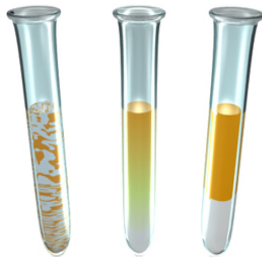


Application-Specific Fault-Tolerant Architecture Synthesis for Digital Microfluidic Biochips

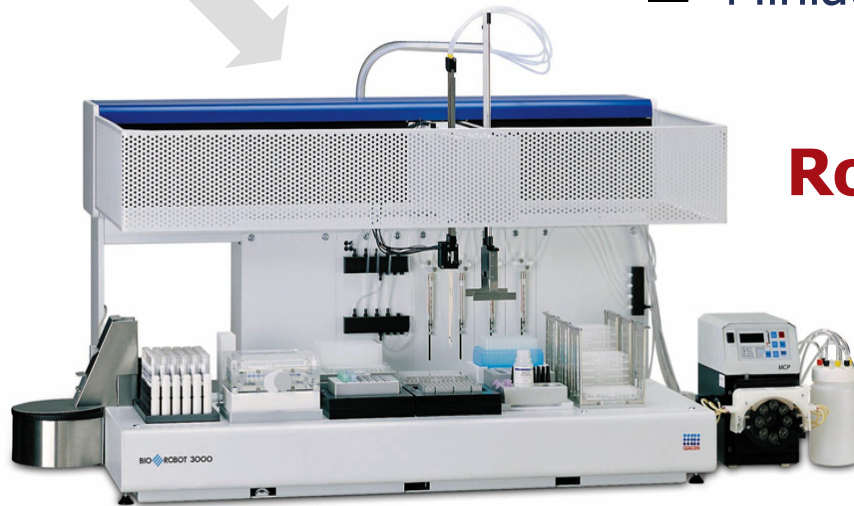
Mirela Alistar, Paul Pop, Jan Madsen
Technical University of Denmark, Lyngby





Test tubes

- Automation
- Integration
- Miniaturization

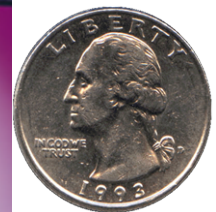
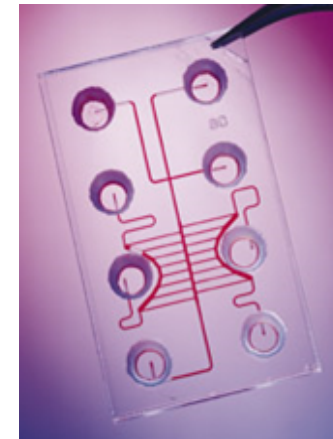


Robotics

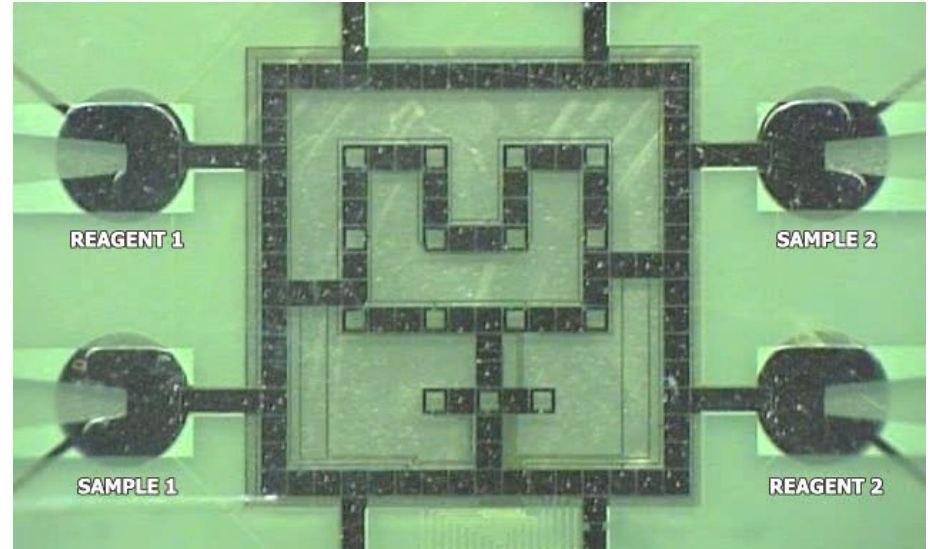
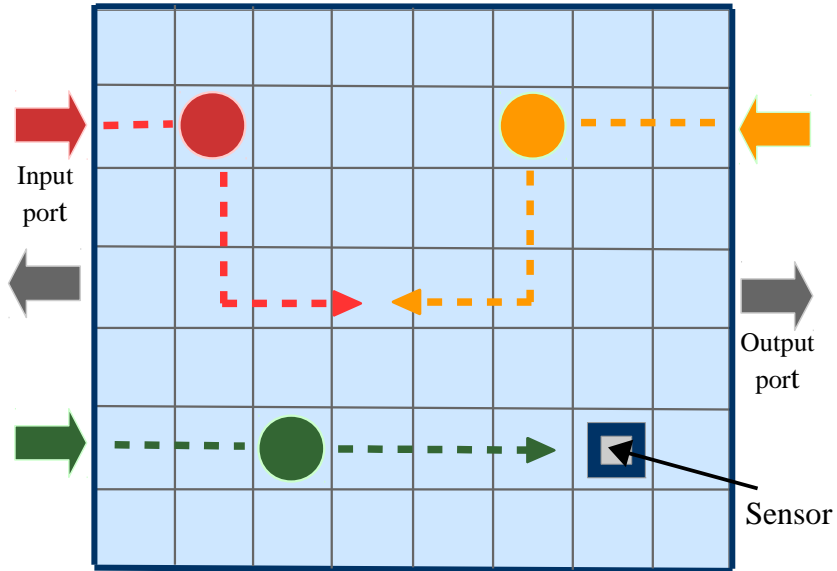
- Automation
- Integration
- Miniaturization

Microfluidics

- Automation
- Integration
- Miniaturization



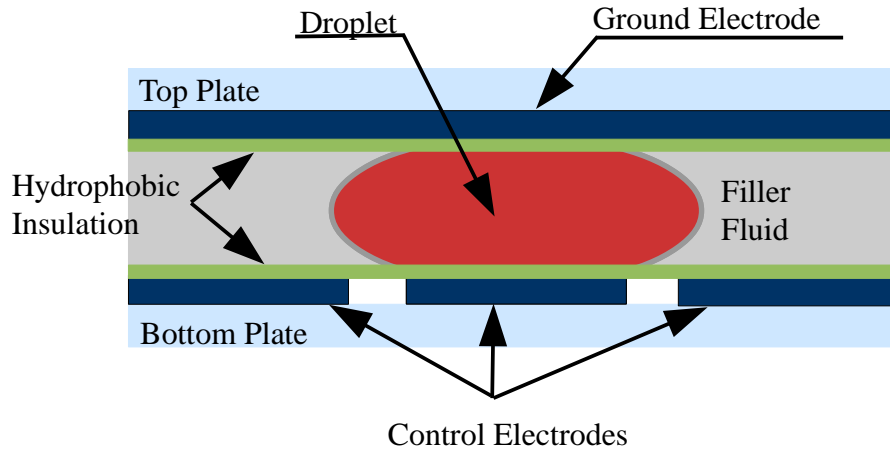
Droplet-based Biochips



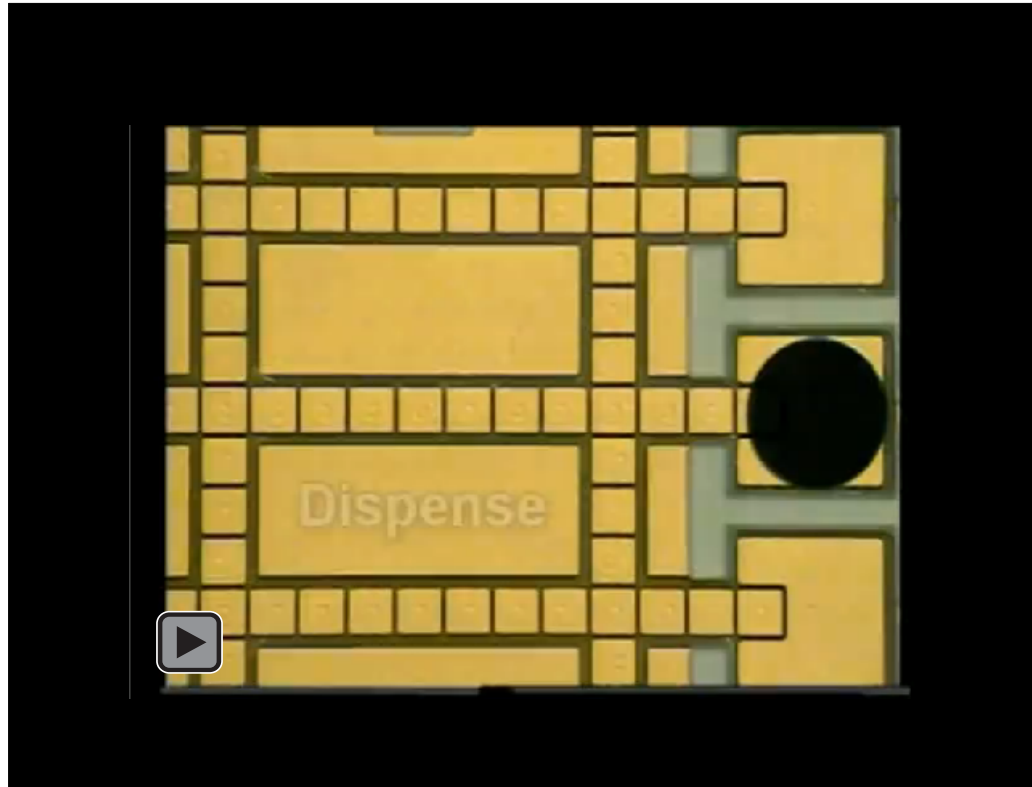
Biochip from Duke University

Digital Microfluidic Biochips (DMB)

Electrowetting on Dielectric



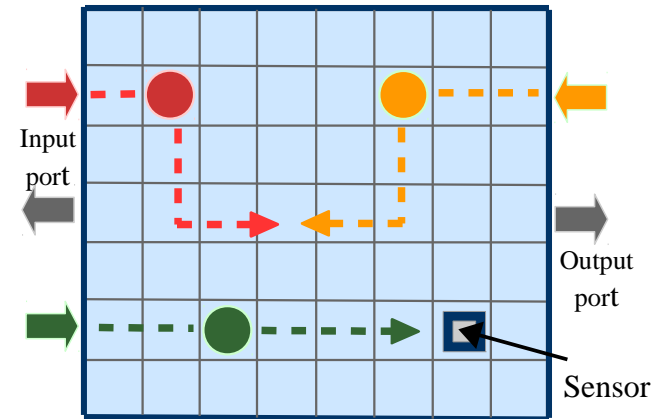
Fluidic Operations



DMB Architecture

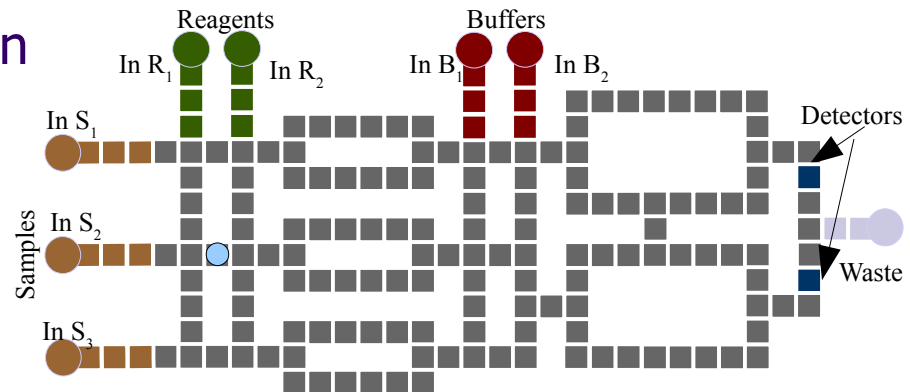
- General-Purpose Architecture

- Reconfigurable
- Versatile
- Fault-tolerant

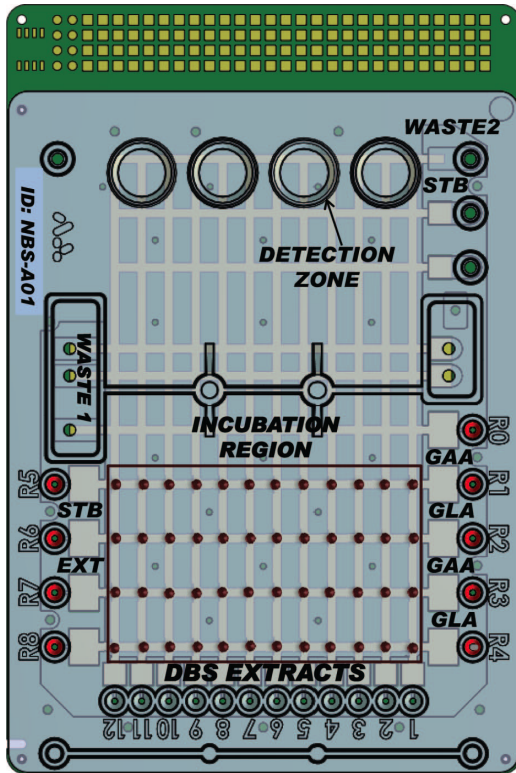


- Application-Specific Architecture

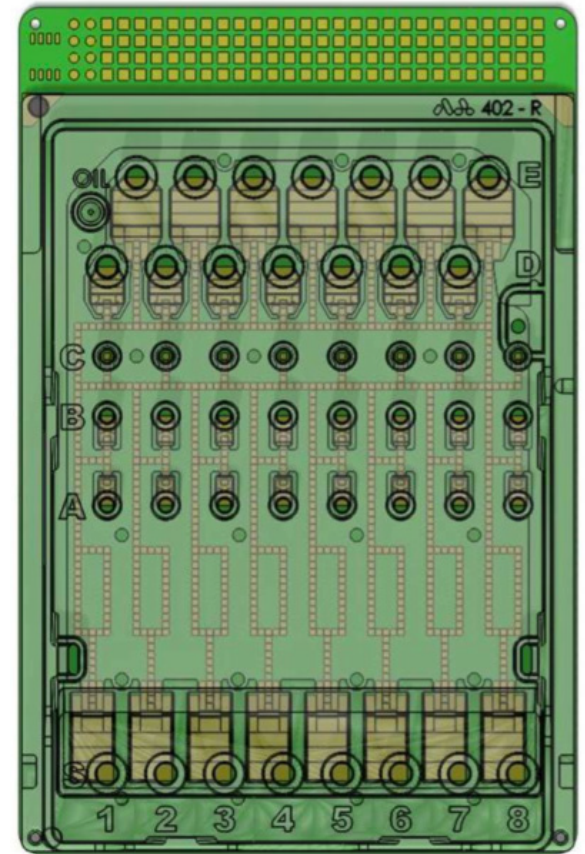
- Designed for one application
- Reduced costs
 - Production costs
 - Reagent costs



Application-Specific Biochips



Biochip for Newborn Screening
<http://www.liquid-logic.com/>



Biochip for Sample Preparation
<http://www.nugeninc.com/>

$$Cost_{\mathcal{A}} = \sum N_{M_i} \times Cost_{M_i}$$

where

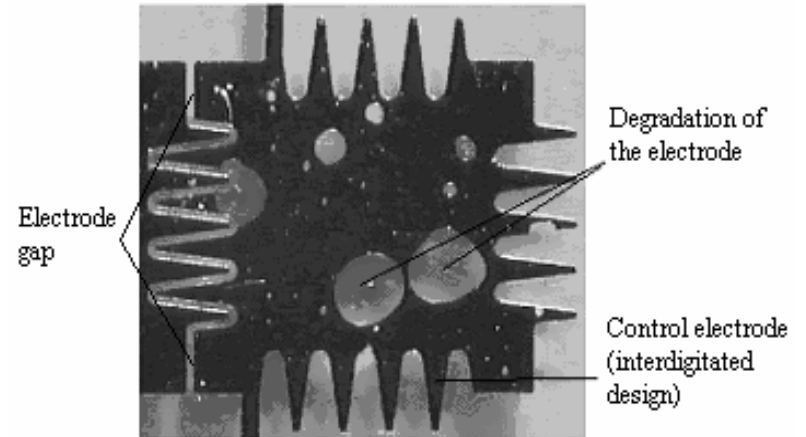
- **A** is the architecture
- **N_{M_i}** is the number of components of type M
- **Cost_{M_i}** is the cost of the physical component M_i

Component Library

Name	Unit cost	Dimensions (mm)	Time (s)
Electrode	1	1.5 × 1.5	N/A
Input Reservoir	3	1.5 × 4.5	2
Waste Reservoir	3	1.5 × 4.5	N/A
Capacitive Sensor	1	1.5 × 4.5	0
Optical Detector	9	4.5 × 4.5	8

Compilation: difficulties

- **Problem:** permanent faults
- **Importance:**
 - Increase the yield of DMBs
 - Improve the batch control



Insulator degradation

Problem: Architecture Synthesis

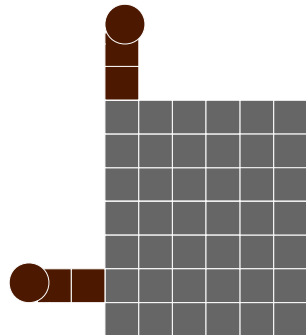


- **Given**
 - Biochemical application
 - Deadline requirements
 - Library of components (physical and virtual)
 - The number k of permanent faults
- **Determine**
 - An application-specific architecture \mathcal{A} , so that
 - the cost is minimized and
 - the application completes within deadline for any occurrence of the k permanent faults

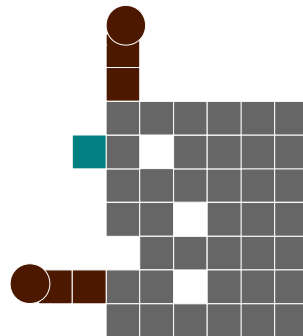
Optimization: SA moves

- Non-reconfigurable components (reservoirs, detectors)
 - Add/Remove
 - Change placement
- Reconfigurable elements (electrodes)
 - Add/Remove a single electrode
 - Add/Remove a row of electrodes on the side

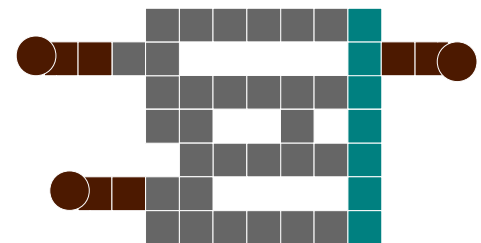
$$Objective(\mathcal{A}) = Cost_{\mathcal{A}} + W \times \max(0, \delta_G^k - D_G)$$



