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# The origin of the plant body axis

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During embryogenesis, the basic body plan of an organism develops from a unicellular zygote. In most flowering plants, the polar zygote divides asymmetrically, making visible the apical–basal axis in the early embryo. The molecular mechanisms governing how the zygote polarizes and how this polarity is linked to embryo axis formation have been obscure, mainly owing to the difficulties to access the zygote that is deeply embedded in the maternal tissue. In this review, we summarize recent findings identifying key regulators in *Arabidopsis* and developing novel approaches in various plant species, which altogether set the stage for unraveling embryo axis formation.

### Addresses

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## Introduction

Initiation of the main body axis is one of the first patterning steps in the development of an organism from a single-celled zygote. In animal and plant model organisms, the zygote undergoes an asymmetric division, converting its unicellular polarity into the main body axis of the early embryo [1,2]. Significant insight into the mechanisms controlling this event in animals has been obtained from studies in *Caenorhabditis elegans*, *Xenopus laevis*, *Drosophila melanogaster*, and mammals [3]. In most flowering plants, a first apical–basal axis is morphologically evident in the egg cell, where the nucleus is localized in the apical half and a large vacuole at the basal end [4,5]. This axis aligns with the micropylar–chalazal axis of the surrounding maternal tissue, raising the question whether maternal factors provide cues for egg cell polarity. A long held view was that the polar localization of the nucleus and vacuoles in the egg cell is contiguous with that in the zygote. However, this appears not to be the case. Upon fertilization, the zygote rapidly undergoes a number of changes, called ‘egg cell activation’, including those affecting the numbers and shapes of

mitochondria, ribosomes, and Golgi vesicles [6,7]. Furthermore, recent studies in *Arabidopsis* and *Nicotiana tabacum* (tobacco) report a symmetrical organization of the zygote after fertilization, with the nucleus in the center and the vacuoles spread throughout the cell (Figure 1) [8,9,10<sup>\*</sup>,11<sup>\*</sup>]. During subsequent zygote elongation, the nucleus relocates to the apical part, and a large vacuole again forms at the basal end of the zygote (Figure 1). In parallel, the microtubule orientation gradually changes from a random orientation in the young *Arabidopsis* zygote into a transverse pattern of cortical microtubules [12]. The regulatory mechanisms that drive these changes in the zygote are unknown. Despite the transient symmetric morphology of the zygote, it is nevertheless possible that polar marks are laid down in the egg cell to provide spatial cues for the re-polarization of the zygote. Investigation of species displaying an opposite polarity in the egg cell (i.e. the nucleus at the basal pole and the vacuole in the apical region), such as *Oryza sativa* (rice) and *Papaver nudicaule* (Iceland poppy), can provide insight into whether transient symmetry and a subsequent symmetry breaking step in the zygote are common in plants [13,14].

