

REVIEW SUMMARY

NEUROSCIENCE

Architectures of neuronal circuits

Liqun Luo

BACKGROUND: The human brain contains about 100 billion neurons, each of which makes thousands of synaptic connections. Although individual neurons can themselves be sophisticated information-processing units, it is their synaptic connection patterns that enable neurons to form specialized circuits for specific functions, making the brain a powerful computational device. Decades of research using anatomical tracing, physiological recording, functional perturbation, and computational modeling in diverse organisms have detailed the connection patterns of neurons and their functions at scales ranging from microcircuits of a few neurons to global organization of millions of neurons. Here, I synthesize these findings from the perspective of circuit architectures and discuss how these architectures might emerge during development and evolution.

ADVANCE: Suppose individual neurons are to the brain what letters are to an article. We can then consider microcircuit motifs as words and larger-scale architectural plans as sentences. At the level of words and microcircuits, specific patterns of connections between excitatory and inhibitory neurons confer elementary information-processing functions. For example, feedforward excitation allows information to propagate across neural regions, with convergent and divergent excitation to integrate signals from multiple upstream

sources and disseminate signals to diverse downstream targets, respectively. Feedforward inhibition and feedback inhibition regulate the duration and magnitude of incoming excitatory signals and often work together to control gain and dynamic range of input signals as well as facilitate synchronous and oscillatory firing. Lateral inhibition selects information to be propagated to downstream circuits by amplifying differences in activity between parallel pathways. Mutual inhibition can produce rhythmic outputs and regulate brain states. These core circuit motifs are almost always used in concert to build up complex signal-processing units.

The next level of organization, the sentences, is more heterogeneous in scale and configuration. Continuous topographic mapping, in which neighboring input neurons connect to neighboring target neurons through orderly axonal projections, provides a way to organize information at successive stages of processing, minimizes wiring length, and can facilitate extraction of local contrast through lateral inhibition. Discrete parallel processing allows signals to be represented and processed in parallel, reducing computational depth and increasing processing speed. Dimensionality expansion enables output neurons to represent different combinations of inputs, facilitating pattern separation by downstream neurons. Biased input and segregated output divide

neuromodulatory systems with broad projections into subsystems, each serving distinct behavioral functions and being differentially regulated by different stimuli. Recurrent loops are abundant in nervous systems and support rich neural activity dynamics. Architectures of many neuronal circuits at this level are awaiting discovery.

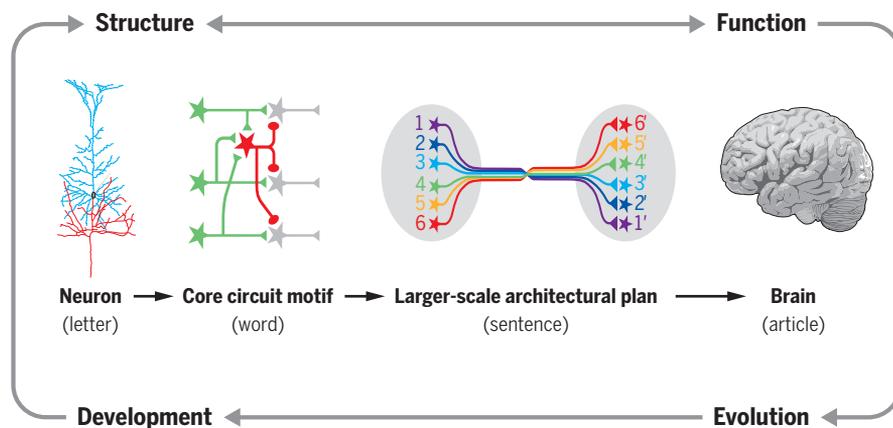
Unlike computer circuits, which are products of top-down design, neuronal circuits are products of hundreds of millions of years of evolution. Some circuit motifs may have originated long ago, having since been conserved across diverse clades and passed on to more recently evolved neural regions. Other architectures have evolved independently in different clades. Duplication and divergence of neuron types and brain regions play important roles in brain evolution and may result in the modularity of brain connections.

Neuronal circuits are also products of development over an individual's life span. Molecular cues hardwire the nervous system using genetic instructions selected by evolution, and neuronal activity and experience fine-tune connectivity. Continuous topographic maps can be constructed by molecular gradients, whereas discrete parallel processing often requires combinatorial cell surface protein codes. These strategies enable a small number of proteins to specify a much larger number of connections.

OUTLOOK: Application of circuit-mapping tools such as serial electron microscopy and transsynaptic tracing to diverse neural regions in different species will generate a wealth of data from which we can distill common principles of neuronal circuit architectures. Activity recording and perturbation of key elements in a circuit can establish their functions in information processing and animal behavior. A key challenge is to investigate how different motifs and architectures operate across scales. A deliberate effort to investigate how letters and words are assembled into sentences in key circuit architectures across species could yield valuable insights. Integrating studies of the structure, function, development, and evolution of neuronal circuits will enable a deeper understanding of nervous system organization beyond the level of individual neurons. Exploration of neuronal circuit architectures will also continue to inspire important advances in artificial intelligence. ■

The list of author affiliations is available in the full article online. Department of Biology and Howard Hughes Medical Institute, Stanford University, Stanford, CA 94305, USA.
Email: lluo@stanford.edu
Cite this article as L. Luo, *Science* 373, eabg7285 (2021).
DOI: 10.1126/science.abg7285

READ THE FULL ARTICLE AT
<https://doi.org/10.1126/science.abg7285>



Between neurons and the brain. Core circuit motifs, such as lateral inhibition depicted here, enable elementary signal processing. Larger-scale architectural plans, such as continuous topographic mapping also depicted here, enable specialized functions. Interrogating the structure and function of neuronal circuits is a central goal of neuroscience; investigating their evolution and development likewise provides crucial insights.

REVIEW

NEUROSCIENCE

Architectures of neuronal circuits

Liqun Luo

Although individual neurons are the basic unit of the nervous system, they process information by working together in neuronal circuits with specific patterns of synaptic connectivity. Here, I review common circuit motifs and architectural plans used in diverse brain regions and animal species. I also consider how these circuit architectures assemble during development and might have evolved. Understanding how specific patterns of synaptic connectivity can implement specific neural computations will help to bridge the huge gap between the biology of the individual neuron and the function of the entire brain, allow us to better understand the neural basis of behavior, and may inspire new advances in artificial intelligence.

Over a century ago, Santiago Ramón y Cajal and his contemporaries proposed that individual neurons are the basic unit of the nervous system. Ramón y Cajal further proposed that information flows from dendrites to cell bodies to axons within individual neurons (Fig. 1) (1). Given that dendrites and axons of most vertebrate neurons are readily distinguishable morphologically, systematic studies of isolated neurons labeled by Golgi staining (2) provided the first overview of how information flows within vertebrate nervous systems (1).

With the advent of modern technologies (Box 1), we have accumulated vast amounts of knowledge of the anatomical, physiological, and functional properties of individual neurons. However, individual neurons do not work in isolation; they work together in neuronal circuits to process information. What is less clear is whether there are generalizable principles about the structural organization of neuronal circuits across different brain regions and animal species. Here, I discuss principles underlying how neurons communicate with each other through specific patterns of synaptic connectivity. Although the importance of activity dynamics in neuronal populations has been increasingly recognized in information processing in diverse systems from invertebrates to mammals (3, 4), synaptic connectivity patterns provide the physical bases on which neuronal dynamics execute their functions. Understanding how these connectivity patterns implement specific computations will allow us to decipher information-processing principles in the nervous system and should inspire new advances in artificial intelligence.

Department of Biology and Howard Hughes Medical Institute, Stanford University, Stanford, CA 94305, USA.
Email: lluo@stanford.edu

Commonly used circuit motifs

If individual neurons are seen as the letters in an alphabet used to write an article that is a brain, then what are the intermediates? In this section, I focus on circuit motifs used across diverse brain regions and animal species (5) (Fig. 2), which can be considered the “words.” In the next section, I explore circuit architectures that might operate at the level of “sentences.” Here, I discuss the most commonly used circuit motifs involving excitatory and inhibitory neurons. Some of these motifs apply not only to neuronal circuits but also to gene-regulatory circuits (6). Architectures based on some of these motifs have also been used in artificial intelligence to great effect (7).

Feedforward excitation

The primary means by which signals flow from one neural region to another is through feedforward excitation, a series of connections between excitatory neurons (Fig. 2A). At each stage, a neuron often receives input from multiple presynaptic partners (convergent excitation) and sends output through branched axons to multiple postsynaptic partners (divergent excitation). Convergent excitation can enable postsynaptic neurons to respond selectively to features not solely or explicitly present in any of the presynaptic neurons. It can also increase the signal-to-noise ratio if multiple input neurons carry the same signal but uncorrelated noise. Divergent excitation allows the same signal to be processed by multiple downstream pathways.

One of the best characterized examples of feedforward excitation is the mammalian visual system, where signals flow from photoreceptors → bipolar cells → retinal ganglion cells → lateral geniculate nucleus (LGN) relay neurons → layer 4 primary visual cortical (V1) neurons → V1 neurons in other layers → neurons in higher cortical areas

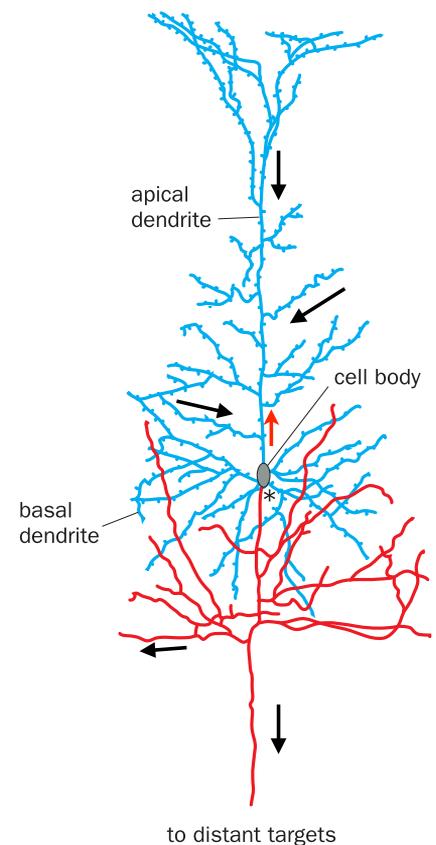


Fig. 1. Information flow within a vertebrate neuron. A pyramidal neuron from rabbit cerebral cortex. Neurons generally use dendrites (blue) to receive information from their presynaptic partners and the axon (red) to send information to their postsynaptic partners. Thus, information flows from dendrites to cell bodies to the axon (black arrows). Asterisk indicates an axon initial segment, where action potentials are generated. Red arrow indicates back-propagating action potentials that can interact with synaptic inputs in complex ways for dendritic computation (115). Whereas axons and dendrites are readily distinguishable by morphological criteria in most vertebrate neurons, most invertebrate central nervous system neurons extend a single process from the cell body that gives rise to both dendrites and the axon. Thus, in invertebrate neurons, information flow is less easily deciphered from morphological criteria alone. Figure was modified and used with permission from (1, 5).