Initial architecture

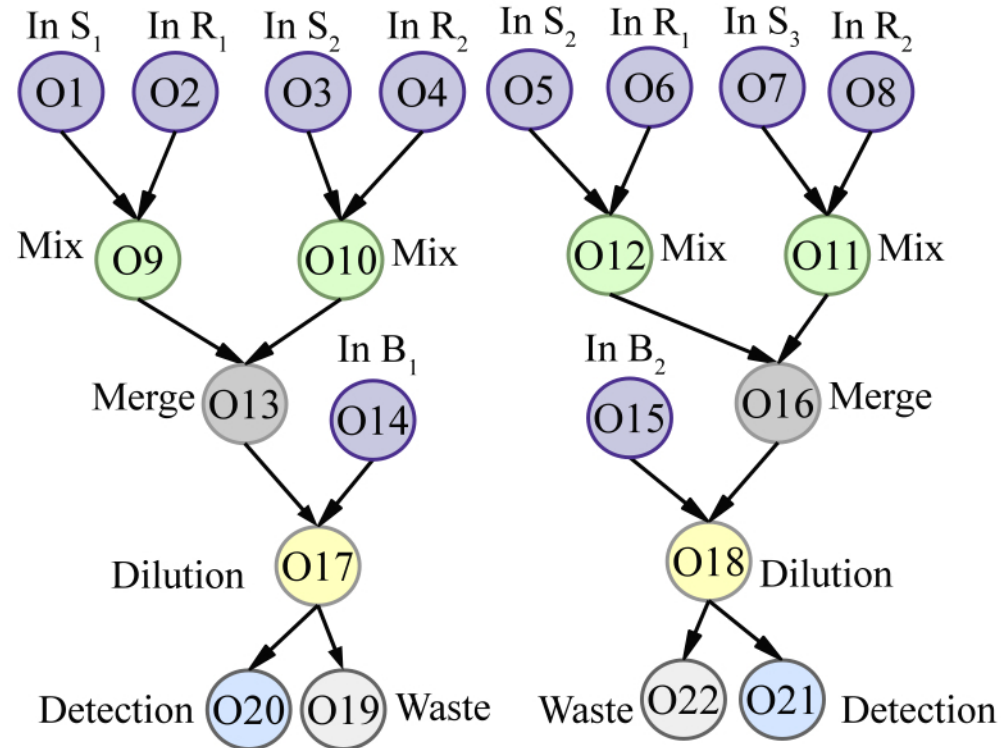


Add single electrode (green)
Remove single electrode (white)

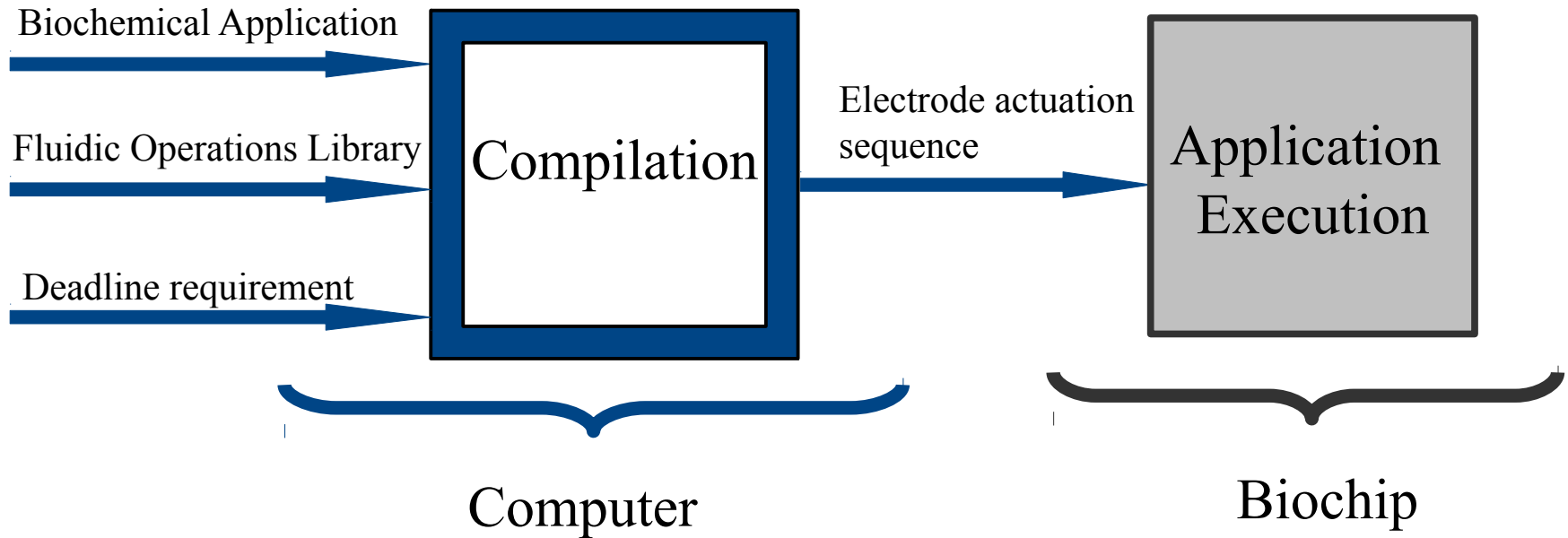


Add row of electrodes (green)
Change placement of reservoir (red)

Biochemical Application Model



Compilation Flow

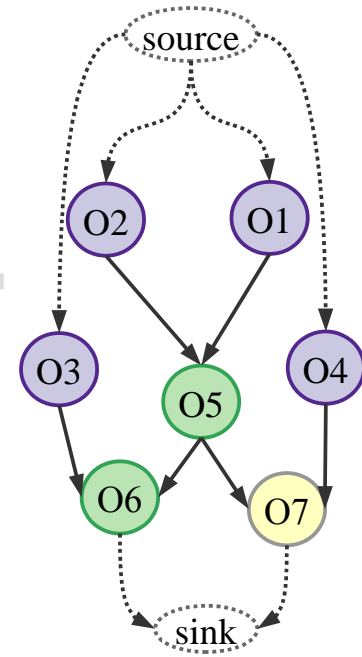


Compilation: Main steps

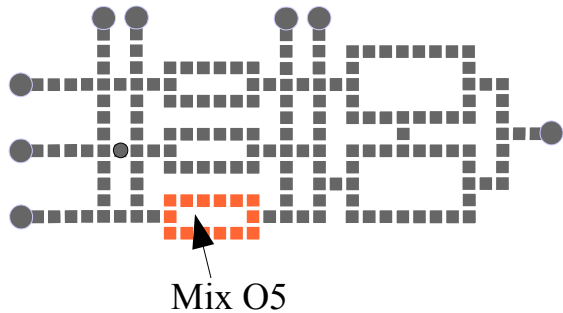
Allocation

Operation	Area	Time (s)
Mix	2x5	2
Mix	2x4	3
Mix	2x2	10
Detection	1x1	30

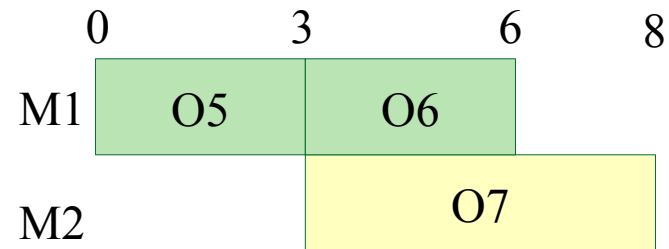
Binding



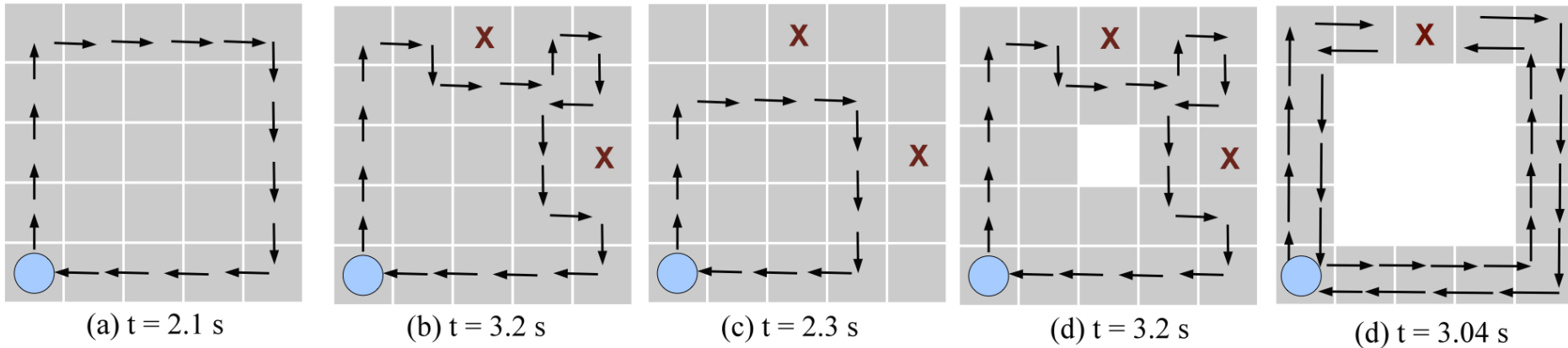
Placement



Scheduling



Fault-Tolerant Overhead



Evaluation of fault-tolerant overhead (faults are marked with X)

