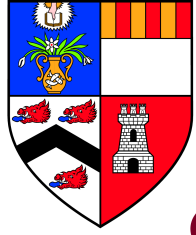


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Heinrich Heine

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UNIVERSITÄT DÜSSELDORF



CEPLAS

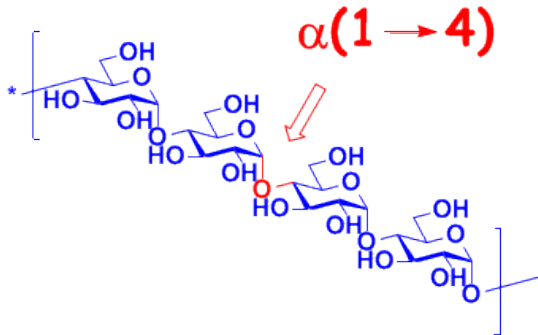
Cluster of Excellence on Plant Sciences

Polysaccharide Metabolism in silico: Conceptual Challenges and Approaches

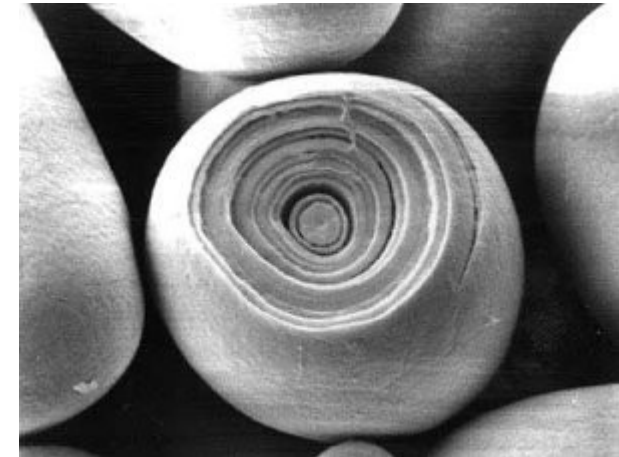
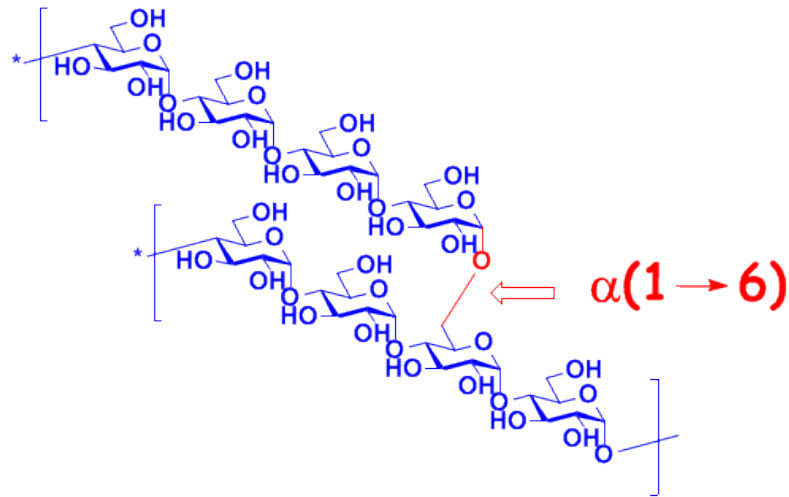
Oliver Ebenhöf, Adélaïde Raguin

