

Notes While Building Computational Context Model

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Abstract

This is an attempt to port Enoch Yeung's Compositional Context Scripts to Julia while expanding them to include the option of adding addition genes, the effect on energetic regions, and to study the effect targeted DNA binding (i.e. Cas effectors) proteins have on the results they observed. This is mainly a notebook so that I can keep track of the math.

1 Introduction

We sought to recreate the work of Yeung et al. in regard to quantification of supercoiling based compositional context effects in regard to synthetic gene networks, to then modify it to include several scenarios. These include the introduction of targeted DNA binding proteins, such as the Cas family, as well as the effect of backbone expression cassettes such as the origin of replication, etc. . .

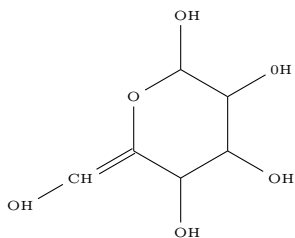
Mainly this will focus on my process in developing these ideas.

2 An Aside Regarding Michaelis-Menten Kinetics and Hill Functions.

2.1 Victory Henri

Henri is generally thought to be the first person to create a mathematical framework to explain enzymatic activity. A Russian physical chemist born in France due to an illegitimate conception, of whom latter returned to Russia (specifically Saint Petersburg). He was the relative of notable control theorist Aleksander Mikhailovich Lyapunov, composer Sergei Mikhailovich Lyapunov and Slavic linguist Boris Mikhailovich Lyapunov.

He was eventually be educated in mathematics and Natural Sciences, and earned two PhDs. The first was in psychology and the second in physical Chemistry.



As a prominent subject of study at the time, **Invertase** became the main focus of Henri's Studies. The reaction of which is included below. Which eventually, with the help of Adrian John Brown, leading him to formulate the equation:

$$\dot{x} = \frac{K * \Phi * (a - x)}{1 + m * (a - x) + n * x}$$

Where $x \equiv$ product formed, and $a \equiv$ initial substrate, and K, Φ, m , and n were empirically derived constants. While under recognized during his time, Henri Victor's work would eventually be picked up 10 years later by Leonor Michaelis and Maud Menten.

2.2 Michaelis-Menten

While not the first to formulate a mathematical framework for enzymatic activity, Leonor Michaelis and Maud Menten were the first to formulate a detailed and bio-physically meaningful model for enzyme activity. The following case describes the activity of an enzyme to produce one molecule count product from one molecule count substrate. This equation recommended by the *International Union of Biochemistry and Molecular Biology*, IUBMB is:

$$\nu = \frac{dp}{dt} = \frac{Va}{K_m + a} \quad (1)$$

Where ν is the rate of product production per time $\frac{dp}{dt}$, V (also written as V_{\max}) is the maximum production rate (more accurately the limit of production as substrate concentration reaches saturation) and K_m is the Michaelis Constant. The physical interpretation of the Michaelis-Constant is the concentration of substrate when $\nu = 0.5V$. This seemingly simplified description of enzymatic activity would go on to be widely used in descriptions of biomolecular reactions.

The assumptions of the Michaelis-Menten model are as follows: