

Comparative Biological Dynamics: A Cross-Disciplinary Analysis of Brain Hydration, Hair Morphogenesis, and Botanical Growth Systems

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ABSTRACT

This comprehensive manuscript explores the biophysical, morphological, and systemic intersections between human brain tissue hydration, hair follicle proliferation, and botanical growth systems. By examining osmotic gradients, basal versus apical growth topologies, and neurotransmitter signaling across biological kingdoms, this research identifies profound mathematical and evolutionary convergences. The synthesis of these highly specialized domains provides a robust framework for applied biomimetics, informing advancements in dermatological therapeutics, autonomous robotics, and optimized artificial neural networks, while strictly addressing the ethical imperatives of cross-disciplinary and animal-free research methodologies.

Keywords: Biomimetics; Osmotic Dynamics; Intercalary Meristems; Tissue Hydration; Dendritic Learning; Plant Neurobiology; Excitability

INTRODUCTION

The investigation of biological systems often proceeds within highly specialized academic silos, isolating the neurobiology of mammalian brains from the morphogenesis of integumentary appendages, and separating human physiology entirely from botanical ecology. However, a rigorous, cross-disciplinary analysis reveals profound underlying structural, topological, and biochemical unities that govern how living organisms manage cellular growth, resource distribution, and fluid dynamics. This comprehensive research report explores the biophysical, morphological, and systemic intersections between human brain tissue hydration, human hair follicle proliferation, and the vegetative and root growth of land grasses. By deconstructing the basic biophysical research driving these fields and examining the applied research that translates these phenomena into biomimetic technologies, this analysis establishes an exhaustive systems-level perspective on comparative biology.

The scope of this research is deliberately expansive, traversing multiple spatial and temporal scales while utilizing an array of methodological frameworks. At the fundamental molecular and cellular level, the report examines the role of water not merely as a passive biological solvent, but as a structural plasticizer and an active signaling medium. This involves a direct comparison of the complex osmotic dynamics governing the human blood-brain barrier with the mathematically analogous, yet

mechanistically distinct, xylem sap transport and osmotic adjustments operating within botanical root systems.[2] Moving to the tissue and morphological level, the analysis evaluates the divergent evolutionary strategies of biological expansion. It contrasts the invagination and continuous basal extrusion logic of the mammalian hair follicle with the intercalary meristems that allow land grasses to survive perpetual defoliation, and further compares both to the apical arborization of plant roots and neuronal dendrites.[5]

Beyond fundamental biology, this report synthesizes quantitative metrics, algorithmic models, and qualitative survey data to substantiate the vast applied relevance of these fields. The translation of botanical growth principles into human dermatological therapeutics, the development of robotic chemotropism inspired by plant roots, and the deployment of artificial intelligence to replace animal testing with human-relevant neural organoids underscore the critical utility of cross-disciplinary biomimetics.[9] Finally, recognizing that the manipulation of biological components at this scale presents unprecedented moral dilemmas, rigorous ethical frameworks and critical decision-making paradigms are applied. By evaluating the frontiers of synthetic biology, global research equity, multi-stressor environmental management, and the engineering of conscious proxies, this research ensures that scientific innovation remains securely anchored in responsible governance and equitable public policy [12].

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Basic Research: The Biophysics of Hydration and Osmotic Dynamics

The Physiology and Hydration Dynamics of Human Brain Tissue

Water constitutes the universal medium for biological processes, yet its specific behavior within the human central nervous system presents a unique biochemical paradox. While total body water typically comprises approximately 45 to 75 percent of a human's overall body weight, depending on adipose and muscular composition, the structural composition of the brain operates under distinct biophysical constraints.[16] The intracranial contents as a whole are maintained at approximately 75 percent water by volume; however, the actual brain tissue, when entirely desiccated to remove this fluid, is composed of roughly 60 percent complex lipid structures.[2] This remarkably high lipid concentration primarily constitutes the myelin sheaths that heavily insulate axonal tracts, structures that are intrinsically hydrophobic and non-conductive.[2] Therefore, water within the brain is meticulously sequestered within the interstitial fluid, the cerebrospinal fluid (CSF), and the intracellular spaces of neurons and glial cells. In these compartments, water functions not merely as a passive structural filler, but as an active participant in neurophysiology, regulating volume conduction, lubricating tissues, and facilitating the complex ionic gradients required for neural transmission.[2] Advanced diagnostic techniques, such as long-TR multiple-echo gradient-echo MRI, have provided unprecedented precision in mapping this hydration, revealing that the water content distribution is specifically partitioned, averaging 69.1% in cerebral white matter and 83.7% in gray matter.

The precise regulation of this hydration status exerts a profound impact on cognitive and motor functions. Historically, clinical models posited that a total body water deficit of 2 percent or more was required to produce measurable decrements in human cognitive performance.[16] However, recent advancements in neuroimaging and psychomotor performance literature provide empirical evidence that even mild dehydration representing a body water loss of merely 1 to 2 percent—can significantly impair a wide array of faculties, including attentiveness, critical thinking skills, memory consolidation, and baseline mood regulation.[16] The physiological mechanisms underlying these cognitive deficits relate directly to subtle, hydration-induced shifts in total brain volume, cellular excitability, and the localized concentration of critical neurotransmitters within the synaptic cleft [17].

