A local activity-dependent learning rule shapes dendritic morphologies and synaptic organization

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Abstract   Dendritic branching and synaptic organization shape single neuron and network computations. How they emerge simultaneously during brain development as neurons become integrated into functional networks is still not mechanistically understood. Here, we propose a computational model in which dendrite growth and the organization of synapses arise from the interaction of local activity-dependent plasticity and activity-independent cues from potential synaptic partners. Consistent with experiments, three phases of dendritic growth – overshoot, pruning, and stabilization – emerge naturally in the model. The model generates biologically-realistic dendritic morphologies under normal and perturbed learning rules, and synapses cluster spatially according to the correlated activity they experience. We demonstrate that activity-dependent and -independent factors influence dendritic growth throughout development, suggesting that early developmental variability can affect mature morphology. Our model generates dendrites with approximately optimal wiring length consistent with experimental measurements. Therefore, a single mechanistic model generates realistic dendritic development and accounts for synaptic organization from correlated inputs. Our work suggests concrete mechanistic components underlying dendritic growth and synaptic formation and removal in function and dysfunction.

Introduction

The dendrites of a neuron are intricately branched structures that receive electrochemical stimulation from other neurons. The morphology of dendrites determines the location of synaptic contacts with other neurons and thereby constrains single-neuron computations. During development, the dendrites of many neurons grow simultaneously and become integrated into neural circuits. Dendrite development is highly dynamic; iterated addition and retraction of branches allow these dendrites to probe various potential synaptic partners before stabilizing (*Cline, 2016*;

Richards et al., 2020). Many intrinsic and extrinsic factors underlie the dynamics of dendritic development. In any given neuron, intrinsic expression of specific genes controls many morphological aspects, including the orientation of the dendrite in the cortex, the general abundance of dendritic branching, and the timing of growth onset (*Puram and Bonni, 2013*). Extrinsic signaling, in contrast, exerts precise control over the detailed dynamics of dendrite development via various mechanisms, including activity-dependent cell-to-cell interactions and molecular signaling (*Polleux et al., 2016*).

While many signaling molecules affect dendrite development, the brain-derived neurotrophic factor (BDNF) and its immature predecessor proBDNF are particularly crucial in the central nervous system (*Lu et al., 2005*). While exposure with BDNF leads to larger dendrites with a higher density of synapses (*McAllister et al., 1995*; *Tyler and Pozzo-Miller, 2001*), exposure with proBDNF leads to smaller dendrites with fewer synapses (*Koshimizu et al., 2009*; *Yang et al., 2014*). Furthermore, the precise balance of BDNF and proBDNF is essential for the emergence of synaptic organization during development (*Kirchner and Gjorgjieva, 2021*; *Winnubst et al., 2015*; *Kleindienst et al., 2011*; *Niculescu et al., 2018*). Interestingly, synaptic activity triggers the cleaving of proBDNF into BDNF (*Je et al., 2012*), providing a mechanistic link between the molecular factors driving dendrite maturation and neural activity.

Activity-dependent factors are equally important in driving dendritic growth. As the sensory periphery is immature during early postnatal development, when many dendrites grow (*Leighton and Lohmann, 2016*), many developing circuits generate their own spontaneous activity. The rich spatiotemporal structure of spontaneous activity instructs the formation, removal, and change in the strength of synaptic inputs (*Sretavan et al., 1988*; *Sakai, 2020*) and triggers the stabilization or retraction of entire dendritic branches (*Riccomagno and Kolodkin, 2015*; *Lohmann et al., 2002*). While blocking spontaneous activity does not result in grossly different dendrite morphology, the density and specificity of synaptic connections are strongly perturbed (*Campbell et al., 1997*; *Ultanir et al., 2007*), highlighting the instructive effect of spontaneous activity on dendritic development (*Crair, 1999*).

One influential hypothesis tying together the extrinsic signaling factors underlying dendritic development is the synaptotrophic hypothesis (*Vaughn, 1989*). According to this hypothesis, a growing dendrite preferentially extends into regions where it is likely to find synaptic partners. Once a dendrite finds such a partner, a synaptic contact forms, anchors the developing dendrite, and serves as an outpost for further dendritic growth. Conversely, loss of synaptic input to the dendrite can lead to retraction unless the remaining synapses stabilize the branch (*Lohmann et al., 2002*; *Niell et al., 2004*; *Haas et al., 2006*; *Cline and Haas, 2008*; *Riccomagno and Kolodkin, 2015*; *Cline, 2016*). However, elaborate dendrites with morphologically defined synapses can also emerge without any synaptic transmission (*Verhage et al., 2000*; *Cijsouw et al., 2014*), suggesting that synaptic activity influences dendritic growth but is not the only driving force. Despite the significant interest in the synaptrophic hypothesis, we still lack a mechanistic understanding of how activity-dependent and -independent factors combine to shape development.

To investigate interactions between known signaling factors and to synthesize different experimental results, computational models of dendrite development provide a fruitful direction to ex-

plore how different mechanisms can generate realistic dendritic morphologies (*Cuntz, 2016*). Previous approaches include modeling dendritic development with random branching (*Kliemann, 1987*) or as a reaction-diffusion system (*Luczak, 2006*), implementing activity-independent growth cones that sense molecular gradients (*van Veen and van Pelt, 1992*; *Torben-Nielsen and De Schutter, 2014*), or constructing dendrites as the solution to an optimal wiring problem (*Cuntz et al., 2010*). While these approaches can generate dendrites that accurately match the statistics of developing and mature biological dendrites (*Koene et al., 2009*; *Cuntz, 2016*), they provide limited insight into how reciprocal interactions between dendrite growth, synapse formation, and spontaneous activity can guide dendrite development which is crucial to our understanding of the link between morphological variability and electrophysiological (*Gouwens et al., 2020*; *Scala et al., 2020*) or functional (*Poirazi and Mel, 2001*; *Poirazi et al., 2003*; *Park et al., 2019*; *Poirazi and Papoutsi, 2020*) properties.

