



The Art of Molecular Programming

Draft Table of Contents

The textbook is divided into three parts. 'Part 0' is foundational material, followed by Part I: Structures, Part II: Circuits and Part III: Interfaces. Each part is divided into chapters (numbered), and chapters into sections (lettered). Each section will be written by an author or team of authors. Bullets represent keywords, or otherwise general guidance on the content of the section.

1. Foreword

2. Introduction & A Brief History of Molecular Computing

3. Foundations

a. Biology for Molecular Programmers

- *Central dogma & cell response to external stimuli*
- *Compartmentalization & reactions?*
- *Common DNA/RNA binding macromolecules*

b. Basic Thermodynamics

- *Concept of entropy, enthalpy, Gibbs energy, Chemical Equilibrium*
- *Thermodynamic energy landscapes, partition function*
- *Example: Thermodynamic models for nucleic acids*

c. Dynamic Models ODEs and Chemical Reaction Networks

- *Stochastic Models*
- *Deterministic Models*
- *Example: Kinetic models for nucleic acids (branch migration)*

d. Experimental Methods to Produce DNA

i. Synthesis of DNA

- *Phosphoramidite, error rates on synthesis*
- *Asymmetric PCR*
- *Cell culture*
- ii. Purification**
 - *Agarose/PAGE purification*
 - *Chromatography (HPLC, LC, maybe capillary?)*
 - *Centrifugation (filter, gradient, ultracentrifugation)*
- e. Experimental Methods for Analysis:**
 - i. Gel electrophoresis (focus on densitometry etc)**
 - *Intercalating dyes vs fluorophores or radiolabeling*
 - *Separation quality as a function of gel % and DNA length?*
 - *Role of buffer conditions*
 - *Pulse field gradient electrophoresis?*
 - ii. Microscopy (AFM, TEM/cryoEM/SEM, fluorescence/superresolution)**
 - *Contrast mechanisms and their drawbacks (stains being inconsistent, AFM always convolutes tip, etc)*
 - iii. Spectroscopy (fluorescence/CD/etc)**
 - *Relative sensitivities*
 - *Background reporter signal?*
 - iv. Sequencing**
- f. Example: Experiments for thermodynamics and kinetics of hybridization**

Part I Structures

4. "From molecules to variables" (*Chemistry*)

- a. What are information bearing molecules? (*biopolymer building blocks*)**
 - *nucleic acids (basic features and geometry)*
 - *proteins/peptides; Protein geometry (alpha helix, ...)*
 - *Other (PNA, L-DNA, new nucleotides, polysaccharides, block-copolymers, other supramolecular programmable molecules)*
- b. How do molecules interact (*the glue*)**
 - *Chemical bonds and interactions (a primer in chemistry)*
 - *Intramolecular forces via covalent bonds: peptide bonds, ester bond*
 - *Intermolecular forces and non-covalent bonds: Van der Waals, hydrogen bond, stacking interactions, electrostatic interactions, Debye screening. (Table with relative bond strengths)*
 - *Entropic forces: depletion, steric, fractionation, crowding, polymers*
- c. How can we represent molecules digitally?**
 - *Structure abstraction layers: Primary, Secondary, Tertiary structure*

- *Sequence abstraction layers: domain level vs nucleotide level*
- *Notations: Dot parens plus notation, DU+ notation, ...*
- *Notions of “valid” conformations (nearest neighbor model)*
- *Intuition behind coarse-grained representations and macrostates*
- *How can we represent molecules graphically*

5. Molecules as construction material

a. DNA properties

- *data sheet to find useful parameters.*
- *Crossover motifs (anti-parallel and parallel)*
- *Special motifs (I motif, G-quads, kissing loops, aptamers)*
- *Biophysical influence of buffer conditions on DNA structure*

b. RNA properties

- *data sheet to find useful parameters.*
- *Special motifs (I motif, G-quads, kissing loops, aptamers)*
- *Biophysical influence of buffer conditions on RNA structure*

c. Protein properties

- *data sheet to find useful parameters.*
- *Enzymatic activity and binding pockets, ...*
- *Special motifs (DNA binding)*
- *Biophysical influence of buffer conditions on enzyme activity and protein structure*