(1, 8, 9). (Note that whereas the discussion here focuses on individual neurons and their synaptic connections, feedforward excitation can also be applied to neural regions in broad strokes, such as retina → LGN → V1.) Along these feedforward pathways, representations of visual information are transformed from light intensity to contrasts, edges, objects, and motion. The feedforward architecture of the mammalian visual system inspired the development of the “perceptron” (10) and

Box 1. Tools for mapping neuronal circuit architecture.

Diverse tools have been used to study the structural organization of the nervous system (94, 95).

Single-neuron tracing

In this approach, a dense library of single neurons within a neural region is created by sparse labeling in individual animals using Golgi staining (2), intracellular dye filling (8), or genetic methods (96–100), such that dendritic morphology and axonal projections are clearly resolved using light microscopy. One can infer that neuron A synapses onto neuron B when A's axon overlaps with B's dendrites. Ramón y Cajal used this approach to chart the coarse organization of the vertebrate nervous system (1). Genetic methods for sparse labeling now allow researchers to infer connectivity between neurons of specific types. A key limitation of this method is that spatial overlap visualized with light microscopy between dendrites and axons is necessary but insufficient to confidently classify two neurons as synaptic partners. Thus, it is only useful for inferring possible connectivity at a coarse level.

Serial electron microscopic reconstruction

Serial electron microscopic (EM) reconstruction is the most comprehensive way of deciphering synaptic wiring diagrams, because it is the only method able to unambiguously visualize synapses. All synapses can be visualized in the same specimen, with the potential of producing a complete wiring diagram. Serial EM reconstruction has been used to decipher the synaptic wiring diagram of the entire *Caenorhabditis elegans* nervous system (101). Recent years have seen rapid progress in the acquisition and partial reconstruction of EM volumes of neural regions from multiple organisms (89, 102–105). A densely reconstructed *Drosophila* hemibrain has been achieved (106), and the entire mouse brain has been proposed as the next ambitious target (107). Limitations include the extensive labor needed to accurately reconstruct connections from EM volumes, especially across large distances, and the difficulty of deciphering cell types or connection signs (excitatory versus inhibitory) unless the region is also well characterized by other means.

Trans-synaptic tracing

Trans-synaptic tracing relies on an event such as gene expression or viral transduction occurring in one neuron to trigger the labeling of its presynaptic partners (retrograde trans-synaptic labeling) or postsynaptic partners (anterograde trans-synaptic labeling). The most widely used methods in mammals use viruses that naturally transduce neurons across synapses, particularly rabies virus for retrograde trans-synaptic tracing from a defined neuron type in a specific location (108, 109). Axon terminal-initiated rabies tracing can reveal inputs to neuronal populations that project to specific targets, allowing inference of input-output architecture (Fig. 4). Anterograde methods have also been reported (110–113). Limitations include poor understanding of trans-synaptic transmission mechanisms, potential biases due to cell type and subcellular locations of synapses, and incomplete characterization of false negatives (synaptic partners not labeled) and false positives (labeling of nonsynaptic partners) for most methods.

Electrophysiological and optical methods

Simultaneous intracellular recordings can reveal synaptic connections between multiple neurons, as well as their sign and strength. This method is mostly limited to in vitro preparations and is therefore mostly used to map local connectivity. However, channelrhodopsin (ChR2)-assisted circuit mapping (114) can map long-range connections between a specific input population (expressing ChR2) and its target neurons in a brain slice, because photostimulating ChR2⁺ axon terminals can often elicit responses in postsynaptic neurons. Because of their low throughput, these electrophysiological and optical methods are mostly used to validate connections suggested by other methods and for detailed analysis of synaptic properties rather than to reveal connectivity within or between neural regions de novo.

“deep neural network” (11) for image recognition and categorization; deep neural networks have also been used in artificial intelligence to solve problems far beyond image analysis (7).

Feedforward and feedback inhibition

Although long-range signals in the nervous system are mostly delivered by excitatory neu-

rons (notable exceptions include basal ganglia and cerebellum circuits), inhibitory interneurons play key roles in sculpting such signals locally (12–14). Two widely used motifs are feedforward and feedback inhibition (Fig. 2B). In feedforward inhibition, an inhibitory neuron receives input from a presynaptic excitatory neuron, and both inhibitory and presynaptic excitatory inputs

converge onto a postsynaptic neuron. In feedback inhibition, an inhibitory neuron receives input from and projects back onto an excitatory neuron, often at its presynaptic terminals. Almost every excitatory connection in the visual pathway described above is accompanied by feedforward inhibition, feedback inhibition, or both. For example, LGN neurons directly excite V1 GABA-releasing neurons to provide feedforward inhibition to layer 4 excitatory neurons, and layer 4 excitatory neurons also activate V1 GABA-releasing neurons to provide feedback inhibition onto themselves (15, 16).

Feedforward inhibition acts more rapidly than feedback inhibition because it reaches the postsynaptic target cell with only one synaptic delay after excitatory signals, whereas feedback inhibition has two synaptic delays (Fig. 2B). Feedforward inhibition is proportional to the strength of the input, whereas feedback inhibition is proportional to the strength of the output; both are used to regulate the duration and magnitude of incoming excitatory signals. For example, limiting the duration of activation in response to sensory input allows circuits to quickly return to their baseline activity levels to maximize their sensitivity to future inputs that signal changes in the environment. Networks of feedforward and feedback inhibitory neurons often act in concert and can perform many interesting functions, such as regulating the gain and dynamic range of input signals and facilitating synchronous or oscillatory firing (14, 17). Feedforward and feedback inhibition also play an essential role in maintaining a “balance” between excitation and inhibition (e.g., strong excitation is accompanied by strong inhibition) to prevent overly excited or inhibited states. Such balanced networks can enhance the speed and signal-to-noise ratio of information processing (18, 19).

Lateral inhibition

Lateral inhibition (Fig. 2C) is a widely occurring circuit motif. It selects information to be propagated to downstream circuits by amplifying differences in activity between parallel pathways. For example, photoreceptor neurons in the vertebrate retina activate horizontal cells, which provide feedback inhibition to many photoreceptor neurons nearby. This action is a major contributor to the classic center-surround receptive field in downstream ganglion cells (20, 21), conferring on these neurons the ability to extract information about spatial or color contrast. Lateral inhibition is also used in other sensory systems (3, 22, 23), with the general purpose of sharpening representations of ethologically relevant information to be processed by downstream circuits.

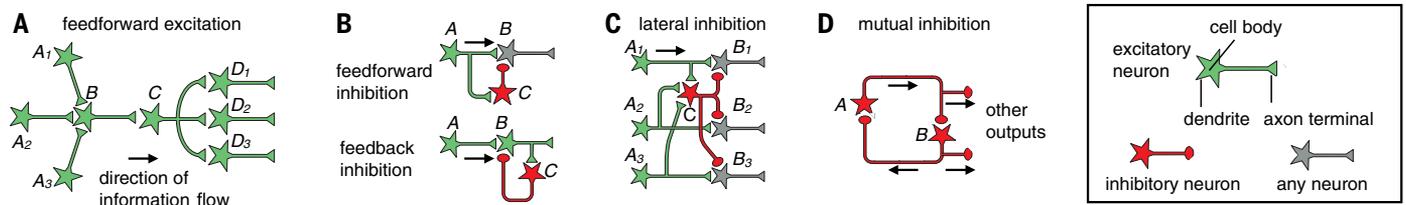


Fig. 2. Commonly used circuit motifs. (A) Feedforward excitation. Information flows through a series of excitatory neurons, A to D . Three different A neurons synapse onto B , exemplifying convergent excitation. C synapses onto three distinct D neurons, exemplifying divergent excitation. (B) In feedforward inhibition (top), inhibitory neuron C receives input from presynaptic excitatory neuron A and sends inhibitory output to postsynaptic neuron B ; in feedback inhibition (bottom), inhibitory neuron C receives input from and sends inhibitory output to postsynaptic excitatory neuron B .

(C) In lateral inhibition, parallel pathways ($A_n \rightarrow B_n$; three are shown) each excite inhibitory neuron C , which in turn sends inhibitory output to all pathways. (D) In mutual inhibition, two inhibitory neurons form reciprocal connections and also provide outputs through branched axons to broadcast their activity states. The inhibitory neurons can also act through intermediary excitatory neurons to inhibit each other (not shown). See box on the right for a key to symbols. Figures in (A) to (C) were modified and used with permission from (5).

Mutual inhibition

Communication between inhibitory neurons can confer circuits with interesting properties. For example, if inhibitory neuron A directly inhibits inhibitory neuron B , then activation of A would disinhibit target neurons of B . If B also inhibits A , then they form the mutual (reciprocal) inhibition motif (Fig. 2D). Mutual inhibition is widely used in circuits that exhibit rhythmic activity, such as those involved in locomotion (24). A classic example is the crustacean stomatogastric ganglion (25). Operating on a longer time scale, mutual inhibition can also be used to regulate brain states such as the sleep–wake cycle (26, 27).

So far, our discussion has involved an alphabet comprising just two letters: excitatory and inhibitory neurons. In reality, the neuronal alphabet is far richer. Both excitatory and inhibitory neurons have many variations because of the heterogeneity in their dendrite morphology, ion channel composition, spiking properties, and the subcellular distribution and strength of their input and output synapses. For example, in the mammalian neocortex, three classes of inhibitory neurons, the Martinotti, basket, and chandelier cells, target their presynaptic terminals to distal dendrites, cell bodies, and axon initial segments of excitatory pyramidal neurons, respectively, and thus control different aspects of how pyramidal neurons integrate synaptic inputs and produce spikes (28, 29). In the stomatogastric ganglion, mutually inhibiting neurons have distinct ion channel compositions and input–output synaptic strengths, which underlie their sequential firing patterns within each rhythmic cycle (30). Finally, the neuronal alphabet also includes many modulatory neuron types to be discussed later.