Here, we propose a new computational model for cortical dendritic development based on reciprocal interactions between dendrite growth, synapse formation, and spontaneous activity. Starting from neuronal somata distributed in a flat sheet of cortex, we grow dendrites by elongating and branching the dendrite in the direction of distributed potential synapses. Upon contact, synaptic connections form and stabilize or disappear according to a local synaptic learning rule inspired by neurotrophin interactions (*Kirchner and Gjorgjieva, 2021*). Consistent with the synaptotrophic hypothesis, the stability of a dendritic branch depends on the stability of its synaptic contacts, with the branch likely retracting after substantial synaptic pruning. The resulting dynamic system naturally leads to the emergence of three distinct phases of dendrite development: 1) an initial overshoot phase characterized by dendrite growth and synapse formation, 2) a pruning phase during which the learning rule prunes poorly synchronized synapses, and 3) a stabilization phase during which morphologically stable dendrites emerge from the balanced reduction of growth and retraction. Varying model parameters in biologically realistic ranges produces dendrite length and synapse density changes consistent with experiments. At the same time, our mechanism leads to the emergence of functional organization and input selectivity as well as approximately optimal wiring. In summary, our model demonstrates that the formation and stabilization of synapses on developing dendrites can explain a range of experimental observations in one unifying framework.

Results

We introduce a computational model of activity-dependent dendrite growth during development based on synapse formation, stabilization, and elimination. We focused on basal dendrites of cortical pyramidal neurons, which primarily extend laterally within a layer of the cortex (*Larkman and Mason, 1990*) and receive numerous feedforward and recurrent inputs (*Rossi et al., 2019*; *Iacaruso et al., 2017*). To investigate the impact of synapse formation on dendrite development, we modeled several neuronal somata and potential synapses in a flat sheet of cortex (*Figure 1*a). Potential synapses represent locations in the cortex where an axon can form a synapse with a nearby dendrite (*Stepanyants and Chklovskii, 2005*). The model consists of two components: An activity-independent component that directly governs branch growth and retraction; and an activity-dependent component that governs synaptic organization and thus indirectly affects branch stability. Inspired

Figure 1. A model of dendritic growth for a cortical pyramidal neuron driven by activity-independent and -dependent mechanisms. (**a**) Schematic of the soma of a pyramidal neuron (orange triangle) with 12 randomly distributed potential synapses from presynaptic axons (circles) with correlated activity patterns indicated by color. (**b**) Schematic of activity-independent and -dependent mechanisms. Soma and synapses correspond to box in **a**. Signaling molecules diffusing from potential synapses (1) attract dendrite growth and promote synapse formation (2) independent of firing pattern (3). Over time, poorly synchronized synapses depress and are pruned from the dendrite (4), while well-synchronized synapses remain stable (5). After a branch retracts, the dendrite is less sensitive to the growth field at that location(5). (**c**) Change in weight of one synapse (green) following the stimulation of itself (green bolt) and of two nearby synapses (purple bolts). Left: Schematic of the developing dendrite from **b** with bolts indicating synaptic activation. Right: Presynaptic accumulator (top), postsynaptic accumulator (middle), and change in synaptic weight (bottom) as a function of time. Dashed line (bottom) indicates zero change.

Figure 1–Figure supplement 1. The growth field is similar to two-dimensional heat diffusion.

Figure 1–Figure supplement 2. Detailed illustration of the dendritic growth mechanism.

by the synaptotrophic hypothesis (*Vaughn, 1989*), we mimic the effect of activity-independent molec-
ular signaling by letting each potential synapse release diffusive signaling molecules that attract
the growing dendrite (*Figure 1*b, *Figure 1–Figure Supplement 1*, *Figure 1–Figure Supplement 2*). In
addition, during development, neural circuits generate patterned spontaneous activity (*Blanken-*
ship and Feller, 2010; *Ackman and Crair, 2014*). Therefore, to model the correlated spontaneous
activity exhibited by different axons (*Scholl et al., 2017*; *Iacaruso et al., 2017*), we divided poten-
tial synapses randomly into activity groups that receive synchronized input (see Methods). Each
group represents either synapses from the same presynaptic neuron or neurons that experience
correlated presynaptic activity.

Because of their attraction to growth-factor releasing synapses and independent of neural ac-
tivity, dendrites in our model grow outward from the soma towards the nearest potential synapse,
where they form a synapse and temporarily stabilize (*Figure 1*b, *Figure 1–Figure Supplement 2*).
We assumed that dendrites could not overlap based on experimental data (*Grueber and Sagasti,*
2010); therefore, dendrites in the model retract, for instance, when further growth would require

self-overlap. Once a synapse is formed, we modeled that its strength changes according to a local, activity-dependent plasticity rule (*Kirchner and Gjorgjieva, 2021*) (*Figure 1*c). The learning rule induces synaptic potentiation whenever presynaptic and local postsnynaptic activity co-occur, and synaptic depression whenever local postsynaptic activity occurs in the dendrite independent of presynaptic stimulation (see Methods),

$$\Delta\text{weight} = \text{post} \times (\text{pre} - \text{offset}). \tag{1}$$

As shown previously, this rule generates synaptic distance-dependent competition, where nearby synapses affect each other more than distant synapses, and correlation-dependent cooperation, where synchronized synapses stabilize. In contrast, desynchronized synapses depress (*Kirchner and Gjorgjieva, 2021*). In our model, we assumed that when a synapse depresses below a threshold, it decouples from the dendrite, and the corresponding branch retracts successively to either the nearest stable synapse, branch point, or the soma (*Figure 1*b, *Figure 1–Figure Supplement 2*). After removal, the vacated synapse turns into a potential synapse again, attracting other growing branches. Thus, a developing dendrite in our model acquires its arborization through the attraction to signaling molecules released by potential synapses and the repeated activity-dependent formation and removal of synapses.