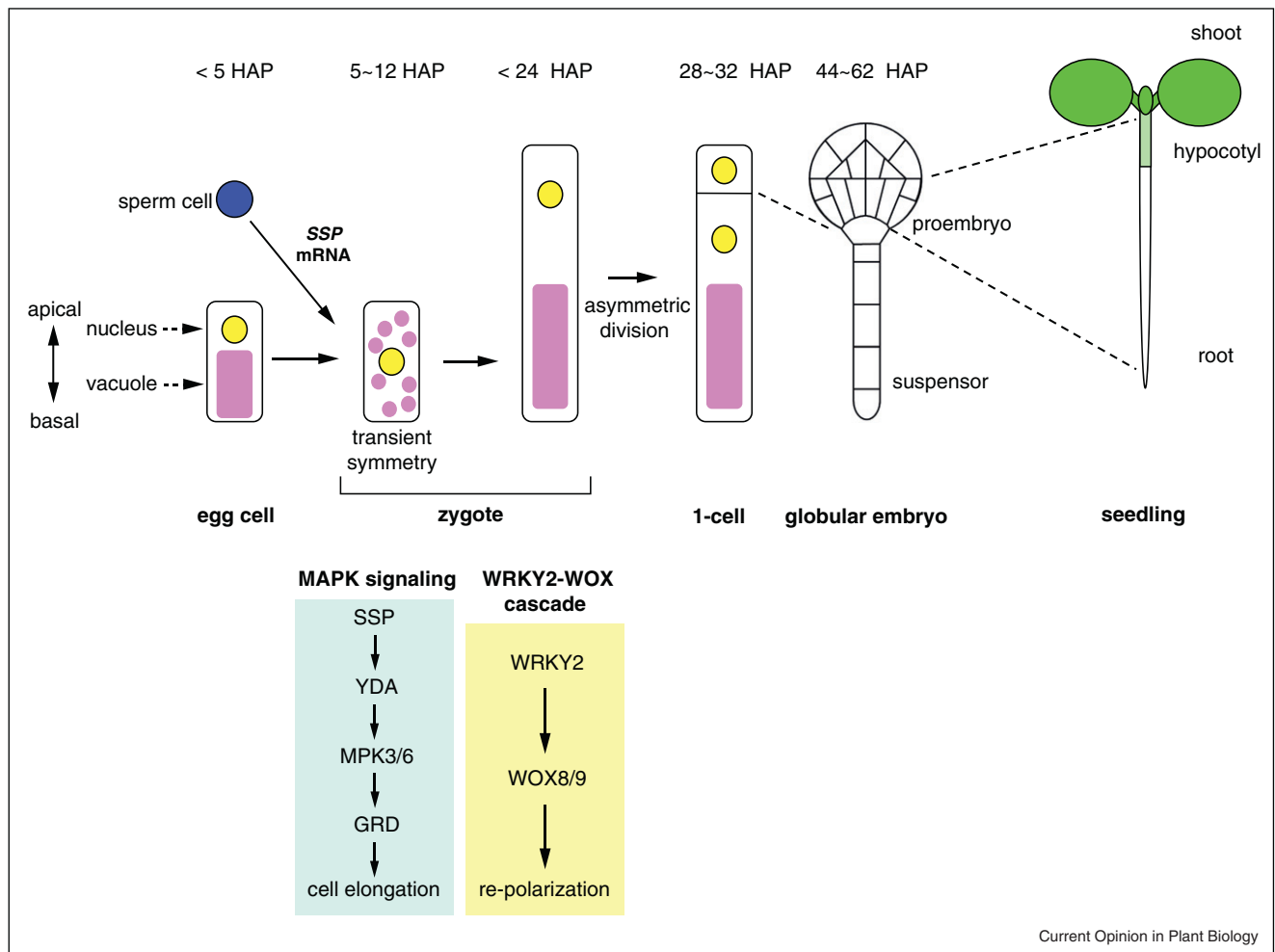
In most plant species, including *Arabidopsis*, the zygote elongates after fertilization and divides asymmetrically into a small cytoplasmic apical daughter cell and a large vacuolated basal daughter cell (Figure 1) [6]. The apical daughter cell gives rise to a spherical proembryo and eventually to most of the plant body (Figure 1) [5]. The initial steps occur by a precise pattern of cell divisions in *Arabidopsis*, whereas divisions are more random in larger embryos such as in maize. The vacuolated basal daughter cell generates the suspensor and, from its uppermost descendant, the hypophyseal cell, a part of the root meristem. The suspensor shapes may vary between different plant species, ranging from a filamentous suspensor in *Arabidopsis* (Figure 1) to the more elaborated branched haustoria as in *Sedum* [15]. It is noteworthy, that in some species, such as *Pelargonium zonale*, the zygotic division generates two cells of similar size [16]. Nevertheless, these species display conspicuous differences between apical and basal cells; for example, small, and circular mitochondria are abundant in the apical cell, whereas the basal cell mainly contains long rod-shaped mitochondria [16].

In summary, the intracellular polarity of the zygote correlates with the apical–basal axis of the embryo.

