Introduction	DRA	FJ-AMVA	FJ-DRA	Efficiency	Results	Conclusion

### Analyzing Queueing Networks with Multiclass Fork-Join Constructs

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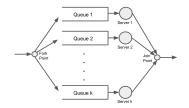
September 2, 2016

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#### What is a fork-join queue?

#### Definition

A fork-join queue is a queue where incoming jobs are split on arrival for service by numerous servers and joined before departure



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We want to come up with an approximation method for multiclass fork-join queueing networks with an implementation that is:

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- Accurate
- Efficient
- Universal

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Overview	V					

Two important building blocks:

- Decay Rate Approximation (DRA)
- Fork-Join Approximate Mean Value Analysis (FJ-AMVA)

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#### Method to approximate multiclass queueing networks

- Does not work with fork-join
- Iteratively solves for the queue lengths and throughputs
- Our method uses the same high level approach/ideas as DRA

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### Decay Rate Approximation

### Obtain initial estimate of throughput, $X_i$ Optimize for min f(X) locally around $X_i$ (using fmincon) def f(X):

- Compute utilization,  $\vec{\rho}$ , for each queue  $\rho_{q,c} = X_c \cdot \theta_{q,c}$
- Compute arrival MMAP[k] into each queue as superposition of departure processes of feeding queues scaled by ρ
- Compute decay rate of each queue when treated as a single MMAP[k]/PH[k]/1 queue
- Use decay rates with product form solver to obtain new estimates for utilization  $\vec{\rho'}$  and queue lengths
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#### • Fork-Join Approximate Mean Value Analysis

- Approximation method for multiclass fork-join queueing networks
- Iteratively solve for the mean queue lengths and throughputs

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- Compute residence time for all queues as  $R'_i(n) = s_i \cdot (1 + \bar{n}_i(n-1))$
- Re-number queues such that  $R'_1(n) \ge R'_2(n) \ge \cdots \ge R'_K(n)$
- Compute residence time for FJ construct as  $\sum_{k=1}^{K} \frac{1}{k} R'_k(n)$
- Compute throughput as  $X(n) = n / \sum_{k=1}^{K} \frac{1}{k} R'_k(n)$

• Compute new mean queue lengths as  $\bar{n}_i(n) = X(n) \cdot R'_i(n)$ Terminate loop when the difference in successive mean queue lengths is less than some  $\epsilon$ 

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Fork-Joi	n DRA	ι				

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- Same high level idea as DRA, but we modify how some steps are performed

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Initializa	tion					

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Two possible methods of initializing the throughput:

- AMVA-FCFS (same as DRA)
- Fork-Join AMVA

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Initializa	tion					

- DRA uses a simple iterative solver, AMVA-FCFS
- Requires the mean service demand,  $\theta_{ir} = v_{ir} \cdot s_{ir}$
- sir : mean service time per visit
- $s_{ir} = \frac{1}{\mu_{ir}}$
- v<sub>ir</sub> : mean number of visits of class r jobs to queue i
- $v_{ir} = \sum_{j=1}^{M} P_{ji} \cdot v_{jr}$
- $P_{ij}$  : routing probability from queue i to queue j
- Visits to queue *i* is the sum of the visits to queues feeding into queue *i*

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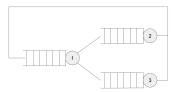
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Initializa	ation					

- For FJ queueing networks, the sum of visits of feeding queues will overcount for queues after the join point
- To deal with that, we need to use the sum of visits of feeding queues into the fork point instead

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- We compared the accuracy of the approximations obtained when initializing with both methods
- We refer to the mean queue length error across all queues and classes
- error =  $\frac{1}{2K}\sum_{i=1}^{M}\sum_{r=1}^{R}|Q_{i,r}-\hat{Q}_{i,r}|$
- Tested on the queueing network below



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#### Initialization - Comparison

$\lambda_{1,1}$	$\lambda_{1,2}$	$\lambda_{2,1}$	$\lambda_{2,2}$	$\lambda_{3,1}$	$\lambda_{3,2}$	<i>K</i> <sub>1</sub>	K <sub>2</sub>	Error (FCFS)	Error (FJ)
2	4	2	4	2	4	1	1	0.1082	0.1082
2	4	2	4	2	4	2	2	0.1196	0.1196
2	4	2	4	2	4	3	3	0.1248	0.1247
2	4	2	4	2	4	4	4	0.1289	0.1289
2	4	2	2	4	8	2	3	0.0476	0.0476
2	4	2	4	3	6	2	3	0.0862	0.0862
2	4	2	4	4	8	2	3	0.0607	0.0607
2	4	2	4	6	12	2	3	0.0467	0.0467