Dendrite development through balancing growth and retraction

After specifying the rules governing the growth of individual dendritic branches, we investigated dendritic development on long timescales. When growing dendrites according to our proposed growth rule in conjunction with signaling molecules attraction and spontaneous activity refinements (*Figure 1*), we found that dendrites form several stems, i.e. branches which start directly at the soma, and rapidly expand outwards (*Figure 2*a). After an initial phase of rapid expansion, we observed that growth rapidly attenuates, and the dendritic length stabilizes (*Figure 2*b). This stability is achieved when the dendrite's expansion and retraction are balanced (*Figure 2*c). To investigate whether the stability in total length also corresponds to stability in dendrite morphology, we quantified morphological stability as the pixel-wise correlation of a dendrite with itself 4.5 hours earlier, which is several orders of magnitude larger than the speed at which dendrites grow and retract in our model (see Table 1). Despite the residual amount of expansion and retraction, we showed that morphological stability increases rapidly, and the dendrite morphology is already stable after the initial expansion phase (*Figure 2*d). Interestingly, rapid stabilization of morphology also occurs in the mouse visual cortex (*Richards et al., 2020*) and the *Drosophila* larvae (*Castro et al., 2020*). The rapid slowing down of growth in our model is not due to a depletion of potential synaptic partners as the number of potential synapses remains high (*Figure 2*e). In summary, by combining activity-independent and -dependent dendrite growth mechanisms, our model produces dendrites that rapidly expand and stabilize by balancing growth and retraction.

Figure 2. Balanced growth and retraction generate morphologically stable dendrites. (**a**) Three example dendrites at five time points from our simulations. For clarity of presentation, connected synapses are not displayed. (**b**) Total length of dendritic tree as a function of time. (**c**) Length of dendrite added (green) and removed (red) as a function of time. (**d**) Morphological stability (correlation between the dendrite shape at time t and $t - 4.5$ hours) as a function of time. (**e**) Number of available potential synapses as a function of time. All lines represent averages across 32 simulations with nine dendrites each. Shaded area indicates two standard deviations.

Figure 2–video 1. Example of a simulation in which several dendrites develop in parallel.

Figure 3. Synapse formation and removal predominate in distinct phases of dendrite development. (**a**) Three examples of dendrites at the beginning ($t = 9$ hours) and end ($t = 72$ hours) of the simulation. Green circles indicate formed synapses. (**b**) Total number of connected synapses as a function of time. Orange arrow highlights overshoot and subsequent pruning. (**c**) Added (green) and pruned synapses (red) as a function of time. (**d**) Average synaptic weights of synapses that ultimately stabilize (solid black; final weight more than 0.5) or are removed (dashed black; final weight less than 0.5) as a function of time. All lines represent averages across 32 simulations with nine dendrites each. Shaded area indicates two standard deviations.

Delayed activity-dependent plasticity produces a rapid increase of synapse density followed by pruning

Since our model couples dendritic growth to the formation and removal of synapses (*Figure 3*a), we next investigated how the number of connected synapses, which are necessary for the dendrite's stabilization, changes over time. As a result of the dendrite's rapid growth, we observed a rapid increase in the number of connected synapses (*Figure 3*b,c). In contrast to the dendrite's length, we found that the initial rapid increase in connected synapses is followed by a brief period of an overall reduction of the number of synapses before additions and removals are balanced (*Figure 3*c). This removal of established synapses resembles the postnatal removal of synapses observed in the mouse neocortex (*Holtmaat et al., 2005*). To understand how the initial overshoot and subsequent removal of synapses emerge in our model, we computed the average synaptic weight of all synapses that eventually stabilize or are pruned (*Figure 3*d). We found that the delayed onset of synapse removal (*Figure 3*c) is due to the slow time scale of the synaptic weight compared to the faster time scale of dendrite growth. Thus, the initial overshoot and subsequent removal of synapses observed in our model (*Figure 3*b) is due to the rapid formation relative to the delayed activity-dependent elimination of synapses.

Activity-dependent competition between synapses produces input selectivity and synaptic organization

Next, we asked if the stabilization of dendrites might be supported by the emergence of organization of connected synapses. First, we compared the synapses connected to dendrites at the apex of the overshoot phase (peak in *Figure 3*b) with those in the stabilization phase (*Figure 4*a,b). While dendrites at the apex do not prefer synapses from any particular input group, in the stabilization phase, they acquire a preference for synapses from only two or three of the activity groups (*Figure 1*b). These dynamics resemble the activity-dependent synaptic competition in the developing visual cortex, where asynchronously activated synapses tend to depress (*Winnubst et al., 2015*). Notably, the remaining synchronized synapses in our model experience correlation-dependent cooperation (*Kirchner and Gjorgjieva, 2021*), and are therefore able to stabilize the dendrite and prevent the total retraction of all branches.

This selective potentiation of synapses according to input correlation also leads to dendritic selectivity for inputs. In particular, synapses on the same dendrite are likely to stem from the same activity group (*Figure 4*c). This selectivity is acquired throughout the simulation, where selectivity starts high (a nascent dendrite is trivially selective to its first synapse; $t = 0 - 1$ hours), drops almost to chance level (indiscriminate addition of synapses; $t = 9$ hours), and finally approaches a value of $\frac{1}{2}$ (two activity groups per dendrite remain after the pruning phase; $t = 72$ hours) (*Figure 4*d). To determine which activity group stabilizes eventually, we computed selectivity for each group early ($t = 9$ hours) and late ($t = 72$ hours). We found that early high (low) selectivity for an activity group translates into even higher (lower) selectivity in the stable state (*Figure 4*e), predicting an outsized impact of early synaptic contacts on continued dendrite function. Finally, we observed that when dendritic trees overlap strongly, they tend to be selective to different activity groups (*Figure 4*f) due

Figure 4. Stable morphology is obtained through selective removal of synapses and dendritic input selectivity. (**a,b**) Dendritic trees before (**a**, 9 hours) and after (**b**, 72 hours) removal of synapses (*Figure 3*). Connected synapses colored corresponding to activity group, which represents activity correlations (*Figure 1*b). (**c**) Left: Schematic illustrating the difference between Euclidean and tree distance. Note that we compute the Euclidean distance between synapses from different trees. Right: Correlation between pairs of synapses as a function of the Euclidean distance (blue) and tree distance (red). (**d**) Input selectivity of dendrites (defined as the fraction of the activity group with the highest representation) as a function of time. Dashed line indicates chance level. All lines represent averages across 32 simulations with nine dendrites each. Shaded area indicates two standard deviations. (**e**) Fraction of connected synapses per activity group early ($t = 9$ hours) and late ($t = 72$ hours) in the simulation. Each dot represents one of the five activity groups on one of the nine dendrites from the 32 simulations, resulting in $5 \times 9 \times 32 = 1440$ data points. (**f**) Left: Schematic of different levels of overlap (rows) between the convex hulls of two dendrites, referring to the smallest convex sets that contain the dendrite. Right: Signal correlation (correlation between fractions of synapses from the same activity groups) for different levels of dendritic overlap. Error bars indicate the standard error of the mean, computed from 1152 pairs of dendrites from 32 simulations.

₂₀₁ to competition for limited potential synapses of any given group. Interestingly, also in the mouse
₂₀₂ visual cortex, neighboring neurons often exhibit different selectivity (*Ohki et al., 2005*), potentially
₂₀₃ reflecting a lasting impact of early competition between different inputs.

₂₀₄   In summary, the emergence of dendrites' selectivity for synapses from specific activity groups
₂₀₅ coincides with and supports the stabilization of dendrites' morphologies.

balance of mature and immature brain-derived neurotrophic factor controls arborization of dendrites

₂₀₈ After establishing that our model can capture some important aspects of the dynamics of den-
₂₀₉ dritic development through the combination of activity-independent and activity-dependent mech-
₂₁₀ anisms, including local plasticity, we asked how changing properties of the plasticity rule might af-
₂₁₁ fect dendritic growth and synaptic organization. Developmentally, the interaction between two
₂₁₂ neurotrophic factors, BDNF and proBDNF (*Figure 5*a), has been found to play a key role in or-
₂₁₃ ganization of synaptic inputs into clusters (*Niculescu et al., 2018*). Therefore, through the previ-
₂₁₄ ously established link between this neurotrophin interaction and synaptic plasticity (*Kirchner and
₂₁₅ Gjorgjieva, 2021*), we investigated the influence of changing the underlying molecular interactions
₂₁₆ on dendritic morphology.

₂₁₇   As we have previously shown, the "offset" term in our plasticity rule (*Equation 1*) represents
₂₁₈ the neurotrophin balance (computed as BDNF/(BDNF+proBDNF)) released upon stimulation of a
₂₁₉ synapse (*Kirchner and Gjorgjieva, 2021*). Consequently, we found that an overabundance of BDNF
₂₂₀ (proBDNF) leads to potentiation (depression) of the synapse (*Figure 5*b), consistent with experi-
₂₂₁ mental data (*Lu et al., 2005*). Furthermore, our plasticity rule acts locally on the dendrite so that
₂₂₂ the strength of individual synapses is affected by interactions with other nearby synapses. Con-
₂₂₃ cretely, a lower (higher) density of nearby synapses tends to lead to potentiation (depression) of
₂₂₄ the synapse (*Kirchner and Gjorgjieva, 2021*).

₂₂₅   To better understand the interactions between the balance of neurotrophins and the density
₂₂₆ of synapses, we analytically derived the maximum density of synapses that can stabilize given a
₂₂₇ balance of neurotrophins (*Figure 5*c, see Methods). We found that an overabundance of BDNF
₂₂₈ (proBDNF) leads to a higher (lower) maximal density of synapses (*Figure 5*c). Indeed, when we
₂₂₉ simulated dendritic development with varying neurotrophin ratios, we found that the density of
₂₃₀ synapses per dendrite increases with increasing neurotrophin ratio (*Figure 5*d,e). Consistent with
₂₃₁ biological evidence (*McAllister et al., 1995*; *Tyler and Pozzo-Miller, 2001*), in our model, developing
₂₃₂ dendrites treated with BDNF tend to grow larger and have a higher density of synapses (*Figure 5*e,g).
₂₃₃ In contrast, over-expression of proBDNF leads to smaller dendrites with fewer synapses (*Koshimizu
₂₃₄ et al., 2009*; *Yang et al., 2014*) (*Figure 5*f,g).