At the level of core motifs, there are also many variations. For example, the mutual inhibition motif often includes intermediary neurons (e.g., inhibitory neuron A inhibits an excitatory neuron that excites inhibitory neuron

B). It is important to note that the core motifs discussed above are almost always used in concert. Indeed, the large-scale architectural patterns discussed in the next section always contain these motifs.

In summary, a rich alphabet of neurons with diverse intrinsic properties can be used to compose words using a set of core motifs and their variations. These words are often used in concert to produce phrases, which together form the basis for sentences, as I discuss next.

Specialized architectures for specific functions

The next level of organization is more heterogeneous in scale and configuration and less readily generalizable. Nevertheless, I attempt here to extract some high-order circuit architectural patterns that have been found in multiple neural regions and diverse species.

Continuous topographic mapping

Continuous topographic mapping is a common organizational scheme for representing information in the nervous system. Neighboring input neurons connect to neighboring target neurons through orderly axonal projections (Fig. 3A). A prime example is retinotopy, in which neighboring retinal ganglion cells synapse onto neighboring LGN neurons, which then connect to neighboring V1 neurons, which in turn connect to neighboring higher-order visual cortical neurons. Retinotopy enables spatial relationships in the outside world captured by the retina to be recapitulated in V1 and higher visual cortical areas. Continuous topographic mapping is also used elsewhere. In the sensory and motor homunculi, somatosensory stimuli from neighboring body parts are coarsely represented in neighboring areas of the primary somatosensory cortex, and motor outputs to neighboring body parts are coarsely controlled by neighboring areas of the motor cortex (31).

Topographic maps provide a convenient way to organize information at successive stages of processing and can be constructed by robust developmental mechanisms (see below). They have a variety of computational advantages. For example, retinotopy facilitates the extraction of local contrast through lateral inhibition for object recognition. Furthermore, by placing circuit elements that are more often functionally connected nearby each other, topographic maps save energy by minimizing wiring length (32). The design of “convolutional neural networks” (7) takes a page from topographic mapping to greatly reduce the number of variables needed to tune an artificial neural network and thus speed up computation.

Discrete parallel processing

Discrete parallel processing (Fig. 3B) allows signals to be represented and processed in parallel by discrete information channels. A prime example is the glomerular organization of the vertebrate olfactory bulb and insect antennal lobe: Olfactory receptor neurons expressing the same odorant receptors send their axons to the same glomerulus to synapse onto the dendrites of their corresponding second-order projection neurons, forming discrete olfactory processing channels (33, 34). Tens to thousands of individual olfactory receptor neurons expressing the same odorant receptor converge their axons onto the same glomerulus, thus enhancing the signal-to-noise ratio. Rather than representing continuous values, different glomeruli represent signals from discrete olfactory receptor neuron types and thus the nature of the chemicals that activate those odorant receptors. Discrete parallel processing also characterizes the mammalian taste system (35).

Discrete parallel processing is often used in conjunction with continuous topographic mapping. In the retina, for example, superimposed on the continuous retinotopy are discrete layers where different bipolar and ganglion cell types

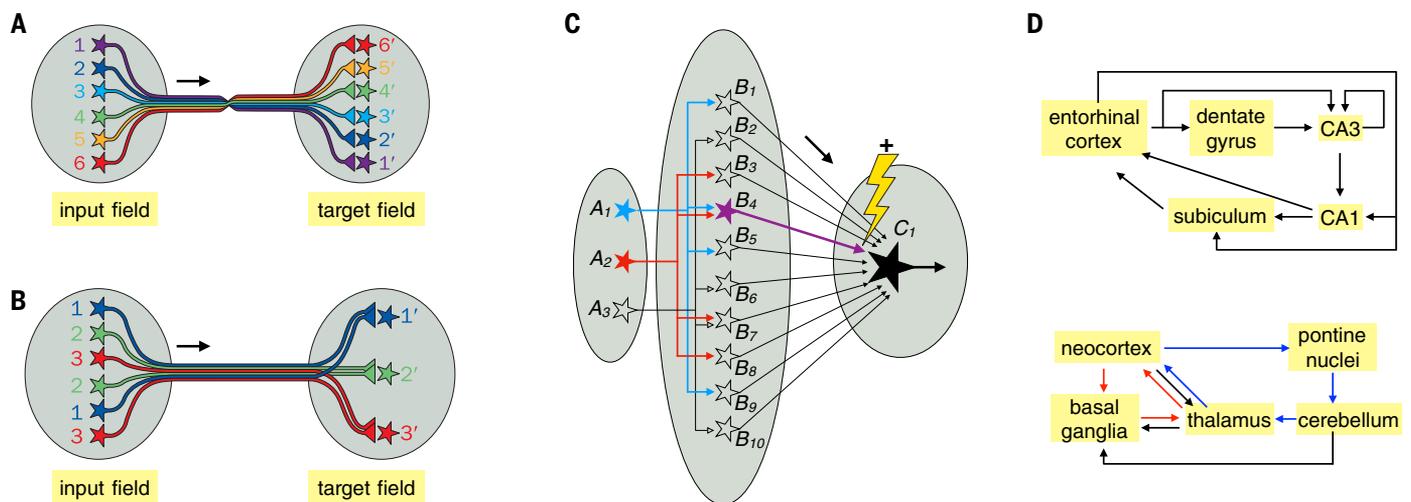


Fig. 3. Specialized architectures for specific functions. (A) Continuous topographic mapping. Neighboring neurons in the input field project their axons in an orderly fashion to connect to neighboring neurons in the target field, preserving their spatial relationships. A prime example is the retinotopic map. (B) Discrete parallel processing. Neurons of a specific type (same color) in the input field, regardless of their spatial locations, connect to the corresponding neuron type in the target field. A prime example is the olfactory glomerular map. Target neurons do not need to be spatially ordered as shown; they could extend their dendrites to connect specifically with the axons of specific input neuron types. (C) Dimensionality expansion. Signals represented by a small number of neurons in field A are represented by a much larger number of neurons in field B, such that activation of B neurons can represent specific combinations of A neurons (e.g., B_4 represents coactivation of A_1 and A_2). Furthermore, the synaptic connections between B and C can be altered by coactivation of a teaching signal (the lightning and plus sign signals

strengthening) and a specific B neuron to modify the synaptic strength between that particular B and C. Thus, only after training would coactivation of A_1 and A_2 reaches the threshold (thick arrow for B_4) for activating C_1 . Likewise, a C_2 neuron (not shown) can be trained to respond to the coactivation of A_2 and A_3 by modifying the strength of $B_7 \rightarrow C_2$. Filled and open symbols represent active and inactive neurons, respectively. (D) Two examples of recurrent loops. In the entorhinal–hippocampal loop (top), arrows indicate direct connections between neurons within the indicated regions. Many connections within these regions are topographically organized (116). In the neocortex–basal ganglia–thalamus (red) and neocortex–pons–cerebellum–thalamus loops (blue), arrows represent connections between these brain regions but not necessarily direct synaptic connections between specific neuron types. Within the basal ganglia and cerebellum, for example, inputs are transformed at intermediary stages to produce outputs. Figures in (A) and (B) were modified and used with permission from (117).

form specific connections to process different types of visual signals such as luminance, color, and motion in parallel. Compared with serial processing, parallel processing reduces computational depth, thus decreasing error rate and increasing processing speed. Indeed, massively parallel processing is a salient feature of complex nervous systems with large numbers of neurons and large numbers of connections per neuron (5). This architecture is increasingly being adopted in computer systems design (11).

Dimensionality expansion

In the dimensionality expansion architecture, signals from a relatively small number of input neurons diverge onto a much larger number of output neurons (Fig. 3C), allowing output neurons to represent distinct combinations of inputs. Similar signals at the input level are more readily distinguished at the output level, facilitating pattern separation by downstream neurons (36–39). Two prime examples are the insect mushroom body (olfactory projection neurons \rightarrow mushroom body Kenyon cells \rightarrow mushroom body output neurons) and the vertebrate cerebellum (mossy fibers \rightarrow cerebellar granule cells \rightarrow Purkinje cells). In both cases, a

relatively small number of inputs (projection neurons or mossy fibers, respectively) synapse onto a much larger number of output neurons (Kenyon cells or granule cells, respectively). Information at the level of the output neurons can thus be represented in a much higher dimensional space, with each dimension representing the firing rate of one cell. Small differences in input firing patterns (e.g., different projection neuron populations representing different odor combinations) can more readily be distinguished by the population firing patterns of their postsynaptic partners. This architecture allows for learning by adjusting the synaptic strengths of the output neurons through “teaching” signals from dopamine neurons in the mushroom body (40, 41) and climbing fibers in the cerebellum (42). After training, the same input can produce different output patterns (Fig. 3C).

Another example of dimensionality expansion is the entorhinal cortex \rightarrow dentate gyrus granule cell \rightarrow CA3 pyramidal neuron circuit (Fig. 3D, top). The large number of dentate gyrus granule cells can perform pattern separation for information from the entorhinal cortex regarding space and objects for further processing by the downstream hippocampal

circuit (43, 44). Unlike in the mushroom body and cerebellar cortex, teaching neurons have not been identified here. This may be because the hippocampal circuit uses unsupervised learning, whereas the cerebellar and mushroom body circuits implement algorithms akin to supervised and reinforcement learning.

Recurrent loops

Nervous systems are full of recurrent loops in which neurons connect back onto themselves, often through intermediary neurons. These recurrent loops are heterogeneous in scale, ranging from within a particular neural region (e.g., mutual inhibition used in the crustacean stomatogastric circuit) to spanning large parts of the brain. In the mammalian visual system, for example, in addition to “bottom-up” projections from LGN \rightarrow V1 \rightarrow higher cortical areas, “top-down” projections from higher cortical areas \rightarrow V1 \rightarrow LGN serve several functions such as attentional control. Long-range recurrent loops may incorporate continuous topographic mapping or discrete parallel processing architectures. Figure 3D illustrates two examples in the mammalian brain at the level of neuronal populations (top) and brain regions (bottom). Recurrent loops generally

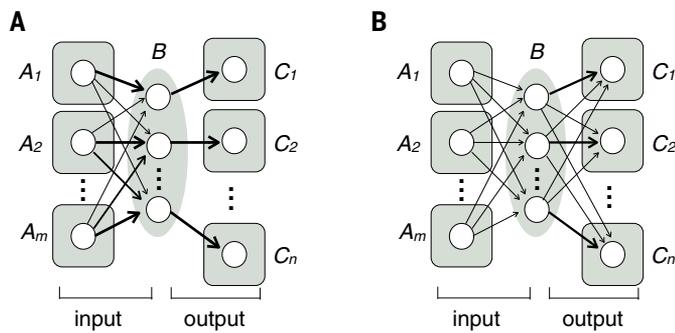


Fig. 4. Input-output organization of neuromodulatory circuits with broad projections.

