A common developmental program can produce diverse leaf shapes

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Introduction

Leaves of eudicots show tremendous morphological diversity (Fig. 1). They can be simple or dissected, that is, partitioned into distinct leaflets, and have margins that are entire (smooth) or have teeth, lobes or sinuses of varying shape and depth (see Supporting Information Fig. S1 for related terminology). In addition, the vascular systems supporting leaf blades may have diverse architectures (Hickey, 1973; Ash et al., 1999). Remarkably different leaf morphologies may occur between closely related species, as within-species variants, or even in the same plant (Kidner & Umbreen, 2010; Nicotra et al., 2011). These differences are illustrated by numerous case studies including the leaves of Pelargonium (Nicotra et al., 2007; Jones et al., 2009), grape vine (Vitis spp.) (Chitwood et al., 2014, 2016), tomato (Solanum lycopersicum) (Nuez et al., 2004), and the poppy family (Gleissberg, 2004). Diverse leaf shapes also emerge in molecular-level studies of reference plants including Arabidopsis thaliana, Cardamine hirsuta and Solanum lycopersicum. The model characterizes leaf development as a product of three interwoven processes: the patterning of serrations, lobes and/or leaflets on the leaf margin; the patterning of the vascular system; and the growth of the leaf blade spanning the main veins. The veins play a significant morphogenetic role as a local determinant of growth directions.

We show that small variations of this model can produce diverse leaf shapes, from simple to lobed to compound.

It is thus plausible that diverse shapes of eudicot leaves result from small variations of a common developmental program.

To better understand the essence of this program and examine how it produces diverse forms, we constructed a parametrized computational model of leaf development. The model integrates three perspectives on leaf development: the growth of the leaf blade viewed as a continuous surface, the morphogenetic role of the leaf margin, and the role of the vascular system.

The continuous surface perspective

The continuous surface perspective has its roots in measurements and a mathematical description of growing tobacco (Nicotiana tabacum) leaves (Avery, 1933; Richards & Kavanagh, 1943). In present terms, the development of the leaf blade is quantified by a growth tensor field (Hejnowicz & Romberger, 1984), which is formally equivalent to the strain tensor field defined in continuum mechanics. Local growth is integrated into a global description of the developing leaf blade using the mechanical notion of stress–strain relations (Boudaoud, 2010). The continuous-surface perspective embedded into the computational framework developed by Kenna way et al. (2011) was applied to characterize the development of entire A. thaliana leaves (Kuchen et al., 2012) and winged-shaped mutant barley (Hordeum vulgare) bracts (Richardson et al., 2016). The latter model explains the emergence of the lobe-like wings in terms of polarizers that define different growth directions within a continuous blade.
Morphogenetic role of the leaf margin

The modelling and understanding of leaf shapes can be facilitated by characterizing them in a more structured way. In particular, the patterning of leaflets, lobes or teeth is largely dependent on the processes that take place at the adaxial–abaxial boundary of a leaf primordium, termed the marginal (leaf) blastozone by Hagemeier & Gleissberg (1996). They pointed out that – even in early development, when the leaf primordium is a three-dimensional bump – the blastozone forms a line that ‘anticipates and circumscribes’, the eventual leaf surface. The blastozone can thus be viewed as a one-dimensional boundary of a two-dimensional leaf, similar to the epidermis of a shoot apical meristem being viewed as a two-dimensional boundary of the three-dimensional meristem (Floyd & Bowman, 2010; Prusinkiewicz & Runions, 2012; Alvarez et al., 2016).

In a growing meristem, feedback between auxin and PIN-FORMED1 (PIN1) proteins – auxin efflux carriers – leads to the emergence of PIN1 convergence points that position new primordia where space is available for them (Reinhardt et al., 2003; Jonsson et al., 2006; Smith et al., 2006). This feedback provides a molecular implementation of Hofmeister’s rule, according to which new primordia emerge at locations that are sufficiently distant from the nearest primordia formed previously (Hofmeister, 1868; see also Kirchoff, 2003). The convergence points also position the endpoints of midveins within the emerging primordia (Bayer et al., 2009). In a similar manner, PIN1 convergence points on the margin of Arabidopsis leaves position leaf serrations (Hay

