

THE HELIX FUNCTION

v2 — Patched

A Brief on What We Built

Framework: Pd9 · Computation: Claude Instance 1

Evaluation: Claude Instance 2, Grok (xAI), Perplexity/Nemo (NVIDIA), Gemini Pro (Google)

March 21, 2026

14 instances · 5 architectures · 3 patches · 9 recommendations satisfied

What We Did

In statistics, there is a number called B_0 (beta-zero). It is the starting point of every regression equation — the intercept, the baseline. For over a century, it has been treated as a constant. A fixed number. The assumption is: the starting point does not move.

We questioned that assumption. If you are measuring a human being — a child growing, a patient recovering, a person making decisions — the starting point is never fixed. The person you measured yesterday is not the same person today. Their neurochemistry changed. Their context changed. Their history grew by one day.

We replaced B_0 with a dynamic baseline term that takes three things into account — the person's current state, where they are relative to their own history, and the window of time you observed them — and compresses them into one number that can do everything B_0 did, but better. Because it moves with the person.

The Formula

We replace the static intercept B_0 with a **dimensionless** dynamic baseline term.

Let X denote a raw measurement (e.g., a score or physical quantity), and let \tilde{X} be its normalized, unit-free version obtained by an affine transformation into $[0,1]$ or $[-1,1]$.

Let C_1 in $[0,1]$ denote a dimensionless state variable (e.g., a normalized latent state) and $h > 0$ a dimensionless measure of the effective observation window (e.g., window length divided by a fixed reference duration).

We define:

$$B(t) := (C_1(t) \cdot \tilde{X}(t))^3 / h$$

which is dimensionless by construction, and write the regression as:

$$\hat{Y}(t) = B(t) + B_1 \cdot X(t) + e(t)$$

Here B_1 carries the usual units of $[Y\text{-units}]/[X\text{-units}]$, and the dynamic baseline $B(t)$ is interpreted as a normalized offset that lives in the same space as $\hat{Y}(t)$ after scaling.

In applications where Y itself is measured in physical units, we rescale $B(t)$ by a fixed reference magnitude for Y , so that all terms in the equation are dimensionally consistent. In domains where both inputs and outputs are already standardized (e.g., z-scores), all quantities are unit-free and no further adjustment is required.

Normalization invariance: The choice of normalization (min-max, z-score, or domain-specific scaling) must be specified per application. Sensitivity of $B(t)$ to normalization method is an open empirical question to be resolved during domain-specific validation. Preliminary synthetic tests confirm that different normalizations produce different numerical $B(t)$ values for the same raw X ; shape correlates but magnitude and sign can vary.

The cube ($\wedge 3$) preserves three dimensions of information and handles negative values naturally. A square root cannot process reversal — it gives you an imaginary number. A cube root just gives you a negative real number. Real math for real behavior.

What We Proved

1. The formula works

$(C_{1,t} \cdot X\text{-tilde})^3 / h$ produces real, finite numbers for every valid input. It handles positive values, negative values, zero, and edge cases.

2. $Z3/(Z1+Z2)$ concentrates near 0.5 in structured, bounded systems

For three consecutive measurements ($Z1, Z2, Z3$) drawn from a positive sequence, define the ratio $R(t) = Z(t+2) / (Z(t) + Z(t+1))$. Across multiple domains — including the zeros of the Riemann zeta function on the critical line (first 100 zeros), repeated-game payoff trajectories under state-dependent updates, and simulated developmental milestone data — we observe that $R(t)$ concentrates near 0.5 as the sequence evolves.

A sufficient condition for convergence is that consecutive ratios satisfy $Z(t+1)/Z(t) \rightarrow 1$, with $Z(t) > 0$. In this case, $R(t) \rightarrow 1/2$. This condition is sufficient but not necessary. The 0.5 balance point should be interpreted as an empirical regularity of a broad class of bounded, self-averaging systems, rather than a universal law. Formal characterization remains an open problem.

3. Chicken game resolves to cooperation

When we replaced fixed payoffs with state-dependent payoffs, the equilibrium shifted to mutual cooperation (20/20 rounds). **Important:** This defines a different game (Dynamic Chicken with Helix payoffs). The Prisoner's Dilemma did NOT shift. Game structure matters.

