

# ECTO Design Notes (SATSA & Dental Cohorts)

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## 1 Purpose & Scope

These notes document the design rationale behind the ECTO pipeline: converting longitudinal psychometric item distributions into an information signal (Shannon entropy, in bits), using that only to *seed initial conditions*, and then evolving a minimal autonomous ODE for two trait states  $N, P$  coupled to a cumulative stress/constraint state  $E_{\text{stress}}$ . The aim is a small, interpretable, falsifiable *phenomenological* law at the cohort level.

## 2 Data $\rightarrow$ Information $\rightarrow$ State

1. **Inputs:** Wave-wise Likert distributions for selected items (per cohort).
2. **Entropy extraction:** For each wave, compute Shannon entropy  $H$  (bits) of the item distribution. Within each dataset, min-max normalize to obtain  $H^* \in [0, 1]$ .
3. **Initialization (seeding only):** Use  $H^*(t_0)$  to set  $N(t_0) = N_0(H^*)$ ,  $P(t_0) = P_0(H^*)$ ; set  $E_{\text{stress}}(t_0) = E_0$ . No time-varying inputs are used in the main ALife runs; the ODE then evolves *autonomously*.

## 3 Time & Units

Integration is in calendar time (years) at the observed timestamps. States  $N, P, E_{\text{stress}}$  and  $H^*$  are dimensionless. Rate parameters  $\mu, \gamma$  have units  $\text{year}^{-1}$ . Parameters  $\alpha, \beta, c_1, c_2, c_3, G, K$  are dimensionless. In the main runs we set  $G = 1$  for comparability;  $K$  is tuned/fitted per run.

## 4 Dynamical Law (ECTO ODE)

$$\begin{aligned} \frac{dN}{dt} &= \mu N - (\alpha N + \beta^2 P) N, & \frac{dP}{dt} &= \mu P - \beta P \left( \frac{c_1 P + c_2 N + c_3 E_{\text{stress}}}{G} \right), \\ \frac{dE_{\text{stress}}}{dt} &= \gamma E_{\text{stress}} \left( \frac{N}{N + K} \right). \end{aligned}$$

**Notes:** Expanding makes the interaction structure explicit:

$$\dot{N} = \mu N - \alpha N^2 - \beta^2 P N, \quad \dot{P} = \mu P - \frac{\beta c_1}{G} P^2 - \frac{\beta c_2}{G} N P - \frac{\beta c_3}{G} E_{\text{stress}} P.$$

## 5 Parameter Roles (phenomenological)

- $\mu$ : innovation/influx baseline sustaining trait expression.
- $\alpha$ : self-limiting constraint on  $N$  (logistic-like).
- $\beta$ : cross-trait coupling scale (squared in  $\dot{N}$  to keep damping nonnegative).
- $c_1, c_2$ : within-/cross-trait cost weights affecting  $P$ .
- $c_3$ : stress sensitivity of  $P$ .
- $G$ : capacity/scale (set  $G=1$  in main runs).
- $\gamma$ : stress amplification (growth) rate.
- $K$ : saturation for the  $N/(N+K)$  kernel (higher  $K$  = slower stress sensitivity onset).

## 6 Interpretation (eco-evo analogues)

Labels such as “pleiotropy,” “selection/constraint,” and “environmental feedback” are cohort-level *phenomenological analogues* rather than biochemical mechanisms. The terms encode: self-limits ( $N^2, P^2$ ), antagonistic trade-offs ( $NP$ ), and a cumulative constraint/stress integrator  $E_{\text{stress}}$  with saturating sensitivity to  $N$ .

## 7 Well-posedness & Invariants (brief)

On the nonnegative orthant  $N, P, E_{\text{stress}} \geq 0$ , the vector field is locally Lipschitz; solutions exist and are unique on finite intervals. With nonnegative initial conditions,  $N, P$  remain nonnegative;  $E_{\text{stress}}$  is monotone nondecreasing with instantaneous rate bounded by  $\gamma E_{\text{stress}}$ . (A relaxing variant with  $-\lambda E_{\text{stress}}$  is a planned ablation, not used in the ALife runs.)

## 8 Validation Protocol

1. **Fit/Calibrate:** Choose/fix parameters to reproduce trajectory *shape* on a subset of waves.
2. **Held-out (LOO):** Hold out one wave, fit on the rest, report held-out RMSE/ $R^2$ .
3. **Artifacts:** Save metrics (`metrics_*.csv`, `loo_*.csv`) and figures (`fig_*.png`); print Python/NumPy/SciPy versions.

## 9 Ablations & Sensitivity (planned)

- **Information measure swap:** Shannon  $\rightarrow$  KL/Fisher/Algorithmic; compare held-out error.
- **Stress relaxation:** add  $-\lambda E_{\text{stress}}$ ; test recovery dynamics.
- **Kernel variants:** replace  $N/(N+K)$  by alternative saturating kernels.
- **Parameter sensitivity/identifiability:** CIs via profile likelihood or bootstrap; Sobol/PRCC.
- **Baselines:** ARIMA/VAR, LV-like systems fit to same normalized trajectories.

## 10 Reproducibility

Runs are deterministic (no stochastic forcing). Use tight solver tolerances and record environment:

Python X.Y.Z | NumPy a.b.c | SciPy d.e.f

Entropies are min–max normalized *within each dataset*; Dental starts near 0 and SATSA near 1 by design to compare *dynamics*, not absolute scale.

## 11 At-a-Glance Pitch for Collaboration:

*One scalar (bits)  $\rightarrow$  tiny autonomous ODE  $\rightarrow$  held-out generalization.* Bring any longitudinal distributions (wave-wise histograms); we will seed initial state with  $H^*$ , evolve the ODE, and report held-out error.