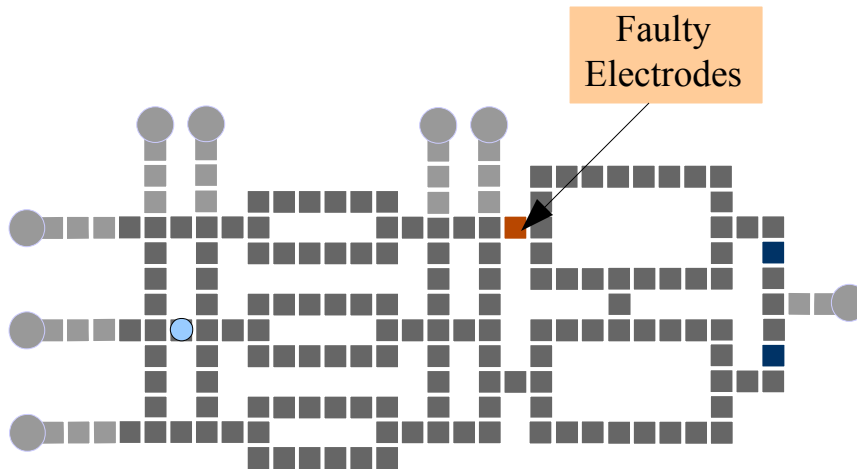
- Fault-tolerant overhead
 - Considers the impact of faults on the operation execution time
 - Routing-based operation execution, *Maftai 2012*

Compilation: difficulties

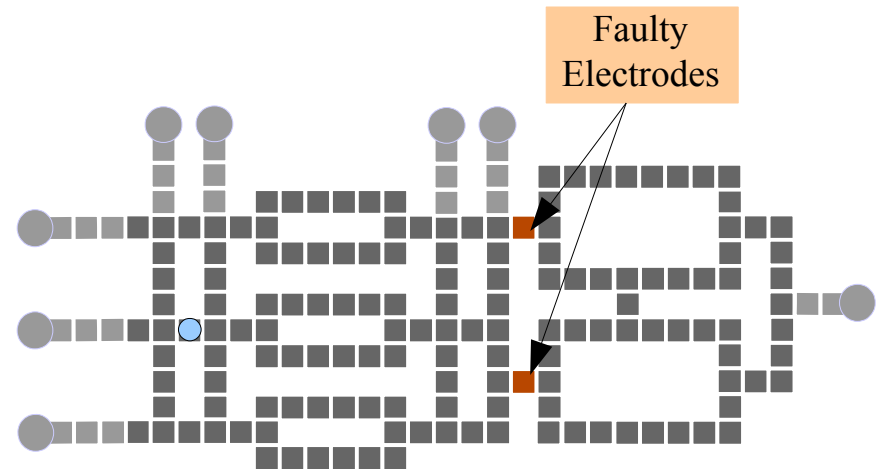


- **Problem:** fast compilation
- **Importance:**
 - It is part of an optimization loop
- **Solution:**
 - List-Scheduling based compilation
 - Routability test
 - Tests if, no matter where k faults are located, there is at least one route between any two electrodes

Routability test



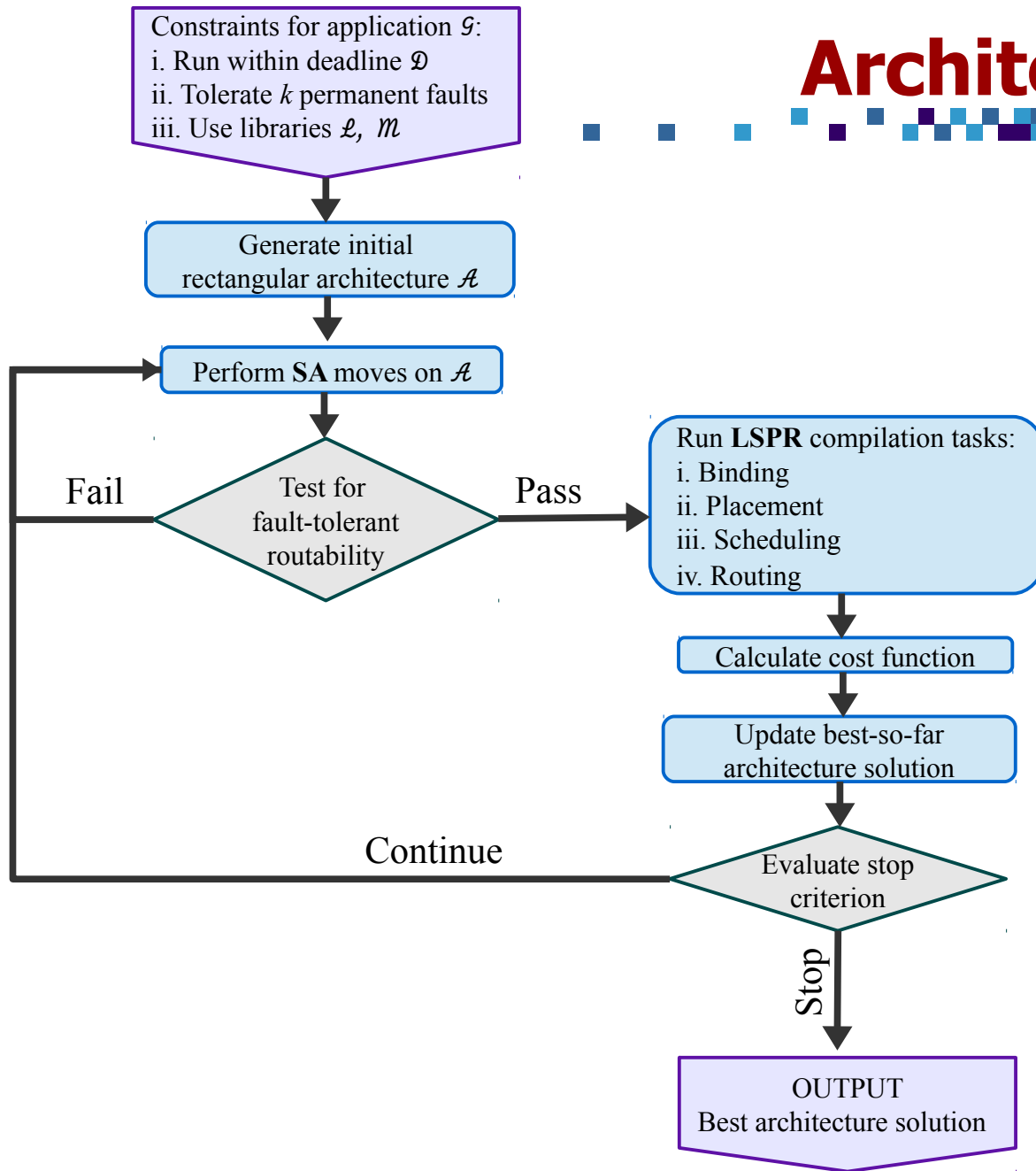
Routable architecture for 1 fault



Non-Routable architecture for 2 faults

- Routability test
 - Tests if, no matter where k faults are located, there is at least one route between any two electrodes
 - Algorithm that tests k -vertex connectivity in a graph, *S. Even (1973)*

Architecture Synthesis



- Simulated Annealing
- Test k-vertex connectivity
- List-scheduling
- Fast-template placement
- Hadlock routing

- Biochemical applications:
 - The mixing stage of polymerase chain reaction (PCR)
 - In-vitro diagnosis on human physiological fluids (IVD)
 - The colorimetric protein assay (CPA)
- Deadlines:
 - PCR – 10 s; IVD – 15 s; CPA – 100 s
- Implementation:
 - Java
- **Evaluation:**
 - Cost-effectiveness of the architectures resulted from our synthesis
 - Pessimism of List-Scheduling based compilation
 - Overhead in execution time due to permanent faults ($k=0,1,2$)

Experiments: Architecture synthesis

App.	$k = 0$				$k = 1$				$k = 2$			
	Arch	Cost	C_{SA}	T_{SA}	Arch	Cost	C_{SA}	T_{SA}	Arch	Cost	C_{SA}	T_{SA}
PCR	7×10 (1,1,1)	79	60	14	7×10 (1,1,1)	79	65	38	9×11 (1,1,1)	108	98	50
IVD	7×10 (2,2,2)	88	62	16	7×10 (2,2,2)	88	70	58	10×8 (2,2,2)	98	85	45
CPA	7×8 (2,1,2)	71	59	10	7×8 (2,1,2)	71	66	20	11×12 (2,1,2)	147	127	30

- Applications are resulted from our synthesis (col. 4, 8, 12)
- Compared with best possible general-purpose architectures (col. 3, 7, 11)
- **Our synthesis produces cheaper architectures**

Experiments: FT Overhead

App.	Cost	δ_G^0 (s)	δ_G^1 (s)	Deviation (%)	δ_G^2 (s)	Deviation (%)
PCR	98	8.42	8.81	4.6	9.43	11.9
IVD	85	12.62	13.11	3.8	14.81	17.3
CPA	129	153.9	169.3	10	190.11	23.5

- $k = 0$ faults (column 2)
- $k = 1$ faults (column 3)
- $k = 2$ faults (column 6)
- Applications are resulted from our synthesis
- Comparison with the case when there are no faults
- **Average deviation is 11.8%**

Experiments: LS Compilation

App. (ops.)	Arch.	$\delta_G^0 (s)$	Exec. time	$\delta_G^{opt} (s)$	Exec. time	Deviation (%)
PCR (7)	9×9	11	25 ms	10	60 min	9
IVD (28)	9×10	77	91 ms	73	60 min	5.4
CPA (103)	11×12	219	498 ms	214	60 min	2.3

- Near-optimal value is obtained with Tabu-Search, *Maftei 2010*
- General-purpose architectures
- No faults
- **Average deviation from near-optimal is 5.5%**

- SA-based architecture synthesis
 - List-Scheduling based compilation (fast)
- Reduced cost architectures
- Fault-tolerant architectures
- Increase the yield of DMBs