Starch

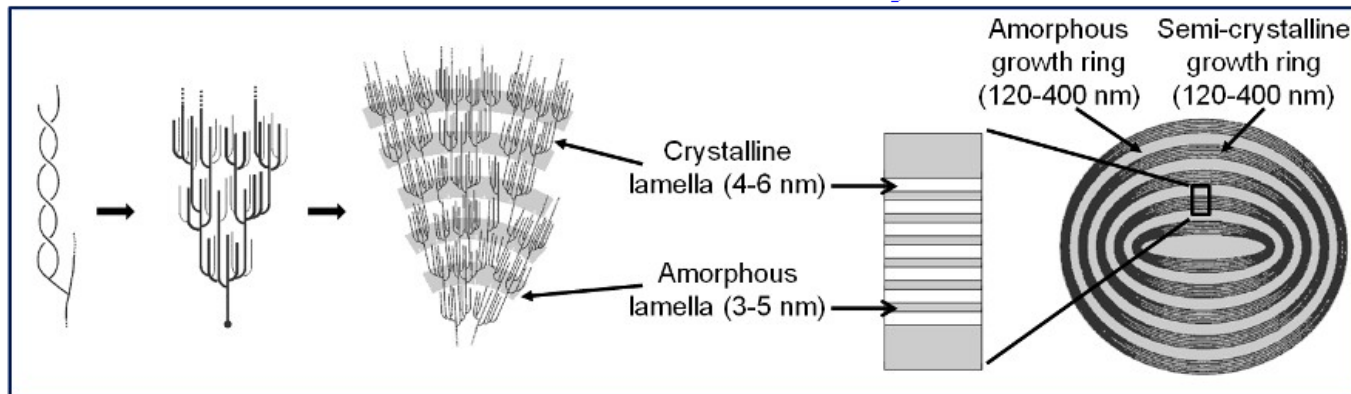
Amylose
(MW 32,000-113,000)



Amylopectin
(MW 10^7 - 10^9)



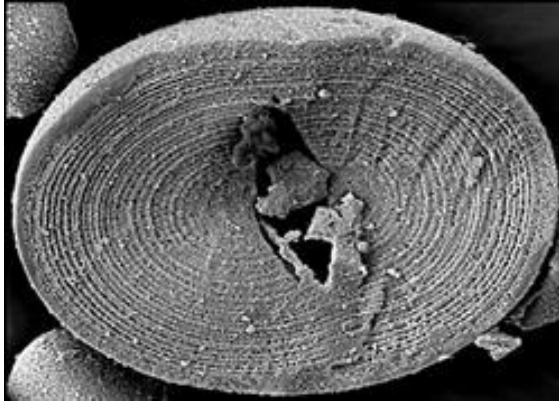
(ianhopkinson.org.uk)



(O'Neill and Field, 2015, Front Bioeng Biotechnol)

To understand the emergence of the complex structure of starch, we need to be able to model it!

Conceptual challenges



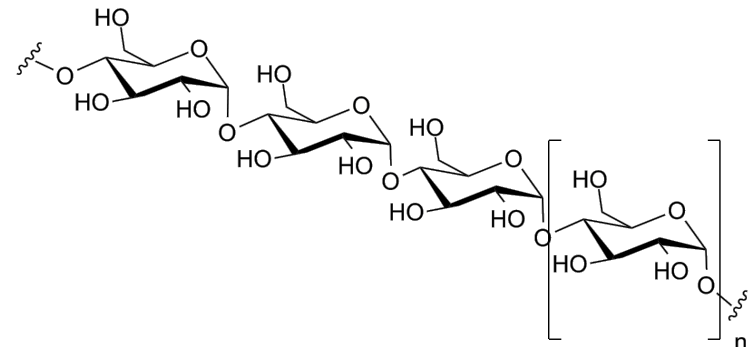
(szeeman-at-ethz.ch)

1

Biochemical reactions on surfaces

2

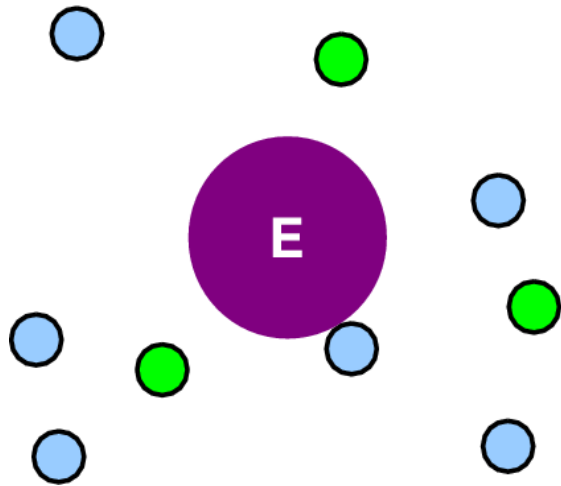
Biochemical reactions on polymers



(by glycoform, commons.wikimedia.org)

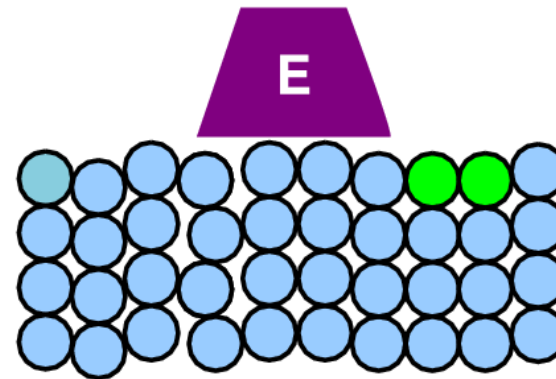
Challenge 1: surface-active enzymes

dissolved substrate



$$v = \frac{V_{\max} S}{K_M + S}$$

**aggregated substrate
(with interfacial reaction space)**

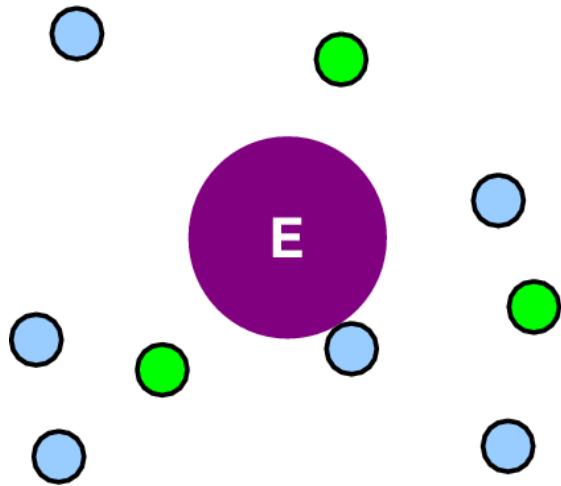


$$v = f(?)$$

Reaction space confined to 2D!

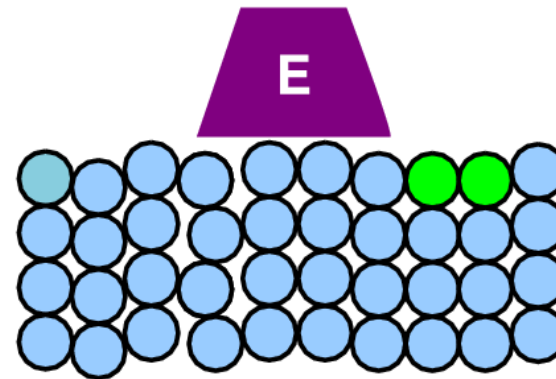
Rate laws for surfactive enzymes

dissolved substrate



$$v = \frac{V_{\max} S}{K_M + S}$$

**aggregated substrate
(with interfacial reaction space)**

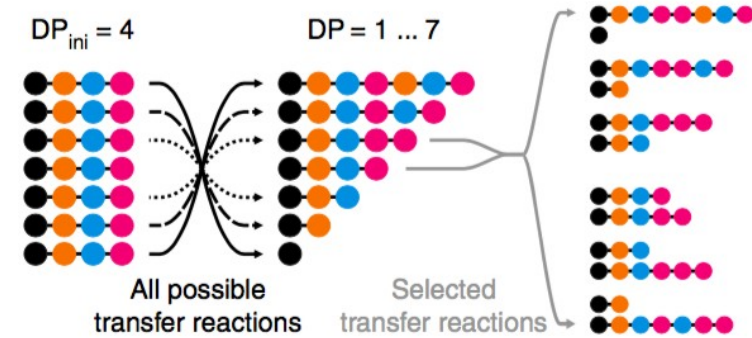
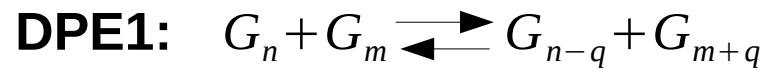


$$v = \frac{k_A a_s \Phi_{\text{eq}} [M] [E_0] (k_S \langle *S \rangle - k_P \langle *P \rangle)}{1 + k_A a_s \Phi_{\text{eq}} [M] \left(1 + \frac{\langle *S \rangle}{K_{mS}} + \frac{\langle *P \rangle}{K_{mP}} \right)} = \frac{V_M^{\text{app}} \frac{[M]}{K_{mM}^{\text{app}}}}{1 + \frac{[M]}{K_{mM}^{\text{app}}}}$$

