, also known as starry night or stan), was largely left out of these models. In our assays, however, it was the only gene needed in both sending and receiving cells, and because its protein product is able to form homodimers from one cell to the next [18], we placed it at the centre of a new model [11]. In our model, the Fmi homodimers act as intercellular bridges. We suggested that, using Fmi to compare its neighbours, each cell points its hair towards the neighbour with the lowest level of Fz activity, and that there is an intercellular feedback via Fmi, which brings the level of Fz activity in one cell towards an average of its neighbours. We argued that Fmi–Fmi homodimers act asymmetrically to convey the level of Fz activity in the sending cell to Vang in the receiving cell. Because information is actually going in both directions—in the wild-type, each cell will both send and receive—it follows Fmi can act in two ways in the same cell depending on whether it sends (with Fz) or receives (with Vang) (Figure 1). A more detailed version of this model was elaborated subsequently [2] and another similar one simulated in silico [19].

Chen et al. [6] recently reported the results of similar experiments to ours [11] but, instead of monitoring hairs, they mostly looked at localisation of the proteins, a concordant indicator of polarity. They reached the same

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conduits for polarising information, or are more passive, for example helping Fz and Vang to contact each other as ligand and receptor.

Chen et al. [6] believe direct contact is implausible because of the wide gulf between the cells. If they are right, Fz might be able to mediate planar cell polarity without the cysteine-rich domain (CRD), part of the protein’s amino-terminal ectodomain, and Chen et al. [6] offer some evidence for this. But their evidence conflicts with previous findings that the Fz ectodomain is essential for planar cell polarity [20]. Also, in contrast to Chen et al. [6], and with stronger evidence, Wu and Mlodzik [8] find once again that the Fz CRD is essential. Extending the disagreement between the two papers, Wu and Mlodzik [8] not only find the interaction of Fz and Vang quite plausible, they actually find direct binding using pulldowns and binding assays in tissue culture cells — but they do not determine whether the binding is in trans (from cell to cell, as they assume) or in cis (which would fit with our evidence that Vang acts in cis to regulate Fz [11]). The data reported by Strutt and Strutt [7] agree with Wu and Mlodzik [8], providing independent evidence from tissue culture experiments that Vang and Fz can bind in trans, a binding that is enhanced in the presence of Fmi.

In spite of the possible significance of direct intercellular binding of Vang and Fz, we know that Fmi is essential for propagation of planar cell polarity information — so how do the three proteins relate? Chen et al. [6] describe evidence from immunoprecipitation that Fz binds to Fmi, and with this Strutt and Strutt [7] agree. But none of the groups shows biochemical evidence of binding between Vang and Fmi. Also, using overexpressed Fmi in vivo, Strutt and Strutt [7] report that Fmi needs both Vang and Fz if it is to stabilise properly in the cell membrane and that it prefers to bind to Fz rather than to Vang. These mixed observations tie Vang, Fz and Fmi closely together but they do not tell us clearly how the information about Fz activity is exchanged, nor what are the structural or functional relationships between the three proteins.

Chen et al. [6] ask whether Fmi acts actively (as we previously proposed [11]) or passively (as now proposed by Wu and Mlodzik [8]). They claim to answer by showing that Vang\(^{-2}\) Fz\(^{-2}\) sending cells do repolarise receiving cells when they strongly overexpress Fmi and argue that this means that Fmi can act instructively. However, this argument is undermined by the conflicting and positive finding of Wu and Mlodzik [8] that Vang\(^{-2}\) Fz\(^{-2}\) sending cells repolarise receiving cells, even without overexpressing Fmi. The contradictory results and conclusions described in the three papers [6–8] show that we are not yet ready to answer this enigmatic question.

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**Figure 1.** Possible intercellular and intracellular relations between Fmi (blue), Fz (red) and Vang (green), according to different authors. Two membranes from neighbouring cells are shown. If a direct physical interaction is suggested, the molecules are shown in contact. In (A), the size of lettering refers to the original gradient of Fz activity, with the strength of color of the molecules indicating the subsequent localisation of the proteins. The arrows show the direction of the flow of information: black is strong, grey is weak signaling. (A) Lawrence et al. [11]; (B) Chen et al. [6]; (C) Strutt and Strutt [7]; (D) Wu and Mlodzik [8].
Animal Personalities: The Advantage of Diversity

Animals do not always perform to the best of their abilities when faced with difficult choices. New findings on foraging honeybees show that co-existing strategies, where some individuals place more emphasis on accuracy and others on speed, can be advantageous to the colony in a variable environment.