Modulatory neurons in region *B* collectively receive inputs from regions A_1 – A_m and send broad output to regions C_1 – C_n . **(A)** Biased input-segregated output architecture. This architecture applies to

several neuromodulatory systems, including midbrain dopamine neurons, dorsal raphe serotonin neurons, and preoptic area galanin neurons. Arrows of different thicknesses represent different input strengths. **(B)** Integration-and-broadcast architecture. Neuronal populations in region *B* that project to a specific output region also send output to other output regions, with the possibility of a quantitative bias; these populations also receive similar inputs. Locus coeruleus norepinephrine neurons approximate this architecture. Each circle symbolizes a neuronal population rather than an individual neuron, as the input-output organization summarized here is based on studies at the population level rather than at the level of individual neurons. Figure was modified and used with permission from (48, 49).

support rich neural activity dynamics, but their exact computational roles are not clear in most cases and are likely to differ on a case-by-case basis. Understanding the general principles of information processing in recurrent loops is a major challenge in modern neuroscience.

Biased input-segregated output

The above discussions have focused on circuits comprising excitatory and inhibitory neurons. Nervous systems also use modulatory neurons for important functions. Modulatory neurons use neurotransmitters such as monoamines and neuropeptides that primarily engage G protein-coupled receptors; therefore, their actions on postsynaptic neurons are slower and last between tens of milliseconds to seconds, in contrast to fast excitatory and inhibitory neurotransmitters, which engage ionotropic receptors and act within a few milliseconds. In addition to acting across the synaptic cleft, modulatory neurotransmitters can also be released at sites without postsynaptic specializations, so-called “volume release,” and can thus influence targets at distances greater than that of a typical synaptic cleft.

Some modulatory neurons in the mammalian brain have cell bodies clustered in small regions but project axons broadly and receive diverse inputs. Viral-genetic tracing in the mouse (Box 1) revealed that midbrain dopamine, dorsal raphe serotonin, and hypothalamic neuropeptide galanin systems all adopt a “biased input-segregated output” architecture at the population level (45–48) (Fig. 4A). Each system can be divided into parallel subsystems defined based on their segregated output projections to distinct target regions that serve different behavioral functions. Each out-

put subsystem receives inputs from similar regions with quantitative biases, allowing these subsystems to be differentially regulated by external and internal stimuli. One exception is the locus coeruleus norepinephrine system. At the population level, locus coeruleus norepinephrine axons projecting to one brain region also project broadly to other regions, even though branching patterns of individual neurons can be idiosyncratic (49, 50). These observations suggest that the locus coeruleus norepinephrine system adopts an integration-and-broadcast architecture (Fig. 4B) that may suit its role in regulating global brain states such as arousal.

Nervous systems also use architectures not discussed above. A prominent architecture in bilaterians is interconnected bilateral symmetry (51), which formal network analysis identified as the top-level organization in forebrain connectivity maps (52). The architectures of many neuronal circuits, such as those of the canonical mammalian neocortex (53) and basal ganglia (54) circuits, do not fit neatly into the categories described above even though they use the aforementioned core circuit motifs and can participate as parts of other architectures such as topographic maps (55) and recurrent loops (Fig. 3D bottom). This may be because we have not yet dug deeply enough into these specific circuits to decipher their computational principles or because our understanding of the nervous system is not broad enough to identify shared architectures. We expect ample future opportunities to explore both the depth and breadth of neural circuit architectures by collecting greater amounts of data with increasingly sophisticated tools (Box 1). Only when we know more about these “sentences” and their numerous variations and complex interactions will we have a deeper

understanding of how they constitute “paragraphs” (e.g., brain regions) and eventually the “article”: the overall organization of an entire nervous system.

Evolutionary and developmental perspectives

Whereas computer circuits are products of top-down design, complex neuronal circuits have evolved over hundreds of millions of years. Neuronal circuits also self-assemble during development using evolutionarily selected genetic instructions and are fine-tuned by experience. Thus, existing neuronal circuit architectures are likely a selection of those that can be readily evolved and assembled during development. Looking at a neuronal circuit in isolation may not tell us what elements are functionally important. Seeing what has been evolutionarily selected, expanded, shrunken, eliminated, or repeatedly produced through convergent evolution can, however, suggest what elements to focus on in functional studies.

Evolution of neuronal circuits

Extant bilaterian nervous systems (including all vertebrate and most invertebrate phyla) likely derived from ancestors through progressive sophistication: those with only myocytes, followed by the sequential evolution of sensorimotor neurons, separate sensory and motor neurons, interneurons, and centralized interneuron networks that gave rise to the central nervous system and brain (51, 56). The ubiquity of some core motifs, such as feedforward excitation and feedforward and feedback inhibition, may have originated early in animals with interneurons and a central nervous system and have since been conserved across diverse species and spread across neural regions within each species because of their utility. Other architectures have evolved independently. The glomerular organization of the insect and vertebrate olfactory systems is likely the result of convergent evolution, because many clades descended from their last common ancestor do not have this organization, and different types of molecules are used as odorant receptors. Visual systems provide striking examples of convergent evolution of many fundamental features from retinotopy to motion detection algorithms in invertebrate and vertebrate lineages (57, 58).

Progressive sophistication of the nervous system requires expansion of neuron numbers (59), neuron types and their connections (60), and brain regions (61). All of these processes must result from changes to DNA. A key mechanism of evolutionary innovation is the duplication and divergence of genes; for example, duplication and divergence of a cone opsin gene conferred trichromacy on some primates (62). Duplication and divergence is also used in the evolution of neuron types (63–65) and

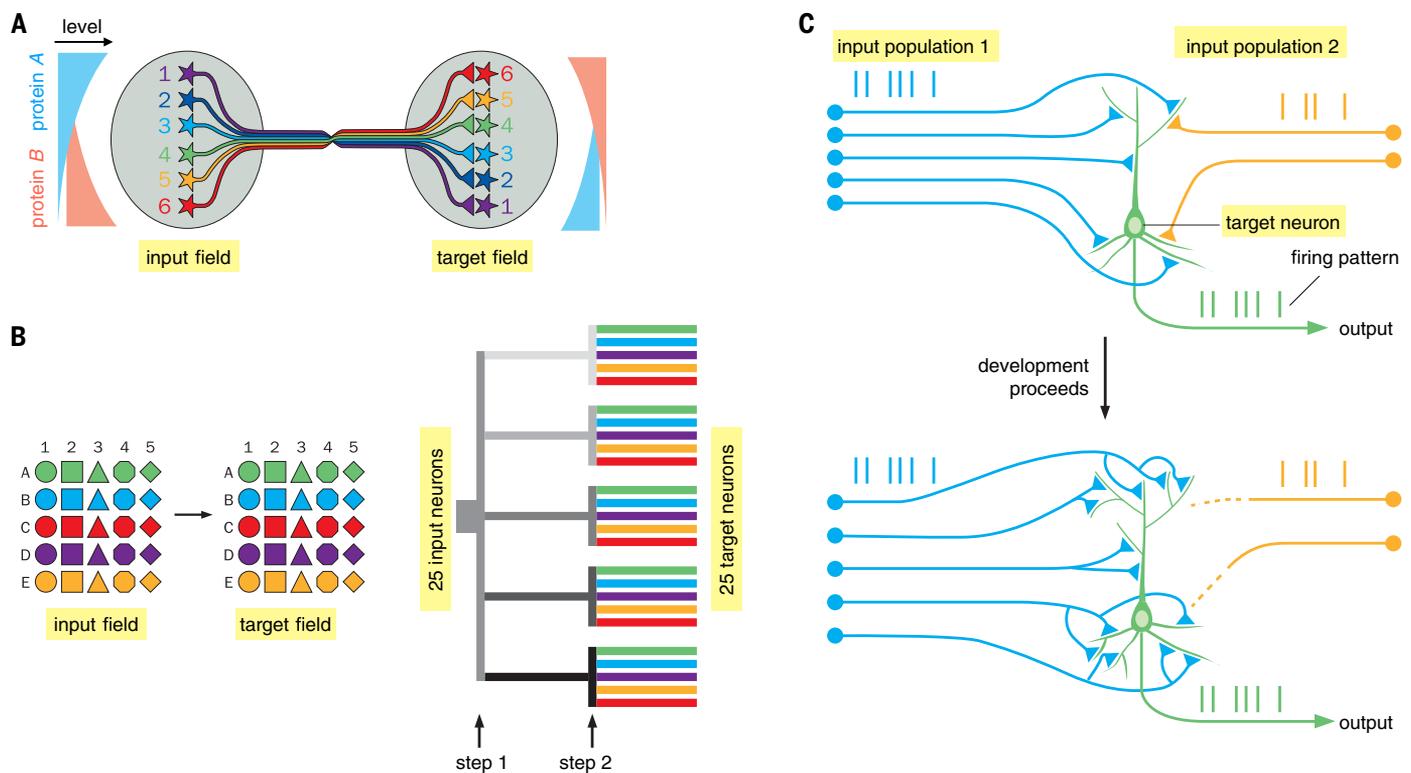


Fig. 5. Wiring up neuronal circuits. (A) Protein gradients can be used to construct continuous topographic maps. In this example, both input and target fields are patterned by opposing gradients of cell surface proteins A and B. Suppose that neuronal processes expressing protein A and protein B mutually repel each other. Because neuron 1 has the highest level of protein A, it seeks a target field with the lowest level of protein B; likewise, neuron 6 seeks a target field with the lowest level of protein A. (B) Illustration of combinatorial strategies to specify connections between 25 discrete cell types in the input and target fields. Left: Suppose that connection specificities between the input and target fields are mediated by homophilic attraction molecules. If each connection is specified by a single molecule, then 25 molecules are needed to specify 25 connections. If each connection is specified by a combination of two molecules (a letter and a number), then only 10 molecules are needed. Right: The

combinatorial strategy is realized by dividing the wiring process into two steps. At step 1, five molecules (represented by different shades of gray) separate five input axons into five groups; at step 2, five more molecules (represented by different colors) are used in each of the five groups to specify the final connections. (C) Schematic illustration of Hebb's rule in instructing wiring. At an early developmental stage, the target neuron is connected to two groups of input neurons with distinct coincident firing patterns (blue and yellow vertical lines). Because stronger connections to the group 1 input drive the target neuron to fire in a pattern (green vertical lines) similar to that of group 1, their connections are strengthened. Synaptic connections between group 2 input and the target neuron weaken because of their dissimilar firing patterns. Eventually, the target neuron is only connected with group 1 input. Figure was modified and used with permission from (5).

brain regions (66). Duplication and divergence for brain region evolution should in principle make neuronal circuits modular: rich connections within a duplicated unit and sparse connections between units (as opposed to all-to-all nonmodular architectures used as the starting conditions in many artificial neural networks). In turn, the modular nature of neuronal circuits might speed up evolution (6) because different modules can evolve independently of each other.