## The *Arabidopsis* sperm triggers zygote elongation and suspensor fate

In plants mutant for the MAPKK kinase YODA (YDA) or its downstream MAP kinases MPK3/MPK6, the zygote

Figure 1



Model for the role of YDA MAPK signaling and the WRKY2-WOX genetic network in *Arabidopsis* zygote polarization. Upon fertilization, the subcellular positions of the nucleus (yellow) and the vacuole (pink) are apolar, and become asymmetrically localized during zygote elongation. The sperm cell provides SSP mRNA that, after translation in the zygote, activates YDA MAP kinase signaling. This indirectly promotes cell elongation of the zygote through activation of the GRD transcription factor. By contrast, zygote re-polarization requires WRKY2, partly via *de novo* transcription of WOX8/9 in the zygote. The asymmetric division of the zygote produces the small apical cell, which generates the proembryo, and the large basal cell, which gives rise to a filamentous suspensor. Subsequently, the proembryo constitutes the entirety of the plant, except for part of the root tip, which is specified by one cell of the suspensor. An approximate time scale in HAP (hours after pollination) based on published reports is indicated [6,9].

fails to elongate and subsequently divides symmetrically, producing a normally sized apical and an abnormally short basal cell [17,18]. Whether these two processes, zygote elongation and asymmetric division are independently affected in *yda* mutants, or whether cell length is a prerequisite for the asymmetry of the zygote division remains an interesting question to be answered. The requirement of MAP kinase signaling for zygote elongation in *Arabidopsis* is reminiscent of the role of the MAP kinase PAR-1 of *C. elegans*, which organizes microtubule dynamics in the zygote [19]. The possibility of a similar link between YODA function and the reorganization of the microtubular network during zygote development can now be addressed.

The direct target of the YDA phosphorylation cascade has not yet been identified. However, Jeong *et al.* show that the RWP-RK-type transcription factor GROUNDED (GRD)/RKD4 is an indirect, but crucial effector of YDA signaling [20]. The *grd* mutant resembles *yda* and *mpk3 mpk6* mutants at all embryo stages and suppresses the effects of a hyperactive form of YDA. Because all of these mutants fail to generate the suspensor properly, YDA signaling plays also an important role to specify suspensor fate. In a parallel study, Waki *et al.* demonstrate that overexpression of GRD in seedlings induces the expression of several embryo-specific genes and the formation of somatic embryos, suggesting that GRD has a potential activity to initiate embryogenesis [21].

In an elegant study, Bayer *et al.* reveal that YDA is activated by the receptor-like cytoplasmic kinase SHORT SUSPENSOR (SSP), whose mRNA is delivered from the sperm cell to be translated in the zygote (Figure 1) [22]. Thus, sperm-triggered activation of YDA MAPK signaling by a paternally provided component connects the fertilization event to the elongation of the zygote and the initiation of embryogenesis. The role of *SSP* mRNA reveals a different type paternal contribution to axis formation compared to model animal species, where symmetry breaking in the zygote is triggered by sperm-derived proteins [23].

### A WRKY2-WOX transcriptional cascade activates zygote re-polarization

Members of the plant-specific WUSCHEL RELATED HOMEBOX (WOX) transcription factors are expressed asymmetrically along the apical-basal axis throughout embryogenesis [24]. Among them, *WOX2* and *WOX8* are co-expressed in the egg cell and in the zygote, but their expression becomes predominant to the apical (*WOX2*) or basal embryo lineage (*WOX8*) after the zygotic division [24]. A recent study detected weak *WOX8* expression also in the apical daughter cell of the zygote [25]. *WOX2* expression is further confined within the apical lineage to the upper half of the globular embryo from which the shoot is derived. There it regulates apical patterning and shoot development, partly through promoting polar transport and asymmetric distribution of a plant hormone auxin [26]. Expression of the closely related *WOX8* and *WOX9* affects not only basal lineage development, but also through *WOX2* the development of the embryo proper [26,27]. Subsequently, expression of the stem-cell regulating *WOX* genes, *WUS* and *WOX5*, is induced in the shoot and root apical meristems, respectively [28,29]. Finally, expression of *WOX1*, *PRESSED FLOWER* (*WOX3*), and *WOX5* in the cotyledons contributes to organ patterning and outgrowth [24,26]. This cascade of *WOX* functions resembles animal *HOX* genes, the key axis regulatory homeobox genes that are sequentially expressed in specific domains of the embryo [30].