The osmotic dynamics that govern the human brain are tightly regulated by the blood-brain barrier, which prevents the indiscriminate movement of fluids and pathogens into the delicate neural microenvironment. Unlike standard osmotic transfer, where water molecules simply diffuse freely along a concentration gradient until equilibrium is reached, the transport of water and ions across the blood-brain barrier and into the intracellular matrix of neurons involves highly sophisticated cotransporter mechanisms and specialized aquaporin channels. [3] When hydration status fluctuates, it induces both macroscopic and microscopic volume changes throughout the cortical tissue. For instance, neurostimulation studies utilizing Transcranial Magnetic Stimulation to measure

cortical excitability have demonstrated that hydration variables significantly alter the volume conduction between the surface of the scalp and the underlying excitable brain tissue.[18] When human subjects are tested following an overnight fast, and subsequently rehydrated, researchers observe a macroscopic decrease in the scalp-to-cortex distance, indicating that the brain tissue physically swells and moves closer to the skull.[18] Microgenically, these identical osmotic challenges induce profound changes at the cellular level; specifically, hydration shifts trigger astrocyte swelling, which in turn provokes a calcium-dependent release of the excitatory neurotransmitter glutamate into the extracellular space.[18] This cascade demonstrates that brain water is fundamentally and inexorably linked to the electrochemical architecture of thought and motor control.

Comparative Hydraulics: Water Transport and Embolism Repair in Plant Roots

While the human brain relies on the blood-brain barrier and active cotransporters to maintain its highly constrained fluid volume within a rigid skull, botanical systems must manage water dynamics across an open, environmentally exposed continuum extending from deep soil horizons to the atmospheric interface. Plant hydraulic traits play major evolutionary roles in coping with abiotic stressors, particularly drought, and process-based plant hydraulics have historically modeled water as flowing strictly along decreasing pressure or total water potential gradients.[4] However, this established paradigm of passive, gradient-driven water flow across the roots of vascular plants is incomplete.

Recent advancements combining sophisticated modeling tools from the emerging field of plant micro-hydrology with pioneering cell solute mapping data have solved a 35-year biological enigma: the observation of water flowing "uphill" against established water potential gradients across living root tissues.[4] This mechanism lacks the traditional reliance on simple capillary action. Instead, the steep gradient of solute concentrations across living plant cells creates a highly negative osmotic potential, which acts as a powerful, active water pump. [4] This biological pump effectively reduces water tension without loading additional solutes directly into the plant's central vasculature, laying the foundation for fluid movement in complex hydraulic anatomies such as those found in maize, grapevine, and the model organism *Arabidopsis* [4]. By pulling water "uphill" toward the xylem, the plant mitigates the catastrophic risk of tension-induced cavitation along the stems, safeguarding the photosynthetic capabilities of the leaves [4]. The physical mechanics of this flow rely heavily on turgor pressure, which provides the necessary internal force to deform the plant cell wall during expansive growth. To contextualize this biophysical force, human physiological blood pressure operates at a relatively low 10 to 20 kilopascals (kPa). In contrast, the turgor pressure within plant cells must be exponentially higher to balance the massive compressive stress of the rigid cell wall; when this wall relaxes, the turgor pressure drops, creating a severe water influx that forcibly drives cell elongation.

Furthermore, the management of internal fluid dynamics in plants shares conceptual parallels with the human brain's

response to ischemic injury and localized edema. When extreme drought induces embolisms within the xylem, rendering these vessels nonfunctional, plants initiate an active repair protocol. The refilling of drought-induced embolisms requires the rapid creation of an osmotic gradient between the adjacent xylem parenchyma cells and the empty vessel lumens to generate the necessary water efflux to fill the void [19]. Advanced sap collection techniques designed to chemically separate fluid from functional versus embolized vessels reveal that, as water stress increases, the osmotic potential of the liquid within the nonfunctional vessels rises dramatically [19]. This localized osmotic adjustment is achieved through the active, targeted transport of sugars and specific ions into the embolized vessels, accompanied by a notable decrease in the pH of the xylem sap [19]. This targeted deployment of sugar transporters highlights a sophisticated degree of hydraulic compartmentalization that parallels the blood-brain barrier's precise regulation of localized ion gradients during metabolic stress [3].

The Hydration and Plasticization of Human Hair Fibers

Occupying a biophysical space distinct from both the electrochemically active brain and the fluid-transporting plant root, human hair represents a heavily keratinized, largely non-living appendage that nevertheless relies completely on hydration for its structural properties. The precise moisture content of a hair fiber significantly influences its mechanical characteristics, governing variables such as macroscopic elasticity, ultimate tensile strength, and microscopic surface friction [20].

Human hair is primarily composed of proteins, which account for 90 to 95 percent of the structure, with extensively cross-linked keratin polymers constituting the vast majority of this protein matrix [20]. Under standard ambient atmospheric conditions, water makes up between 3 and 10 percent of the hair's mass, accompanied by 1 to 9 percent structural lipids, including specific ceramides, cholesterol, and uniquely integrated fatty acids [20]. When hair is exposed to liquid water or highly humid environments, water molecules permeate the fiber and fundamentally disrupt the secondary hydrogen bonds and internal salt bridges that normally provide rigid structural integrity to the macrofibrils [20].