₂₃₅   In our model, these changes in length and density are explained by a change in selectivity of
₂₃₆ synapses (*Figure 5*h). Concretely, an increase in BDNF erases all synaptic competition, reducing
₂₃₇ the selectivity to chance level, while an increase in proBDNF greatly amplifies synaptic competition
₂₃₈ and thus selectivity. These differences in competition determine the number of pruned synapses
₂₃₉ and thus the length at which the dendrite stabilizes. Thus, our model predicts that biologically-
₂₄₀ relevant differences in dendritic morphology may arise from different neurotrophin ratios due to

Figure 5. Dendritic arborization is controlled by the ratio of neurotrophic factors. (**a**) Interactions between molecular factors underlie a local activity-dependent plasticity rule for synaptic change (*Equation 1*, (*Kirchner and Gjorgjieva, 2021*)). Neurotrophins (BDNF and proBDNF) bind to different neurotrophin receptors, and a cleaving protease (MMP9) converts proBDNF to BDNF in an activity-dependent manner. (**b**) Schematic illustrating the impact of different concentrations of BDNF on synaptic change. Upon stimulation of a synapse (top), proBDNF and BDNF is released into extracellular space (middle), where proBDNF can be cleaved into BDNF by MMP9. Depending on the neurotrophin ratio, computed as BDNF/(BDNF + proBDNF), the synapse is stabilized (left) or depressed and hence eventually removed (right). (**c**) Maximal possible stable density of synapses as a function of the initial concentration of BDNF. Stable (no pruning; green) and unstable (pruning occurs; red) areas are indicates. (**d**) Three examples of dendrites with superimposed synapses (green) with high initial BDNF concentration (49%), the baseline concentration (45%, same as *Figure 1-Figure 3*) and low initial BDNF (40%). Symbols correspond to locations marked in panel c. (**e-f**) Averages for density of synapses on the dendrite (**e**), number of connected synapses (**f**), total length of dendrite (**g**) and global selectivity (**h**) as a function of time for dendrites from the three conditions shown in **d**. All lines represent averages across 32 simulations with nine dendrites each.

²⁴¹ the maximal density of synapses that stabilizes the dendrite.

Different impacts of activity-dependent and -independent factors on dendritic development

²⁴⁴ We next aimed to understand the different roles of activity-dependent and -independent mech-
²⁴⁵ anisms on dendritic morphology. To this end, we varied either only activity-dependent factors
²⁴⁶ (firing patterns of potential synapses) or only activity-independent factors (stochasticity in den-
²⁴⁷ drite branching and growth) across a set of simulations (*Figure 6*a,b). Consistent with experiments
²⁴⁸ (*Scala et al., 2020*), dendrites produced by our model exhibit substantial variability in morphology
²⁴⁹ (*Figure 6*a), length (*Figure 6*c), and number of synapses (*Figure 6*d). Comparing dendrites that expe-
²⁵⁰ rienced either identical activity-dependent or -independent factors allowed us to compute the per-
²⁵¹ centage of change in morphology attributable to each factor as a function of developmental time
²⁵² (*Figure 6*e,f). We found that while activity-independent factors tend to lead to large differences
²⁵³ in morphology early on, activity-dependent factors affect dendrite morphology with a substantial
²⁵⁴ delay. These differences can be explained by the delay in synaptic pruning (*Figure 3*d).

²⁵⁵ Despite substantial variability, there are predictive factors for the final length of the dendrite. In
²⁵⁶ particular, we found a positive relationship between the number of major branches, i.e. branches
²⁵⁷ starting from the soma, and the final length (*Figure 6*g). Interestingly, this is consistent with re-
²⁵⁸ constructed dendrites from multiple regions of the mouse cortex (*Figure 6–Figure Supplement 1*).
²⁵⁹ Furthermore, our model predicts that dendrites that have a high (low) total length early on will, on
²⁶⁰ average, retain a (high) low total length throughout development (*Figure 6*e).

²⁶¹ Thus, our model suggests that while activity-independent factors affect dendritic morphology
²⁶² early on during development, activity-dependent factors dominate later. Properties like the num-
²⁶³ ber of major branches or the length of dendrites during early development might be predictive of
²⁶⁴ the dendrite's morphology throughout the animal's lifetime.

Coupled dendrite growth and synapse formation leads to approximately optimal wiring

²⁶⁷ Since space is limited in the cortex and maintaining complex morphologies is costly, it is beneficial
²⁶⁸ for a neuron to connect to its synapses with the minimum possible dendrite length (*Cuntz et al.,*
²⁶⁹ *2012*) (*Figure 7*a). In our model, dendrites are assumed to grow towards the nearest potential
²⁷⁰ synapse. Thus, we investigated how the final length in our model compares to the optimal wiring
²⁷¹ length. The optimal length (L) of dendrites in a plane scales with the square root of the number
²⁷² of synapses (N) times the area over which the synapses are distributed (A), divided by a scalar:
²⁷³ $L = \sqrt{N\frac{A}{\pi}}$ (*Cuntz et al., 2012*). In contrast, the length of a dendrite when synapses are connected
²⁷⁴ randomly scales with the number of connected synapses times the average distance of two random
²⁷⁵ points on a circle (*Uspensky, 1937*), $L = N\frac{128}{45\pi}\sqrt{\frac{A}{2\pi}}$ which differs from the optimal result by a square
²⁷⁶ root in the number of synapses. Using the convex hull that circumscribes the stabilized synapses
²⁷⁷ as the area over which the synapses are distributed (*Figure 7*b), we compared the actual dendrite
²⁷⁸ length with the optimal and the random wiring length (*Figure 7*c). We found that our simulations'
²⁷⁹ dendritic lengths are shorter than random wiring and longer than the theoretical optimal length.

