



# The Art of Molecular Programming

## *Draft Table of Contents*

*The textbook is divided into three parts. 'Part 0' is preliminary 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). 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

### 1. "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*

## **2. 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*

## **3. 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*

#### **4. 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 work<sup>1</sup>*
- *Nanomotors*

##### **e. Other phases of DNA structures (Physical properties)**

- *DNA hydrogels*
- *DNA liquids*

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

#### **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**

### **1. Introduction to Computation**

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

### **2. Programming molecular behaviors over time (CRNs)**

- Introduction**
- 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*
- Compute with distributions**
  - *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*

**d. Advanced topics in CRNs**

**3. 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*

**4. 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?*
- 5. 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**
- 6. 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)**
- 7. Conclusion**

## **Part III Interfaces**

- 1. 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*
- 2. 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*

### **3. 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 work<sup>2</sup>*
- *Jaeseung Hahn's work<sup>3</sup>*
- *More examples<sup>45</sup>*

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

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

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

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



**f. Electronic driven redox power sources**

- *Electrically driven process<sup>6</sup>*

**g. Future**

**4. 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?*
  - *How are drugs loaded onto DNA structures? Various examples available.*
  - *Mechanisms for release of drugs in particular areas of the body.*

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

#### **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)<sup>7</sup>*
- *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*

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