Table: Error for Different Initialization Methods

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- Both methods are O(IMK) where I is the number of iterations
- Runtime of both methods are orders of magnitude smaller than the overall solver
- FJ-AMVA produces initial estimations that are much closer to the actual values
- FJ-AMVA is easier to use, no extra work required to prepare input
- No discernible difference in accuracy or performance
- We chose FJ-AMVA as the initialization method for ease of use

Initializa	ation -	Compar	rison				
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Computing Arrival Processes

- For queues after the join point, we need a new way to compute the arrival process
- For all other queues, arrival process is unaffected and nothing new is required

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#### Synchronizing Fork-Join Queues

Existing method to approximate the departure process from join point by assuming finite length synchronization queues

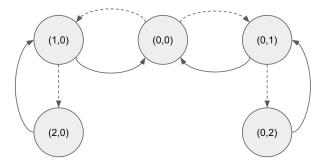


Figure: MAP representing departure process with sync queue length = 2

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Synchronizing Fork-Join Queues

- Consider the service process for class r at queue i
- We approximate the departure process as the service process multiplied by ρ<sub>i,r</sub>
- Use the method to generate a new  $D_0$  matrix in exactly the same way as before
- Consider each class one at a time and generate new D<sub>1,i</sub> matrices
- Set  $D_1 = \sum_{i=1}^{M} D_{1,i}$
- Normalize the MMAP, ensuring  $D_0 + D_1$  is a valid transition rate matrix

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- Part of the input to the product form (PF) solver is the job population for each class
- Mean queue lengths from PF solver will be based on those populations
- $c_i = \sum_{q=1}^M Q_{q,i}^{(PF)}$
- where  $c_i$  is the population of class i
- Q<sup>(PF)</sup><sub>q,i</sub> is the mean queue lengths for class i at queue q obtained from the PF solver
- However, when we have fork-join queues we have:
- $c_i < \sum_{q=1}^M Q_{q,i}$
- because one job splits into multiple jobs at the fork point

- Part of the input to the product form (PF) solver is the job population for each class
- Mean queue lengths from PF solver will be based on those populations

• 
$$c_i = \sum_{q=1}^{M} Q_{q,i}^{(PF)}$$

- where *c<sub>i</sub>* is the population of class *i*
- Q<sub>q,i</sub><sup>(PF)</sup> is the mean queue lengths for class i at queue q obtained from the PF solver
- However, when we have fork-join queues we have:
- $c_i < \sum_{q=1}^M Q_{q,i}$
- because one job splits into multiple jobs at the fork point



- To deal with this, we provide the PF solver a modified set of populations
- $c'_i = \sum_{q=1}^M Q_{q,i}^{\text{init}}$
- where  $Q_{q,i}^{\text{init}}$  is the initial approximation of mean queue length for queue q and class i
- $\bullet$  Doing this scales  $Q_{q,i}^{(PF)}$  to be a more accurate approximation

Introduction	DRA	FJ-AMVA	FJ-DRA	Efficiency	Results	Conclusion
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Efficienc	у					

- Profiled code using MATLAB's built in profiler
- Investigate how runtime scales as we increase:
  - Number of fork-join queues
  - Length of synchronization queue

Introduction	DRA	FJ-AMVA	FJ-DRA	Efficiency	Results	Conclusion
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Efficienc	v - Pi	ofiling				

Fork-Join Queues	Total Time (s)
2	2.412
3	5.536
4	35.643
5	814.599

Table: Increasing Number of Fork-Join Queues and Runtime

Sync Queue Length	Total Time (s)
1	4.501
2	8.092
3	14.925
4	33.174
5	88.88

Table: Increasing Synchronization Queue Length and Runtime

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#### Efficiency - Resizing MMAP

### Generated MMAPs become extremely large as we increase the two factors

• Resize the MMAP once it becomes larger than a certain size

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• We investigated resizing to MMAP with one state



#### Efficiency - Resizing MMAP

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• We investigated resizing to MMAP with one state



#### Efficiency - Resizing MMAP

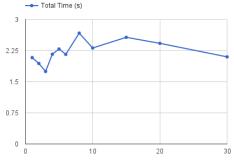
- Generated MMAPs become extremely large as we increase the two factors
- Resize the MMAP once it becomes larger than a certain size