Fig. 1 Examples of diverse shapes and features of eudicot leaves. (a) Quercus imbricaria. (b) Garcinia spicata. (c) Catalpa bignonioides. (d) Populus tremula. (e) Crataegus marshallii. (f) Fatsia japonica. (g) Quercus rubra. (h) Acer saccharinum. (i) Pueraria montana var. lobata. (j) Cannabis sativa. (k) Handroanthus sp. (l) Robinia pseudoacacia. Photograph sources and credits (see also Supporting Information Notes S1): (a) Jan De Langhe, courtesy of Foundation Arboretum Wespelaar (www.arboretumwespelaar.be); (b) Irène Bouguerra and Nicolas Lagarrigue, Pitchandikulam Forest Virtual Herbarium (www.pitchandikulam-herbarium.org), licensed under CC-Attribution-NonCommercial-ShareAlike 4.0; (c) W. Mark and J. Reimer, courtesy of SelectTree (www.selectree.calpoly.edu); (d) commons.wikimedia.org, licensed under CC-Attribution-Share Alike 3.0; (e) John Pickering, courtesy of Discover Life (www.discoverlife.org); (f) commons.wikimedia.org, licensed under CC-Attribution-Share Alike 3.0; (h) Renn Tumlison, Henderson State University (www.hsu.edu/Academics/ARNatureTrivia/), used with permission; (i, l) J. K. Marlow, courtesy of Native and Naturalized Plants of the Carolinas and Georgia (www.namethatplant.net); (j) http://commons.wikimedia.org, image in public domain; Karen Blixen (www.flickr.com), licensed under CC-Attribution-NonCommercial-ShareAlike 2.0.
whether serrations, lobes or leaflets will develop. For instance, *CUP-SHAPED COTYLEDON* (*CUC*) genes specify the boundary between serrations in wild-type *A. thaliana* (Bilsborough et al., 2011) (Fig. 2b), but induce lobes by deepening the sinuses when overexpressed (Nikovics et al., 2006). In compound *C. hirsuta* leaves CUC2 marks the boundaries between leaflets (Blein et al., 2008; Hasson et al., 2011; Rast-Somssich et al., 2015) (Fig. 2b,c). The postulated interaction between auxin, PIN1 and CUC2 on the growing leaf margin has been included in the computational model of wild-type and mutant *A. thaliana* leaves (Bilsborough et al., 2011). An even simpler model, focused on the feedback between auxin and PIN1 alone, faithfully reproduced lobed ivy (*Hedera* sp.) leaves (Prusinkiewicz & Lane, 2013). In these models, auxin concentration controlled the outward expansion of the leaf margin: faster at the PIN1 convergence points and slower between them. A possible tangential component of growth and the displacement of distal leaf parts by growing proximal parts were ignored. Consequently, the geometry of complex leaf shapes, with a hierarchy of growth axes, could only be captured with limited accuracy (Nakamasu et al., 2014).

Morphogenetic role of the vascular system

Growth directions are inherently accounted for in the third perspective on leaf development, which treats leaves as modified shoots (Arber, 1950). This perspective is related to telome theory, according to which leaves evolved by connecting, or webbing, the free-standing branching structure of ancestral plants (Zimmermann, 1952; see also Beierling & Fleming, 2007). The local alignment of growth directions with main veins—the present-day counterpart of ancestral branching structures—is evident in multifid leaves, such as the *Pelargonium graveolens* leaf in Fig. 2(d). Its blade consists of narrow segments, which can only acquire their form by growing faster in the direction of midveins than perpendicular to them. Examination of variegated leaves, in which the dominant directions of growth are indicated by groups of cells with contrasting colours, indicates that they are also aligned with the major veins in leaves with a broad lamina (Dolan & Poethig, 1998) (Fig. 2e). The relative rigidity of veins (Hagemann, 1999; Bar-Sinaï et al., 2016) predisposes them mechanically to act as the main axes of growth, with the leaf blade spanning the skeleton of veins. As observed by Dengler & Kang (2001) commenting on an earlier hypothesis by Van Volkenburgh (1999), vascular parenchyma may act as a driver of leaf development, that is, provide the motive force for overall leaf expansion, while mechanical resistance is offered by the epidermis. At the molecular level, the patterning of veins is commonly explained in terms of the canalization theory (Sachs, 1981), which posits that veins differentiate along self-organizing paths of auxin flow. In this process, a positive feedback between polar auxin transport and the distribution of auxin transporters in cells creates narrow canals of auxin transport in a manner analogous to the carving of rivers by flowing water (Sachs, 2003). The canalization theory is consistent with experimental data and computational models, in which new vascular strands emerge as gradually refined conduits of auxin, connecting convergence points with the existing vasculature.
Models with the aim of capturing the essence of the canalization process at a higher, geometric level of abstraction have also been proposed (Rodkaew et al., 2003; Runions et al., 2005; Owens et al., 2016).

**Description**

To better understand the development of leaf form and the basis of leaf diversity, we have constructed a computational model of leaf development that builds upon the three perspectives discussed in the previous section. The model is specified in the L+C extension of the C++ programming language (Karwowski & Prusinkiewicz, 2003). The source code and parameter files are available from the authors’ website (http://algorithmicbotany.org/papers/leaves2017.html). Parameters of all models are also collected in Table S1. Simulations can be executed and visualized using the Virtual Laboratory software environment (http://algorithmicbotany.org/virtual_laboratory). The structure and operation of the model are described in the remainder of this section; further details are given in Notes S2.