Backup slides



Optimization: Simulated Annealing

\mathcal{A}^0 - initial architecture

T^0 - initial temperature

T^L - temperature length

eps - cooling rate

temp = T^0 ;

$\mathcal{A} = \mathcal{A}^0$;

repeat

 while (temp < T^L) do

$\mathcal{A}^{new} = \text{moves}(\mathcal{A})$; *//generate new architecture*

 delta = Objective(\mathcal{A}) - Objective(\mathcal{A}^{best});

 if (delta < 0)

$\mathcal{A}^{best} = \mathcal{A}^{new}$;

 elseif (Math.random < $e^{-\text{delta}/\text{temp}}$) *//accept bad solutions with low probability*

$\mathcal{A}^{best} = \mathcal{A}^{new}$;

 endif

 endwhile

 temp = temp * eps;

until stop criterion is true

$$\text{Objective}(\mathcal{A}) = \text{Cost}_{\mathcal{A}} + W \times \max(0, \delta_{\mathcal{G}}^k - D_{\mathcal{G}})$$

ListScheduling($Graph, \mathcal{C}, \mathcal{B}, \mathcal{P}$)

1 **CriticalPath**($Graph$)

2 **repeat**

3 $List = \text{GetReadyOperations}(Graph)$

4 $O_i = \text{RemoveOperation}(List)$

5 $t_i^{start} = \text{Schedule}(O_i, \mathcal{B}(O_i), \mathcal{C}, \mathcal{P})$

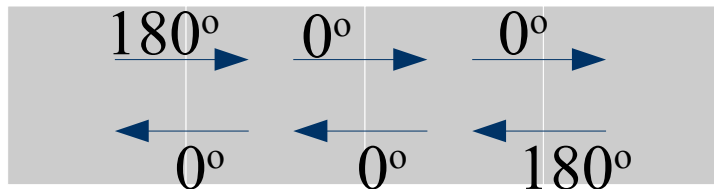
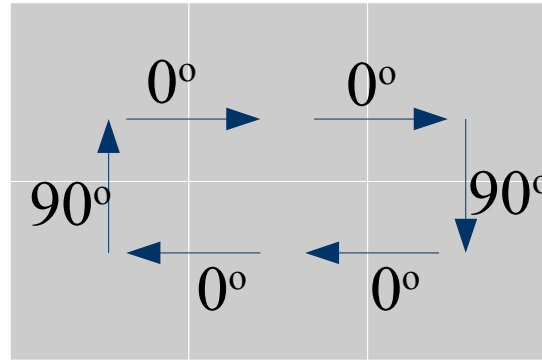
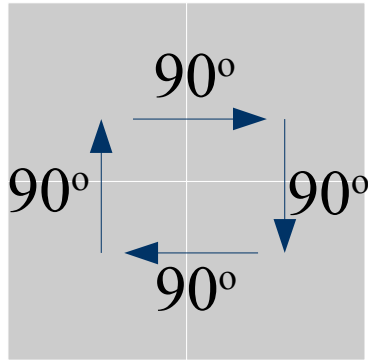
6 $t =$ earliest time when a scheduled operation terminates

7 $\text{UpdateReadyList}(Graph, t, List)$

8 **until** $List = \emptyset$

9 **return** \mathcal{S}

Routing-based Operation Execution



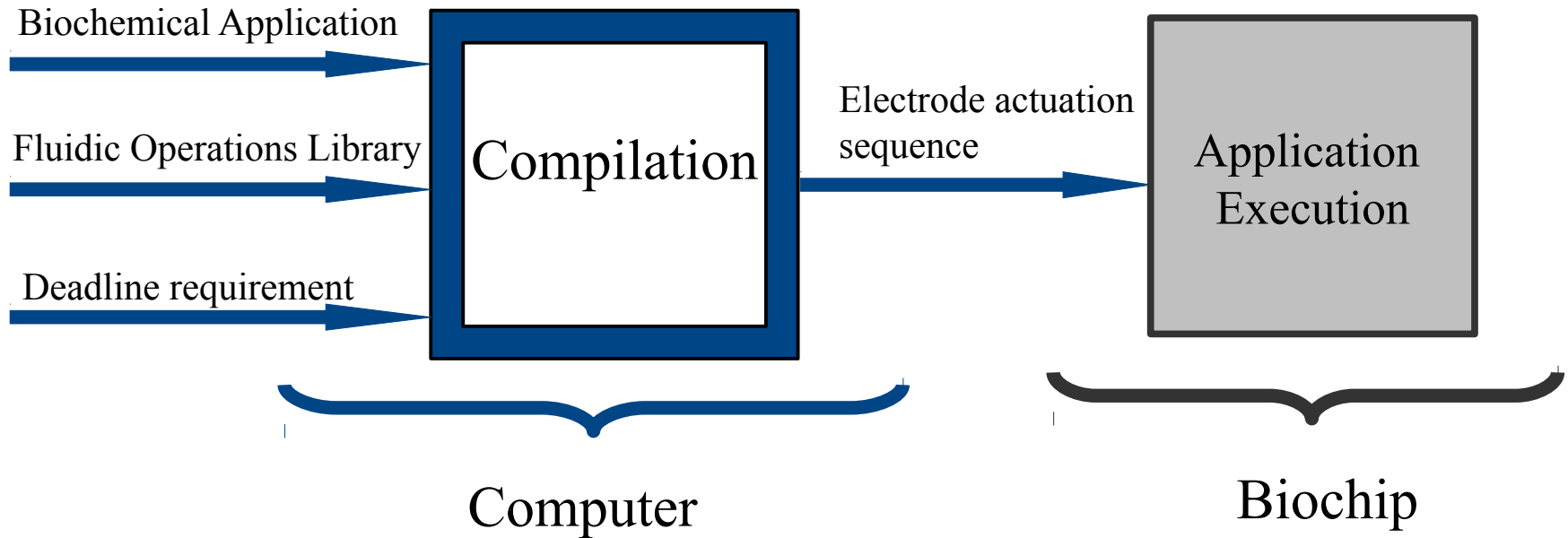
Operation	Area (cells)	Time (s)
Mix	2 x 2	10
Mix	2 x 3	6
Mix	1 x 4	5

$$p^{90} = 0.1\%$$

$$p^0 = 0.29\% \quad p^{00} = 0.58\%$$

$$p^{180} = -0.5\%$$

Compilation Flow

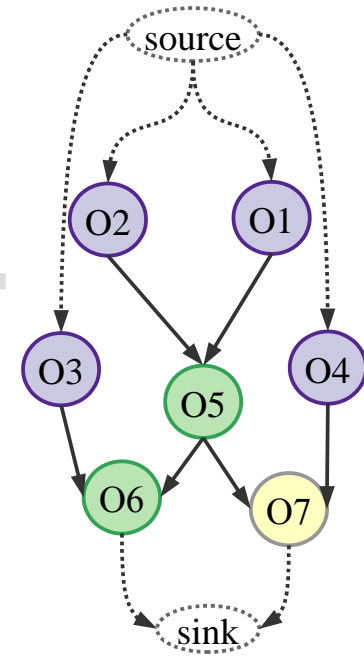


Compilation: Main steps

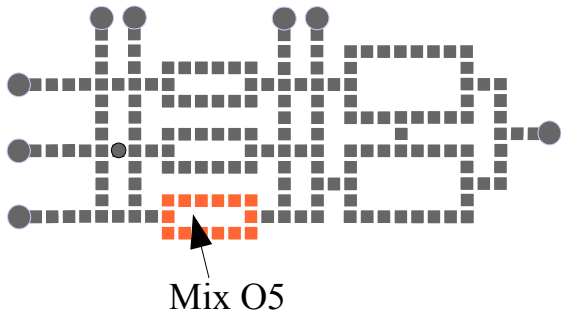
Allocation

Operation	Area	Time (s)
Mix	2x5	2
Mix	2x4	3
Mix	2x2	10
Detection	1x1	30

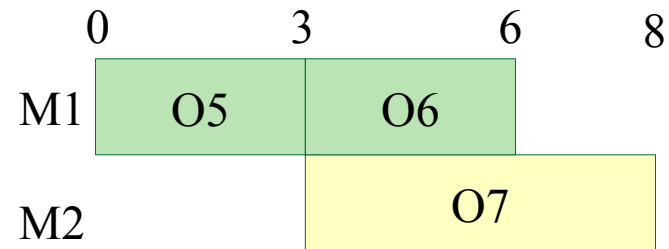
Binding



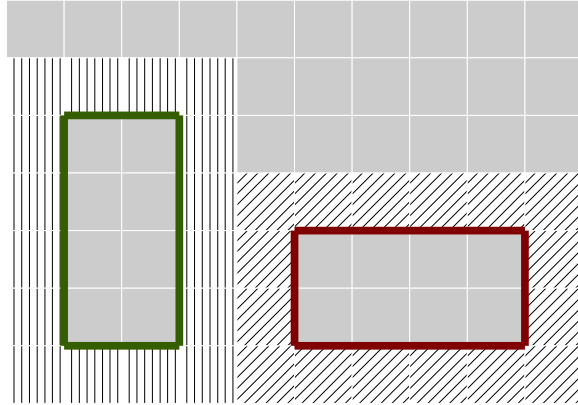
Placement



Scheduling

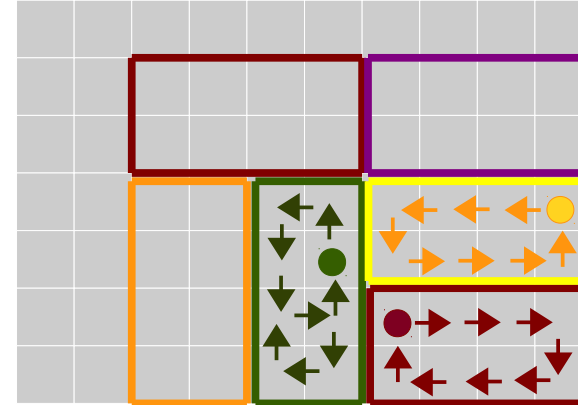


Droplet vs. Module Compilation



Module based

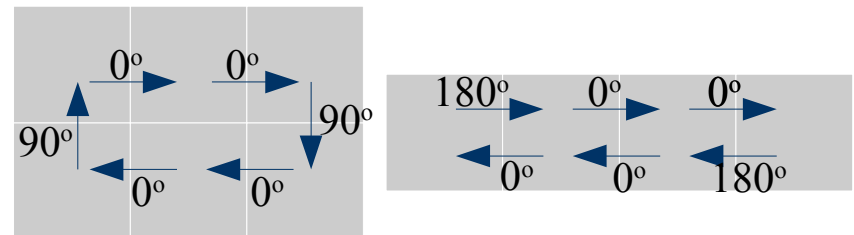
- module library
- black boxes
- protection borders