Development of neuronal circuits

Evolution exerts its influence on neuronal circuits primarily by modifying genes involved in circuit wiring during development. A key question is how a limited number of genes (~20,000 across many animal species) can construct nervous systems with much larger numbers of synaptic connections (~ 10^7 in fruit

flies, ~ 10^{11} in mice, and $>10^{14}$ in humans) with specific motifs and architectures.

Extracellular cues and their cell surface receptors enable recognition of specific targets by axonal and dendritic growth cones. These molecules are the predominant force for establishing a coarse organization of the nervous system and can also specify synaptic connectivity with great precision in certain circuits and organisms (67–69). One strategy to establish specificity of a large number of connections with a limited number of genes is to use different expression levels of the same protein to specify different connections. This strategy is readily used in constructing continuous topographic maps (70, 71) (Fig. 5A), perhaps contributing to the prevalence of this circuit architecture (Fig. 3A). Graded expression of cell surface molecules is also used in the early steps of constructing discrete maps (72, 73).

However, discrete parallel processing (Fig. 3B) requires distinguishing between discrete cell types and often uses combinatorial cell surface protein codes such that a small number of proteins can specify many more connections (Fig. 5B, left). An efficient way to implement combinatorial coding is to divide the wiring process into distinct spatiotemporal steps (Fig. 5B, right); in addition to conserving molecules, this strategy can also enhance robustness because growth cones are faced with few simultaneous choices at each step. The same wiring molecules can be used at different times and places, sometimes in different parts of the same circuit, through elaborate spatiotemporal regulation of their expression patterns (74–76).

Neuronal activity, both spontaneous and experience driven, refines synaptic wiring diagrams. Activity-dependent wiring, often through competition between neurons with

different activity levels, has been well documented (77–81). A prominent mechanism by which neuronal activity influences wiring is by implementing Hebb's rule: Synapses at which firing of presynaptic neurons causes firing of postsynaptic neurons are strengthened, or colloquially, “fire together, wire together” (82, 83) (Fig. 5C). Non-Hebbian mechanisms such as homeostatic synaptic plasticity also contribute to activity-dependent circuit wiring (84). These activity-dependent mechanisms continue to operate in the adult nervous system, enabling animals to change their synaptic connectivity patterns as a consequence of experience throughout life.

Many synaptic connections are not completely specified. In vertebrate neuromuscular systems, for example, whereas the connections between motor neuron pools and muscles are precisely specified (85), the specific connection patterns between individual motor neurons and muscle fibers within a motor pool are highly variable (86). Likewise, in the fly olfactory circuit, synaptic connections between specific olfactory projection neuron types and mushroom body Kenyon cells (Fig. 3C) are mostly random (87–89). In both cases, it is not necessary, or even desirable, to have more stereotyped connectivity. As more synaptic connectomes are mapped (Box 1), more examples of wiring variability will surely emerge.

In summary, two broad kinds of mechanisms are used to establish wiring patterns of neuronal circuits: molecular cues hardwire the nervous system and neuronal activity and experience fine-tune connectivity. There is also interplay between neuronal activity and molecular cues; for example, neuronal activity can regulate the expression of molecular cues or complement their action (90, 91). However, apart from the limited examples discussed above, most developmental studies have not focused on addressing how specific circuit motifs and architectures are established, and most investigations of circuit function have not considered developmental constraints. There is thus ample opportunity for cross-fertilization of developmental and functional studies of neuronal circuits.

Outlook

Applications of circuit-mapping tools such as serial electron microscopy and trans-synaptic tracing (Box 1) to diverse neural regions and organisms will surely generate a wealth of data from which we can distill common principles of structural organization of neuronal circuits. Relating structures to the functions they implement will be an important next step. This can be done by leveraging powerful tools that have been developed and applied to functionally interrogate neuronal circuits in the context of animal behavior (92–94). Such interrogation is essential for identifying the functions of each

circuit elements. In addition, a key challenge is to investigate how these motifs and architectures interact with each other across scales. Understanding how different architectures cooperate in an individual nervous system should also inspire new artificial neural networks that might someday achieve general purpose artificial intelligence.

We are still only beginning to gain insights into the evolutionary and developmental processes that give rise to circuit architecture in complex nervous systems. We still do not know, for example, whether and to what degree algorithmic changes in the wiring process and operation of neuronal circuits can account for the increased complexity of the mammalian brain. A deliberate effort to investigate how letters are assembled into words and words into sentences in key circuit architectures across different species could yield valuable insights. Comparative study of neuron-type composition of homologous brain regions using single-cell transcriptomics (63–66) is a useful first step. This can be followed by investigation of the mechanisms that establish their connectivity patterns and that underlie their functional operations. Integrating studies of the structure, function, development, and evolution of neuronal circuits will enable a deeper understanding of nervous system organization beyond the level of individual neurons.

