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Factorization Based Dilution of Biochemical Fluids with Micro-Electrode-Dot-Array Biochips

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Session 6C: New Trends in Biochips

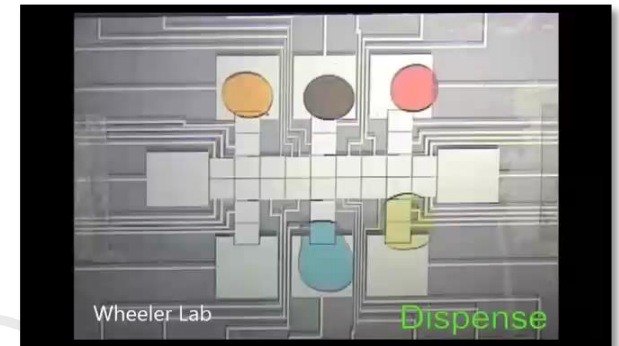
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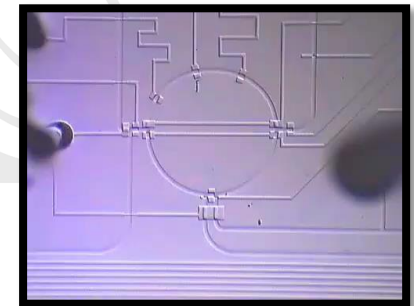
- Introduction to Microfluidic Biochips
- Sample Preparation using Microfluidic Biochips
- Proposed Approach: *FacDA*
- Conclusions

Introduction: Microfluidic Biochips

- Microfluidic biochips manipulate sub-milliliter scale volumes of fluid to perform different bioassays
- Microfluidic biochips are broadly classified as:
 - Digital Microfluidic biochips (DMF):
 - Electrowetting-on-dielectric (EWOD) based technology
 - Manipulates droplets of fluids
 - Continuous-Flow Microfluidic biochips (CMF):
 - Micro-pump and micro-valve based technology
 - Manipulates fluids in micro-channels



Working of DMF biochip [1]



Working of CMF biochip [2]

Sample Preparation: Automation and Optimization



- It refers to the generation of droplets of a particular concentration factor (CF) of a sample fluid OR volumetric ratio of some fluids
- In many biological assays, it is an inherent step that needs longer time using macroscopic biological systems
- Dilution: a particular biological sample and a buffer are involved
- Mixing: multiple sample and reagent fluids are involved
- **Inputs:** sample (S, 100% conc.), buffer (B, 0% conc.), target-concentration (OR volumetric ratio) C_t , concentration error-tolerance (\mathcal{E}), mixing volume constraint (M)
- **Objectives:** minimize sample or reagent count (n_s), waste count (n_w), mixing steps (n_m)

Sample Preparation using DMF and CMF

- Several algorithms proposed for both DMF and CMF biochips
- Different algorithms have different optimization criteria
- Dilution algorithms for CMF can be used for DMF as well, if $(n : n)$ mixing model is followed
- Sample preparation algorithms:

DMF

- *twoWayMix* [2]
- *DMRW* [3]
- *GORMA* [4]
- *NFSP* [5]

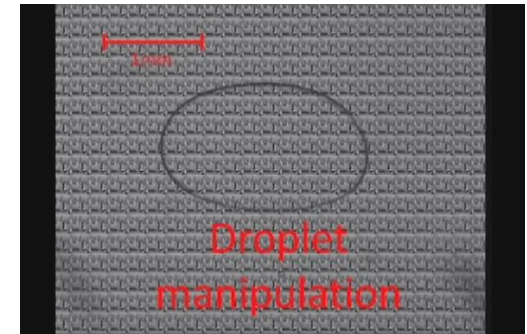
CMF

- *NWayMix* [2]
- *TPG* [6]
- *VOSPA* [7]
- *FloSPA* [8]

Properties	DMF	CMF
Mixing model	(1:1)	Depends on the size of the mixer
Reconfigurability	Yes	No
Mixing time	Relatively more	Less