6. Introduction to self assembly. Basics of molecular self-assembly (theory)

a. Sequence based models of intramolecular self assembly

- *Levinthal’s paradox. Why can we engineer DNA/RNA but proteins are still hard?*
- *DNA/RNA Sequence design. Given a sequence, what is the structure? Given a structure, design a sequence?*
- *Proteins engineering*

b. Principles of natural multi-component self-assembly processes

- *Examples (viruses, lipid micelles, nanowires, crystals, self-assembled monolayer, hydrogels, lipids, block co-polymers)*
- *Biophysical models: The role of thermal energy driving self-assembly, nucleation, kinetic/thermodynamic control, stoichiometry, cyclisation*
- *Stochasticity — having equilibrium close to thermal energy (kT) means there will be equilibrium population of defects*

c. Tile self-assembly

- *Tiling theory (Wang tiles, Penrose tiles, ...)*
- *What shapes can be assembled in which tile model?*

- *aTAM, kTAM*
 - *Sequence design (Box OK)*
- d. Deeper into computation and self-assembly (Fundamental limits of computation — reference chapter 8)**
- *Finite state automata*
 - *Tiles and algorithmic self assembly*

7. Programmed molecular self-assemblies (*experiments*)

a. Scaffold-less DNA assemblies

- *ss Tiles,*
- *dx Tiles*
- *HCR*
- *Simple polyhedra: Seeman cube, tetrahedron, Yamuna's icosahedron, Mao's Bucky ball and octahedron.*

b. DNA Origami

- *The concept*
- *Design principles, cooperativity*
 - *Sequence design constraints*
- *Design and simulation tools*
- *"Wireframe origami" examples 1D / 2D / 3D structures*
- *Production and purification*
 - *Custom scaffold design*
 - *Thermo/stoichiometry*

c. Periodic and Multi component assemblies [Higher complexity structures, DNA LEGO, Conformational changes of origami]

- *Shape complementary, base stacking*
- *Self-limiting assemblies (rings)*
- *Fractal assemblies*
- *Lattices, ribbons, nanotubes, and crystals*
- *Interlocked assemblies (e.g. origami rotaxanes) → connection to mechanics*

d. Dynamic rearrangements of structures

- *DNA tweezers, DNA Walkers, burnt-bridge motor*
- *Nanomechanical devices. Mechanical constructs / active components/ machines / walkers (comparison with molecular motors/enzymes)*
 - *Castro's work¹*
- *Nanomotors*

e. Other phases of DNA structures (Physical properties)

- *DNA hydrogels*

¹<https://www.pnas.org/content/112/3/713/tab-figures-data>

- *DNA liquids*
- f. Hybrid DNA objects**
 - *Colloids*
 - *Gels*
 - *Polymer/DNA brushes (surface/colloid coatings)*
 - *Coacervates*
 - *Grafted nanoparticles*
- g. Other biopolymer building materials — Beyond DNA**
 - *RNA*
 - *Protein engineering (David Baker, U Washington)*
 - *Polysaccharide, spider silk, ...*
 - *DNA-chimera assemblies (protein bioconjugation, eg the work of Nicholas Stephanopoulos)*

Part II Circuits

8. Introduction to Computation

- a. Intro to computation / information processing
- b. Conventional computation
- c. Background: Boolean logic, Turing machines, register machines
- d. Examples of non-conventional computing
- e. Examples of natural computing
- f. Guiding principles? Molecular programming “languages”, modular/circuits intro
- g. Advanced: limits on computation / computability frameworks?