Kartal and Ebenh oh (2013) FEBS Letters
(centenary issue commemorating Michaelis-Menten 'Kinetik der Invertinwirkung')

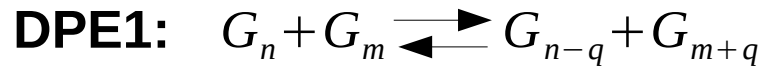
Challenge 2: polymer-active enzymes

How do you model enzymes catalysing an infinite number of reactions?



Challenge 2: polymer-active enzymes

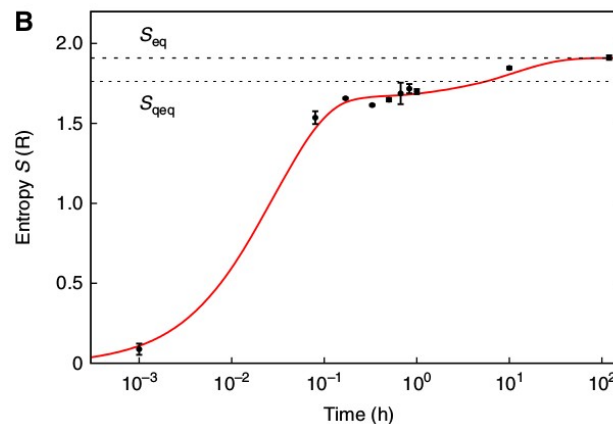
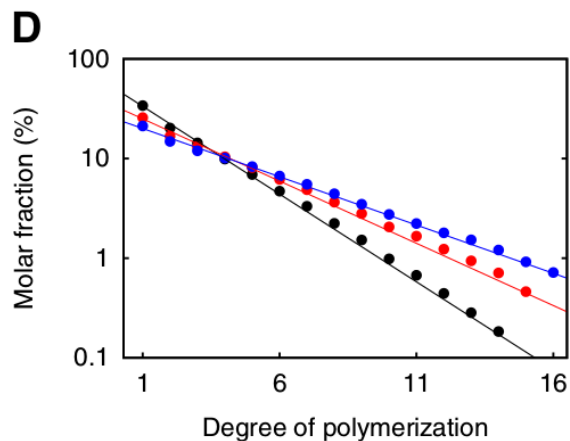
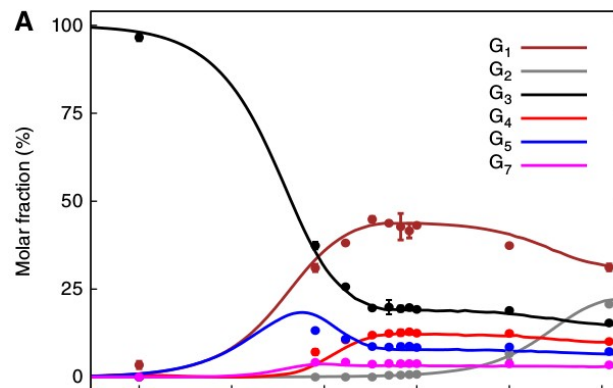
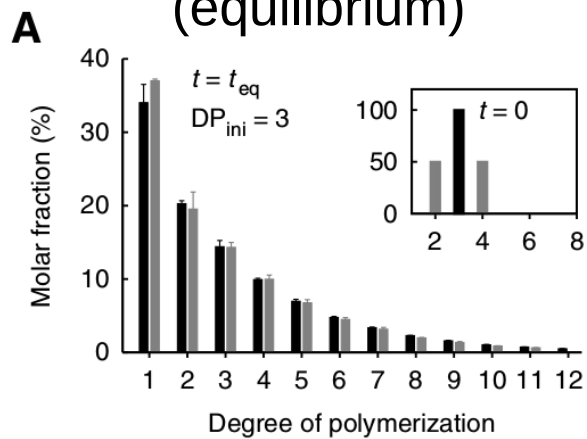
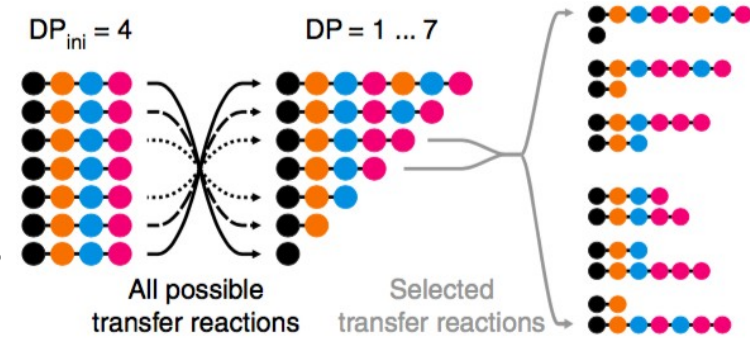
How do you model enzymes catalysing an infinite number of reactions?