Micro-Electrode-Dot-Array (MEDA) DMF

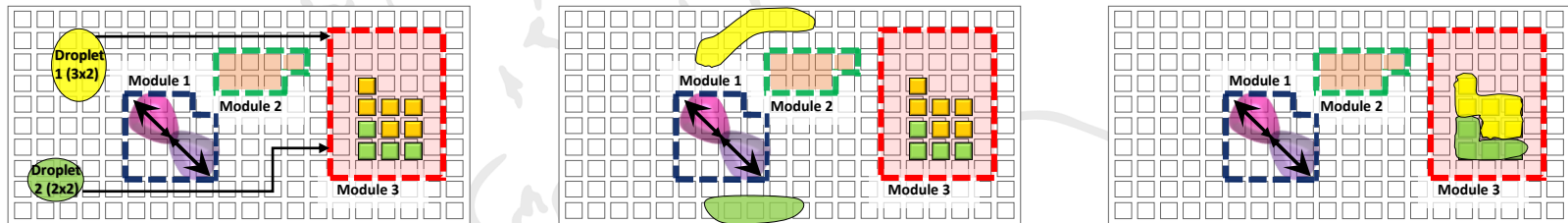
- MEDA-DMF [9, 10] is EWOD-based, sea-of-microelectrodes technology
- Many advantages over DMF and CMF:
 - Manipulates droplets with higher granularity
 - More reconfigurable on-chip components
 - Addition droplet routability and more flexibility in droplet-shape
 - Equipped with in-built sensors and hence suitable for cyber-physical error management
- New EDA tools needed for mapping biochemical assays on MEDA-DMF
- Sample preparation with MEDA-DMF possesses new challenges



Working of MEDA biochip [9]

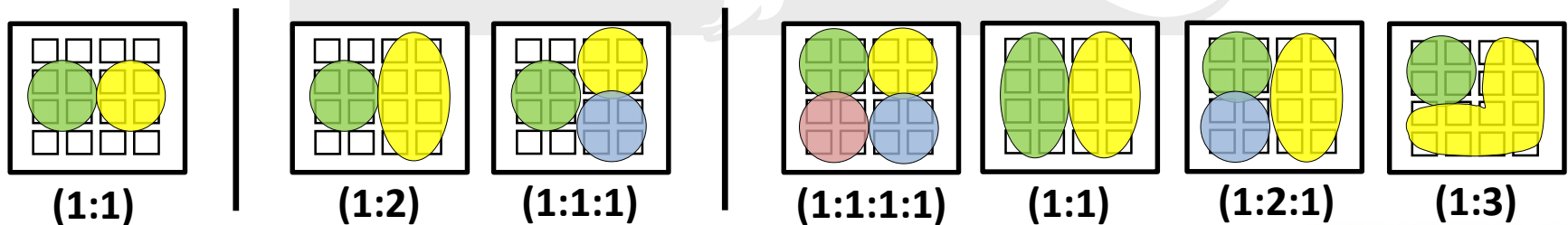
Sample Preparation in MEDA

- Advantages of sample preparation in MEDA-DMF biochips:
 - Significant reduce in mixing time due to SAR (split-and-rotate) based laminar mixing
 - More mixing models that can reduce mixing steps/time
 - More reliable sample preparation due to integrated sensors with each micro-electrode



Fluid flow on MEDA

- Mixing models with a mixing volume constraint, $M = 4$ for MEDA



Previous Work

- **WSPM** [11]: – Dilution algorithm exploiting various mixing models of MEDA

Weighted
Sample
Preparation
Method

- *WSPM* works as follows:
 - Generates a set of primary droplets (*PDs*) of intermediate *CFs*
 - Determines dilution graph for all *PDs*
 - Obtain minimum cost dilution graph

- **FloSPA-D** [8]: – Dilution algorithm for CMF biochip that can be tuned for MEDA