REFERENCES AND NOTES

- S. R. Cajal, *Histology of the Nervous System of Man and Vertebrates* (Oxford Univ. Press, 1995).
- C. Golgi, Sulla struttura della sostanza grigia del cervello. *Gazzetta medica lombarda* **IV** (1873).
- G. Laurent *et al.*, Odor encoding as an active, dynamical process: Experiments, computation, and theory. *Annu. Rev. Neurosci.* **24**, 263–297 (2001). doi: [10.1146/annurev.neuro.24.1.263](https://doi.org/10.1146/annurev.neuro.24.1.263); pmid: [11283312](https://pubmed.ncbi.nlm.nih.gov/11283312/)
- K. V. Shenoy, M. Sahani, M. M. Churchland, Cortical control of arm movements: A dynamical systems perspective. *Annu. Rev. Neurosci.* **36**, 337–359 (2013). doi: [10.1146/annurev-neuro-062111-150509](https://doi.org/10.1146/annurev-neuro-062111-150509); pmid: [23725001](https://pubmed.ncbi.nlm.nih.gov/23725001/)
- L. Luo, *Principles of Neurobiology* (CRC Press/Garland Science, ed. 2, 2020).
- U. Alon, *An Introduction to Systems Biology* (CRC Press, ed. 2, 2020).
- Y. LeCun, Y. Bengio, G. Hinton, Deep learning. *Nature* **521**, 436–444 (2015). doi: [10.1038/nature14539](https://doi.org/10.1038/nature14539); pmid: [26017442](https://pubmed.ncbi.nlm.nih.gov/26017442/)
- C. D. Gilbert, T. N. Wiesel, Morphology and intracortical projections of functionally characterised neurones in the cat visual cortex. *Nature* **280**, 120–125 (1979). doi: [10.1038/280120a0](https://doi.org/10.1038/280120a0); pmid: [552600](https://pubmed.ncbi.nlm.nih.gov/552600/)
- D. J. Felleman, D. C. Van Essen, Distributed hierarchical processing in the primate cerebral cortex. *Cereb. Cortex* **1**, 1–47 (1991). doi: [10.1093/cercor/1.1.1](https://doi.org/10.1093/cercor/1.1.1); pmid: [1822724](https://pubmed.ncbi.nlm.nih.gov/1822724/)
- F. Rosenblatt, The perceptron: A probabilistic model for information storage and organization in the brain. *Psychol. Rev.* **65**, 386–408 (1958). doi: [10.1037/h0042519](https://doi.org/10.1037/h0042519); pmid: [13602029](https://pubmed.ncbi.nlm.nih.gov/13602029/)
- A. Krizhevsky, I. Sutskever, G. Hinton, Imagenet classification with deep convolutional neural networks. *In Proc. Adv. Neural Inf. Process. Syst.* **25**, 1090–1098 (2012).
- C. S. Sherrington, Remarks on some aspects of reflex inhibition. *Proc. R. Soc. London Ser. B* **97**, 519–545 (1925). doi: [10.1098/rspb.1925.0017](https://doi.org/10.1098/rspb.1925.0017)
- J. S. Coombs, J. C. Eccles, P. Fatt, The inhibitory suppression of reflex discharges from motoneurons. *J. Physiol.* **130**, 396–413 (1955). doi: [10.1113/jphysiol.1955.sp005414](https://doi.org/10.1113/jphysiol.1955.sp005414); pmid: [13278907](https://pubmed.ncbi.nlm.nih.gov/13278907/)
- J. S. Isaacson, M. Scanziani, How inhibition shapes cortical activity. *Neuron* **72**, 231–243 (2011). doi: [10.1016/j.neuron.2011.09.027](https://doi.org/10.1016/j.neuron.2011.09.027); pmid: [22017986](https://pubmed.ncbi.nlm.nih.gov/22017986/)
- K. A. Martin, P. Somogyi, D. Whitteridge, Physiological and morphological properties of identified basket cells in the cat's visual cortex. *Exp. Brain Res.* **50**, 193–200 (1983). doi: [10.1007/BF00239183](https://doi.org/10.1007/BF00239183); pmid: [6641854](https://pubmed.ncbi.nlm.nih.gov/6641854/)
- X. Y. Ji *et al.*, Thalamocortical innervation pattern in mouse auditory and visual cortex: Laminar and cell-type specificity. *Cereb. Cortex* **26**, 2612–2625 (2016). doi: [10.1093/cercor/bhv099](https://doi.org/10.1093/cercor/bhv099); pmid: [25979090](https://pubmed.ncbi.nlm.nih.gov/25979090/)
- L. Roux, G. Buzsáki, Tasks for inhibitory interneurons in intact brain circuits. *Neuropharmacology* **88**, 10–23 (2015). doi: [10.1016/j.neuropharm.2014.09.011](https://doi.org/10.1016/j.neuropharm.2014.09.011); pmid: [25239808](https://pubmed.ncbi.nlm.nih.gov/25239808/)
- C. van Vreeswijk, H. Sompolinsky, Chaos in neuronal networks with balanced excitatory and inhibitory activity. *Science* **274**, 1724–1726 (1996). doi: [10.1126/science.274.5293.1724](https://doi.org/10.1126/science.274.5293.1724); pmid: [8939866](https://pubmed.ncbi.nlm.nih.gov/8939866/)
- M. N. Shadlen, W. T. Newsome, The variable discharge of cortical neurons: Implications for connectivity, computation, and information coding. *J. Neurosci.* **18**, 3870–3896 (1998). doi: [10.1523/JNEUROSCI.18-10-03870.1998](https://doi.org/10.1523/JNEUROSCI.18-10-03870.1998); pmid: [9570816](https://pubmed.ncbi.nlm.nih.gov/9570816/)
- S. W. Kuffler, Discharge patterns and functional organization of mammalian retina. *J. Neurophysiol.* **16**, 37–68 (1953). doi: [10.1152/jn.1953.16.1.37](https://doi.org/10.1152/jn.1953.16.1.37); pmid: [13035466](https://pubmed.ncbi.nlm.nih.gov/13035466/)
- H. B. Barlow, Summation and inhibition in the frog's retina. *J. Physiol.* **119**, 69–88 (1953). doi: [10.1113/jphysiol.1953.sp004829](https://doi.org/10.1113/jphysiol.1953.sp004829); pmid: [13035718](https://pubmed.ncbi.nlm.nih.gov/13035718/)
- V. B. Mountcastle, Modality and topographic properties of single neurons of cat's somatic sensory cortex. *J. Neurophysiol.* **20**, 408–434 (1957). doi: [10.1152/jn.1957.20.4.408](https://doi.org/10.1152/jn.1957.20.4.408); pmid: [13439410](https://pubmed.ncbi.nlm.nih.gov/13439410/)
- G. M. Shepherd, W. R. Chen, C. A. Greer, “Olfactory bulb,” in *The Synaptic Organization of the Brain*, G. M. Shepherd, Ed. (Oxford Univ. Press, 2004), pp. 165–216.
- S. Grillner, Biological pattern generation: The cellular and computational logic of networks in motion. *Neuron* **52**, 751–766 (2006). doi: [10.1016/j.neuron.2006.11.008](https://doi.org/10.1016/j.neuron.2006.11.008); pmid: [17145498](https://pubmed.ncbi.nlm.nih.gov/17145498/)
- E. Marder, D. Bucher, Understanding circuit dynamics using the stomatogastric nervous system of lobsters and crabs. *Annu. Rev. Physiol.* **69**, 291–316 (2007). doi: [10.1146/annurev.physiol.69.031905.161516](https://doi.org/10.1146/annurev.physiol.69.031905.161516); pmid: [17009928](https://pubmed.ncbi.nlm.nih.gov/17009928/)
- C. B. Saper, P. M. Fuller, N. P. Pedersen, J. Lu, T. E. Scammell, Sleep state switching. *Neuron* **68**, 1023–1042 (2010). doi: [10.1016/j.neuron.2010.11.032](https://doi.org/10.1016/j.neuron.2010.11.032); pmid: [21172606](https://pubmed.ncbi.nlm.nih.gov/21172606/)
- F. Weber, Y. Dan, Circuit-based interrogation of sleep control. *Nature* **538**, 51–59 (2016). doi: [10.1038/nature19773](https://doi.org/10.1038/nature19773); pmid: [27708309](https://pubmed.ncbi.nlm.nih.gov/27708309/)
- H. Markram *et al.*, Interneurons of the neocortical inhibitory system. *Nat. Rev. Neurosci.* **5**, 793–807 (2004). doi: [10.1038/nrn1519](https://doi.org/10.1038/nrn1519); pmid: [15378039](https://pubmed.ncbi.nlm.nih.gov/15378039/)
- Z. J. Huang, A. Paul, The diversity of GABAergic neurons and neural communication elements. *Nat. Rev. Neurosci.* **20**, 563–572 (2019). doi: [10.1038/s41583-019-0195-4](https://doi.org/10.1038/s41583-019-0195-4); pmid: [31222186](https://pubmed.ncbi.nlm.nih.gov/31222186/)
- A. A. Prinz, D. Bucher, E. Marder, Similar network activity from disparate circuit parameters. *Nat. Neurosci.* **7**, 1345–1352 (2004). doi: [10.1038/nrn1352](https://doi.org/10.1038/nrn1352); pmid: [15558066](https://pubmed.ncbi.nlm.nih.gov/15558066/)
- W. Penfield, T. Rasmussen, *The Cerebral Cortex of Man: A Clinical Study of Localization of Function* (Macmillan, 1950).
- I. E. Wang, T. R. Clandinin, The influence of wiring economy on nervous system evolution. *Curr. Biol.* **26**, R1101–R1108 (2016). doi: [10.1016/j.cub.2016.08.053](https://doi.org/10.1016/j.cub.2016.08.053); pmid: [27780051](https://pubmed.ncbi.nlm.nih.gov/27780051/)
- R. Axel, The molecular logic of smell. *Sci. Am.* **273**, 154–159 (1995). doi: [10.1038/scientificamerican1095-154](https://doi.org/10.1038/scientificamerican1095-154); pmid: [7481719](https://pubmed.ncbi.nlm.nih.gov/7481719/)
- L. B. Vosshall, R. F. Stocker, Molecular architecture of smell and taste in *Drosophila*. *Annu. Rev. Neurosci.* **30**, 505–533 (2007). doi: [10.1146/annurev.neuro.30.051606.094306](https://doi.org/10.1146/annurev.neuro.30.051606.094306); pmid: [17506643](https://pubmed.ncbi.nlm.nih.gov/17506643/)
- D. A. Yarmolinsky, C. S. Zuker, N. J. Ryba, Common sense about taste: From mammals to insects. *Cell* **139**, 234–244 (2009). doi: [10.1016/j.cell.2009.10.001](https://doi.org/10.1016/j.cell.2009.10.001); pmid: [19837029](https://pubmed.ncbi.nlm.nih.gov/19837029/)
- D. Marr, A theory of cerebellar cortex. *J. Physiol.* **202**, 437–470 (1969). doi: [10.1113/jphysiol.1969.sp008820](https://doi.org/10.1113/jphysiol.1969.sp008820); pmid: [5784296](https://pubmed.ncbi.nlm.nih.gov/5784296/)
- J. S. Albus, A theory of cerebellar function. *Math. Biosci.* **10**, 25–61 (1971). doi: [10.1016/0025-5564\(71\)90051-4](https://doi.org/10.1016/0025-5564(71)90051-4)
- A. Litwin-Kumar, K. D. Harris, R. Axel, H. Sompolinsky, L. F. Abbott, Optimal degrees of synaptic connectivity. *Neuron* **93**, 1153–1164.e7 (2017). doi: [10.1016/j.neuron.2017.01.030](https://doi.org/10.1016/j.neuron.2017.01.030); pmid: [28215558](https://pubmed.ncbi.nlm.nih.gov/28215558/)
- N. A. Cayco-Gajic, R. A. Silver, Re-evaluating circuit mechanisms underlying pattern separation. *Neuron* **101**, 584–602 (2019). doi: [10.1016/j.neuron.2019.01.044](https://doi.org/10.1016/j.neuron.2019.01.044); pmid: [30790539](https://pubmed.ncbi.nlm.nih.gov/30790539/)