**Leaf representation**

Consistent with previous models of leaf shape (Bilsborough et al., 2011; Prusinkiewicz & Lane, 2013; Nakamasu et al., 2014), we disregard the adaxial–abaxial leaf dimension and model the developing leaf as a two-dimensional structure. We further simplify models by considering development as a planar process, thus ignoring possible three-dimensional wrinkling and folding of the leaf. We follow Hagemann & Gleissberg (1996) in abstracting from the cellular-level details of leaf development and focusing on larger components. The leaf is thus represented as three coupled data structures: an open polygon representing the leaf margin (marginal blastozone), a two-dimensional tree representing the main veins (synonymous with growth axes), and a triangle mesh representing the leaf blade (Fig. 3a). The margin polygon is defined by a sequence of vertices, referred to as the sample points, which begins and ends at the bottom of the petiole. The vascular system is approximated as an open branching structure with straight branches (we do not consider reticulate venation patterns). The root of this system is located at the leaf base, and the terminal points coincide with the convergence points on the margin. The points in the neighbourhood of each convergence point define the margin segment that is associated with this point and with the vein that terminates at it. Each margin point can be characterized further by the presence or absence of one or more morphogens that affect the patterning of subsequent convergence points and/or the growth of the margin (Fig. 3b). Please note that we use the noun ‘morphogen’ as a general term denoting a substance involved in the production of form (Turing, 1952). We do not require that morphogens form gradients.

A sample point is thus specified by its position in space, type (convergence point or not), pointer to the convergence point/vein it is associated with, and one or more flags representing the presence or absence of morphogens. The leaf blade is represented by a triangle mesh, constrained such that all vertices and edges of the leaf margin and the vascular system coincide with vertices and edges of the mesh. During simulation this mesh is dynamically refined to allow for faithful representation of details of the leaf shape as it develops (see Notes S2.1).

**Simulation outline**

Leaf form emerges as an outcome of development, simulated as a feedback loop of growth, patterning of the margin and the

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**Fig. 3** Key elements of the model. (a) Data structures representing a leaf: a polygon representing the leaf margin (black), a tree representing the main veins (blue), and a triangular mesh representing the lamina (green). Yellow circles indicate convergence points, black dots are additional sample points on the margin, and blue dots indicate the vertices at which the mesh coincides with the veins. (b–d) Example of patterning on the margin. (b) Margin fragment with two convergence points, CP1 and CP2, and the associated margin segments (green bars). Red bars indicate the presence of a morphogen. (c) Distance relations change as a result of margin growth. An interval with red morphogen reaches threshold length λ. (d) A new convergence point CP’ forms in the middle of the red interval and initiates a new vein. Sample points at distance δ or less from CP’ become the margin segment associated with CP’. The red morphogen is excluded from points closer than ρmin to CP’. Another morphogen (blue bars) is introduced at distances greater than ρmin and less than ρmax from CP’.
insertion of new veins. The initial shape of the leaf (leaf primordium with a midvein) and the distribution of morphogens on its margin are specified explicitly at the beginning of the simulation (Fig. 4a). The simulation proceeds iteratively. Each iteration begins with a growth step, driven by expansion of the vascular system. The expansion increases distances measured along the margin and modifies the distribution of morphogens that affect patterning or growth. These changes may trigger the insertion of new convergence points followed by a further modification to the distribution of morphogens (Fig. 4b). The insertion of new convergence points leads to the creation of veins that connect these points to the existing vascular structure and define new growth axes. This step closes the feedback loop: the system is now ready for the next growth step (Fig. 4c,d). The leaf blade interior, spanning the space between the branching vascular system and the leaf margin, expands passively, following the growth of veins and propagation of the leaf margin.

### Growth of the vascular system

Leaf growth is modelled as the superposition of two processes: uniform expansion of the blade, which represents the isotropic component of growth, and growth driven by veins, which introduces an anisotropic component. The growth of veins is modelled, in turn, as the sum of two factors: the addition of new vein segments of a given length at vein tips, and the elongation of existing segments. The latter factor is defined in terms of the relative elementary rate of growth (RERG) (Richards & Kavanagh, 1943; Hejnowicz & Romberger, 1984):

\[
\text{RERG}(P) = \frac{\Delta l}{\Delta t}, \quad \text{Eqn 1}
\]

where \(l\) is the length of an (infinitesimal) vein segment at point \(P\), and \(\Delta l\) is the increase of this segment’s length over time \(\Delta t\). The elongation of a finite-length vein segment between points \(P_1\) and \(P_2\) is thus the line integral:

\[
\Delta l_{P_1 P_2} = \int_{P_1}^{P_2} \text{RERG}(P(s))\,ds. \quad \text{Eqn 2}
\]

In general, the RERG may be a highly nonlinear function of position \(P\) (Kuchen et al., 2012). For simplicity, we assume that:

![Fig. 4 Overview of the model operation. (a–d) Two iterations of simulated development. Coloured lines along the margin indicate morphogens that control development. In this case, the red morphogen enables formation of convergence points and defines sinuses, and the blue morphogen defines the petiole. (a) A leaf at the beginning of the simulation. (b) The midvein grows, driving the expansion of the margin and the blade. New convergence points (yellow circles) emerge on the margin, causing the insertion of new veins and dividing the leaf into regions associated with each vein. (c) Beginning of the next iteration: the veins grow again, causing differential expansion of the leaf in the directions associated with veins. (d) The shape of the margin is further affected by the margin propagation (webbing). (e) Vascular system expansion. Vein elongation is limited to the bottom portion of the leaf (shown in red). This portion does not propagate with the growing leaf and is limited to the vasculature points within the same distance from the leaf base in each simulation step. Vein segments \(AB\) and \(BP\) are outside the growth zone and do not elongate (e.g. \(|AB| = |A'B'|\) and \(|BP| = |B'P'|\)). Although distal parts of the leaf do not grow, they are displaced by the growing bottom part; for instance, point \(P\) moves to \(P'\). (f) Margin growth. Points within a margin segment (e.g. \(A, B\)) are projected perpendicularly onto the corresponding veins (\(A', B'\)). A boundary point (\(C\)) is projected onto the branching point between veins (\(C'\)). Following vein expansion, point \(D\) is moved by the same vector as its projection: \(DE = D'E'\).](image-url)
Vein insertion

Consistent with molecular data (Scarpella et al., 2006; Bayer et al., 2009; O’Connor et al., 2014), a new convergence point on the leaf margin induces a vein that connects this point to the existing vasculature. The attachment point \( P \) at which this vein will meet the vasculature is computed using two heuristics (Fig. 5d). The first heuristic is motivated by the observation that vascular strands tend to provide short and straight connections between sources and sinks (Bayer et al., 2009), and the hypothesis that these connections may minimize resistance to the transport of water and photosynthates in the leaf (Sack & Scoffoni, 2013). According to this heuristic, the attachment point \( P \) is found by minimizing the expression \( \tau = b \| CP−P \| + \nu \| P−B \| \), where \( \| CP−P \| \) is the length of the inserted vein (the distance between the new convergence point \( CP \) and the attachment point \( P \)), \( \| P−B \| \) is the length of the path from the attachment point to the leaf base \( B \) (the arc-length distance from \( P \) to \( B \), measured along the veins), and parameters \( b \geq \nu > 0 \) are the resistances to transport per unit distance in the leaf blade and in the veins, respectively. Variable \( \tau \) thus represents the total resistance to transport from the convergence point to the leaf base. It can be shown that any vein minimizing \( \tau \) meets an existing vein at a constant branching angle \( \Theta = \arccos(v/b) \) (Notes S3) or is attached to an existing branching point. Assuming that resistance to the transport of water and sugars is anticipated by resistance to the transport of auxin that patterns the vascular system, the resistance-based model is qualitatively supported by the observed reduction of the angle at which secondary veins meet the midvein in plants treated with an auxin transport inhibitor (Mattsson et al., 1999). The second heuristic constrains the angle between the new vein and the normal to the margin to an interval \( [−\sigma_{\text{max}}, \sigma_{\text{max}}] \). It prevents veins from approaching the margin at unnaturally small grazing angles.

Results

The parameter values for the leaf forms and developmental sequences discussed in this section are listed in Table S1. These values were found in two types of \textit{in silico} experiment. In the first type (theoretical morphospace exploration; McGhee, 1999), we systematically modified one or two parameters while keeping the remaining parameters constant to visualize the space of resulting forms. In the second type (synergistic human-computer problem solving;icklidi, 1960), we modified and refined parameter values interactively to approximate select real leaves. When needed, this process also involved adjustments to the model code, gradually yielding the current implementation.

Plausible patterns of leaf development emerge from a self-organizing process

A basic pattern of development produced by the model is illustrated in Fig. 6(a–f) (Movie S1). A heart-shaped leaf with branching venation is produced in a self-organizing process, in which convergence points determine veins, veins determine growth...
directions, and growth of the margin leads to the formation of new convergence points. The only leaf part excluded from this loop is the petiole, where a locally expressed morphogen suppresses lateral growth and inhibits the formation of convergence points. The overall form is not defined globally, but results from the integration of local subprocesses.

Growth distribution controls both the shape and venation pattern of simple leaves

The impact of additive marginal growth at vein tips is illustrated in Fig. 6(g,h,i–m). If the marginal growth is small, the vascular system expands uniformly. Lateral veins, inserted after the mid-vein, are relatively short, and this proportion is maintained throughout the subsequent growth of the leaf. The result is an elongated leaf with a strictly or approximately pinnate venation (Fig. 6g,h,i,j). A uniform increase in the marginal component of growth gradually reduces the relative differences in vein lengths, producing leaves with broader blades (Fig. 6k,l). With even stronger marginal growth, higher order veins emerge near the leaf base, yielding cordate leaf forms (Fig. 6f,m).