The co-expression of *WOX2* and *WOX8/9* implies that the zygote contains a mixture of basal and apical regulators that becomes separated through the asymmetric division. In a study to reveal how the asymmetric expression patterns are linked to the structural polarity of the zygote, Ueda *et al.* show that the plant-specific zinc-finger transcription factor WRKY2 is a direct activator of *WOX8/9* transcription [10\*\*]. In the *wrky2* mutant, egg cell polarity and the transiently symmetric zygote are normal, but re-polarization of the zygote does not take place and the zygote divides symmetrically [10\*\*]. This reveals the different genetic requirements between the polarization processes of the egg cell and the zygote. Forcing *WOX8* expression partially restores all defects in the *wrky2* zygote, indicating that WRKY2 polarizes the zygote partly

through activating *WOX8* (Figure 1) [10\*\*]. One important observation from this study is that *WRKY2* provided from the pollen is sufficient to activate *WOX8* transcription and to establish zygote polarity. This indicates that transcription after fertilization functions in zygote polarity, in contrast to animal models where early pattern information is derived from maternally supplied products [31]. Several expression profiling experiments corroborate this difference and indicate transcription in the plant zygote on a large scale (see below).

In contrast to the *yda* mutant, *wrky2* does not severely affect zygote elongation, indicating that cell elongation is not sufficient to establish zygote polarity and asymmetric division, and the specific requirement of WRKY2 in the mechanism to establish zygote polarity [10\*\*]. Furthermore, when *yda* or *grd* mutation are combined with the *wox8 wox9* double mutant, the triple mutant embryos display synergistic phenotypes and arrest the development at the zygote stage or shortly thereafter [20\*\*,26]. Therefore in the early embryo patterning, a WRKY2-WOX cascade and YDA MAPK signaling seem to regulate independently zygote polarity and elongation, respectively (Figure 1).