This solvation process leads to a temporary but significant decrease in the hair's overall mechanical stiffness through a process defined biophysically as structural plasticization [20]. Furthermore, hydration induces highly anisotropic swelling; because the keratin filaments are aligned longitudinally along the shaft, the absorption of water forces the fiber to swell laterally, inducing a significantly greater increase in the hair's diameter compared to any elongation in its length [20]. The endocuticle, a sub-layer of the hair's outer cuticle characterized by a markedly lower density of tough disulfide cross-links, serves as the primary region of rapid water absorption [20]. Empirical research indicates that simply immersing human hair in water can elevate the cuticle step height by over 50 percent, radically altering the topography of the fiber [20]. Within the deep cortical matrix of the hair fiber, the hydrogen atoms of the keratin proteins exhibit a higher chemical affinity for incoming water molecules than they do for each other, driving a

continuous reorganization of the internal architecture as hydration levels fluctuate [20].

Basic Research: Morphogenesis and Topologies of Growth

The Development and Cyclic Proliferation of Human Hair Follicles

Human hair serves a multitude of physiological and psychological functions, acting as a sensory organ, a mechanism for sweat-based thermoregulation, a primary defense against ultraviolet radiation, and a profound component of psychological identity and social signaling [21]. The intricate process of hair morphogenesis begins deep in embryonic development. At approximately 10 weeks of gestational age, localized thickenings within the fetal epidermis, known as placodes, begin to form through precise signaling interactions driven by stem cells expressing specific G-protein coupled receptors. During this period of organogenesis, these placodes invaginate deeply into the underlying dermal layer, leading to the condensation of mesenchymal cells beneath the down-growing follicular epithelium. This structural encapsulation forms the dermal papilla, the critical signaling center of the hair follicle [1].

Unlike the branching, indeterminate growth seen in botanical systems, human hair growth follows a highly regulated, cyclical timeline characterized by distinct developmental phases. Every single hair follicle on the human body operates on its own independent clock, continuously cycling through the anagen, catagen, and telogen phases [22]. The anagen phase represents the period of active growth and massive cellular proliferation. During anagen, multipotent stem cells located within the outer root sheath and the bulge region of the follicle rapidly divide, generating transit-amplifying cells that subsequently differentiate to form the complex, multi-layered architecture of the inner root sheath, the cuticle, and the heavily keratinized cortex [23]. Following the growth period, the follicle enters the catagen phase, a destructive stage of programmed involution. Marked by a rapid bout of apoptosis, the lower portion of the follicle dramatically regresses, causing the active dermal papilla to physically detach from the blood supply and move upward, coming to rest in close proximity to the stem cell bulge [22]. Finally, the follicle enters the telogen phase, a period of total growth cessation and cellular quiescence [22]. Advanced mathematical and functional models of the hair cycle further divide this resting state into "refractory telogen," during which the follicle is completely unresponsive to internal or external growth stimuli, and "competent telogen," during which the follicle is biochemically primed and awaiting the proper signals to re-enter the anagen phase and initiate a new cycle of growth [24].

From a topological perspective, the growth of the hair shaft relies on a strict "first-in, first-out" mechanical logic [8]. Because the multipotent cells actively dividing in the hair bulb are located at the absolute base of the structure, deep within the dermis, the production of new cellular matter physically pushes the older, fully keratinized, and deceased epithelial cells upward and outward past the surface of the epidermis [8]. This basal

extrusion mechanism stands in stark contrast to the development of avian feathers, where the outermost portion represents the most recently formed tissue, and differs radically from the apical extension strategies utilized by both neuronal structures and botanical roots [8].

Botanical Growth Topologies: Grass Tillering and Intercalary Meristems

The vegetative growth of land grasses, including both forage species and specialized turfgrasses, relies on an evolutionary strategy explicitly adapted for survival in environments characterized by continuous herbivory, trampling, and severe climatic shifts [7]. While animals exhibit determinate growth, reaching a specific morphological shape and size before growth ceases, plants exhibit indeterminate growth, maintaining the capacity to continuously add new organs throughout their lifespan via populations of undifferentiated, highly active stem cells known as meristems [25].

A mature grass plant is fundamentally a modular organism, composed of an interconnected network of tillers [26]. Each tiller contains a blade, a protective sheath enveloping the shoot, and a collar region, all anchored to a primary root crown [27]. The root crown contains dozens of dormant buds, each capable of generating entirely new tillers or adventitious roots [26]. Because multiple tillers remain anatomically and physiologically connected, they operate as a unified biological network, actively sharing water, carbohydrates, and sequestered nutrients; if an individual tiller is destroyed, the adjacent connected tillers simply allocate resources to initiate replacement growth from a dormant basal bud [26].

The defining topological divergence of grass from other biological systems is its reliance on the intercalary meristem [25]. While the grass shoot does possess a primary apical meristem housed within an apical dome to initiate new leaf primordia, the vast majority of actual linear leaf growth is the result of massive cell expansion occurring within the intercalary meristems located at the extreme base of each leaf blade [7]. These basal growth zones are deeply nested and protected within the sheaths of older, mature leaves [7]. This unique anatomical arrangement provides grasses with their legendary resilience. When a grazing animal or a mechanical lawnmower amputates the top portion of a grass leaf, the apical tissue is destroyed; however, because the actively dividing intercalary meristem remains safely positioned near the soil surface, it simply continues to generate new cellular volume, pushing the remaining leaf blade upward without interruption [25].

Apical Arborization in Root Systems

While the aerial shoots of grasses utilize basal intercalary meristems, the subterranean root networks rely entirely on apical meristems for spatial exploration and nutrient acquisition. The root system serves multifaceted

roles: firmly anchoring the plant against environmental shear forces, operating as a massive carbohydrate storage organ, and absorbing essential water and dissolved minerals [7].