Limiting growth to basal portions of the leaf prevents elongation of veins in more distal positions, which results in a transition of leaf shape from elliptic to ovate to oblong (Fig. 6n–p). The aspect ratio (width: length) of a leaf also depends on the branching angle \( \theta \) between the veins: as this angle increases, the leaf becomes wider (Fig. 6q,r). Decreasing the range \( \sigma_{\text{max}} \) of angles that a vein can form with respect to the margin prevents veins from reaching the margin tangentially (compare the veins near the base of the leaf in Fig. 6q with those in Fig. 6s,t). If clamping to \( \pm \sigma_{\text{max}} \) overrides other criteria of vein insertion, small values of \( \sigma_{\text{max}} \) may induce a variation in the branching angles \( \theta \) between veins. This variation is reflected in variable vein orientations and a less regular leaf margin (Fig. 6s,t).

Differences in webbing control the margin of simple leaves

Webbing plays a critical role in defining the leaf margin. Strong webbing, characterized by a significant resistance to stretching and bending, results in a smooth margin (Figs 6i–p, 7a). With reduced webbing, the parts of the margin further from the extending tips lag behind those near the tips, resulting in the
The shape of these teeth depends on the resistance of the leaf margin to bending (Fig. 7b,c). Different resistance for stretching at the proximal and distal segments of the polygonal approximation of the margin, meeting at a convergence point, results in asymmetric serrations (Fig. 7d).

Uniform expansion promotes the emergence of compound teeth

In the examples discussed so far, different parts of the leaf blade expand anisotropically following the growth of veins in their proximity. A margin segment spanning the endpoints of two parallel growing veins may thus not expand at all. By contrast, an isotropic expansion of the entire leaf increases all distances along its margin uniformly. This may lead to the recursive insertion of intercalary convergence points between those formed previously, inducing a hierarchy of teeth (Fig. 7e–g; Movie S2). Similar hierarchies occur in many leaves in nature; for instance, compare the model in Fig. 7(g) with the photograph of a *Crataegus marshallii* (parsley hawthorn) leaf in Fig. 1(e). Another example is given in Fig. 7(h,i), which compares a photograph and model of a *Platanus occidentalis* (American sycamore) leaf. Note the similarities in both the shape of the leaf and the structure of its vascular system.
Inhibition of convergence point formation in sinuses yields leaves with lobes

Decreasing the strength of webbing deepens the indentations between teeth. If, in addition, a morphogen inhibits the formation of new convergence points in these indentations, a lobed leaf results (Fig. 8a–e; Movie S3). The formation of new convergence points, and thus new veins and higher order lobes, is then limited to the distal (subapical) part of each lobe. With different parameter values, this model captures the shape of many common leaves. For instance, Fig. 8(f–h) shows the palmately lobed leaves of three maple species.

In Fig. 8(a–h), first-order lobes are initiated close to the leaf base. If the window of morphogenetic competence is moved upward from the leaf base, a more elongated leaf blade supported by a pinnate vascular system results (Fig. 8i). A similar dependence of leaf type (palmate vs pinnate) on the position of the window of morphogenetic competence was observed in the model study by Jeune & Lacroix (1993).

Elongated pinnately lobed leaves also emerge when the growth of lateral veins is delayed with respect to the insertion of convergence points that induce these veins. Conversely, broader palmately lobed leaves emerge if the growth of lateral veins is accelerated (Movie S4). The morphospace in Fig. 9 illustrates the combined effect of this delay/acceleration and the rate of webbing. The latter factor controls a progression of leaf shapes from simple to recursively lobed. Select forms in this morphospace resemble leaves of different Pelargonium and Chrysanthemum species, although the full spectrum of their diversity is larger (Jones et al., 2009; Kim et al., 2014).

Spatio-temporally limited competence to create convergence points yields leaves with simple lobes

The progression from leaves with compound teeth (Fig. 7e–i) to leaves with compound lobes (Figs 8, 9) was modelled by introducing a morphogen that suppressed the emergence of convergence points and growth in sinuses. An additional developmental
control is needed to suppress the formation of higher order lobes. It can be effected by a spatio-temporal window of morphogenetic competence that limits the formation of sinuses to early stages of leaf development and to locations near the leaf base (Fig. 10a; Movie S5). Small changes in the duration of competence yield leaves with different numbers of lobes (Fig. 10b,c). The simple, quantitative nature of these changes may explain the lability of lobe numbers common in some plant species, for example *Brachychiton acerifolius* (flame tree; Fig. 10d). Moreover, small differences in the initial development of the left and right sides of the leaf, for example as a result of differences in the distribution of morphogens, may be amplified by further development producing asymmetric leaves with different numbers of lobes on the left and right sides (Fig. 11).