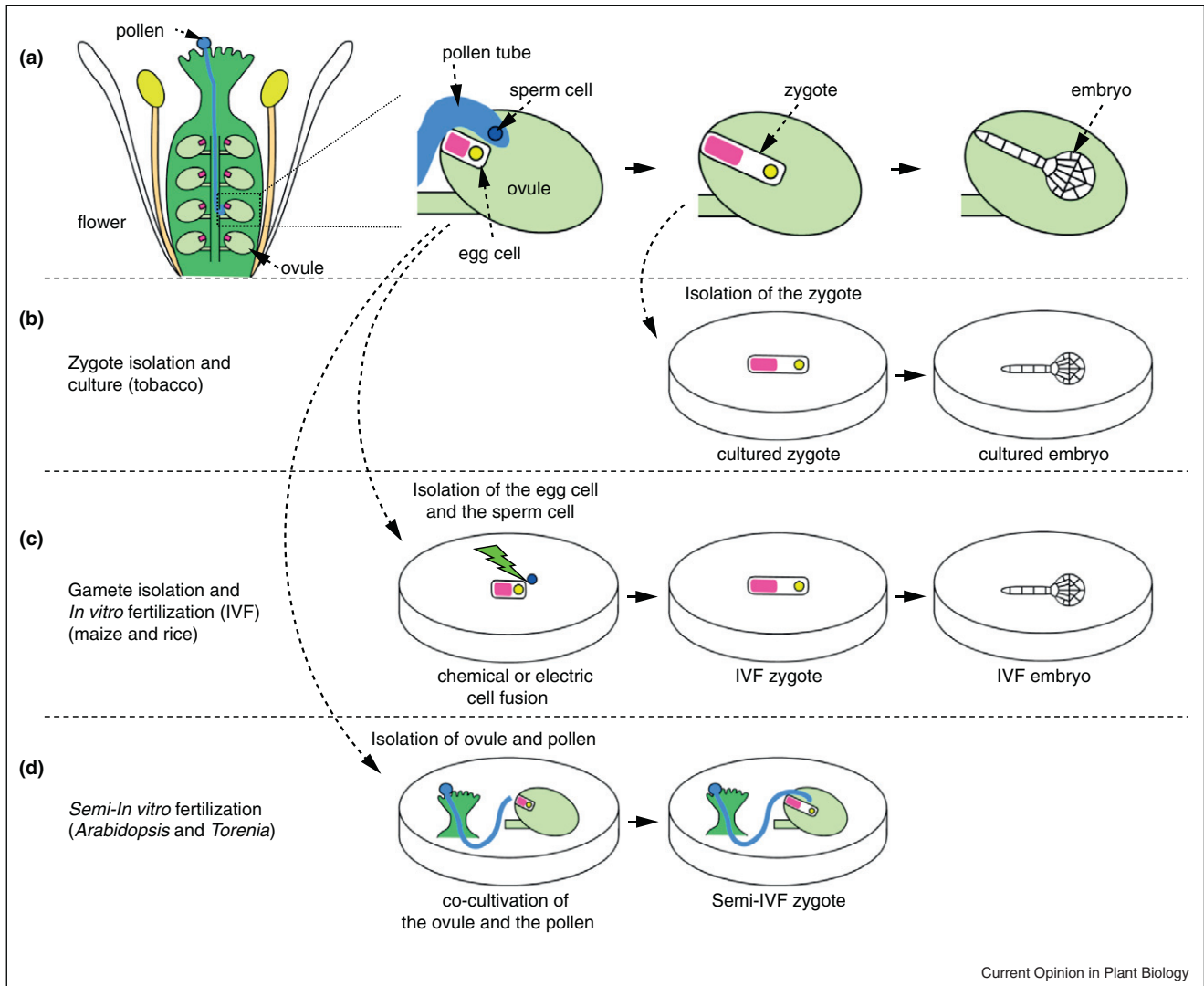
### Transcriptome analysis of the zygote

When transcription is activated after fertilization and whether parental imprinting plays an important role in plants has been a matter of debate. Xiang *et al.* collected *Arabidopsis* zygotes and young embryos from various developmental stages and studied the expression profiles [32\*]. They find that mRNA for 58% of all genes in the *Arabidopsis* genome are present in the zygote, with an over-representation of transcription factor transcripts compared to subsequent embryo stages [32\*]. Autran *et al.* report that maternal transcripts dominate early stages in plant embryogenesis, similar to the situation in animals [33\*\*]. Initially, paternal alleles are repressed via DNA-methylation and histone-methylation and only become activated at later stages through the maternally supplied histone chaperone complex CAF1. In contrast to that study, the transcriptome analysis by Nodine and Bartel of 1-cell or 2-cell embryos provides evidence that paternal and maternal derived mRNA are equally represented, with only very few transcripts preferentially derived from a specific parent [34\*\*]. Widespread transcription in the zygote is also inferred by studies in maize and tobacco [11\*,35,36]. Altogether, these results indicate that plants have a different transcriptional strategy in early embryo development compared to model animals, where *de novo* transcription is largely repressed after fertilization until the late embryo stage [31].

### Cellular analysis and *in vitro* culture of the zygote

Because it is deeply embedded in the ovule (Figure 2A), the development of *in vitro* culture protocols is of great

Figure 2



Methods of zygote studies. Schematic representations of: fertilization and embryogenesis in a flower **(A)**, development of a cultured zygote **(B)**, *in vitro* fertilization (IVF) of an egg cell **(C)**, and *semi-IVF* of an ovule **(D)**. Currently available species for each protocol are designated. In the flower, the pollen grain germinates on the stigmatic surface of the pistil, and the pollen tube delivers the two sperm cells to the ovule. Only one of the two sperm cells is shown to simplify the drawing. The pollen tube that grows through a cultured pistil section is shown in **(D)**.