Droplet based

- routing based operation execution
- the position of the droplet is tracked
- better use of space

Operation	Area (cells)	Time (s)
Mix	2 x 4	3
Mix	2 x 2	4
Dilution	2 x 4	4



Component Library

Name	Unit cost	Dimensions (mm)	Time (s)
Electrode	1	1.5 × 1.5	N/A
Input Reservoir	3	1.5 × 4.5	2
Waste Reservoir	3	1.5 × 4.5	N/A
Capacitive Sensor	1	1.5 × 4.5	0
Optical Detector	9	4.5 × 4.5	8

Virtual Devices Library

Op.	Shape	Time (s) no faults	Time (s) $k = 1$	Time (s) $k = 2$
Mix	3 × 6	2.52	2.71	3.77
Mix	5 × 8	2.05	2.09	2.3
Mix	4 × 7	2.14	2.39	2.51
Mix	5 × 5	2.19	2.28	2.71
Mix	5 × 5 × 1	2.19	2.73	3.92
Mix	5 × 5 × 2	3.98	5.82	7.56
Dilution	3 × 6	4.4	4.67	4.11
Dilution	5 × 8	3.75	4.76	6.3
Dilution	4 × 7	3.88	4.22	4.46
Dilution	5 × 5	3.98	4.12	4.67
Split	1 × 1	0	0	0
Storage	1 × 1	N/A	N/A	N/A

Test for k-vertex connectivity

Algorithm 1:

- (1) For every i and j such that $1 \leq i < j \leq k$ check whether there are k vertex-disjoint paths between them. If for some i and j the test fails, then G 's connectivity is less than k .
- (2) For every j , $k+1 \leq j \leq n$, form \tilde{G} and check whether there are k vertex-disjoint paths between a and j . If for some j the test fails, then G 's connectivity is less than k .
- (3) The connectivity of G is at least k .

Complexity $O(kn^3)$

Capacitive Sensor

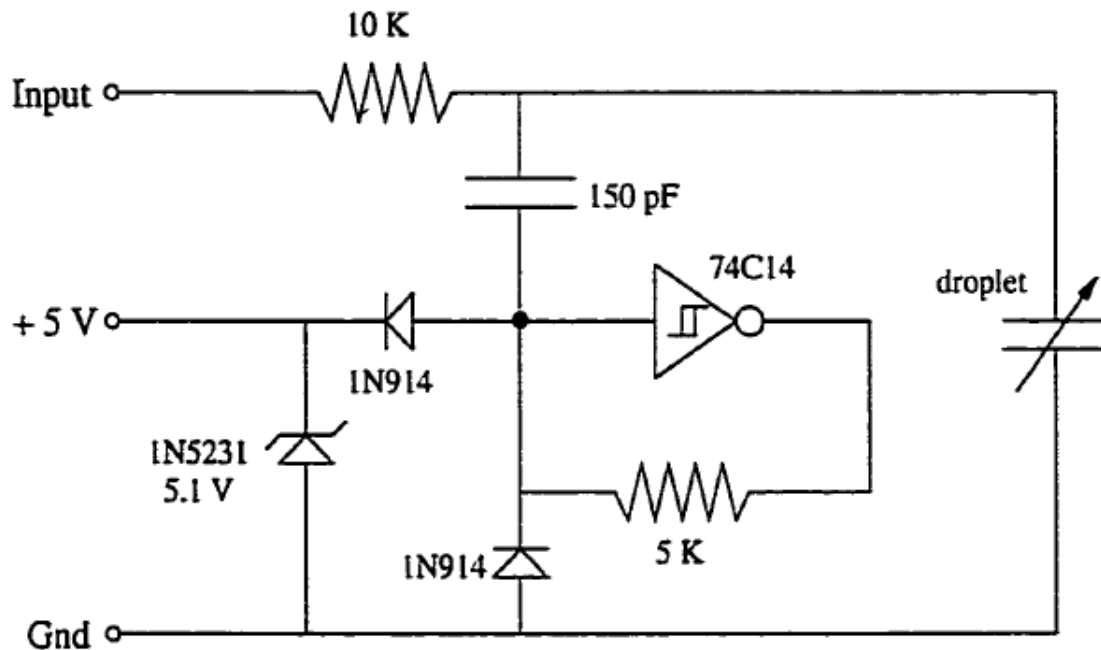


Figure 5.7: Ring-oscillator circuit for capacitance measurements of droplets [Pollack*]

Biochip Parameters

Parameter	Range or conditions
Gap height, h	0.19 – 1.04 mm, with $L = 1.5$ mm and $D/L \approx 1.3$
Droplet diameter, D	1.31 – 2.18 mm, with $L = 1.5$ mm and $L/h \approx 5$
Electrode pitch, L	0.15 – 1.50 mm, with $L/h \approx 5$ and $D/L \approx 1.3$
Electrode shape	interdigitated, square, circular, wedge
Medium type	air, silicone oil
Droplet viscosity	1.0 – 58.4 cP
Oil viscosity	0.8 – 9.4 cP
Droplet volume, V_D	3 nl – 3 μ l
Velocity, v	0 – 10 cm/s

Table 4.1: Parameter space for droplet dynamic investigations [Pollack*]

Glucose detection

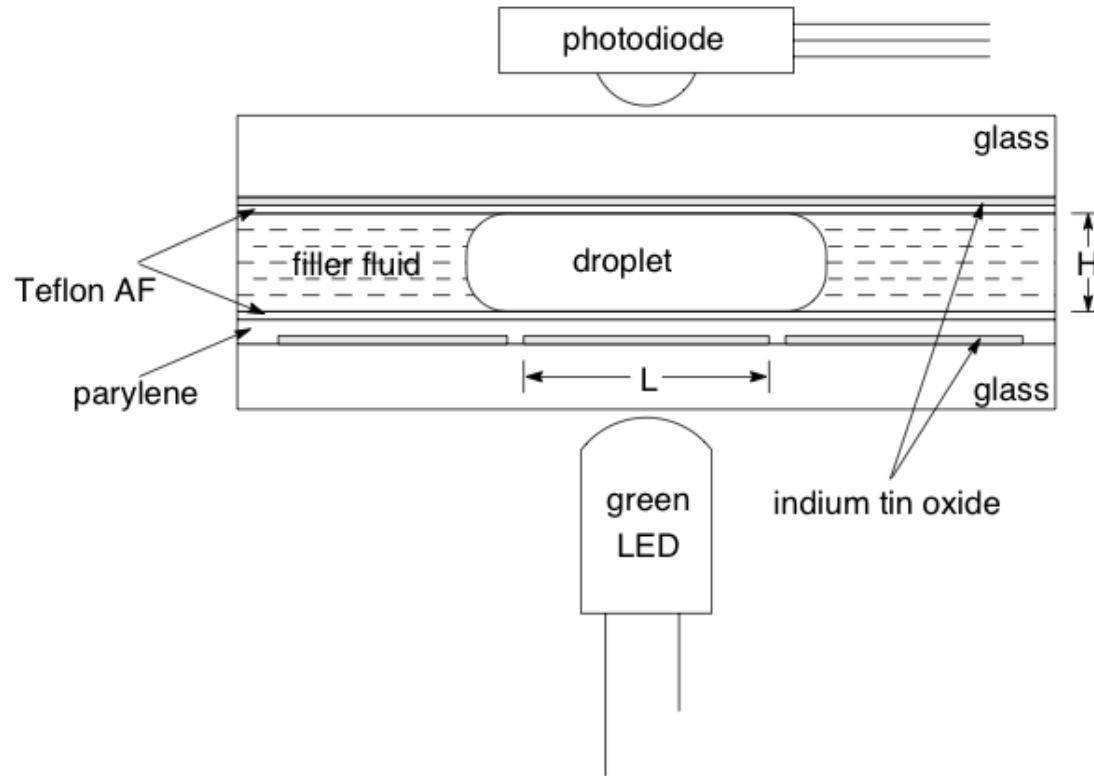
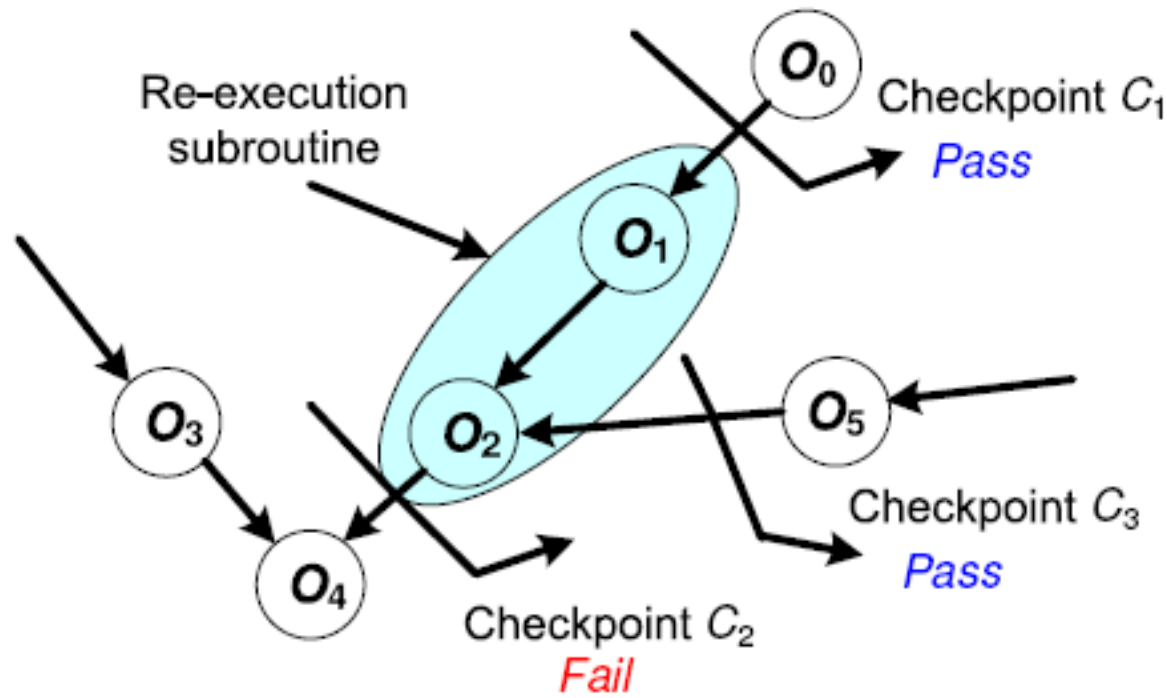