9. Programming molecular behaviors over time (CRNs)

- a. Introduction
- b. Computing with Stochastic CRNs
 - *Theory example and analysis: min/max and Boolean logic programming with CRNs*
 - *Biology example: something simple*
 - *Computational Power of Stochastic CRN*
 - *Time complexity of stochastic CRN*
- c. Computing with distributions
- d. Computing with Deterministic CRNs
 - *Theory examples: circuits, boolean circuits, oscillators, bistability, etc.*
 - *Well-mixed CRNs as example of analog computing*
 - *Computing functions (e.g. $y = kx$)*

- *Approximate majority*
- *Dynamic system: Oscillators, bistability*
- *Biology example: (predator prey / ecology models)*
- *Computational power of deterministic CRNs*

e. Advanced topics in CRNs

10. Nucleic acids as a universal substrate for molecular programming

a. Introduction (reference Foundations for overlapping topics)

- *Reference to Structures section on background: nucleic acid hybridization and thermodynamics*
- *Background: nucleic acid branch migration*
- *Toehold-mediated strand displacement*
- *Experiment examples: toehold exchange reaction*

b. Strand displacement cascades

- *Theory & experiment: DNA universal substrate for CRNs*
- *Boolean circuits: seesaw circuit theory*
- *Box: discuss considerations and strategies in e.g. designing mutually orthogonal sequences*

c. Theory & Experiment examples: 3wDSD circuits

- *Oscillators*
- *Bistable memory*
- *Approximate majority*
- *Amplification, catalysis, and thresholding*
- *Seesaw circuits*
- *Neural networks*

d. Advanced designs in DNA strand displacement cascades

- *Toehold activation*
- *Mutation detection; use mismatch to control kinetics?*
- *Theory & Experiment: 4wDSD circuits*

e. Leakless circuits

- *Why do circuits leak and why leaks are problematic*
- *How to avoid leak. Examples of leakless circuits*

11. Incorporating enzymes into nucleic acid cascades

a. Introduction

- *Background: enzymatic behaviors*
- *Early example: traveling salesman?*

b. DNA polymerase-based circuits

- *PEN circuits, predator prey*

- *APR, PER circuits (whiplash PCR?)*
- *Shah et al. work on logic gates*

c. Transcribed RNA-based circuits

- *Toehold switches*
- *Conditional CRISPR*
- *Oritatami*
- *Genelets*

d. Translated protein-based circuits

- *Repressilator*
- *References to good syn bio resources?*

12. Spatially-Organized Circuits

a. Introduction: advantages of spatial structures

b. Background: compartmentalization in biology

c. Surface CRNs

- *Surface DNA circuits and DNA walkers*

d. Droplet-based computing / “synthetic compartmentalization”

e. Reaction diffusion circuits

f. Microfluidics Breakout Box

13. Advanced topics in tile assembly (algorithmic self-assembly)

a. Computational complexity

b. Wang tile Turing machine implementation?

c. CRN-Tile hybrid systems?

d. Relationships between DSD and TAM etc, e.g. TBNs (Thermodynamic Binding Networks)

14. Conclusion

Part III Interfaces

15. Introduction to Interfaces & Applications

a. Intro

b. General preview of interfaces (and the techniques used) with other fields through

- *Electronic interfacing*
- *Chemical and physical interactions*
- *Interacting with biology*

16. The Interface between Traditional and Molecular Computers

a. Background/Foundations

- *Sequencing & Synthesis (reference to Foundations); NGS*
- *Nanopores*
- *Microarrays (mention how big of an industry this is...multi billion...e.g. 23 and Me uses it exclusively)*

b. DNA Computation

- *Why use molecules to do traditional computations?*
 - *Pros: lower energy usage, parallelism , physical stability, data density, etc.*
 - *Cons: expensive at high throughput, etc*

c. DNA Data Storage

- *Encoding data from digital data to DNA*
- *Synthesizing DNA*
- *Storing DNA*
- *Recovering stored DNA*
- *Sequencing the DNA*
- *Decoding the DNA back to digital data*

