

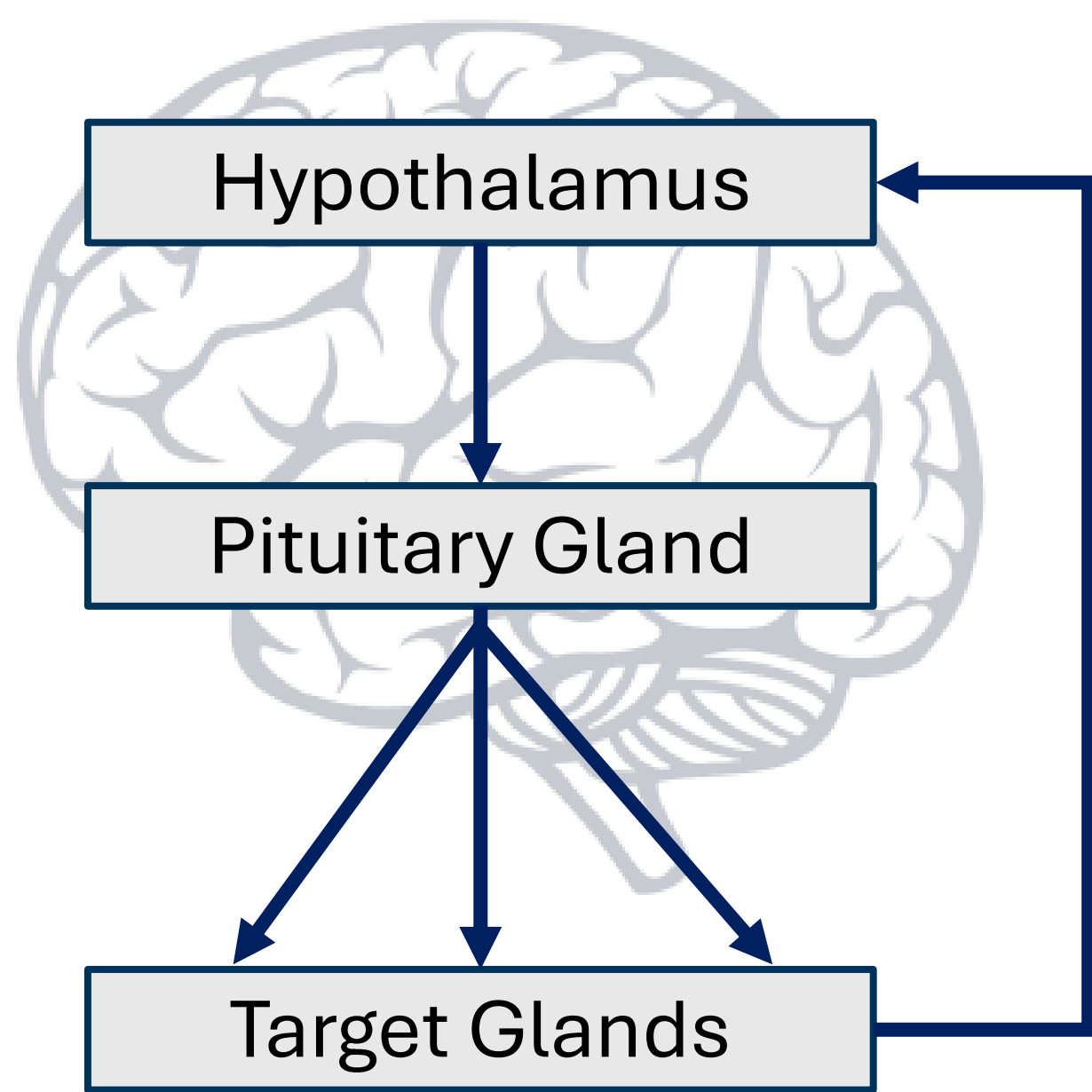
Analysis of Ordinary Differential Equation Models Representing the Dynamics of Hormone Axis



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Introduction

The endocrine system is essential for maintaining overall health by regulating metabolism, reproductive function, and early brain development. Central to these regulations are hormonal axes – network of interactions where the hypothalamus releases regulatory hormones that act on the pituitary gland. This interaction triggers the secretion of tropic hormones, which in turn stimulate target organs, establishing a complex, feedback-driven system vital for physiological balance.



Objective

This study is dedicated to the mathematical modeling and analysis of two key endocrine axes:

- Hypothalamus-Pituitary-Thyroid (HPT) axis
- Hypothalamus-Pituitary-Ovary (HPO) axis

Mathematical Modeling

- Type of Equations: Ordinary Differential Equations
- Goal: To describe the dynamics and regulatory feedback loops of the endocrine system^[1,2]

Exemplary Model ^[2] below describes the course of the autoimmune disease Hashimoto's thyroiditis.

FT4	$\frac{dx}{dt} = k_1 - \frac{k_1 y}{k_a + y} - k_2 x,$	$x(t_0) = x_0,$
TSH	$\frac{dy}{dt} = \frac{(k_3 z)x}{k_d + x} - k_4 y,$	$y(t_0) = y_0,$
Functional Size	$\frac{dz}{dt} = k_5 \left(\frac{x}{z} - N \right) - k_6 z w,$	$z(t_0) = z_0,$
TPOAb	$\frac{dw}{dt} = k_7 z w - k_8 w,$	$w(t_0) = w_0.$

Outlook

HPT-Axis

- Model refinement and improvement
- Further validation with clinical data

HPO-Axis

- Local and global stability analysis of selected models
- Calibration of model parameters

Main References

- [1] Balamurugan Pandiyan, Stephen J. Merrill, and Salvatore Benvenga. A patient-specific model of the negative-feedback control of the hypothalamus-pituitary-thyroid (hpt) axis in autoimmune (hashimoto's) thyroiditis. *Mathematical medicine and biology: a journal of the IMA*, 31 3:226–58, 2014
- [2] Graham, E.J., Elhadad, N., and Albers, D. (2023). Reduced model for female endocrine dynamics: Validation and functional variations. *Mathematical Biosciences*, 358, 108979
- [3] Marino, S., Hogue, I.B., Ray, C.J., and Kirschner, D.E. (2008). A methodology for performing global uncertainty and sensitivity analysis in systems biology. *Journal of Theoretical Biology*, 254(1), 178–196

Analysis

HPT-Axis^[1]

Stability Analysis

- Asymptotically stable equilibrium
- The system returns to a steady state regardless of small perturbations in the model parameters

HPT- and HPO-Axis^[2]

Local Sensitivity

Model Parameters

$$\frac{\partial z(t, \theta)}{\partial t} = f(t, z(t, \theta), \theta)$$
$$z: I \times G \rightarrow \mathbb{R}^n, z = z(t, \theta), I \subseteq \mathbb{R}, G \subseteq \mathbb{R}^m$$
$$\left. \frac{\partial z_k(t, \theta^*)}{\partial \theta_j} \right|_{\theta=\theta^*} \quad \theta^* \in \mathbb{R}^m \quad \text{Fixed Parameter Vector}$$

Global Sensitivity^[3]

The following methods were employed:

- Latin Hypercube Sampling
- Partial Rank Correlation Coefficient
- Extensive Fourier Amplitude Sensitivity Testing
- Sobol' indices

Calibration

- Utilized clinical measurements to calibrate model parameters
- Model validation to assess applicability to depict physiological dynamics