Fig. 2 Setup for glucose detection

Srinivasan 2003, Clinical diagnostics on human whole blood, plasma, serum, urine, saliva, sweat, and tears on a digital microfluidic platform

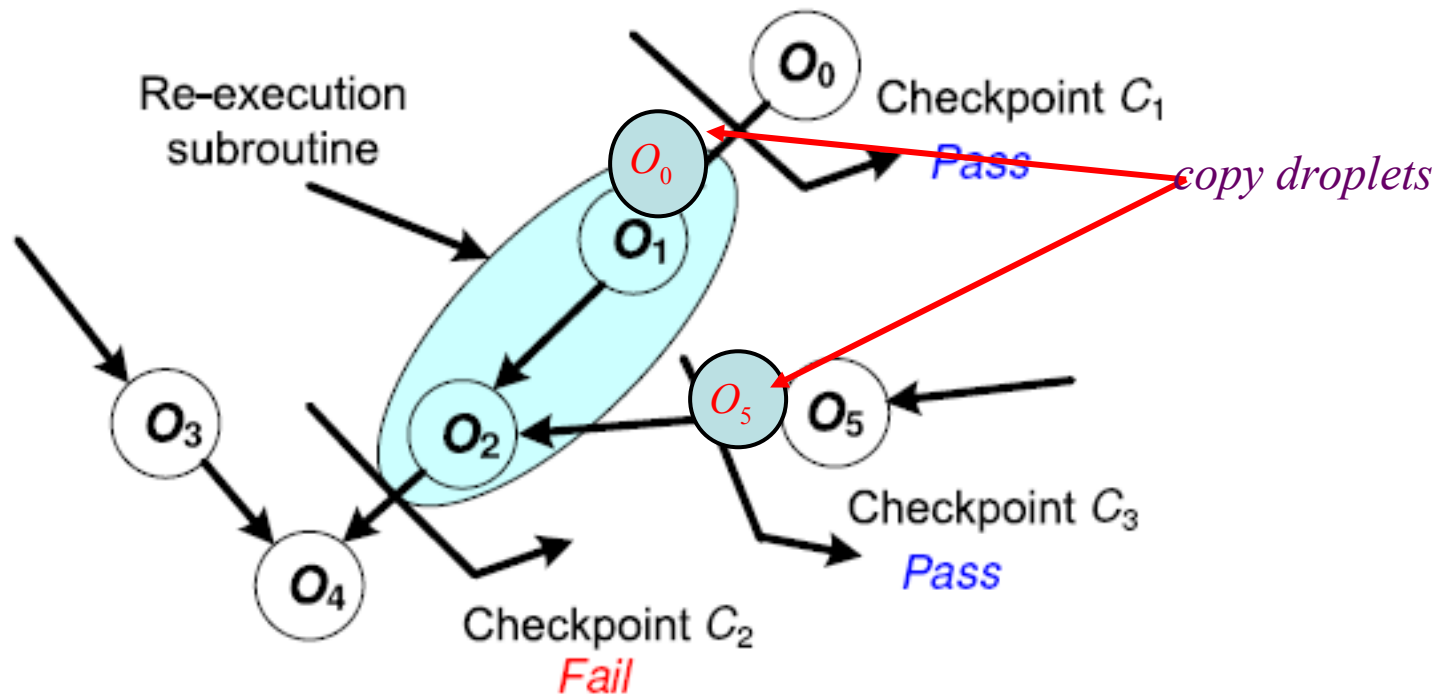
Control-Path Design

- Add *checkpoints* to monitor outcomes of fluidic operations
 - **Checkpoint**: storage of the intermediate product droplet
- Assign each checkpoint a *re-execution subroutine*
 - **Subroutine**: fluidic operations between checkpoints



Control-Path Design

- Extra **copy droplets** needed
- **Checkpoints:** where ?
- Costs:
 - Time
 - Area

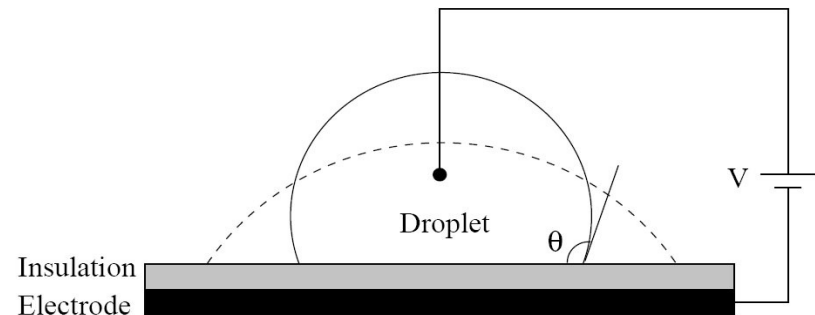
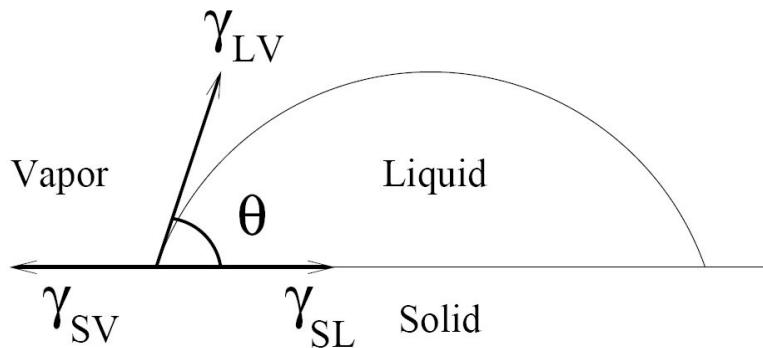


And the droplet moves!

- Electrowetting on dielectric principle (EWOD)
 - Electrical modulation of the solid-liquid interfacial tension

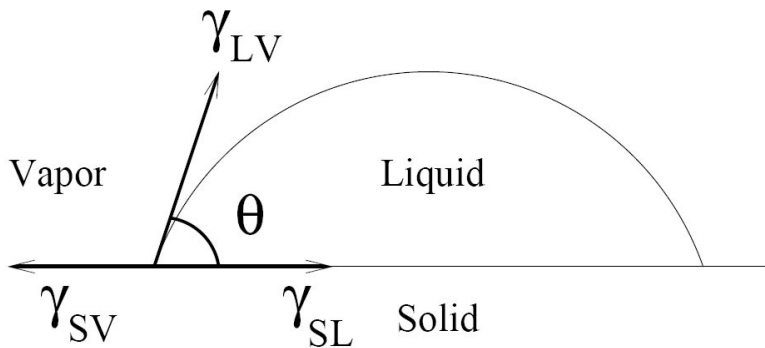
Young equation

$$\gamma_{LV} \cos(\theta) = \gamma_{SV} - \gamma_{SL}$$



Electrowetting: Physical Principles (I)

- Motion of droplets is based on the differences between contact angles in the advancing and receding lines of a droplet.
- When a droplet rests on a non-wetting solid surface, the forces acting at the solid-liquid-vapor interface equilibrate and result in a contact angle θ between the droplet and solid, as described by Young's equation,

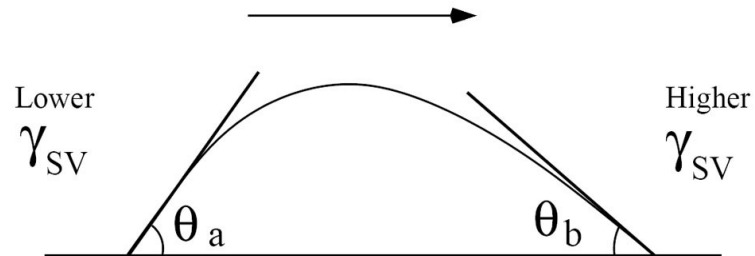


$$\gamma_{LV} \cos(\theta) = \gamma_{SV} - \gamma_{SL}$$

γ_{LV} , γ_{SV} and γ_{SL} are the liquid-vapor, solid-vapor and solid-liquid surface energies

Electrowetting: Physical Principles (II)

- When an imbalance in these surface energies occurs (as in the case of a droplet resting on a surface with a gradient surface energy), a net force is induced
 - Initiate droplet motion
- Imbalance can be induced by chemical, thermal, or electrostatic means
 - In the case of thermally-induced droplet motion, a surface tension gradient can be induced by differentially heating the ends of a droplet, since the surface tension of a liquid decreases with temperature.



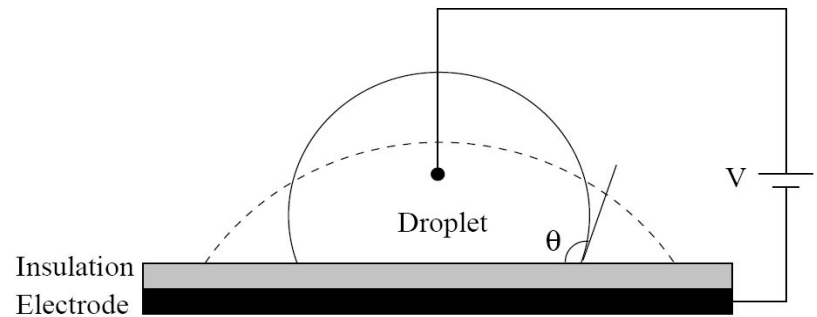
Electrowetting: Physical Principles (III)



- Electrowetting-based actuation of droplets: electrical fields used to induce surface tension gradients.
 - Electrowetting effect \Rightarrow the surface energy can be directly modified by the application of an electric field
- Consider a droplet resting on a electrode separated by a hydrophobic insulator
 - A potential is applied between the droplet and the electrode, resulting in a capacitive energy E stored in the insulator. The resulting energy is:

$$E = \frac{\epsilon_0 \epsilon_r A}{2d} V^2$$

$$\Rightarrow \gamma_{SL}(V) = \gamma_{SL}(0) - \frac{\epsilon_0 \epsilon_r A}{2d} V^2$$



Contact angle

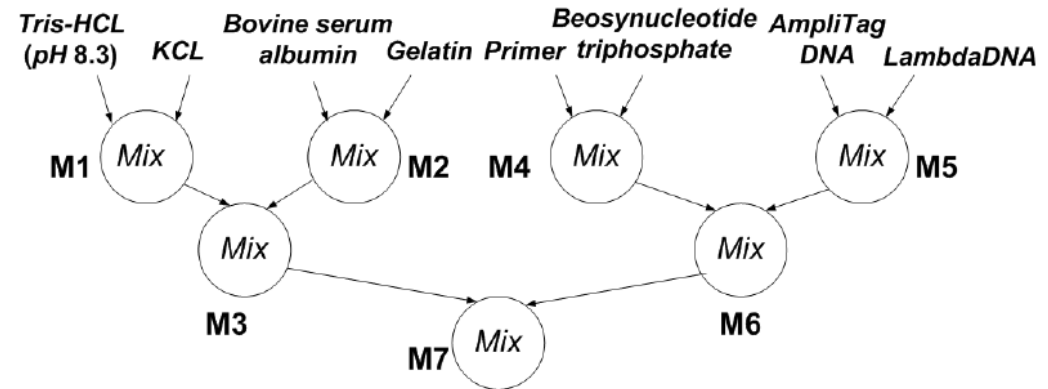
change:

$$\cos \theta(V) = \cos \theta(0) - \frac{\epsilon_0 \epsilon_r A}{2d \gamma_{LV}} V^2$$

Reference: P. Y. Paik, V. K. Pamula and K. Chakrabarty, "Adaptive Cooling of Integrated Circuits using Digital Microfluidics", Artech House, Norwood, MA, 2007.

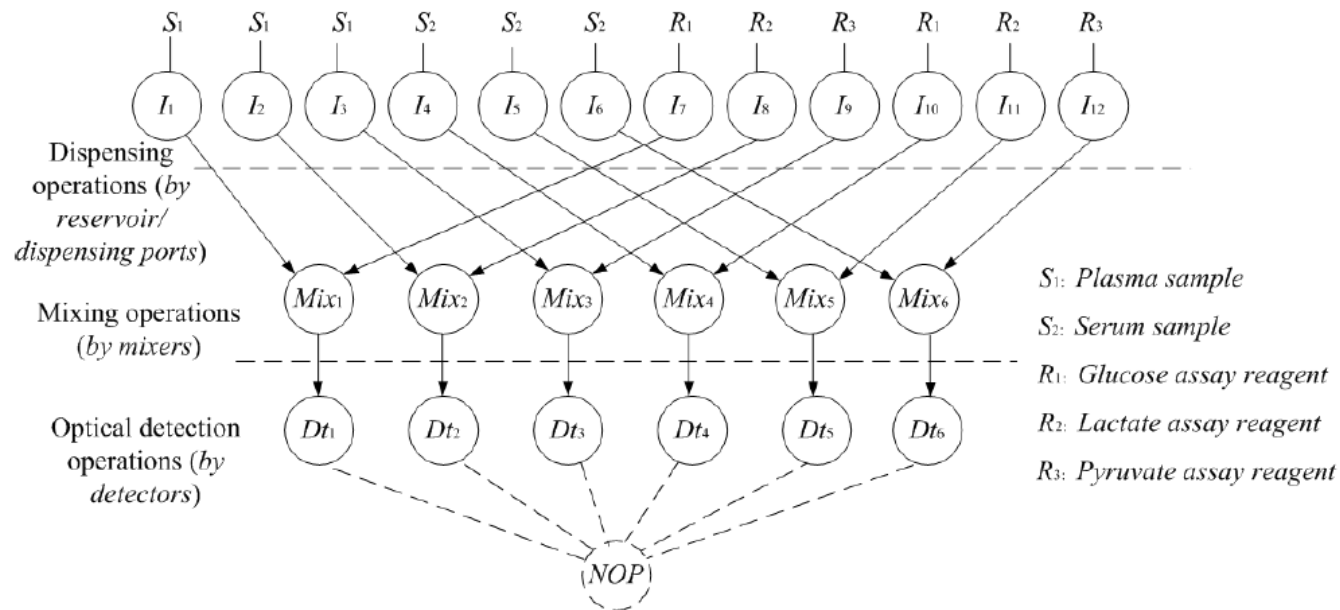
Benchmarks: PCR

- Mixing stage for Polymerase Chain Reaction
- Electrode pitch: 1.5 mm, Gap height: 600 μm



Operation	Hardware*	Module	Mixing time
mixing	2x2 electrode array	4x4 cells	10s
	4-electrode linear array	3x6 cells	5s
	2x3 electrode array	4x5 cells	6s
	2x4 electrode array	4x6 cells	3s

- Multiplexed in-vitro diagnosis
- Electrode pitch: 1.5 mm, Gap height: 600 μm



- Colorimetric Protein Assay
- Electrode pitch: 1.5 mm, Gap height: 600 μm

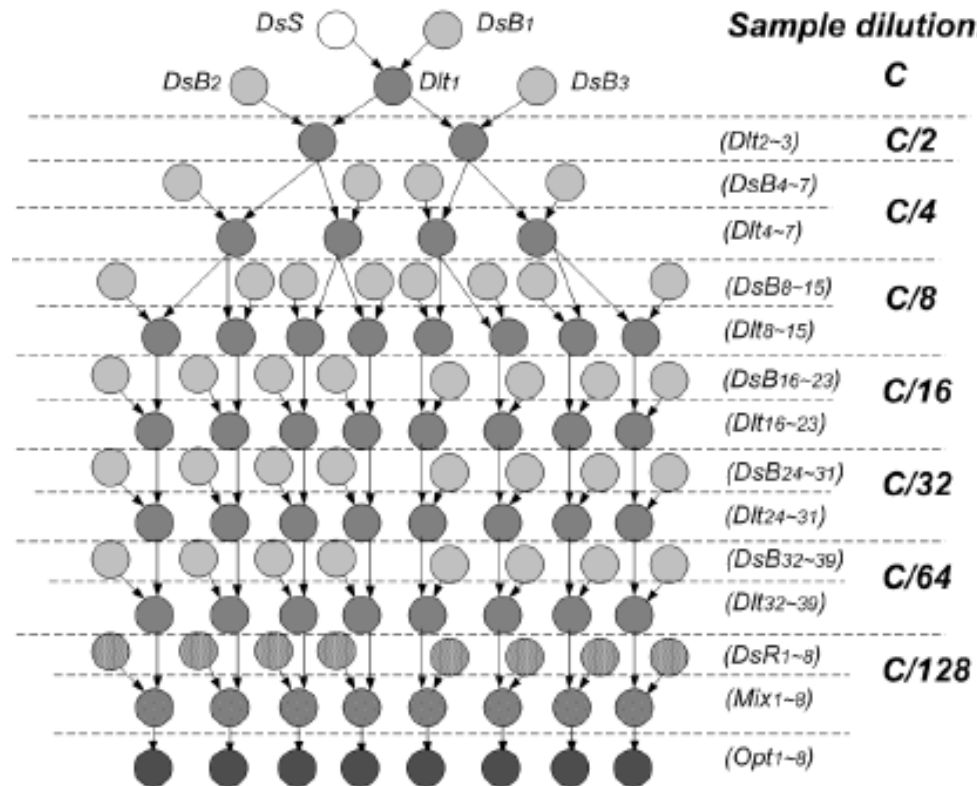


Figure 3-2. Sequencing graph for a protein assay.