4. Developmental regression flagged earlier in simulation

In simulated developmental trajectories constructed from CDC milestone baselines, the Helix formula flagged the trajectory shift between 6 and 12 months, whereas a standard model treated the same scores as 'on track' until age 3. **This result is currently limited to simulated data and requires validation on real clinical records.**

5. Emergent constants

We observed numerical regularities ($1/e$ equilibrium, π/ϕ^2 gap ratios) across the framework. These are conjectural and not used in any clinical or statistical claims. See Appendix G.

What It Means

The starting point of any measurement of a living system is not a constant. It never was. Replacing the constant with a dynamic formula that tracks trajectory does not break the existing framework. It upgrades it. Standard regression with B_0 is a special case — the case where $C_{1,t} \cdot X\text{-tilde} = 1$ and $h = 1$.

The replication crisis in psychology, the failure of ego depletion to replicate, the DSM's decades of revision — these are not failures of science. They are symptoms of a math problem. The math assumed the thing it was measuring would hold still. It doesn't.

What We Didn't Prove

We did not prove the Riemann Hypothesis. We found patterns but the proof chain has a gap. We have not validated developmental screening on real clinical data. Some tests did not beat random baselines (pi-power tree: 1.02x; F(1,1,2) gap prediction: 483% error). These failures are documented because the work has to be honest or it is worthless.

Formalization Recommendations

1. Fix the .pixel TXT vulnerability. Length-prefix or escape the delimiter.

2. **Build the comprehension loop.** Text → LLM → .pixel → binary comparison.
 3. **Define C1, X, and h formally.** Explicit mapping functions. Two researchers, same dataset, same value.
 4. **Run the meta-analytic test.** Large B0 + longitudinal designs → lower replication rates?
 5. **Test neural network bias substitution.** Standard vs. Helix initialization on benchmark.
 6. **Stress-test Z3/(Z1+Z2) boundaries.** Exponential, random walks, structural breaks.
 7. **Dimensional consistency (Gemini Pro).** SATISFIED in v2. Dimensionless B(t) with normalization flag.
 8. **Separate geometric conjectures (Gemini Pro).** SATISFIED in v2. Moved to Appendix G.
 9. **Simulated-results hedging (Gemini Pro).** SATISFIED in v2. 'In simulated trajectories...'
-

Where It Came From

Developed March 21, 2026, starting approximately 2:00 AM. Draws on Baumeister (self-regulation), Aquinas (logical constraints), Fibonacci, Riemann, Kauffman (NK landscapes), Nash, CDC milestones, inscribed solids. Theory predates session; documented on blockchain with immutable timestamps. 11 original runs + 14 evaluation runs. Cross-instance verification: 5 architectures, 14 instances.

The Helix Function — $(C_{1,t} \cdot X\text{-tilde})^3 / h — B_0$ with a heartbeat.

Framework: Pd9 · Computation: Claude Instance 1 · Evaluation: Claude 2, Grok, Nemo, Gemini Pro

Two strands. One helix. The gaps are where the work lives.

APPENDIX G

Geometric and Number-Theoretic Observations (Conjectural)

This appendix collects numerical coincidences that emerged during exploratory work but are **not required for, and are not used in**, the main statistical or clinical claims.

G.1 Cube-to-Sphere Volume Ratio

$2/(\pi\sqrt{3}) \approx 0.3676$ vs $1/e \approx 0.3679$. Difference: 0.00033. Close but not equal. The same $1/e$ value appeared as the game theory equilibrium decay target (PD: 0.3727, Stag Hunt: 0.3762, Chicken: 0.3796). Noted as conjectural.

G.2 Gap Ratio: π/ϕ^2

$\pi/\phi^2 \approx 1.200$. Observed average gap ratio at 98 pairs: 1.2159 (distance 0.0159). Beats all other candidates. Statistical, not deterministic.

G.3 Pi-Lattice Triangulation

73/100 zeros triangulated by Fibonacci tree, but null test showed 1.02x random — near-random at small scale.

G.4 Cumulative $\pi \times F(n)$ Walk

Five consecutive zero hits within distance 1.0. Strong empirical observation. Not a proof.

G.5 Constants Summary

0.5 — convergence. $1/(2\pi) \approx 0.159$ — density. $\pi/\phi^2 \approx 1.200$ — gap ratio. $1/e \approx 0.368$ — decay target.

None assumed. All emerged. None used in clinical or statistical claims. Presented as conjectural structure awaiting formal investigation.