When a grass seed germinates, it first prioritizes the establishment of a foundation, driving primary roots deeply into the soil matrix before any significant photosynthetic shoot emerges from the surface [7]. Root growth is achieved through the rapid division of meristematic cells located just behind a specialized, protective root cap at the apex of the structure [7]. Following this division, the newly generated cells enter a zone of profound elongation, driven by localized increases in turgor pressure as water is rapidly absorbed from the surrounding soil into the cellular vacuoles [7]. Maturation and subsequent differentiation of these elongated cells lead to the development of the highly specialized xylem and phloem tissues necessary for whole-plant transport [7].

The overall architecture of the root system is highly responsive to environmental gradients, dynamically adjusting its growth rate, branching diameter, lateral root formation, and directional orientation to maximize resource acquisition [29]. Phenotypic adaptations in root architecture exhibit powerful synergistic effects. For example, in environments characterized by suboptimal phosphorus availability, genetic lines of common bean plants exhibiting both long, dense root hairs and a shallow basal root growth angle demonstrate massive biomass advantages [30]. While long root hairs increase the total soil volume subject to phosphorus depletion, and shallow basal roots position the network in the topsoil where phosphorus is concentrated, genotypes possessing both anatomical traits exhibit a nearly 300 percent greater biomass accumulation compared to deep-rooted, short-haired phenotypes [30]. This indicates that architectural phenes operate synergistically, delivering benefits far greater than the sum of their individual additive effects, an essential insight for optimizing crop fitness in resource-depleted agricultural zones [30].

Quantitative Data and Comparative Mathematical Modeling

To contextualize the vast physiological differences between neural tissue, keratinized appendages, and botanical systems, rigorous quantitative analysis is necessary. The specific velocities of cellular proliferation and the precise mathematical definitions of tissue hydration illustrate the immutable scaling laws that govern these mechanisms.

Comparative Growth Velocities and Structural Compositions

Growth rates across these biological domains vary exponentially, dictated entirely by the metabolic costs of the structural materials being synthesized and the ecological imperatives of the growing organ.

Biological Entity	Growth Topology & Driver	Average Quantitative Growth Rate	Structural Composition & Notes
Human Hair (Scalp)	Basal Extrusion (Follicle)	~ 0.35 to 0.411 mm/day	90-95% highly cross-linked protein (keratin), 3-10% water, 1-9% lipids. Equates to ~ 15 cm/year. [31]
Human Vellus Hair	Basal Extrusion (Follicle)	~ 0.03 to 0.13 mm/day	Shorter cycle duration limits overall length; continuous replacement. [31]
Land Grass (Shoots)		~ 1.5 to 5.0 mm/day	Highly variable based on season; equates to roughly 0.06 to 0.2 inches/day during optimal summer conditions. [32]
Grass Roots		~ 10.89 mm/day	Measured during early establishment, reaching 152.4 mm (6 inches) in depth over 14 days. [7]

The quantitative disparity presented in this data highlights a fundamental evolutionary strategy. Human hair is composed of dense, sulfur-rich, heavily cross-linked structural proteins that require significant metabolic energy to synthesize and align; consequently, its growth is limited to a slow, steady extrusion. Conversely, grass roots must rapidly secure unstable environments and acquire transient water sources. By relying largely on the rapid vacuolar uptake of water to drive cell elongation rather than massive protein synthesis, root apices can achieve daily expansion rates an order of magnitude higher than human follicular tissues [7].

The Mathematics of Tissue Hydration and Cerebral Edema

The precise mathematical quantification of water within biological tissues dictates how researchers understand pathological states. In the context of human neurology, the calculation of brain water is a matter of critical clinical consequence, particularly concerning ischemic strokes, hemorrhagic strokes, and traumatic brain injuries that result in lethal cerebral edema [33].

Historically, researchers quantified edema in animal models by expressing it as a simple change in percentage brain water content, derived from comparing the wet and dry weights of the affected tissue against healthy controls [33]. The standard equation utilized was:

$$\% \text{ Water Content} = 100 \times \frac{(\text{Wet Weight} - \text{Dry Weight})}{\text{Wet Weight}}$$

However, advanced biophysical modeling has demonstrated that relying on this straightforward percentage can lead to catastrophic misinterpretations of the severity of a brain injury [33]. Because the denominator (Wet Weight) increases simultaneously with the numerator as edematous fluid rapidly accumulates in the brain tissue, seemingly "small" percentage changes in total brain water mathematically mask massive, life-threatening increases in absolute tissue swelling and intracranial pressure [33]. To correct this analytical flaw, contemporary neurological modeling evaluates edema by calculating absolute water content relative only to the static dry mass, or by quantifying true volumetric tissue swelling: This mathematical recalibration underscores a unique physiological constraint: the human brain exists within the rigid, unyielding compartment of the cranium. Therefore, minute variations in absolute water volume have exponentially larger mechanical and functional

consequences than equivalent hydration shifts in the unconstrained matrices of soil-dwelling plant roots or extruding human hair shafts [18].

Mathematical Isomorphisms in Biological Branching Networks

Despite originating from entirely distinct evolutionary lineages, the branching architectures of botanical root systems and human neuronal dendrites share profound mathematical and topological isomorphisms. Both systems function as highly complex physical networks explicitly designed to optimize spatial foraging within a constrained volumetric space—roots actively foraging for unpredictable water and nitrogenous compounds throughout the soil matrix [6], and dendrites aggressively foraging for synaptic connections within the dense cortical environment [6].