**Strong inhibition of webbing in sinuses yields dissected leaves**

Increasing the inhibition of webbing by a morphogen acting in sinuses shifts leaf form from moderately to strongly lobed to pinnately compound (Fig. 12a–c). Development of pinnately compound leaves is more complex. Not only must the webbing be suppressed along the midvein to produce the petiole and rachis, but the elongation of the rachis must also be suppressed at the points of leaflet attachment to prevent excessive widening of the leaflet bases. These requirements can be satisfied by the coordinated action of two morphogens (Fig. 12d; Movie S6). The first morphogen suppresses lateral growth in sinuses, as in Fig. 12(c). It does not prevent, however, the formation of new convergence points (a) (b) (c) (d) (e) (f) (g) (h) (i) Fig. 8 Modelling lobed leaves. (a–e) Simulation of the development of a generic palmately lobed leaf (bars indicate constant reference length). The red morphogen inhibits the formation of intercalary convergence points and deepens the sinuses. (a) At the beginning of the simulation, the red morphogen is present near the base of the incipient leaf. (b) As the leaf grows, new convergence points emerge in the morphogen-free intervals, and new intervals of the red morphogen form between the old and new convergence points (arrows). (c–e) Iteration of this process elaborates leaf shape while preventing the formation of new convergence points in the sinuses. As in the previous examples, the blue morphogen defines the petiole. (f–h) Modelling leaf diversity: photographs and models of (f) *Acer macrophyllum* (big leaf maple), (g) *Acer campestre* (field maple) and (h) *Acer grandidentatum* (bigtooth maple). All models result from small changes in the parameter values of the generic palmate leaf model illustrated in (e). (i) A photograph and model of a pinnately compound *Quercus rubra* (northern red oak) leaf. The transition from palmately to pinnately compound form results from a larger window of morphogenetic competence for the red morphogen, enabling the insertion of primary lobes further from the leaf base, compared with the palmately lobed leaves. Photograph sources and credits: (f) Dan Anderson (www.tree-species.blogspot.com), used with permission; (g) Middle European Woods data set (Novotný & Suk, 2013) (http://zoi.utia.cas.cz/node/662), licensed under CC-Attribution-ShareAlike 3.0; (h) Adapted from a photograph by Dean Hueber (http://www.pbase.com/deanhueber/image/90024562; downloaded 20 July 2016).
points. The growth of the petiole and rachis can thus induce new convergence points and leaflets. The second morphogen is generated near each convergence point and defines the leaflet boundary at its point of attachment to the rachis. It inhibits longitudinal growth of the rachis, as required to properly form leaflet bases, and divides the margin into intervals within which distances are measured independently. This division stabilizes the development of leaflets by isolating them from the processes that initiate new leaflets along the midvein. Through varying model parameters, both sessile and non-sessile leaf forms arise (Fig. 12e,f). In addition, the model can capture subtle asymmetries in leaflet shape along the proximo-distal leaf axis, as a result of slightly different initial conditions on the two sides of the petiolules (Fig. 12g).

Discussion

The central question addressed in our paper is the developmental origin of leaf diversity. Following the inferences of Hagemann & Gleissberg (1996), molecular data, and previous computational models outlined in the Introduction, we have attributed patterning of protrusions and indentations to morphogenetic processes taking place on the leaf margin. Furthermore, based on observations of multifid and variegated leaves, and taking into account the mechanical rigidity of veins, we hypothesized that main veins play an important morphogenetic role by defining local growth directions, that is locally polarizing growth. Consistent with the hypotheses of Van Volkenburgh (1999) and Dengler & Kang (2001), we also assumed that the intervening leaf blade tissue locally follows these directions. With different parameters, our model captures the essential aspects of the development and shape of a wide range of eudicot leaves, which supports its plausibility and leads to the following conclusions.

Leaf development is a self-organizing process

Molecular-level processes apparently act by establishing the ‘rules of the game’ that integrate growth, dynamic patterning on the leaf margin, and the formation of the vascular pattern into a self-organizing system characterized by several feedback loops (Fig. 13). In particular, auxin-driven interactions on the leaf margin establish a metric (distance measure) for patterning protrusions and indentations, and the vascular system complements the morphogenetic role of the margin by specifying local growth directions.

A common mechanism can produce widely diverse leaf forms

It is known that small modifications to a self-organizing process can fundamentally change its outcome (Wolfram, 1984). This does not preclude different molecular implementations of the same developmental program, or, conversely, the recruitment of the same molecular process for different morphogenetic purposes. The self-organizing character of leaf development is probably essential to the diversity of leaf forms. For example, the frequently observed transitions between simple, lobed and recursively lobed leaves (e.g. Hareven et al., 1996; Hay & Tsiantis, 2006; Jones et al., 2009; Efroni et al., 2010; Bar & Ori, 2014) (Fig. 9) may be attributed to the feedback loop in which weaker webbing produces deeper sinuses, and the resulting increase in the length of the leaf margin creates space for new convergence points, veins and lobes. Such changes can be plausibly attributed to small modifications of the plant genotype, differences in developmental context (heteroblasty), or environmental factors (Nicotra et al., 2011).