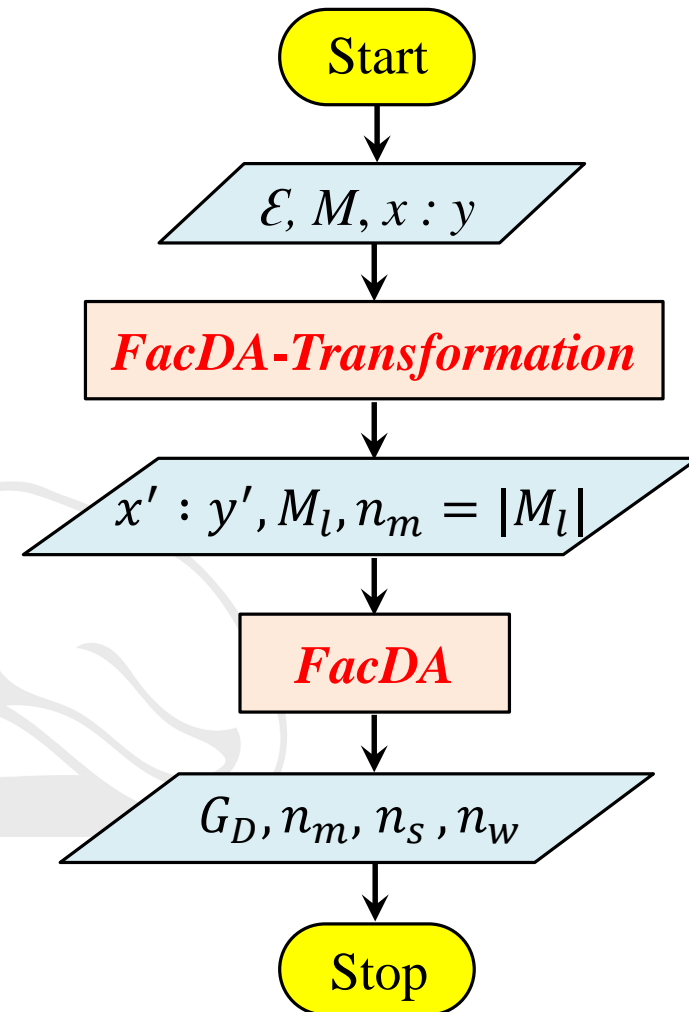
Flow-based
Sample
Preparation
Algorithm for
Dilution

- Mixing tree with minimum number of mixing operations generated by *NWayMix* [8]
- SMT-based approach reduces the sample or reagent count, n_s

Proposed Approach

- **Factorization based Dilution Algorithm for MEDA-DMF biochip (***FacDA***)**

- Exploits all possible mixing models for a mixer of size M
- Assures a dilution graph with minimum height that reduces mixing steps/time, n_m
- Uses a Satisfiability Modulo Theories (SMT) based approach to reduce the sample or reagent count, n_s

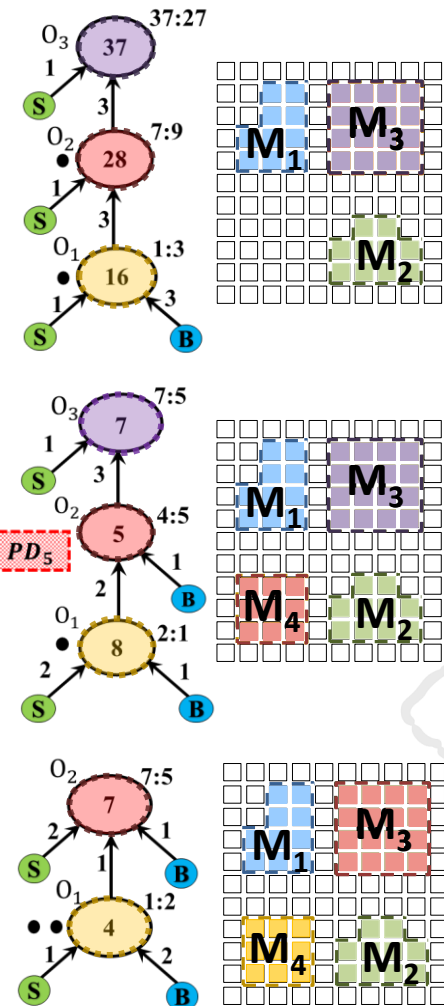


Motivating Example

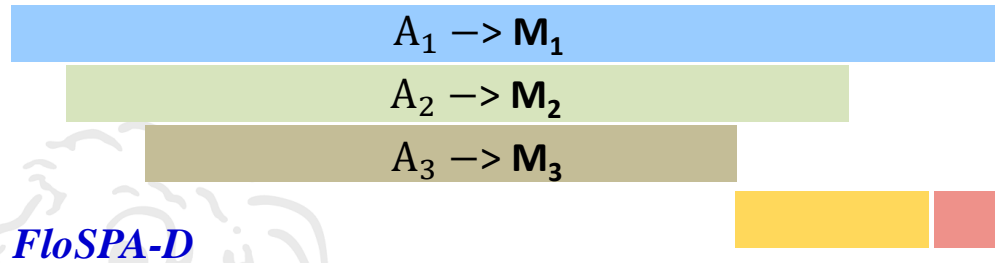
$$C_t = \frac{7}{12}, \mathcal{E} = 0.02, M = 4, \text{transformed } C_t = \frac{37}{64}$$

● : waste droplet

Time →



Previous Operations



WSPM



FacDA



Module	Size	M
M ₁	10	3
M ₂	10	3
M ₃	16	4
M ₄	9	3

FacDA-Transformation

Problem Formulation:

Inputs: Target ratio *sample* : *buffer* = $x : y$, where $x + y = N$, concentration error-tolerance \mathcal{E} , mixing volume constraint M .

Output: Transformed ratio $x' : y'$, feasible mixer-set (M_ℓ).

- Find all possible factors of N
- Sort the factor sets in non-increasing order of set-cardinalities
- Choose the first factor set whose every element is less than M , in many cases this will give us an exact solution

$$\begin{aligned} x : y &= 233:7, \\ \epsilon &= 0.000005, \\ M &= 5 \end{aligned}$$

$$240 = \{ \dots, \boxed{12} * 4 * 5, \dots, 2 * 4 * 5 * \boxed{6}, \boxed{3 * 4 * 4 * 5}, \dots \}$$

×
×
✓
 M_ℓ

- If no such factor set exists, then approximate the given ratio maintaining the error-tolerance by increasing N and repeating the above steps

- Order of the factors in the final mixer-set decided by a greedy approach called as *Division-by-Factor* method
- A set of constraints is imposed on the skeleton of the final dilution graph and a *SMT-based approach* reuses the fluids of intermediate CFs

$$C_t = \frac{233}{240}$$

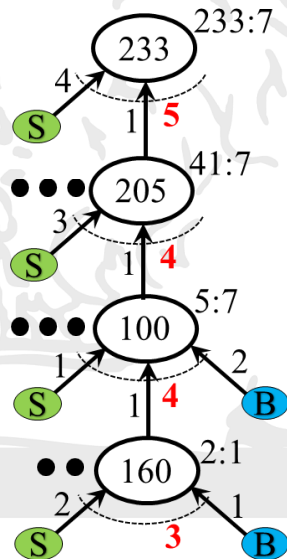
$$M_\ell = \{3, 4, 4, 5\}$$

Factor ($\in M_\ell$)	Remainder		Quotient	
	#S	#B	233	7
3	2	1	77	2
4	1	2	19	0
4	3	0	4	0
5	4	0	0	0

Total = 10

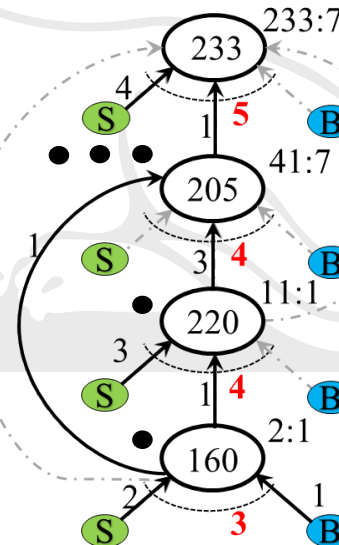
$$n_m = 4, n_s = 10, n_w = 8$$

Division-by-Factor



$$X_i = \frac{\frac{X_{i+1}}{\prod_{j=d}^{i+1} f_j} + k + (\dots)}{\prod_{j=d}^{i+1} f_j} = \frac{X_{i+1} + k \prod_{j=d}^{i+1} f_j + (\dots)}{\prod_{j=d}^{i+1} f_j}$$

$$X_i = \frac{\frac{X_{i+1}}{\prod_{j=d}^{i+1} f_j} + k \frac{X_p}{\prod_{j=d}^p f_j} + (\dots)}{f_i} = \frac{X_{i+1} + k X_p \prod_{j=p-1}^{i+1} f_j + (\dots)}{\prod_{j=d}^i f_j}$$



$$n_m = 4, n_s = 9, n_w = 5$$

SMT-based Approach

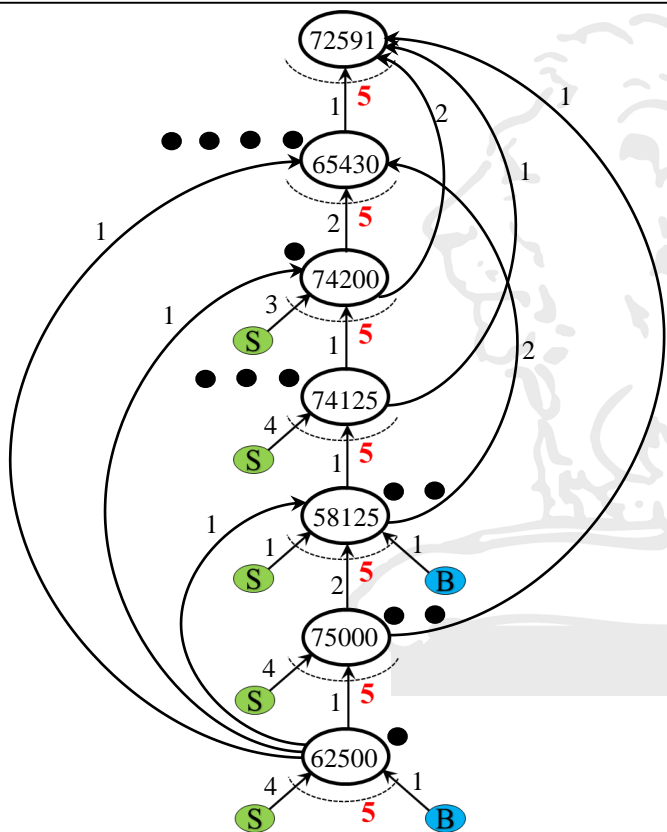
Comparative Example

$C_t = \frac{233}{240}$, $\mathcal{E} = 0.000005$, $M = 5$, transformed $C_t = \frac{72591}{57}$ • : waste droplet

$S = \{78125/5^7\}$, $B = \{0/5^7\}$

$n_m = 7$

$n_s = 16, n_w = 13$



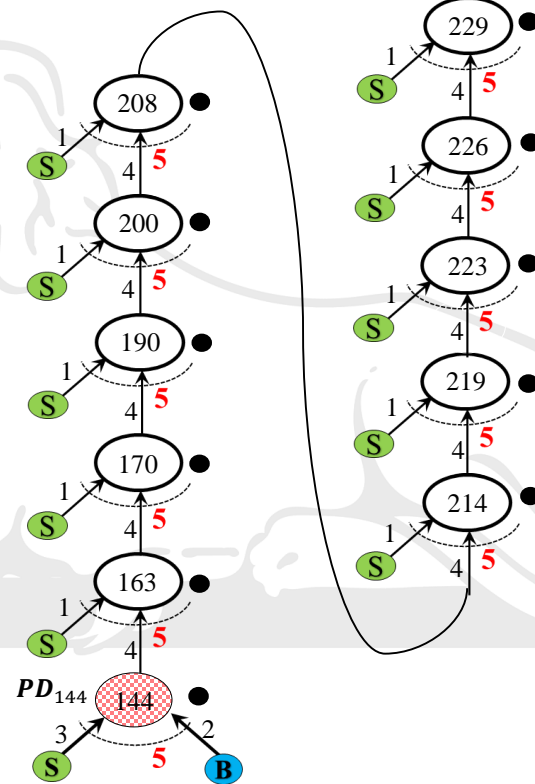
FloSPA-D

$S = \{240/240\}$

$B = \{0/240\}$

$n_m = 13$

$n_s = 15, n_w = 12$



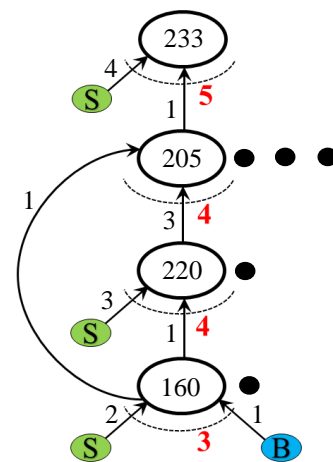
WSPM

$S = \{240/240\}$

$B = \{0/240\}$

$n_m = 4$

$n_s = 9, n_w = 5$

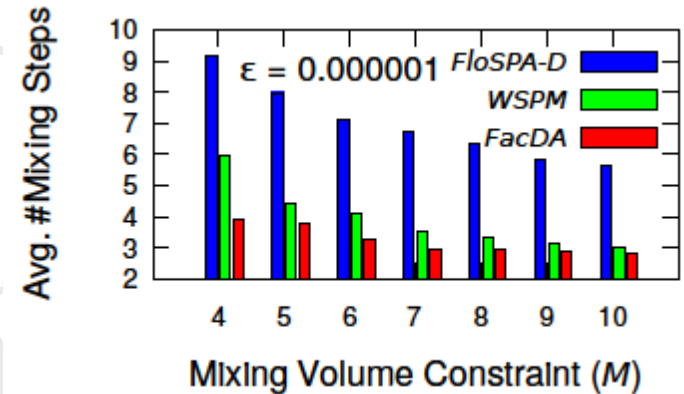
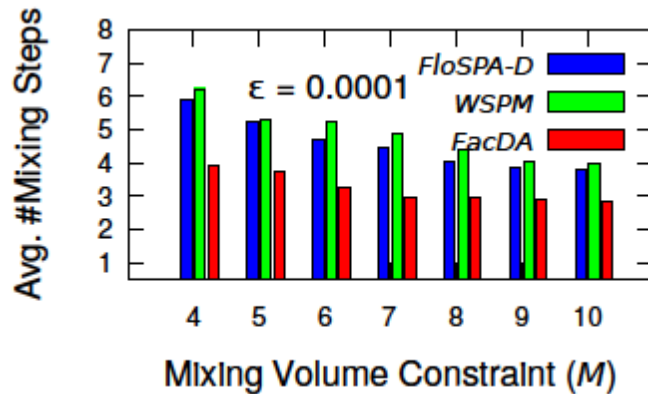


FacDA

Simulation Results

- A set of total 19,947 unique target ratios of *sample* and *buffer* considered, where the ratio-sum varies between 2 and 256
(i.e., $1:1, 1:2, 2:1, 1:3, 3:1, 1:4, 2:3, 3:2, 4:1, \dots, 1:255, \dots, 255:1$)

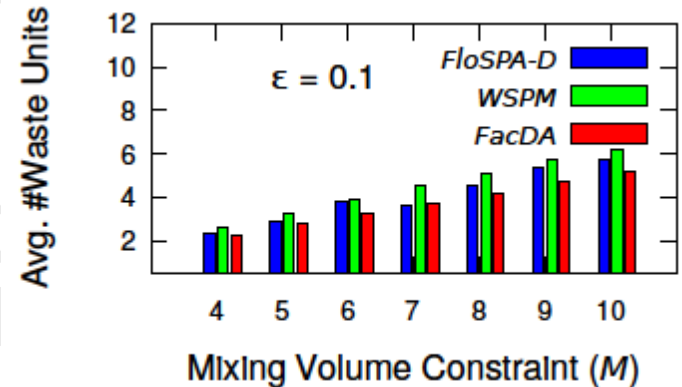
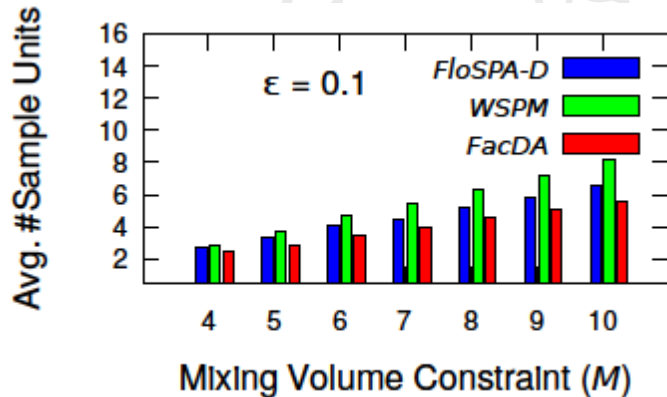
2
3
4
5
...
256
- M varies in the range of 4 to 10
- Average number of mixing steps compared among three methods



Distributions of avg. #mixing steps for three algorithms with $\epsilon = 0.0001$ and $\epsilon = 0.000001$

Simulation Results

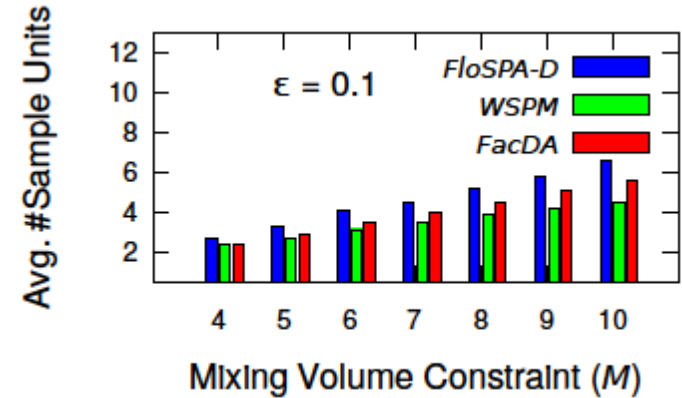
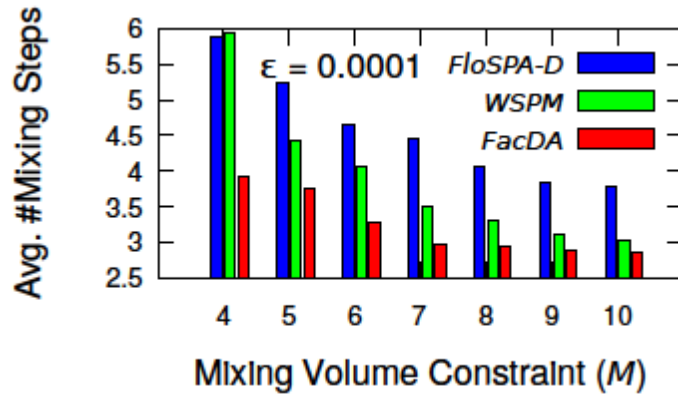
- Target volume V_t is one of the input parameters for *WSPM* [11]
- Average sample and waste count compared for two cases:
 - $V_t = 2$
 - $V_t > 2$ (final volume of target concentration is same as of *FacDA*)
- Case I: $V_t = 2$



Distributions of avg. #sample units and avg. #waste units by three algorithms with $\epsilon = 0.1$

Simulation Results

- Case II: $V_t > 2$



Distributions of n_m and n_s , when V_t is same as the output volume by *FacDA* with $\epsilon = 0.0001$ and $\epsilon = 0.1$

- Average CPU time comparison among three algorithms, where M varies in the range of 4 to 8 and $\epsilon = 0.05$

Algorithm	Avg. CPU time (in seconds)				
	$M = 4$	$M = 5$	$M = 6$	$M = 7$	$M = 8$
<i>FloSPA-D</i>	0.0173	0.0288	0.0231	0.0166	0.0126
<i>WSPM</i>	0.0085	0.0320	0.0732	0.1708	0.2816
<i>FacDA</i>	0.0745	0.0919	0.0505	0.0253	0.0267

Simulation Results

- Comparison between *FloSPA-D* [8] and *FacDA* for the number of test-cases **without ratio transformation**
- Considered a subset of the target ratio set that includes the ratios whose ratio-sum lies between M^{d-1} to M^d

ε	M	#Test-cases	#Test-cases without transformation by <i>FloSPA-D</i>	#Test-cases without transformation by <i>FacDA</i>	% Improvement of <i>FacDA</i> over <i>FloSPA-D</i>
0.001	3	2914	105	193	29.53
	4	30624	255	1069	61.4
	5	187250	625	4762	76.82
	6	250042	0	9540	100.0
0.0001	3	2914	105	193	29.53
	4	30624	591	1069	28.79
	5	187250	776	4762	71.97
	6	250042	255	9540	94.79

Conclusions

- Sample preparation on MEDA-DMF based architecture has many new scopes due to the availability of wide range of mixer models
- Unlike the conventional approach of $(m_1 : m_2)$ here we assume the mixer models to be $(m_1 : m_2 : \dots : m_n)$ for n distinct input fluids in case of MEDA-DMF
- *FacDA* manages to exploit the architectural advantages of MEDA-DMF and assures a dilution graph with minimum height
- The theory behind earlier methods *twoWayMix* and *FloSPA-D* follows as corollary to *FacDA*
- As a future work mixing algorithm can be designed for MEDA-DMF

References



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Thank you for your attention!