Solution

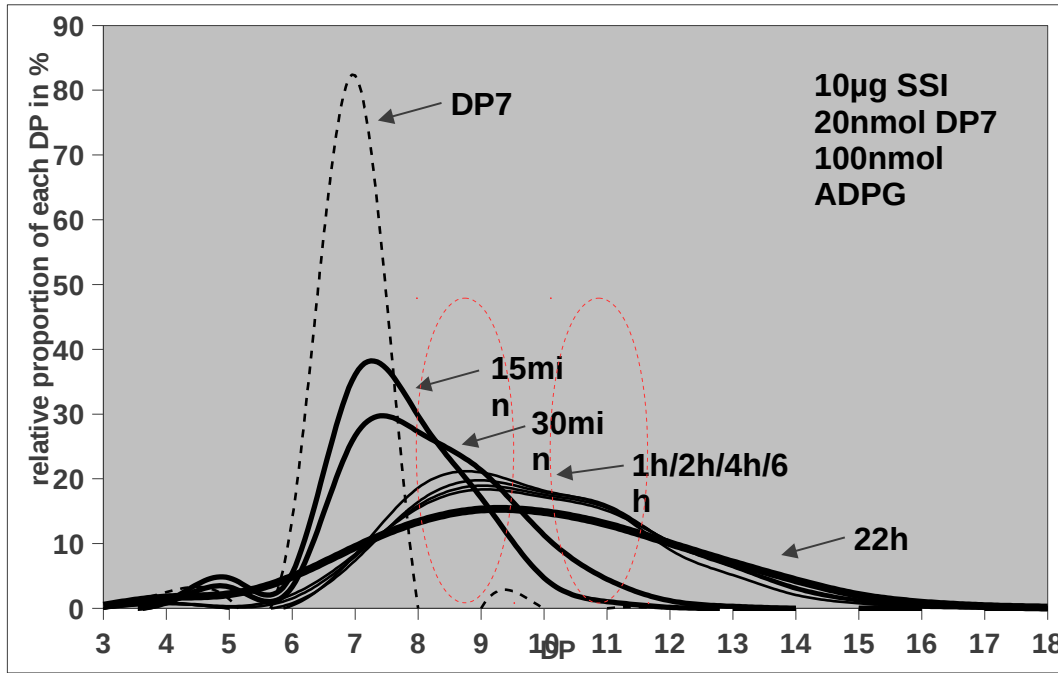
Statistical thermodynamics
(equilibrium)

Stochastic simulations
(dynamics)

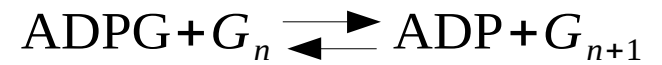


(Kartal et al, 2011, Mol Syst Biol)