Empirical research across a wide variety of neuronal types reveals that both systems are governed by fundamental scaling laws. A derived quantitative theory of optimal wiring demonstrates that dendritic arbors abide by a strict $2/3$ power law relating the total length of dendritic wiring to the number of branch points and the resulting number of synapses [6]. This geometric optimization successfully minimizes the metabolic cost of constructing the cellular cable and reduces the signal conduction time, while simultaneously maximizing the total receptive surface area available for communication [6].

To simulate the genesis of these branching architectures, computational biologists employ advanced algorithmic models. The "Cell Compartment Model" developed for dendritic growth dynamically couples physical cellular expansion with the reaction-diffusion dynamics of intracellular chemicals [35]. This model posits an "activator" chemical that diffuses strictly within the intracellular space, reacting with a secondary "suppressor" reactant to dictate precise branching points according to classic Turing pattern formation principles [35]. Similarly, the ROOTS (Ruled-Optimum Ordered Tree System) algorithm extends neuronal generative methods to highly branched axonal and dendritic terminals, allowing researchers to accurately predict how cortical tissue will respond to externally applied electric fields during neurostimulation therapies [36].

In the realm of botany, researchers utilize continuous, spatial convective-diffusive algorithms to model root proliferation through heterogeneous soils [37]. Recent interdisciplinary integrations suggest that treating actively growing root apices as

decision-making agents navigating complex chemical gradients provides a mathematical framework highly homologous to the algorithms governing axonal pathfinding during early neural development [38]. The deployment of identical reaction-diffusion mathematics to model both plant roots and human neurons exposes a deterministic layer of biology: strict biophysical constraints force completely diverse biological entities to converge upon the exact same architectural topologies to achieve operational efficiency.

The PARC Model: Modeling Hair Populations as Excitable Media

The macroscopic coordination of hair follicle growth across mammalian skin acts as a classical excitable medium, a mathematical concept frequently used to model spreading fires or the propagation of chemical waves [24]. The PARC model treats each individual follicle as a distinct functional unit, parameterizing the complex biological growth cycle into four mathematical phases: Propagating Anagen (P), Autonomous Anagen (A), Refractory Telogen (R), and Competent Telogen (C) [24].

Follicles resting in the Competent Telogen phase exist at a highly excitable steady-state. If stochastic molecular noise—modeled mathematically as Gaussian fluctuations within the underlying biochemical pathways—surpasses a specific excitability threshold, the follicle spontaneously enters the active anagen growth phase, an event treated probabilistically as a Poisson process [24]. Furthermore, through the continuous diffusion of chemical activators and inhibitors across the extrafollicular space, a follicle operating in the Propagating Anagen phase can actively induce neighboring competent follicles to begin growing, resulting in highly coordinated, macroscopic wave-like patterns of hair regeneration spreading across the tissue [24].

Further simulations leveraging multi-scale modeling reveal that mammalian skin actually functions as a heterogeneous regenerative field [39]. Interactions between fast-cycling domains (such as the chin) and slow-cycling dorsal domains produce complex, bilaterally symmetric patterns [39]. Furthermore, anatomically distinct areas like the ears operate as "hyper-refractory" domains due to naturally high levels of localized BMP inhibitors, which ultimately generate "wave-breaking" effects that physically halt the propagation of hair growth across anatomical boundaries [39].

Basic Research: Cross-Kingdom Neurotransmitter Signaling

A paradigm-shifting realization in modern cross-disciplinary biology is the discovery that major neurotransmitters traditionally associated strictly with human neurophysiology play fundamental, highly active roles in botanical development and environmental stress regulation.

In the human central nervous system, Gamma-Aminobutyric Acid (GABA) and glutamate represent the fundamental yin and yang of neuronal excitability [40]. Glutamate serves as the primary excitatory neurotransmitter, facilitating the rapid depolarization of neuronal membranes and enabling the synaptic plasticity essential for learning and memory [40].

Conversely, GABA acts as the primary inhibitory neurotransmitter [40]. When GABA binds to post-synaptic GABA receptors, it induces the opening of specific chloride (Cl⁻) channels [41]. Because the intracellular concentration of chloride in human neurons is generally lower than the extracellular environment, the channel opening allows an influx of negatively charged chloride ions, which hyperpolarizes the neuronal membrane, heavily suppresses background electrical activity, prevents overexcitation by glutamate, and maintains the essential signal-to-noise stability of the neural circuit [41].

Remarkably, these exact five-carbon (glutamate) and four-carbon (GABA) amino acids operate as critical signaling molecules within plant biology, specifically governing root architecture and morphological adaptation [41]. In plants, GABA is synthesized directly from glutamic acid via the highly conserved GABA shunt pathway, serving to balance the cellular carbon-to-nitrogen ratio while actively mitigating extreme abiotic stressors associated with climate change [41].