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For decades, researchers in animal behaviour have been largely concerned with the accuracy, and not the speed, of decision making, and have measured choice percentages when animals faced multiple options in terms of foraging, mates or predation risk. In human psychophysics, however, it has long been known that decision accuracy and speed are interrelated, and accuracy can only be understood in a meaningful way if decision time is also quantified [1,2]. This is because, in noisy or uncertain conditions, accurate decisions require a higher sampling time [3]. In research on nonhuman animals, this interaction has received more attention since two studies in 2003 examined the possibility of speed-accuracy trade-offs in olfactory discrimination by rats [4] and colour discrimination by bees [5]. Far from simply copying concepts from human psychophysics, behavioural ecologists have since explored several new dimensions of such tradeoffs, including their ecological and evolutionary relevance — for example, in house-hunting ants [6], spatial exploration by passerine birds [7], predators choosing between aposematic prey [8], and predator avoidance behaviour by pollinators [9].

Burns [10] recognized the potential implications for research on animal ‘personality’. Rather than the typical scenario where speed-accuracy trade-offs are evaluated within subjects, there might also be consistent between-individual differences in terms of whether an animal places greater emphasis on speed or precision [2,5]. Some individuals might consistently be meticulous and slow, while others choose a ‘fast-and-sloppy approach’ — and perhaps such ‘impulsive’ individuals might not be selected against, because despite their high number of errors, their strategy can be advantageous, if the temporal costs of accurate decisions exceed those of errors [10].

One of the major challenges in research on the individuality of animals, from insects to humans, is understanding its adaptive significance [11,12]. How can multiple ‘personalities’ persist, side-by-side, in the same environment, when one might expect that one particular configuration of traits might outperform all others, and should therefore be favoured by selection? One possibility is that variation is selectively neutral [11], but in many cases, spatial and temporal heterogeneity in the environment might play important roles in maintaining diversity [12]. As they report in this issue of Current Biology, Burns and Dyer [13] set out to identify the kinds of environmental conditions that might favour ‘fast-and-sloppy’ individuals among honeybees, as well as the conditions that might give ‘slow-and-precise’ individuals an edge.

Bees typically obtain their entire diet — nectar and pollen — from flowers. In doing so, they operate in a ‘pollination market’, where they must choose adaptively between multiple flower species that differ in reward profitability, handling costs, densities and predation threat — and memorise these features by associating them with flower signals such as colours [9,11]. The complexity of this interaction makes the collection of meaningful data in field conditions difficult, and so Burns and Dyer [13] used artificial flowers with precisely controlled rewards and colours. The authors assessed the ‘personality’ of 12 freely-flying worker bees by evaluating choice precision and times spent in flight between flower visits, and the consistency of these parameters over time and experimental conditions. They then quantified their individual nectar collection rates by testing each bee in two conditions (Figure 1). In both conditions, the bees foraged on a patch containing two ‘flower species’ with two similar colours, with one flower type containing nectar and the other containing water. In condition 1, there were as many rewarding as unrewarding flowers. Therefore, the cost of inaccuracy was relatively low, as bees had a one-in-two chance to find nectar by random choice. In condition 2, however, the rewarding flowers were outnumbered by the similarly coloured unrewarding flowers by a factor of two, so that bees had a 33% chance of finding nectar.

Burns and Dyer [13] found that individuals fell along a continuum from slow-accurate to fast-inaccurate strategies. Moreover, they discovered that, when there were equal numbers of both flower types, fast-inaccurate bees collected slightly more nectar than slow-accurate bees. Conversely, when the accumulating cost of mistakes was higher, slow-accurate bees clearly out-competed the fast-inaccurate bees (Figure 1). Therefore, these findings support a differential advantage for