importance for studying the zygote at the cellular level. The *in vitro* protocol of He *et al.* supports development from a dissected, elongated tobacco zygote to a fertile plant (Figure 2B) [37]. However, if the zygote is not yet elongated, developmental arrest takes place [37]. This could be owing to the requirement of maternal tissue for the very first steps of embryogenesis, or alternatively, a higher sensitivity of the young zygote to the dissection procedure. Interestingly, an elongated zygote lacking the cell wall also fails to develop although a new wall is regenerated on culture media [37]. This finding is reminiscent to pioneering studies in zygotes of the brown algae *Fucus*, where attachment of the cell wall has been shown to be instrumental for specifying cell fate [38]. In a

new study analyzing the transcriptome of zygotes after inhibition of Arabinogalactan proteins (AGPs), Hu *et al.* report that AGPs are required for normal expression of genes involved in various cellular processes including vesicle transport and tip growth [39]. AGPs were originally found to localize preferentially at the apical cell surface of the tobacco zygote, and AGP inhibition forced the zygote to divide symmetrically [40]. Together with the finding that AGPs promote cell elongation in the tip-growing cells in the moss *Physcomitrella patens* [41], a role of AGPs in promoting zygote elongation appears plausible. It will be of interest to understand how this function intersects with the role of the YDA pathway in zygote elongation.

## Live imaging of fertilization

In order to study processes during or immediately after fertilization, *in vitro* fertilization (IVF) protocols have been established in grass species (Figure 2C). Here, the isolated gametes are fused electrically or in the presence of polyethylene glycol (PEG) or bovine serum albumin (BSA), and the resulting cell can be cultured into a mature, fertile plant [42–44]. These studies reveal that sperm entry causes a calcium spike to promote cell wall synthesis in the maize zygote [45], and that the entry site is not crucial to set the cell division plane in the rice zygote [46]. In dicots, however, *in vitro* fertilization has not been successful yet. Chemical fusion of tobacco gametes results in a zygote that arrests without any division [43]. What could cause the differential effectiveness of IVF in grasses and dicotyledons? In grasses, the sperm cell formed in the pollen grains is at G1 and is ready to fuse with the egg cell that is also in G1 [43]. Therefore, sperm isolated from grass pollen is at the right stage for fertilization. The sperm of *Arabidopsis* and tobacco is also at G1 in the pollen grain, but completes S phase after the pollen tube enters the ovule and before fusion with the egg cell [47]. Thus, fertilization-competent sperm may have to be isolated from the discharged pollen tube in the ovule, which seems technically challenging.

As an alternative to IVF, *semi*-IVF has been established in *Torenia fournieri* and *Arabidopsis* (Figure 2D) [48\*\*,49,50]. In this procedure, ovules are co-cultured with germinating pollen from a cut pistil, and fertilization occurs in almost intact ovules [48\*\*]. Hamamura *et al.* report live-imaging of ovules fertilized by this approach revealing that the sperm cell invariably fuses with apical tip of the egg cell [48\*\*]. Using a similar approach, Ingouff *et al.* show that sperm-specific and egg-specific HISTONE 3 variants are degraded shortly after fertilization and are rapidly replaced by newly synthesized variants in the zygote [51]. Because HISTONE 3 variants carry epigenetic information, this finding suggests that significant epigenetic restructuring of both parental genomes takes place in the zygote [51].

## Conclusions

The mechanism of plant zygote polarization has long been obscure, but recent studies identify principal components of this process. Furthermore, experimental advances set the stage for significant and new discoveries at the cellular level of the zygote. Fundamental questions can now be addressed. What are the cues that polarize the zygote? If the sperm entry site is not crucial for zygote polarity, does the egg cell establish stable polarity marks that are used in the zygote? How is zygote polarity connected to embryo patterning? Specifically, does the zygotic division distribute cell fate determinants in an unequal manner to its daughter cells?

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