17. Chemical and Physical Interactions

a. Material science

- *Patterning non-DNA molecules*
- *Surface assisted methods*

b. Chemical control

- *pH [primer on DNA triplexes]*
- *Ions*
- *crowding agents*

c. Magnetic control

- *swimmers*
- *beads (for purification and/or computation)*

d. Optical control

- *DNA plasmonic and photonic circuits*
- *DNA scaffolding of non-DNA reaction networks*
- *Dynamic control of optical DNA devices*

e. Thermal control

- *Vinothan Manoharan's work²*
- *Jaeseung Hahn's work³*

²<https://science.sciencemag.org/content/347/6222/639.abstract>

³<https://academic.oup.com/nar/article/47/20/10968/5580905>

- *More examples*⁴⁵

f. Electronic driven redox power sources

- *Electrically driven process*⁶

g. Future

18. Interacting with Biology, Medical Diagnostics and Therapeutics

a. Biology toolbox primers — topics not included in Foundations primer

- *Cutting DNA (e.g., restriction enzymes, CRISPR, ribozymes?; talk about syn-bio gene editing using CRISPR and other restriction enzymes)*
- *DNA Synthesis (e.g., ligation, overhang PCR)*

b. Diagnostics

- *How do current diagnostics work [primer on various techniques, biomarkers]*
 - *Which particular diseases will benefit from in-depth diagnostics? Why do we need to know about so many biomarkers?*
 - * *Shotgun / Microarrays*
- *Why are some molecules difficult to detect? What are the advantages of DNA nanotech methods over the standard techniques?*
 - *Methods*
 - *Aptamers — How can we program DNA to bind with targets other than DNA/RNA? [primer/external reference on selex]*
 - *toeholds*
 - *electrochemical/fluorescence*
 - *SPR (surface plasmon resonance) and QCM (quartz crystal microbalance) and FET (field effect transistor)*
 - *etc.*

c. Therapy

- *Programmable nucleic acids-based therapeutics*
 - *Protein production silencing or activation — mRNA mimics, siRNA etc. (Pfizer/Moderna vaccines are real-world examples of this) [primer/external reference on transcription/translation]*
 - *Aptamer therapeutics*
- *Protection of therapeutic agents e.g. tetrahedron structures.*
 - *Different protection methods for nucleic acid nanostructures (e.g., polylysine, protein, lipid, etc.)*
- *Examples of approved or under-trial DNA nanotech drugs.*
- *Drug Delivery*
 - *Why is it beneficial to have targeted delivery of drugs in the body?*

⁴<https://onlinelibrary.wiley.com/doi/abs/10.1002/adfm.201706410>

⁵<https://pubs.acs.org/doi/10.1021/nn4030543>

⁶<http://www.francescoricciab.com/wp-content/uploads/92.pdf>

- *How are drugs loaded onto DNA structures? Various examples available.*
- *Mechanisms for release of drugs in particular areas of the body.*

d. Burgeoning biological interfaces

- *Cellular interactions Communicating with cells and cell surfaces*
 - *Why do we want to create artificial cell signal controllers? [primer/external reference on cell signalling systems]*
 - *DNA nanotech channel receptor mimics (various versions using DNA origami; include lipid bilayer section here)⁷*
- *Replicating cellular components*
- *Proteinosomes (?)*

e. Challenges in working in a biological medium

- *Dealing with toxicity*
- *Preventing degradation — [primer or external links to well-known DNA modifications and their functionality, use of XNAs?]*
- *Examples of Chemical modifications, ligand attachments and encapsulating protection (nanoparticles)*
- *How to scale up “bio factories,” using cells to manufacture molecules is difficult*
- *Working with RNA (mRNA, RNA origami, etc.)*
 - *Why RNA?*
 - *What’s different from DNA?*

f. Regulation & Ethics — breakout box

- *High level overview of the general international medical regulatory processes, with some specific examples from, e.g., the FDA approval process*
- *Maybe briefly mention other regulatory processes for, e.g., synthetic crops*
- *DNA Data Storage — chemical waste*

⁷[https://www.cell.com/trends/biotechnology/fulltext/S0167-7799\(21\)00034-2](https://www.cell.com/trends/biotechnology/fulltext/S0167-7799(21)00034-2)