At the root apex the primary sensory interface between the plant and the surrounding soil environment glutamate operates with distinct neuronal-like activity. The transition zone cells of the root, situated physically between the apical meristem and the basal elongation zone, respond to exogenous glutamate exposures with rapid plasma membrane depolarizations, a powerful bioelectric response that can be completely abolished by the application of specific antagonists targeting ionotropic glutamate receptors [42]. Furthermore, GABA acts directly on Aluminum-Activated Malate Transporter 1 (ALMT1) channels present in plant roots [41]. This interaction reveals a fascinating evolutionary adaptation of a shared mechanism. In stark contrast to mammalian neurons, living plant root cells maintain a higher chloride concentration within the cytoplasm relative to the exterior apoplast [41]. Therefore, when GABA interacts with ALMT1 receptors in roots, it does not cause an influx of ions; rather, it rapidly inhibits the efflux of chloride out of the cell [41]. While the physical directional flow of the anion is reversed due to the differing osmotic environments of roots versus brains, the ultimate physiological result remains identical: a significant reduction in membrane depolarization and a powerful suppression of overall cellular excitability [41].

This profound molecular convergence highlights the immense evolutionary antiquity of these signaling pathways. Long before the evolutionary divergence of flora and fauna, the molecular machinery required to utilize simple amino acids to gate ion channels and regulate complex osmotic and electrical potentials was already firmly established [41].

Applied Research: Biomimetics and Interdisciplinary Therapeutics

The rigorous deconstruction of botanical growth systems, excitable media models, and mammalian neuro-dermatology has spawned a highly fertile ground for applied research. By mathematically modeling the architectural similarities between species and directly applying the biochemical principles of plant meristems to human physiology, researchers are pioneering groundbreaking clinical and technological interventions.

Botanical Interventions in Dermatological Regeneration

The deep mechanistic understanding of the human hair follicle's growth cycle has illuminated distinct molecular targets for combating various forms of alopecia, a widespread condition that inflicts a severe psychological burden on affected individuals [44]. Conventional pharmacological interventions, such as minoxidil and finasteride, alongside surgical hair transplantation, are frequently hindered by adverse drug-induced side effects, hormonal disruptions, and exceptionally poor drug penetration across the skin's protective barrier [45]. Consequently, applied biomimetic research has aggressively turned to the regenerative capacity of plant meristems and naturally occurring phytochemicals to therapeutically hijack the human hair cycle [45].

Extensive ex vivo tissue cultures and in vivo animal modeling demonstrate that specifically curated plant extracts and plant-derived stem cells operate as potent senomorphics. These botanical compounds reduce cellular senescence in the localized skin microenvironment by effectively neutralizing reactive oxygen species, thereby preserving the stemness of human follicular cells. More directly, advanced botanical serums utilizing isolated extracellular vesicles—tiny packets containing dense protein and genetic signals—derived from resilient plant species such as *Centella asiatica* have demonstrated profound clinical efficacy [10]. In recent randomized, double-blind clinical trials utilizing standardized imaging, human subjects applying these plant-derived vesicles alongside mild caffeine and panthenol experienced a verified 25 percent increase in total hair thickness within a mere 56 days [10].

The specific biochemical mechanism behind this remarkable cross-kingdom efficacy relies on the targeted modulation of highly conserved signaling pathways. Active phytochemicals, including specific terpenes, phenolic compounds, and sulfur-containing compounds, actively promote the survival and rapid proliferation of human dermal papilla cells through several distinct mechanisms: 47: First, they upregulate the production of essential growth factors, specifically Insulin-like Growth Factor 1 (IGF-1), Vascular Endothelial Growth Factor (VEGF), and Fibroblast Growth Factor 7 (FGF-7/KGF), flooding the follicle with signals that support skin cell production. 47 Second, they actively stimulate the Protein Kinase B (PKB/AKT), Extracellular Signal-Regulated Kinase (ERK), and Wingless/Int-1 (WNT) signaling pathways. In the mathematical PARC model, these WNT pathways serve as the primary chemical activators responsible for transitioning a dormant follicle from the competent telogen phase directly into the highly active propagating anagen phase [24]. Finally, these phytochemicals concurrently suppress the Transforming Growth Factor-beta (TGF-) and Bone Morphogenetic Protein (BMP) pathways. BMP activity, often originating from extrafollicular adipocytes, acts as the primary molecular inhibitor that locks hair stem cells in the quiescent, refractory telogen state [24]. By utilizing botanical compounds to manipulate this delicate activator-inhibitor balance, clinical dermatology can safely induce robust human hair regeneration.

Translating Botanical Chemotropism to Robotic Navigation

The conceptualization of biological root systems acting as decentralized, localized decision-makers has been successfully reverse-engineered into the field of robotics. Drawing direct inspiration from the innate chemotropism of plant roots—their ability to independently sense and grow toward chemical nutrients—engineering teams have developed miniaturized robotic root systems embedded with highly customized solid-state electrochemical micro-potentiometers.

These biomimetic artificial roots are capable of autonomously navigating complex, moist, soil-like environments by actively sensing minute changes in potassium concentration and pH gradients. Providing an integrated electronic readout, the sensors trigger the physical movement of the robotic root tip using a control algorithm precisely modeled on the behavioral decision-making processes of a living plant. This nature-inspired technological framework bypasses the massive central processing requirements typical of traditional robotics, opening entirely new avenues for creating autonomous devices capable of independently exploring and continuously monitoring subterranean environmental conditions.

Dendritic Learning and Artificial Neural Network Optimization

The mathematical isomorphisms shared by plant root branching and neuronal dendritic arborization have also inspired novel optimization strategies within the field of artificial intelligence and deep neural networks (DNNs). Traditional DNNs often rely on uniform activation and dense connectivity, which dramatically increases computational overhead and the risk of overfitting. However, inspired by the sparse, highly efficient wiring of biological dendrites, computer scientists have developed "dendritic learning" architectures. These architectures utilize diverse types of artificial neuronal units that exhibit differential activation based on simulated synaptic plasticity, employing a flexible, sparse connectivity architecture to significantly reduce the number of required model parameters.