Hofmeister’s rule extends to leaf development

The insertion of new convergence points when the distances to previously formed points exceed a threshold plays a
prominent role in leaf development. In the context of phyllo-tactic patterning, this distance-based criterion is known as Hofmeister’s rule. Hofmeister noticed that ‘the appearance of new lateral organs in the largest of the spaces between the nearest older organs of the same type on the same axis is a phenomenon of almost complete universality’ (quoted after Kirchoff, 2003). We observe that the universality of this rule exceeds even its author’s expectations: it applies to the emergence of new outgrowth not only ‘on the same axis’, but along the leaf margin as well.
The extension of Hofmeister’s rule from phyllotaxis to leaf formation has its source in the similarities between the molecular processes governing the two phenomena. They include the creation of convergence points through the interaction between PIN1 and auxin, the role of additional morphogens, such as CUC proteins, as factors shaping boundaries, and the formation of vascular strands by auxin-driven canalization beginning at convergence points (Floyd & Bowman, 2010; Alvarez et al., 2016). These similarities may reflect the common nature of the developmental problem solved by plants in each case: how to place a number of elements (flowers, floral organs, leaves, leaflets and lobes) within the constraints of available space. The strikingly different appearance of spiral phyllotactic patterns and leaves results ‘not from fundamentally different morphogenetic processes, but from different geometries on which they operate: an approximately paraboloid SAM (shoot apical meristem) dynamically maintaining its form vs a flattening leaf that changes its shape and size’ (Prusinkiewicz & Runions, 2012).

The telome theory and the notion of blastozone are related to each other

Zimmerman’s telome theory postulates that leaves evolved by webbing the branching structures of early land plants. It does not specify, however, whether the control of development has
remained with the branching structure – corresponding to the leaf vasculature – or has been transferred to another part of the leaf. Hagemann and Gleisberg’s blastozone theory implies that the control of development has been transferred to the leaf margin. Over the last decade, it found strong experimental support in molecular studies of reference plants. From an evolutionary perspective, the transfer of control from the branching structure to the leaf boundary has the advantage of creating a vascular scaffolding in concert with the available space on the leaf margin. This phenomenon can be observed, for example, in the gradual proliferation of veins near the base of cordate leaves (e.g. Figs 6a–f, 7hi,j) and the emergence of small veins supporting higher order protrusions in recursively lobed leaves (compare leaves in the right column of Fig. 9).

Geometric models provide a framework for interpreting molecular mechanisms of leaf development

Geometric terms are an abstraction that highlights the morphogenetic role of specific molecular-level processes (Prusinkiewicz & Runions, 2012; Runions et al., 2014). For instance, the interaction between auxin and PIN proteins in the epidermis of the shoot apical meristem or on the leaf margin can be characterized as a mechanism for distance measurement. This characterization furnishes an explanation for the initiation of convergence points in regular patterns (Jönsson et al., 2006; Smith et al., 2006; Bilsborough et al., 2011; O’Connor et al., 2014), although additional biomechanical (Hamant et al., 2008) or biochemical factors (e.g. CUC proteins; Bilsborough et al., 2011) are also relevant.

The current understanding of CUC proteins suggests that they are involved in shaping the boundaries of protrusions, such as serrations in *A. thaliana* leaves (Nikovics et al., 2006; Blein et al., 2008; Bilsborough et al., 2011), or leaflets in the compound leaves of tomato (Berger et al., 2009) and *C. hirsuta* (Rast-Somssich et al., 2015). However, CUC also plays an important role in enabling the formation of convergence points (Bilsborough et al., 2011). This role was not conserved in all the models devised in our work, suggesting that the coordination between the initiation of protrusions and the sculpting of the indentations between them may be more diversified than studies of current reference plants indicate.

Other genes and proteins can also be interpreted in the context of our models. The recently discovered growth inhibitor REDUCED COMPLEXITY (RCO) contributes to leaf dissection (Sicard et al., 2014; Vlad et al., 2014) and appears to play a critical role in defining linear elements – the rachis and petiolules – of compound leaves (Vlad et al., 2014). A similar role may be played by several auxin response factors (ARFs) in tomato (Ben-Gera et al., 2016). The development of compound leaves also involves class I *KNOTTED-like homeobox* (*KNOX*) genes (Bharathan et al., 2002; Bar & Ori, 2014), which delay the progression of leaf maturation, and appear to extend the spatio-temporal window of marginal patterning. By contrast, *TCP* expression delays maturation and increases the number of protrusions initiated at the leaf margin (Barkoulas et al., 2007; Bar & Ori, 2014). With research in progress on the molecular underpinnings of leaf shape, computational modelling is likely to continue to play a significant role in verifying whether the spatio-temporal patterns of gene expression and morphogen distribution are consistent with the morphogenetic roles attributed to them, synthesizing our understanding of leaf development, and providing a framework for considering this development from an evolutionary perspective.