40. T. Hige, Y. Aso, M. N. Modi, G. M. Rubin, G. C. Turner, Heterosynaptic plasticity underlies aversive olfactory learning in *Drosophila*. *Neuron* **88**, 985–998 (2015). doi: [10.1016/j.neuron.2015.11.003](https://doi.org/10.1016/j.neuron.2015.11.003); pmid: 26637800
41. R. Cohn, I. Morantte, V. Ruta, Coordinated and compartmentalized neuromodulation shapes sensory processing in *Drosophila*. *Cell* **163**, 1742–1755 (2015). doi: [10.1016/j.cell.2015.11.019](https://doi.org/10.1016/j.cell.2015.11.019); pmid: 26687359
42. C. I. De Zeeuw, S. G. Lisberger, J. L. Raymond, Diversity and dynamism in the cerebellum. *Nat. Neurosci.* **24**, 160–167 (2021). doi: [10.1038/s41593-020-00754-9](https://doi.org/10.1038/s41593-020-00754-9); pmid: 33288911
43. D. Marr, Simple memory: A theory for archicortex. *Philos. Trans. R. Soc. Lond. B Biol. Sci.* **262**, 23–81 (1971). doi: [10.1098/rstb.1971.0078](https://doi.org/10.1098/rstb.1971.0078); pmid: 4399412
44. J. P. Neunuebel, J. J. Knierim, CA3 retrieves coherent representations from degraded input: Direct evidence for CA3 pattern completion and dentate gyrus pattern separation. *Neuron* **81**, 416–427 (2014). doi: [10.1016/j.neuron.2013.11.017](https://doi.org/10.1016/j.neuron.2013.11.017); pmid: 24462102
45. K. T. Beier *et al.*, Circuit architecture of VTA dopamine neurons revealed by systematic input-output mapping. *Cell* **162**, 622–634 (2015). doi: [10.1016/j.cell.2015.07.015](https://doi.org/10.1016/j.cell.2015.07.015); pmid: 26232228
46. T. N. Lerner *et al.*, Intact-brain analyses reveal distinct information carried by SNC dopamine subcircuits. *Cell* **162**, 635–647 (2015). doi: [10.1016/j.cell.2015.07.014](https://doi.org/10.1016/j.cell.2015.07.014); pmid: 26232229
47. J. Ren *et al.*, Anatomically defined and functionally distinct dorsal raphe serotonin sub-systems. *Cell* **175**, 472–487.e20 (2018). doi: [10.1016/j.cell.2018.07.043](https://doi.org/10.1016/j.cell.2018.07.043); pmid: 30146164
48. J. Kohl *et al.*, Functional circuit architecture underlying parental behaviour. *Nature* **556**, 326–331 (2018). doi: [10.1038/s41586-018-0027-0](https://doi.org/10.1038/s41586-018-0027-0); pmid: 29643503
49. L. A. Schwarz *et al.*, Viral-genetic tracing of the input-output organization of a central noradrenergic circuit. *Nature* **524**, 88–92 (2015). doi: [10.1038/nature14600](https://doi.org/10.1038/nature14600); pmid: 26131933
50. J. M. Kebschull *et al.*, High-throughput mapping of single-neuron projections by sequencing of barcoded rna. *Neuron* **91**, 975–987 (2016). doi: [10.1016/j.neuron.2016.07.036](https://doi.org/10.1016/j.neuron.2016.07.036); pmid: 27545715
51. L. W. Swanson, *Brain Architecture* (Oxford Univ. Press, ed. 2, 2012).
52. L. W. Swanson, J. D. Hahn, O. Sporns, Structure-function subsystem models of female and male forebrain networks integrating cognition, affect, behavior, and bodily functions. *Proc. Natl. Acad. Sci. U.S.A.* **117**, 31470–31481 (2020). doi: [10.1073/pnas.2017733117](https://doi.org/10.1073/pnas.2017733117); pmid: 33229546
53. R. J. Douglas, K. A. Martin, Neuronal circuits of the neocortex. *Annu. Rev. Neurosci.* **27**, 419–451 (2004). doi: [10.1146/annurev.neuro.27.070203.144152](https://doi.org/10.1146/annurev.neuro.27.070203.144152); pmid: 15217339
54. C. R. Gerfen, D. J. Surmeier, Modulation of striatal projection systems by dopamine. *Annu. Rev. Neurosci.* **34**, 441–466 (2011). doi: [10.1146/annurev-neuro-061010-113641](https://doi.org/10.1146/annurev-neuro-061010-113641); pmid: 21469956
55. J. Lee, W. Wang, B. L. Sabatini, Anatomically segregated basal ganglia pathways allow parallel behavioral modulation. *Nat. Neurosci.* **23**, 1388–1398 (2020). doi: [10.1038/s41593-020-00712-5](https://doi.org/10.1038/s41593-020-00712-5); pmid: 32989293
56. J. H. Kaas, *Evolutionary Neuroscience* (Elsevier, ed. 2, 2020).
57. L. V. Salvini-Plawen, E. Mayr, On the evolution of photoreceptors and eyes. *Evol. Biol.* **10**, 207–263 (1977).
58. D. A. Clark *et al.*, Flies and humans share a motion estimation strategy that exploits natural scene statistics. *Nat. Neurosci.* **17**, 296–303 (2014). doi: [10.1038/nrn.3600](https://doi.org/10.1038/nrn.3600); pmid: 24390225
59. J. H. Lui, D. V. Hansen, A. R. Kriegstein, Development and evolution of the human neocortex. *Cell* **146**, 18–36 (2011). doi: [10.1016/j.cell.2011.06.030](https://doi.org/10.1016/j.cell.2011.06.030); pmid: 21729779
60. D. Arendt *et al.*, The origin and evolution of cell types. *Nat. Rev. Genet.* **17**, 744–757 (2016). doi: [10.1038/nrg.2016.127](https://doi.org/10.1038/nrg.2016.127); pmid: 27818507
61. M. Chakraborty, E. D. Jarvis, Brain evolution by brain pathway duplication. *Philos. Trans. R. Soc. Lond. B Biol. Sci.* **370**, 20150056 (2015). doi: [10.1098/rstb.2015.0056](https://doi.org/10.1098/rstb.2015.0056); pmid: 26554045
62. G. H. Jacobs, J. Nathans, The evolution of Primate color vision. *Sci. Am.* **300**, 56–63 (2009). doi: [10.1038/scientificamerican0409-56](https://doi.org/10.1038/scientificamerican0409-56); pmid: 19363921
63. M. A. Tosches *et al.*, Evolution of pallium, hippocampus, and cortical cell types revealed by single-cell transcriptomics in reptiles. *Science* **360**, 881–888 (2018). doi: [10.1126/science.aar4237](https://doi.org/10.1126/science.aar4237); pmid: 29724907
64. Y. R. Peng *et al.*, Molecular classification and comparative taxonomies of foveal and peripheral cells in primate retina. *Cell* **176**, 1222–1237.e22 (2019). doi: [10.1016/j.cell.2019.01.004](https://doi.org/10.1016/j.cell.2019.01.004); pmid: 30712875
65. R. D. Hodge *et al.*, Conserved cell types with divergent features in human versus mouse cortex. *Nature* **573**, 61–68 (2019). doi: [10.1038/s41586-019-1506-7](https://doi.org/10.1038/s41586-019-1506-7); pmid: 31435019
66. J. M. Kebschull *et al.*, Cerebellar nuclei evolved by repeatedly duplicating a conserved cell-type set. *Science* **370**, eabd5059 (2020). pmid: 33335034
67. A. L. Kolodkin, M. Tessier-Lavigne, Mechanisms and molecules of neuronal wiring: A primer. *Cold Spring Harb. Perspect. Biol.* **3**, a001727 (2011). doi: [10.1101/cshperspect.a001727](https://doi.org/10.1101/cshperspect.a001727); pmid: 21123392
68. W. Hong, L. Luo, Genetic control of wiring specificity in the fly olfactory system. *Genetics* **196**, 17–29 (2014). doi: [10.1534/genetics.113.154336](https://doi.org/10.1534/genetics.113.154336); pmid: 24395823
69. J. R. Sanes, S. L. Zipursky, Synaptic specificity, recognition molecules, and assembly of neural circuits. *Cell* **181**, 536–556 (2020). doi: [10.1016/j.cell.2020.04.008](https://doi.org/10.1016/j.cell.2020.04.008); pmid: 32359437
70. U. Drescher *et al.*, In vitro guidance of retinal ganglion cell axons by RAGS, a 25 kDa tectal protein related to ligands for Eph receptor tyrosine kinases. *Cell* **82**, 359–370 (1995). doi: [10.1016/0092-8674\(95\)90425-5](https://doi.org/10.1016/0092-8674(95)90425-5); pmid: 7634326
71. H.-J. Cheng, M. Nakamoto, A. D. Bergemann, J. G. Flanagan, Complementary gradients in expression and binding of ELF-1 and Mek4 in development of the topographic retinotectal projection map. *Cell* **82**, 371–381 (1995). doi: [10.1016/0092-8674\(95\)90426-3](https://doi.org/10.1016/0092-8674(95)90426-3); pmid: 7634327
72. T. Imai, M. Suzuki, H. Sakano, Odorant receptor-derived cAMP signals direct axonal targeting. *Science* **314**, 657–661 (2006). doi: [10.1126/science.1131794](https://doi.org/10.1126/science.1131794); pmid: 16990513
73. T. Komyama, L. B. Sweeney, O. Schuldiner, K. C. Garcia, L. Luo, Graded expression of semaphorin-1a cell-autonomously directs dendritic targeting of olfactory projection neurons. *Cell* **128**, 399–410 (2007). doi: [10.1016/j.cell.2006.12.028](https://doi.org/10.1016/j.cell.2006.12.028); pmid: 17254975
74. W. J. Joo, L. B. Sweeney, L. Liang, L. Luo, Linking cell fate, trajectory choice, and target selection: Genetic analysis of Sema-2b in olfactory axon targeting. *Neuron* **78**, 673–686 (2013). doi: [10.1016/j.neuron.2013.03.022](https://doi.org/10.1016/j.neuron.2013.03.022); pmid: 23719164
75. J. Cang, D. A. Feldheim, Developmental mechanisms of topographic map formation and alignment. *Annu. Rev. Neurosci.* **36**, 51–77 (2013). doi: [10.1146/annurev-neuro-062012-170341](https://doi.org/10.1146/annurev-neuro-062012-170341); pmid: 23642132
76. D. T. Pedrick *et al.*, Reciprocal repulsions instruct the precise assembly of parallel hippocampal networks. *Science* **372**, 1068–1073 (2021). doi: [10.1126/science.abg1774](https://doi.org/10.1126/science.abg1774); pmid: 34083484
77. T. N. Wiesel, D. H. Hubel, Single-cell responses in striate cortex of kittens deprived of vision in one eye. *J. Neurophysiol.* **26**, 1003–1017 (1963). doi: [10.1152/jn.1963.26.6.1003](https://doi.org/10.1152/jn.1963.26.6.1003); pmid: 14084161
78. M. Constantine-Paton, M. I. Law, Eye-specific termination bands in tecta of three-eyed frogs. *Science* **202**, 639–641 (1978). doi: [10.1126/science.309179](https://doi.org/10.1126/science.309179); pmid: 309179
79. M. P. Stryker, W. A. Harris, Binocular impulse blockade prevents the formation of ocular dominance columns in cat visual cortex. *J. Neurosci.* **6**, 2117–2133 (1986). doi: [10.1523/JNEUROSCI.06-08-02117.1986](https://doi.org/10.1523/JNEUROSCI.06-08-02117.1986); pmid: 3746403
80. A. A. Penn, P. A. Riquelme, M. B. Feller, C. J. Shatz, Competition in retinogeniculate patterning driven by spontaneous activity. *Science* **279**, 2108–2112 (1998). doi: [10.1126/science.279.5359.2108](https://doi.org/10.1126/science.279.5359.2108); pmid: 9516112
81. M. Buffelli *et al.*, Genetic evidence that relative synaptic efficacy biases the outcome of synaptic competition. *Nature* **424**, 430–434 (2003). doi: [10.1038/nature01844](https://doi.org/10.1038/nature01844); pmid: 12879071
82. D. O. Hebb, *The Organization of Behavior* (Wiley, 1949).
83. G. S. Stent, A physiological mechanism for Hebb's postulate of learning. *Proc. Natl. Acad. Sci. U.S.A.* **70**, 997–1001 (1973). doi: [10.1073/pnas.70.4.997](https://doi.org/10.1073/pnas.70.4.997); pmid: 4352227
84. G. G. Turrigiano, The dialectic of Hebb and homeostasis. *Philos. Trans. R. Soc. London Ser. B* **372**, 20160258 (2017). doi: [10.1098/rstb.2016.0258](https://doi.org/10.1098/rstb.2016.0258); pmid: 28093556
85. C. Lance-Jones, L. Landmesser, Pathway selection by embryonic chick motoneurons in an experimentally altered environment. *Proc. R. Soc. London Ser. B* **214**, 19–52 (1981). doi: [10.1098/rspb.1981.0080](https://doi.org/10.1098/rspb.1981.0080); pmid: 6121329
86. J. Lu, J. C. Tapia, O. L. White, J. W. Lichtman, The interscutalaris muscle connectome. *PLOS Biol.* **7**, e32 (2009). pmid: 19209956
87. J. Caron, V. Ruta, L. F. Abbott, R. Axel, Random convergence of olfactory inputs in the *Drosophila* mushroom body. *Nature* **497**, 113–117 (2013). doi: [10.1038/nature12063](https://doi.org/10.1038/nature12063); pmid: 23615618
88. F. Li *et al.*, The connectome of the adult *Drosophila* mushroom body provides insights into function. *eLife* **9**, e62576 (2020). doi: [10.7554/eLife.62576](https://doi.org/10.7554/eLife.62576); pmid: 33315010
89. Z. Zheng *et al.*, A complete electron microscopy volume of the brain of adult *Drosophila melanogaster*. *Cell* **174**, 730–743.e22 (2018). doi: [10.1016/j.cell.2018.06.019](https://doi.org/10.1016/j.cell.2018.06.019); pmid: 30033368
90. S. Serizawa *et al.*, A neuronal identity code for the odorant receptor-specific and activity-dependent axon sorting. *Cell* **127**, 1057–1069 (2006). doi: [10.1016/j.cell.2006.10.031](https://doi.org/10.1016/j.cell.2006.10.031); pmid: 17129788
91. T. McLaughlin, C. L. Torborg, M. B. Feller, D. O'Leary, Retinotopic map refinement requires spontaneous retinal waves during a brief critical period of development. *Neuron* **40**, 1147–1160 (2003). doi: [10.1016/S0896-6273\(03\)00790-6](https://doi.org/10.1016/S0896-6273(03)00790-6); pmid: 14687549
92. M. Z. Lin, M. J. Schnitzer, Genetically encoded indicators of neuronal activity. *Nat. Neurosci.* **19**, 1142–1153 (2016). doi: [10.1038/nrn.4359](https://doi.org/10.1038/nrn.4359); pmid: 27571193
93. C. K. Kim, A. Adhikari, K. Deisseroth, Integration of optogenetics with complementary methodologies in systems neuroscience. *Nat. Rev. Neurosci.* **18**, 222–235 (2017). doi: [10.1038/nrn.2017.15](https://doi.org/10.1038/nrn.2017.15); pmid: 28303019
94. L. Luo, E. M. Callaway, K. Svoboda, Genetic dissection of neural circuits: A decade of progress. *Neuron* **98**, 256–281 (2018). doi: [10.1016/j.neuron.2018.03.040](https://doi.org/10.1016/j.neuron.2018.03.040); pmid: 29673479
95. W. M. Cowan, The emergence of modern neuroanatomy and developmental neurobiology. *Neuron* **20**, 413–426 (1998). doi: [10.1016/S0896-6273\(00\)80985-X](https://doi.org/10.1016/S0896-6273(00)80985-X); pmid: 9539119
96. T. Lee, L. Luo, Mosaic analysis with a repressible cell marker for studies of gene function in neuronal morphogenesis. *Neuron* **22**, 451–461 (1999). doi: [10.1016/S0896-6273\(00\)80701-1](https://doi.org/10.1016/S0896-6273(00)80701-1); pmid: 10197526
97. G. Feng *et al.*, Imaging neuronal subsets in transgenic mice expressing multiple spectral variants of GFP. *Neuron* **28**, 41–51 (2000). doi: [10.1016/S0896-6273\(00\)00084-2](https://doi.org/10.1016/S0896-6273(00)00084-2); pmid: 11086982
98. H. Zong, J. S. Espinosa, H. H. Su, M. D. Muzumdar, L. Luo, Mosaic analysis with double markers in mice. *Cell* **121**, 479–492 (2005). doi: [10.1016/j.cell.2005.02.012](https://doi.org/10.1016/j.cell.2005.02.012); pmid: 15882628
99. J. Livet *et al.*, Transgenic strategies for combinatorial expression of fluorescent proteins in the nervous system. *Nature* **450**, 56–62 (2007). doi: [10.1038/nature06293](https://doi.org/10.1038/nature06293); pmid: 17972876
100. A. Nern, B. D. Pfeiffer, G. M. Rubin, Optimized tools for multicolor stochastic labeling reveal diverse stereotyped cell arrangements in the fly visual system. *Proc. Natl. Acad. Sci. U.S.A.* **112**, E2967–E2976 (2015). doi: [10.1073/pnas.1506763112](https://doi.org/10.1073/pnas.1506763112); pmid: 25964354
101. J. G. White, E. Southgate, J. M. Thomson, S. Brenner, The structure of the nervous system of the nematode *Caenorhabditis elegans*. *Philos. Trans. R. Soc. London Ser. B* **314**, 1–340 (1986). doi: [10.1098/rstb.1986.0056](https://doi.org/10.1098/rstb.1986.0056); pmid: 22462104
102. S. Y. Takemura *et al.*, A visual motion detection circuit suggested by *Drosophila* connectomics. *Nature* **500**, 175–181 (2013). doi: [10.1038/nature12450](https://doi.org/10.1038/nature12450); pmid: 23925240
103. J. S. Kim *et al.*, Space-time wiring specificity supports direction selectivity in the retina. *Nature* **509**, 331–336 (2014). doi: [10.1038/nature13240](https://doi.org/10.1038/nature13240); pmid: 24805243
104. W. C. Lee *et al.*, Anatomy and function of an excitatory network in the visual cortex. *Nature* **532**, 370–374 (2016). doi: [10.1038/nature17192](https://doi.org/10.1038/nature17192); pmid: 27018655
105. D. G. C. Hildebrand *et al.*, Whole-brain serial-section electron microscopy in larval zebrafish. *Nature* **545**, 345–349 (2017). doi: [10.1038/nature22356](https://doi.org/10.1038/nature22356); pmid: 28489821
106. L. K. Scheffer *et al.*, A connectome and analysis of the adult *Drosophila* central brain. *eLife* **9**, e57443 (2020). doi: [10.7554/eLife.57443](https://doi.org/10.7554/eLife.57443); pmid: 32880371
107. L. F. Abbott *et al.*, The mind of a mouse. *Cell* **182**, 1372–1376 (2020). doi: [10.1016/j.cell.2020.08.010](https://doi.org/10.1016/j.cell.2020.08.010); pmid: 32946777
108. I. R. Wickersham *et al.*, Monosynaptic restriction of transsynaptic tracing from single, genetically targeted neurons. *Neuron* **53**, 639–647 (2007). doi: [10.1016/j.neuron.2007.01.033](https://doi.org/10.1016/j.neuron.2007.01.033); pmid: 17329205
109. E. M. Callaway, L. Luo, Monosynaptic circuit tracing with glycoprotein-deleted rabies viruses. *J. Neurosci.* **35**, 8979–8985 (2015). doi: [10.1523/JNEUROSCI.0409-15.2015](https://doi.org/10.1523/JNEUROSCI.0409-15.2015); pmid: 26085623