Figure 6. Morphological variability emerges from the interaction of activity-dependent and -independent factors. (**a**) Example of three dendrites with identical initial conditions but different random seeds. The colors illustrate that initial growth is governed by activity-independent factors, while later growth is governed by activity-dependent factors. (**b**) Schematic illustrating how variability is introduced into model: activity-dependent via the patterns of spontaneous activity (orange), and activity-independent via fluctuations in both the extrinsic growth stimulating field (purple 1) and the intrinsic mechanisms underlying dendrite growth (purple 2; see Methods). (**c,d**) Total length (**c**) and number of synapses (**d**) as a function of time for dendrites with identical initial conditions. Each gray line corresponds to one dendrite from one of 32 simulations. Bold line represents average. (**e,f**) Percentage in change of morphological similarity (**e**) and similarity of connected synapses (**f**) as a function of time for simulations where activity-dependent (orange) or -independent (purple) factors vary. Lines represent averages across 32 simulations with nine dendrites each. Shaded area indicates two standard deviations. (**g,h**) Final length as a function of number of major branches (**g**) and maximal length in the first 18 hours of the simulation (**h**). Lines indicate linear regression.

Figure 6–Figure supplement 1. Total tree length increases with the number of stems.

Figure 7. Dendritic morphology approximately minimizes cable length. (**a**) Schematic illustrating optimal (top) and random (bottom) wiring to connect a given set of synapses. (**b**) The convex hull of all synapses connected to a dendrite with the proportionality of optimal length (bottom). (**c**) Total tree length as a function of convex hull area in the optimal, simulated and random scenario. Each dot corresponds to one of 288 dendrites from 32 simulations. Lines correspond to analytic predictions for the average density across all simulations. (**d**) Total tree length against the number of branch points in log scale, both for data and theoretical optimum. (data extracted from (*Cuntz et al., 2012*)). (**e**) Total tree length in the data (black, average of n=13112), our simulations (blue, average of 288 dendrites from 32 simulations), and the random baseline (green, analytically computed) relative to theoretical optimum (pink, analytically computed).

²⁸⁰   We were interested to see if the deviation from optimality might quantitatively match the one

²⁸¹ observed in real dendrites. To investigate this question, we reanalyzed a published dataset (*Cuntz*

²⁸² *et al., 2012*) containing the total lengths and the number of branch points of 13,112 dendrites

²⁸³ pooled across 74 sources. When computing the fold change between the real and the optimal

²⁸⁴ dendritic length in the dataset, we confirmed that real dendrites tend to be consistently larger

²⁸⁵ than the theoretical optimum (*Figure 7*d). Interestingly, the fold change between the length of real

²⁸⁶ dendrites and the theoretical optimum is similar to the fold change of our simulated dendrites and

²⁸⁷ the theoretical optimum (*Figure 7*e). This deviation is expected given the heterogeneous structure

²⁸⁸ of neuronal tissue that hampers diffusion of signaling molecules (*Nicholson et al., 2000*; *Motta*

²⁸⁹ *et al., 2019*), which mirrors the fluctuations in activity-independent factors in our model. There-

²⁹⁰ fore, activity-dependent dendrite growth produces morphologies with a total length close to the

²⁹¹ theoretically possible minimum.

Discussion

²⁹³ Dendrite growth, synapse formation, and spontaneous activity influence each other during early

²⁹⁴ development, but the nature of these interactions and the implications for dendritic function through-

²⁹⁵ out life remain largely unexplored. In this study, we proposed a model of dendritic growth dur-

²⁹⁶ ing development based on the formation and removal of synapses as well as growth and retrac-

²⁹⁷ tion of dendritic branches. In this model, three distinct phases of synapse development – over-

²⁹⁸ shoot, pruning, stabilization – and stable dendritic trees emerge naturally from the interactions be-

²⁹⁹ tween activity-independent release of signaling molecules from potential synapses and the activity-

³⁰⁰ dependent synaptic weight changes. The stable morphology of dendrites in our model is enabled

³⁰¹ by input selectivity, which implies local organization of connected synapses on the dendrite. Our

³⁰² model allows us to vary activity-independent and -dependent influences on dendrite morphology

³⁰³ individually and to predict the impact of early perturbations on the mature morphology. Further,

304 we provide a mechanistic explanation for how mature dendrite length might approximate optimal
305 wiring length since dendritic growth is guided towards potential synapses. Thus, our model pro-
306 vides a new perspective on activity-dependent dendrite growth and suggests that the formation
307 and removal of synapses might exert powerful control over the growth process.

Comparison with the synaptotrophic hypothesis

The synaptotrophic hypothesis, originally proposed three decades ago (*Vaughn, 1989*), has pro-
vided a useful framework for interpreting the effect of neural activity and synapse formation on
dendrite development. Our proposed model is inspired by the synaptotrophic hypothesis but dif-
fers from it in a few key aspects. (1) The synaptotrophic hypothesis postulates that synaptic activity
is necessary for dendrite development (*Cline and Haas, 2008*). In contrast, our model contains an
activity-independent component that allows dendrites to grow even in the absence of synaptic
activity. Our model is thus consistent with the finding that animals lacking all neurotransmitter
secretion still form connected circuits of neurons with morphologically defined synapses (*Verhage
et al., 2000*) and computational models that produce dendrites with many relevant morphological
properties without relying on activity (*Cuntz, 2016*). (2) The synaptotrophic hypothesis does not
specify the exact molecular actors that underlie the information exchange between pre- and post-
synapse. Informed by recent experiments that identify central molecular candidates (*Winnubst
et al., 2015*; *Kleindienst et al., 2011*; *Niculescu et al., 2018*; *Lu et al., 2005*), our model proposes
a concrete mechanistic implementation based on neurotrophic factors (*Kirchner and Gjorgjieva,
2021*). (3) The synaptotrophic hypothesis postulates that pre- and postsynapse can rapidly evalu-
ate whether a potential contact is appropriate. Consistent with experiments (*Lohmann et al., 2002*;
Niell et al., 2004), the fate of a synapse in our model is determined only within tens of minutes or
hours after it is formed. This is due to the slow timescale of synaptic plasticity (*Figure 3*d).