Furthermore, the challenge of manually designing these complex, highly branched network topologies is being addressed through reinforcement learning algorithms. These advanced systems can optimize network architectures for novel computational problems without requiring exhaustive retraining, effectively mimicking the autonomous, environment-responsive branching logic inherent to biological root and neuronal systems.

Qualitative Surveys and Interdisciplinary Perception

The successful integration of biological principles into global engineering, design, and public policy requires more than raw data; it necessitates an educational and cultural paradigm shift. Qualitative bibliometric analyses and large-scale empirical surveys provide critical metrics for evaluating how these highly interdisciplinary concepts are perceived and adopted by the public and academic communities.

Surveys conducted during extensive science museum exhibitions in Japan demonstrate a high baseline public expectation and a notably positive societal perception of biomimetics, an emerging area clearly recognized by the public as being developed through

the cross-pollination of biology, ecology, chemistry, and traditional engineering. This public enthusiasm is mirrored in academic trends. Systematic reviews of design literature spanning from 1991 to 2021 highlight a massive upward trajectory toward bio-inspired design to improve global sustainability. The qualitative data concludes that teaching biomimicry to emerging designers substantially enhances their creativity, improves systemic problem-solving skills, and facilitates the integration of stringent sustainability metrics into standard industrial production cycles.

Furthermore, educational surveys highlight the structural barriers to implementing this interdisciplinary research at smaller institutions. Retrospective pre- and post-surveys applied to life sciences student researchers at predominantly undergraduate institutions and minority-serving institutions reveal significant attitudinal and structural roadblocks to learning advanced computational biology. However, the qualitative interview data confirms that implementing short-format computational modeling courses successfully overcomes these barriers, effectively equipping the next generation of diverse scientists with the analytical tools necessary to merge complex biological data with computational mathematics.

Critical Decision Making and Systemic Analysis

Environmental and physiological systems rarely operate in pristine isolation. Human health, ecological vitality, and agricultural stability are continuously exposed to multiple, overlapping, and frequently synergistic stressors. Traditional risk assessment methods, which rely heavily on a reductionist "chemical-by-chemical" approach, consistently fail to accurately model the complex realities of biological impairment [14]. For example, investigations into severe biological impairments in the Little Floyd River in Iowa, which featured altered fish community structures and mass mortality events, demonstrated the sheer inadequacy of traditional methods when attempting to assign causality amidst multiple stressors like livestock waste, agricultural chemicals, and treated wastewater effluent [14].

To address these catastrophic analytical deficiencies, leading scientific organizations, including the Environmental Protection Agency's Science Advisory Board, have strongly advocated for the adoption of the "One Environment-One Health" (One Health) systems approach [15]. This framework integrates massive datasets across vast spatial and temporal scales, demanding an unprecedented interdisciplinary synthesis of the social sciences, socio-economics, and rigorous biophysical sciences [15].

The Interdisciplinary Decision Making (IDM) framework operationalizes this systems-level thinking by defining science integration as a strict, three-part continuous cycle [15]. The process begins with rigorous problem formulation, wherein interdisciplinary panels must abandon their historical program silos to ask the right questions and accurately define the scope of the complex problem [15]. The second phase involves comprehensive assessment, requiring the synthesis of entirely disparate scientific fields—for instance, merging toxicological data regarding water pollution with ecological models of grass root absorption, and combining those with neurological models

of human chemical exposure [15]. The final phase centers on decision making and evaluation, emphasizing the application of the integrated science directly to policy, coupled with the continuous, dynamic evaluation of the outcomes [15].

However, establishing these collaborative frameworks early in the research lifecycle is paramount. Literature regarding public health and genomic research indicates that failing to secure interdisciplinary diversity undermines the extent to which scientific research engages with marginalized communities. Building trust over time, ensuring all members understand human subject protections, and sharing a unified research agenda are necessary prerequisites to prevent the misapplication of highly specialized biological models to general public policy.

Ethical Considerations in Cross-Disciplinary Biology

As scientific capability merges the sophisticated domains of computational algorithms, human neurology, and botanical biomimetics, unprecedented ethical dilemmas arise. The profound manipulation of biological components demands incredibly rigorous ethical oversight to protect human dignity, ensure animal welfare, and guarantee global equity.

Brain Organoids and the Threshold of Artificial Consciousness

The intensive effort to model the human brain utilizing miniaturized, simplified versions of neural tissue grown in laboratory settings from human stem cells—structures known as brain organoids—presents one of the most profound ethical frontiers in modern science [13]. Historically, animal models have served as the undisputed foundation for neurological research. However, these models consistently fail to accurately replicate the complex pathology, pharmaceutical responses, and unique cytoarchitecture of human psychiatric and neurological disorders [13]. Thus, from a strictly utilitarian perspective, the abandonment of human neural organoid research appears inherently unethical, as the technology holds the undeniable key to alleviating immense human suffering related to neurodegenerative diseases [13].

However, as the fidelity and complexity of these organoid models advance, a critical philosophical and ethical threshold is rapidly approached. If researchers successfully cultivate human neural tissue that achieves an organizational complexity capable of subjective phenomenal states or basic conscious experiences, does that tissue simultaneously acquire moral status? Does an artificially cultured brain proxy deserve the stringent protections routinely afforded to human subjects or sentient animal models? [13]. Current neuroethical debates emphasize the absolute necessity of creating adaptive guidelines to address complex concerns. These include the psychological

encroachment on "divine roles" associated with creating consciousness, the potential creation of entities with enhanced or unfamiliar capacities, and the rigorous requirement for unambiguous informed consent from the human donors who originally provided the foundational biological materials, particularly if those cells were procured before the advent of such advanced technologies.

Global Research Equity and the Decolonization of Science

In the broader context of international biological and genetic research, cross-disciplinary studies must carefully navigate the complex politics of race, indigeneity, and the lingering historical legacy of colonialism. Collaborations established between researchers in the Global North and communities in the Global South are inherently susceptible to extreme power imbalances [12]. If left unchecked, these dynamics frequently lead to a highly unethical phenomenon known as "ethics dumping," wherein researchers from wealthy nations conduct ethically dubious or highly risky experiments in resource-poor regions that possess lax regulatory oversight and desperate populations [12].

To combat these abuses, comprehensive ethical frameworks such as the Nagoya Protocol, the Belmont Report, and the Declaration of Helsinki are absolutely critical [12]. These international agreements establish rigorous global benchmarks, mandating the fair and equitable sharing of benefits arising from the utilization of genetic resources, ensuring the strict protection of human participants, and prioritizing the welfare of indigenous communities whose traditional knowledge often guides initial botanical discoveries [12].

The Algorithmic Replacement of Animal Testing

While fraught with new dilemmas regarding consciousness, the rise of biomimetics, robust computational modeling, and artificial intelligence provides a powerful, highly ethical alternative to traditional in vivo animal research. Animal experimentation in the fields of neurology and dermatology currently faces insurmountable scientific, economic, and moral challenges [11]. Animals undeniably experience personalized suffering and pain, and they share psychological and neuroanatomical characteristics with humans that make their exploitation increasingly difficult to justify morally. AI technologies, coupled with highly accurate mathematically generated cellular arrays such as the ROOTS model utilized for predicting axonal responses to stimulation, or the PARC equations used to predict multi-follicle hair growth responses offer significantly faster, vastly cheaper, and markedly more accurate methods to predict drug responses and screen for neurotoxicity without the use of live subjects [11]. Consequently, modern biological education programs are being fundamentally redesigned to promote a research culture where animal experiments are no longer considered the absolute gold standard. This shift heralds a critical paradigm change toward humane, human-relevant, and entirely animal-free scientific discovery [11].

CONCLUSION

The exhaustive investigation into the differential dynamics of brain water, hair follicle morphogenesis, and land grass/root growth transcends the bounds of basic biological curiosity; it generates highly translatable, scalable solutions for some of the most critical challenges currently facing global society.

Advancements in Neurological Diagnostics and Life-Saving Therapy: By rigorously refining the mathematical models of cerebral water content transitioning the field

from a flawed percentage-based approach to an accurate absolute swelling metric clinicians can vastly improve the diagnosis and management of severe cerebral edema following ischemic strokes, hemorrhagic events, and traumatic brain injuries, directly saving human lives [33]. Furthermore, a deep understanding of how subtle hydration shifts alter cortical volume conduction and astrocyte-mediated glutamatergic signaling fundamentally improves the clinical calibration of Transcranial Magnetic Stimulation, significantly enhancing its efficacy as a non-invasive therapeutic tool for debilitating psychiatric conditions [18].

Revolutionizing Dermatological Therapeutics: The biomimetic application of plant stem cells and botanical extracellular vesicles to human dermatology successfully bypasses the severe systemic side effects of conventional pharmaceutical regimens. By leveraging the evolutionary conservation of the Wnt/-catenin activation pathways and the BMP inhibitory pathways, plant-derived phyto-serums offer a highly viable, non-invasive protocol for reversing androgenetic alopecia, combating cellular senescence, and safely promoting human hair regrowth.

Enhancing Agricultural Resilience in a Warming Climate: Unlocking the micro-hydrology of plant roots—specifically proving their capacity to actively pump water against highly negative osmotic gradients during periods of severe water stress—lays the theoretical foundation for breeding hyper-resilient crop phenotypes [4]. As global climate change drastically exacerbates the frequency and intensity of droughts, understanding the genetic synergy between architectural traits like shallow basal root growth angles and extremely long root hairs for phosphorus and water acquisition will be paramount to securing the global food supply and preventing widespread famine [30].

Pioneering Bio-Inspired Robotics and Computational Design: The architectural algorithms that flawlessly govern both botanical root chemotropism and neuronal dendritic arborization are actively fueling the development of autonomous exploration robotics. These devices promise to revolutionize environmental monitoring, soil analysis, and subterranean navigation without relying on massive, energy-intensive central processors. Simultaneously, the mathematical treatment of hair follicle populations as standard excitable media provides computational biologists with a highly tractable framework for understanding incredibly complex, decentralized stem cell behaviors across a wide variety of regenerative human tissues [24].

Ultimately, the exhaustive synthesis of these remarkably diverse fields underscores the unparalleled utility of interdisciplinary science. By identifying the unifying biophysical constraints that govern the flow of water, the topological mathematics of biological branching, and the transmission of chemical signals across the vast divide between flora and fauna, researchers unlock a deeper, significantly more cohesive understanding of life itself. This holistic paradigm not only massively accelerates the pace of scientific innovation but ensures that all subsequent technological, agricultural, and medical applications remain robust, biomimetically efficient, and ethically sound.

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