Open problems

The mechanism by which the veins extending from a convergence point find their target location is unclear (Bayer et al., 2009) and, consequently, so is the developmental mechanism that determines the branching angles between veins of different orders. The heuristics used in this paper provide a working hypothesis, but also highlight the need for research exposing the molecular mechanisms through which veins find their attachment point and create particular branching angles. Moreover, the assumption of an open venation system is conspicuously violated in brachidodromous patterns, in which secondary veins form pronounced loops. Although some models can create closed loops (Couder et al., 2002; Runions et al., 2005), the mechanisms that control the development of closed venation patterns remain largely unknown (Scarpella et al., 2010). Even in open venation systems, the assumption of straight veins is a simplification, as different processes – for example, nonuniform expansion of the lamina with the embedded veins – may yield curved veins. In addition, some
leaves include veins that penetrate indentations rather than protrusions. A detailed investigation of the paths of veins and the mechanisms that produce them remains an intriguing open problem, bridging molecular biology and differential geometry.

Our model postulates that growth directions are locally aligned with the veins. With veins running in different directions, as dictated by their branching pattern, the growth directions and magnitudes may form a highly nonuniform field. A detailed multiscale analysis of growing leaves is needed to reveal whether this mathematical possibility holds for real leaves. The growth tensor would then be scale dependent, in the same way as distance measures are scale dependent in fractal objects. The fractal nature of the growth field may be obscured when considering leaves at larger scales. This would explain why the average values of the growth tensor inherent in continuous-surface models suffice when characterizing and modeling simple or broadly lobed leaves (Kuchen et al., 2012; Richardson et al., 2016).

Recent experimental results obtained by Das Gupta & Nath (2016) indicate that different leaf growth patterns, at a coarse level manifested by the distinction between basipetal, uniform and acropetal growth, occur across eudicot species, contributing to the diversity of their leaves. We have acknowledged this source of diversity in our paper (Fig. 6n–p), but its interplay with other morphogenetic factors and impact on the final leaf forms is presently unclear and awaits further study.

Our model ignores departures of leaves from planarity. The molecular basis of leaf curvature – for example a delayed arrest of growth near the leaf margin – is increasingly well understood (Nath et al., 2003; Sharon et al., 2007; Efroni et al., 2008; Prusinkiewicz & de Reuille, 2010), but modelling the morphogenesis of leaves that are curved as well as lobed or serrated remains an open problem. Extending the model described in this paper to curved leaves would probably entail incorporating a biomechanical component, as their forms can most readily be expressed as states of minimum energy of surfaces resisting stretching and bending. Challenges are posed by the nonhomogeneous structure of leaves, with veins embedded within a thin leaf blade (Hong et al., 2005), and the need to replace straight vein segments with their more complicated counterparts defined on curved surfaces: the geodesic curves.

Many leaves are folded as they develop. This folding may play an important morphogenetic role, especially in the case of leaves that develop within the confines of a bud (Couturier et al., 2009, 2011, 2012). Reconciling models described in this paper with those attributing a significant morphogenetic role to folding remains an open problem, although a link between them can be foreseen (for instance, convergence points may define how the leaf is folded).

While the leaves discussed so far can be approximated as flat or curved surfaces (formally, two manifolds), some succulent leaves are fully three-dimensional, volumetric structures (three manifolds) and have a correspondingly three-dimensional vasculature (Korn, 2011; Ogburn & Edwards, 2013). It would be most interesting to determine whether our model can be extended to this case as well, possibly shedding light on the evolutionary path between surface-like and volumetric leaves.

Finally, our models are supported by visual comparisons of generated forms with photographs of mature leaves. This brings into focus the lack of adequate data concerning the development of diverse leaves from the earliest stage of leaf primordia to maturity. Acquiring such data using current methods is a tedious process (cf. Kuchen et al., 2012); however, recent results indicate that the diversity of growth patterns is not well represented by the limited spectrum of current reference plants (Das Gupta & Nath, 2015, 2016). In addition, visual comparisons should be complemented by measurements and comparison criteria rooted in leaf morphometrics (for example, see Biot et al., 2016 and references therein). An examination of models in light of quantitative data will provide an opportunity to refine the models and validate them objectively; conversely, we expect that the models will provide a useful theoretical framework for interpreting new experimental results.

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Author contributions

P.P. and A.R. designed the research; A.R. performed the research; P.P., A.R. and M.T. analyzed and interpreted the results; P.P. and A.R. wrote the paper with input from M.T.

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Bharathan G, Goliber T, Moore C, Kessler S, Pham T, Sinha N. 2002. GOBLET specifies leaflet boundaries in compound tomato leaves. The NAC-domain transcription factor.
Movie S2 Simulation of the development of a leaf with compound teeth (corresponds to Fig. 7g).

Movie S3 Simulation of a generic palmately lobed leaf (corresponds to Fig. 8a–e).

Movie S4 Simulation of the development of a representative leaf from the 2D morphospace in Fig. 9 (row 3, column 4).

Movie S5 Simulation of the development of a generic palmate leaf with sequential emergence of lobes (corresponds to Fig. 10a).

Movie S6 Simulation of the development of a pinnately compound leaf (corresponds to Fig. 12d).

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