Relationship between dendritic form and function

While previous studies focused on how dendritic morphology affects function, e.g. through non-
linear signal transformation (*Poirazi and Papoutsi, 2020*) or dynamic routing of signals (*Payeur
et al., 2019*), we propose that dendrite form and function reciprocally shape each other during de-
velopment. While the dendrite's morphology constrains the pool of available potential synapses,
synaptic activity determines the dendritic branch's stability (Fig. 1). As a result, the dendritic tree
self-organizes into a well-suited shape to support a limited number of functionally related synapses.
These initial synaptic contacts might then serve as a scaffold around which additional, functionally
related synapses cluster to form the building blocks to support the powerful computations of ma-
ture dendrites (*Kirchner and Gjorgjieva, 2021*).

Dynamics of dendritic development

338 Here we focus on the early developmental period of highly dynamic growth and retraction. How-
339 ever, dendritic morphology remains remarkably stable in later development and throughout adult-
340 hood (*Richards et al., 2020*; *Castro et al., 2020*; *Koleske, 2013*). This stability is achieved despite
341 substantial increases in overall size of the animal (*Richards et al., 2020*; *Castro et al., 2020*) and

ongoing functional and structural plasticity of synapses (*Kleindienst et al., 2011*; *Winnubst et al., 2015*; *Kirchner and Gjorgjieva, 2021*). While it is still unclear how exactly synaptic organization is established during early development and how synapses are affected by the overall increase in dendrite size, somatic voltage responses to synaptic activity are largely independent of dendrite size (*Cuntz et al., 2021*). It has been shown that dendrite stability plays a crucial role in enabling the correct function of the adult brain and is disrupted in many psychiatric disorders and neurodegenerative diseases. In particular, the release of BDNF, which is connected to synaptic activity, affects structural consolidation of dendrites and, thus, long-term stability (*Koleske, 2013*). With our mechanistic model, we can perturb the balance of neurotrophic factors and observe detrimental effects on dendrite stability as a result of extreme or non-existent input selectivity, providing insight into functional consequences of disrupted dendrite growth in neurodevelopmental disorders (*Johnston et al., 2016*).

Interneurons and inhibitory synapses

In addition to the excitatory neurons and synapses that are the focus of this study, there are also inhibitory interneurons and inhibitory synapses that play an important role in brain development (*Naskar et al., 2019*). Interneurons fall into genetically-distinct subtypes, which tend to target different portions of pyramidal neurons (*Rudy et al., 2011*; *Kepecs and Fishell, 2014*). In particular, somatostatin-expressing (SST) interneurons preferentially target the dendrites of pyramidal neurons, while parvalbumin-expressing (PV) interneurons preferentially target the soma. Furthermore, the dendrites of inhibitory neurons have a complex morphology that likely allows them to perform intricate transformations of incoming signals (*Tzilivaki et al., 2019*, *2021*). Investigating whether and how inhibitory interneurons and synapses might interact with excitatory ones during dendritic development is an exciting direction for future research.

In summary, by proposing a mechanistic model of dendritic development which combines activity-independent and dependent components, our study explains several experimental findings and makes predictions about the factors underlying variable dendritic morphologies and synaptic organization. Interestingly, the stable morphologies it generates are approximately optimal in terms of wiring length and experimental data. Finally, our model provides the basis for future exploration of different learning rules and cell types which could differ across brain regions, species and healthy vs./ disease states.

Methods and Materials

Activity-independent synaptic signals. In the synaptotrophic hypothesis, dendrite growth is directed towards potential synaptic partners. In our model, we capture this aspect by introducing a *growth field* of activity-independent synaptic signals, $T(\mathbf{p})$, over all positions $\mathbf{p}$ in our sheet of cortex. This field contains point sources at the positions of potential synapses, $\mathbf{p}_i$, and evolves over time according according to a diffusion equation,

$$T(\mathbf{p})^{t+1} = T(\mathbf{p})^t * D + \mu \sum_i \mathbf{p}_i + \sigma \mathbf{N}. \tag{2}$$

The growth field at time point $t + 1$ is therefore given by the sum of the growth field at time t convolved with a diffusion filter D, a constant input of size μ from all potential synapses, which are currently not connected to a dendrite, as well as independent Gaussian noise, $\mathbf{N}$, with standard deviation σ. We chose a two dimensional Gaussian for the diffusion filter D, making the field $T(\mathbf{p})$ mathematically equivalent to a heat diffusion in two dimensions (*Figure 1–Figure Supplement 1*).

Asynchronous dendrite growth and retraction. Dendrite development critically depends on resources from the soma (*Ye et al., 2007*). Consequently, we modeled the growth of dendrites to depend on *scouting agents* that spread outward from the soma at regular time intervals, t_{scout}, and that traverse the dendritic tree at speed v_{scout} (*Figure 1–Figure Supplement 2*). These scouting agents resemble actin-blobs that control dendrite growth (*Nithianandam and Chien, 2018*). When a scouting agent encounters a branch point, there is a 0.5 chance for it to continue in any direction. This means it can go in one direction, but it can also duplicate or disappear completely. We further model these scouting agents to detect the growth field's value – a scouting agent stops at a position on the dendrite where this field is locally maximal and marks this location for growth. The dendrite will then expand at the marked positions in the direction of the gradient of the growth field, and the scouting agent moves to this new position. If the dendrite grows to the location of a potential synapse, this synapse is then realized, and its initial weight is set to $w_{init} = \frac{1}{2}$. Two branches of the same neuron may never become adjacent; however, branches from other neurons may be crossed freely. If a scouting agent reaches the end of a branch without finding a local maximum of the growth field along its path, the scouting agent initiates the retraction of this branch. Depending on proximity, a branch then retracts up to the nearest stable synapse, the next branch point, or the soma. Because our simulations are a spatially discrete approximation of a biological flat sheet of cortex, we had to ensure that growth behaves appropriately in cases where the discretization scheme becomes relevant (*Figure 1–Figure Supplement 2*).