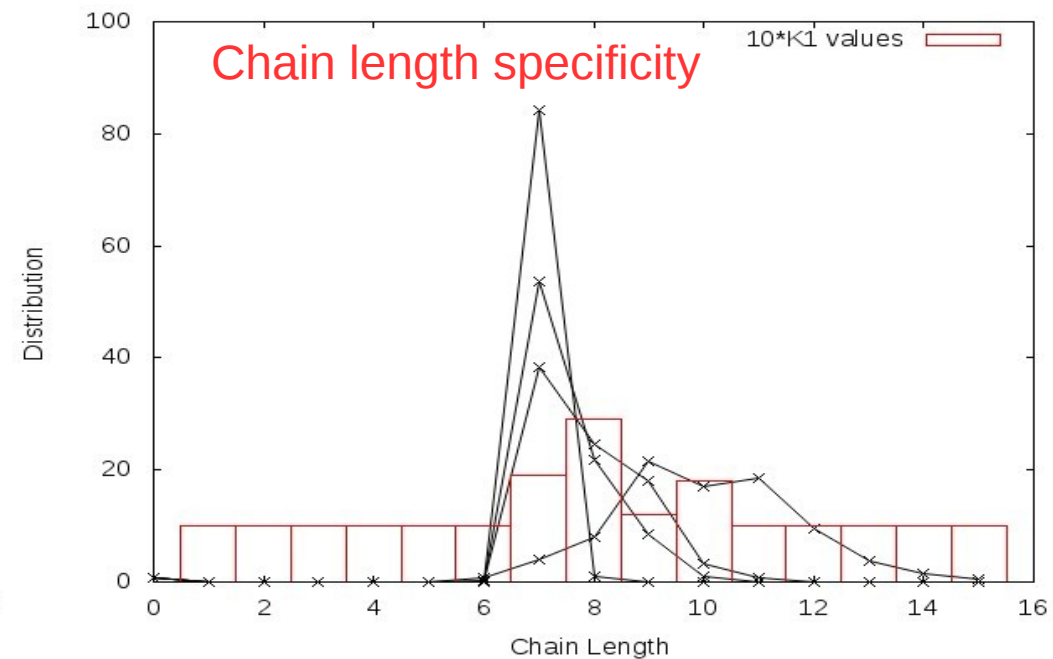
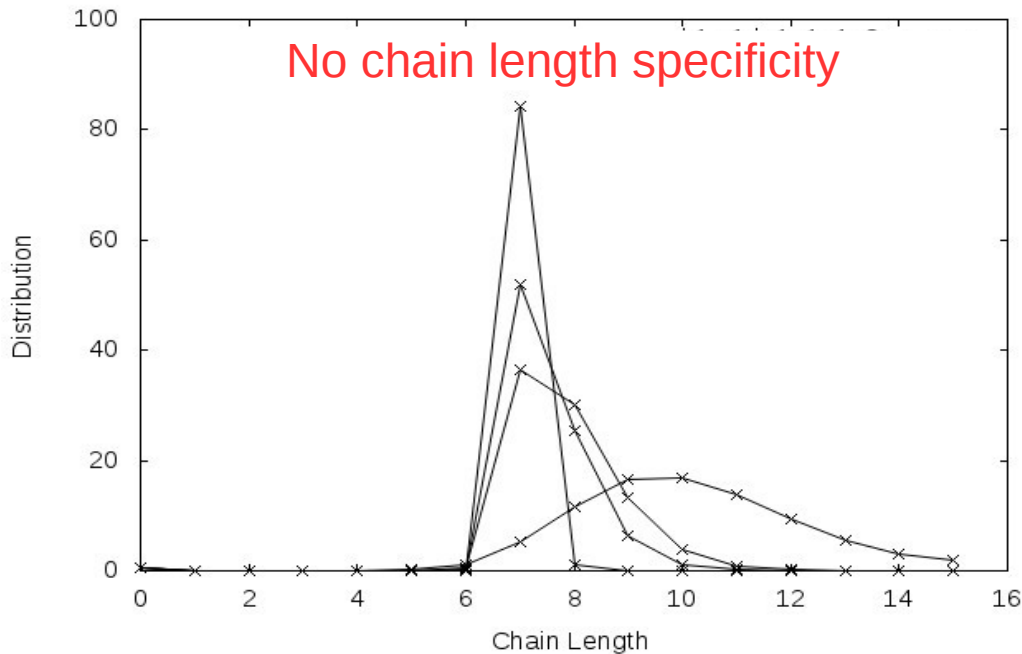
What can we learn from models?



SS1 (Starch Synthase I):



data from Henrike Brust, U Potsdam
(see also Brust et al, 2013, J Appl Glycosci 60)



MATHEMATICAL MODELS



A SOPHISTICATED
DATA ANALYSIS
METHOD



UNDERSTANDING!