110. L. Lo, D. J. Anderson, A Cre-dependent, anterograde transsynaptic viral tracer for mapping output pathways of genetically marked neurons. *Neuron* **72**, 938–950 (2011). doi: [10.1016/j.neuron.2011.12.002](https://doi.org/10.1016/j.neuron.2011.12.002); pmid: [22196330](https://pubmed.ncbi.nlm.nih.gov/22196330/)
111. B. Zingg *et al.*, Aav-mediated anterograde transsynaptic tagging: Mapping corticocollicular input-defined neural pathways for defense behaviors. *Neuron* **93**, 33–47 (2017). doi: [10.1016/j.neuron.2016.11.045](https://doi.org/10.1016/j.neuron.2016.11.045); pmid: [27989459](https://pubmed.ncbi.nlm.nih.gov/27989459/)
112. M. Talay *et al.*, Transsynaptic mapping of second-order taste neurons in flies by trans-tango. *Neuron* **96**, 783–795.e4 (2017). doi: [10.1016/j.neuron.2017.10.011](https://doi.org/10.1016/j.neuron.2017.10.011); pmid: [29107518](https://pubmed.ncbi.nlm.nih.gov/29107518/)
113. T. H. Huang *et al.*, Tracing neuronal circuits in transgenic animals by transneuronal control of transcription (*TRACT*). *eLife* **6**, e32027 (2017). doi: [10.7554/eLife.32027](https://doi.org/10.7554/eLife.32027); pmid: [29231171](https://pubmed.ncbi.nlm.nih.gov/29231171/)
114. L. Petreanu, D. Huber, A. Sobczyk, K. Svoboda. Channelrhodopsin-2-assisted circuit mapping of long-range callosal projections. *Nat. Neurosci.* **10**, 663–668 (2007). doi: [10.1038/nrn1891](https://doi.org/10.1038/nrn1891); pmid: [17435752](https://pubmed.ncbi.nlm.nih.gov/17435752/)
115. M. London, M. Häusser, Dendritic computation. *Annu. Rev. Neurosci.* **28**, 503–532 (2005). doi: [10.1146/annurev.neuro.28.061604.135703](https://doi.org/10.1146/annurev.neuro.28.061604.135703); pmid: [16033324](https://pubmed.ncbi.nlm.nih.gov/16033324/)
116. N. M. van Strien, N. L. Cappaert, M. P. Witter, The anatomy of memory: An interactive overview of the parahippocampal-hippocampal network. *Nat. Rev. Neurosci.* **10**, 272–282 (2009). doi: [10.1038/nrn2614](https://doi.org/10.1038/nrn2614); pmid: [19300446](https://pubmed.ncbi.nlm.nih.gov/19300446/)
117. L. Luo, J. G. Flanagan, Development of continuous and discrete neural maps. *Neuron* **56**, 284–300 (2007). doi: [10.1016/j.neuron.2007.10.014](https://doi.org/10.1016/j.neuron.2007.10.014); pmid: [17964246](https://pubmed.ncbi.nlm.nih.gov/17964246/)

ACKNOWLEDGMENTS

I thank W. Allen, T. Clandinin, L. Fei-Fei, M. Feller, J. Kebschull, J. Lui, J. Ren, M. Scanziani, A. Shuster, L. Stryer, and M. Wagner for helpful critiques and Taylor & Francis Group, LLC for granting permission to adapt figures from *Principles of Neurobiology* (5). **Funding:** This work was supported by the National Institutes of Health, the National Science Foundation, the Howard Hughes Medical Institute, and the Wu Tsai Neurosciences Institute at Stanford University. **Competing interests:** The author declares no competing interests.

10.1126/science.abg7285

Architectures of neuronal circuits

Liqun Luo

Science, 373 (6559), eabg7285. • DOI: 10.1126/science.abg7285

General principles of neuronal communication

The explosion of modern technologies over the past years has produced vast amounts of knowledge about the molecular, anatomical, and physiological properties of individual neurons. However, individual neurons do not function in isolation; they interact with each other in circuits to process information. Luo attempted to synthesize different types of investigations to extract generalized principles underlying how neurons communicate with each other through specific patterns of synaptic connectivity. Although different elements of the projection motifs and architectural plans have been discussed separately in the past, these motifs and architectures can now be brought together in the same intellectual framework. —PRS

View the article online

<https://www.science.org/doi/10.1126/science.abg7285>

Permissions

<https://www.science.org/help/reprints-and-permissions>

Use of think article is subject to the [Terms of service](#)

Science (ISSN) is published by the American Association for the Advancement of Science. 1200 New York Avenue NW, Washington, DC 20005. The title *Science* is a registered trademark of AAAS.

Copyright © 2021 The Authors, some rights reserved; exclusive licensee American Association for the Advancement of Science. No claim to original U.S. Government Works