Minimal plasticity model. When a synapse k forms on the dendrite, its weight w_k evolves according to a previously proposed minimal plasticity model for interactions between synapses on developing dendrites (*Kirchner and Gjorgjieva, 2021*). This model can be linked to a full neurotrophin model that interprets the parameters in terms of the neurotrophic factors BDNF, proBDNF, and the protease MMP9. In this model, the k-th synapse is stimulated within input event trains x_k

$$x_k(t) = \int_0^\infty \sum_f \delta(s - s_k^f)(H(t - s) - H(t - x_{\text{dur}} - s))\mathrm{d}s \tag{3}$$

408 with events at times t_k^f and where the Heaviside step function $H(t)$ is 0 when t is less than 0 and

409 1 when t is greater or equal than 0, so that events have duration x_{dur} (50 time steps). The minimal

410 plasticity model consists of a synapse-specific presynaptic accumulator v_k,

$$\tau_v \frac{\mathrm{d}v_k}{\mathrm{d}t} = -v_k(t) + \phi x_k(t), \tag{4}$$

411 and a postsynaptic accumulator u_k that averages over nearby synapses in a weighted and distance-

412 dependent manner,

$$\tau_u \frac{\mathrm{d}u_k}{\mathrm{d}t} = -u_k(t) + \sum_{l=1}^{N} s_{kl} w_l(t) x_l(t). \tag{5}$$

413 The multiplicative factor ϕ is an MMP9 efficiency constant that determines how efficiently MMP9

414 converts proBDNF into BDNF per unit of time and the proximity variables s_{kl} between synapses k

415 and l on the same dendrite are computed as $s_{kl} = e^{-\frac{d_{kl}^2}{2\sigma_s^2}}$, where σ_s determines the spatial postsynap-

416 tic calcium spread constant. The equation governing the weight development of w_k (***Equation 6***) is a

417 Hebbian equation that directly combines the pre- and postsynaptic accumulator with an additional

418 offset constant ρ,

$$\tau_w \dot{w}_k = u_k(t)(v_k(t) + \rho), \tag{6}$$

419 with $\rho = \frac{2\eta-1}{2(1-\eta)}$ and $\tau_w = \tau_W \frac{1}{2(1-\eta)}$. Here, η is the constitutive ratio of BDNF to proBDNF and

420 $\tau_W = 3000\,\text{ms}$ This model is minimal in the sense that it cannot be reduced further without losing

421 either the dependence on correlation through the link to the BTDP rule, or the dependence on

422 distance.

423 To model structural plasticity, we implemented a structural plasticity rule inspired by ref. (***Holt-***

424 ***maat and Svoboda, 2009***) where each synapse whose efficacy falls below a fixed threshold W_{thr} is

425 pruned from the dendrite.

426 **Simulations and parameters.** For all simulations in this report, we distributed nine somata at

427 regular distances in a grid formation. We used 1500 potential synapses and divided them into five

428 groups of equal size, with each group receiving Poisson input with rate r_{in}. Therefore, all synapses

429 in the same group are perfectly correlated, while synapses in different groups are uncorrelated.

430 ## Figures and Tables

431 ## Acknowledgments

432 ## Competing interests

433 No competing interests declared.

434 ## References

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Table 1. Parameters of the minimal plasticity model (*Kirchner and Gjorgjieva, 2021*) and the synaptotrophic growth model.

Parameter	Variable	Value
Synaptic efficacy time constant	τ_W	6000 time steps
Postsynaptic accumulator time constant	τ_u	300 time steps
Presynaptic accumulator time constant	τ_v	600 time steps
Constitutive percent of BDNF of total neurotrophins	η	45%
MMP9 efficiency constant	ϕ	$\frac{3}{50}$ per time step
Heterosynaptic offset	ρ	$\rho = \frac{2\eta-1}{2(1-\eta)}$
Minimal model synaptic efficacy time constant	τ_w	$\tau_w = \tau_W \frac{1}{2(1-\eta)}$
Standard deviation of calcium spread	σ_c	$200\,\mu m$
Turnover threshold below which a synapse is replaced	W_{thr}	0.02
Firing rate of synapses	r_{in}	$0.116\,min^{-1}$
Scout intervals and speed	t_{scout}, v_{scout}	$10\,min, 0.18\,\mu m\,min^{-1}$

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Figure 1–Figure supplement 1. The growth field is similar to two-dimensional heat diffusion. By iteratively convolving the potential synapses in the growth field with a Gaussian filter, the growth field reaches a steady-state that resembles a two-dimensional heat diffusion with point sources. Over time, individual point sources disappear and reappear to mimic when the corresponding synapses are connected or pruned from the dendrite.

Figure 1–Figure supplement 2. Four scenarios of asynchronous dendritic growth can be modeled as a scout agent representing the tip of a dendrite exploring a two-dimensional grid. A scout agent (yellow dot) has reached the location of a potential synapse (blue dot). Scenario 1: The scout agent will extend the dendrite and form a new synapse if nothing else happens. Scenario 2: To prevent overlap, if a second branch from the same dendrite blocks the path to the potential synapse, the original branch retracts. Scenario 3: If a branch from another dendrite reaches the potential synapse first, the original branch retracts. Scenario 4: If a new potential synapse becomes available adjacent to the dendrite (1) so that growth is not possible (since the branch cannot form immediately adjacent to two other parts of the dendrite), the corner flips (2), and the synapse forms.

Figure 6–Figure supplement 1. Total tree length increases with the number of stems. Average (solid black) and individual (gray circles) total tree lengths as a function of the number of stems. $N = 701$. Data from the Allen Cell Types Database (2015).