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• We investigated resizing to MMAP with one state

## Introduction DRA of FJ-AMVA of FJ-DRA conclusion of the state MMAP

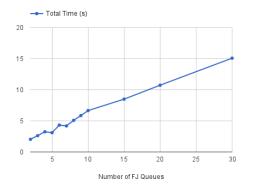
- We compute the rate for each class as  $\lambda_c = \operatorname{sum}(\pi \cdot D_{1,c})$
- where  $\pi$  is the equilibrium distribution of the MMAP
- So the new matrices are  $D_{1,c} = [\lambda_c]$  and  $D_0 = -\sum_{c=1}^{K} D_{1,c}$



Sync Queue Length



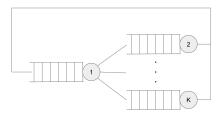
 Scales better - No discernible increase with synchronization queue length and approximately linear increase with number of FJ queues



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Introduction	DRA	FJ-AMVA	FJ-DRA	Efficiency	Results	Conclusion
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Results -	Overv	view				

- Compare our method against the FJ-AMVA method
- Vary the following factors:
  - Number of FJ queues
  - Heterogeneity of FJ queues
  - Complexity of service distributions
- Homogeneous FJ queues with exponential service distribution (unless directly testing that factor)

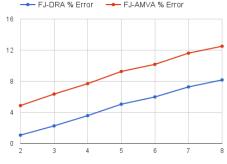


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#### Results - Number of FJ Queues

- Our FJ-DRA method is more accurate than the FJ-AMVA method for all our tests
- Trend suggests that it will continue to be more accurate even as we increase the number of FJ queues
- Overall error increases as number of FJ queues increases



Introduction	DRA	FJ-AMVA	FJ-DRA	Efficiency	Results	Conclusion
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Results -	Heter	rogeneity				

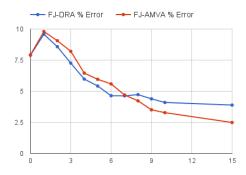
We fixed the number of FJ queues at 4 and used the following parameters:

Parameter	Value
<i>c</i> <sub>1</sub>	2
<i>c</i> <sub>2</sub>	3
$\lambda_{1,1}$	1
$\lambda_{1,2}$	2
$\lambda_{k,1} \; (\forall k  eq 1)$	1+0.1(k-1)h
$\lambda_{k,2} \; (orall k  eq 1)$	2+0.2(k-1)h

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Results	- Hete	erogeneit	V			

- The performance of both approximation methods is quite close
- FJ-DRA is better at lower levels of heterogeneity
- Overall, error is trending down



Introduction	DRA	FJ-AMVA	FJ-DRA	Efficiency	Results	Conclusion			
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Results -	Results - Erlang-2 Service Distribution								

- Used Erlang-2 as service distribution for all queues
- Used 2 FJ queues
- Fix some parameters as in the table below:

Parameter	Value		
$\lambda_{1,1}$	2		
$\lambda_{1,2}$	4		
<i>c</i> <sub>1</sub>	1		
<i>c</i> <sub>2</sub>	2		

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#### Results - Erlang-2 Service Distribution

- Performance is quite close, FJ-DRA is slightly better in most cases
- Error is less than 5% for all tests
- No clear trend as  $\lambda$ 's change

$\lambda_{k,1}$	$\lambda_{k,2}$	FJ-DRA % Error	FJ-AMVA % Error
2	4	4	4
3	4	3.12	3.16
4	4	2.63	4.36
3	5	2.76	3
3	6	2.56	3.11
4	8	3.85	3.39
5	10	4.27	3.68
6	12	4.23	3.72
10	20	3.57	3.18

Introduction	DRA	FJ-AMVA	FJ-DRA	Efficiency	Results	Conclusion		
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Results - Summary								

- Tested three factors:
  - Number of FJ Queues
  - Heterogeneity of FJ Queues
  - Complexity of service distributions
- In most of our tests, FJ-DRA is at least approximately as accurate as the FJ-AMVA method and there are some cases where FJ-DRA clearly outperforms FJ-AMVA
- Only tests where FJ-AMVA performs better is where FJ queues are very heterogeneous
- Overall, FJ-DRA method had less than 10% error in all our tests

Introduction	DRA	FJ-AMVA	FJ-DRA	Efficiency	Results	Conclusion
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Conclusi	on					

- We presented the FJ-DRA method to approximate multiclass FJ queues
- We investigated how resizing the MMAP to a single state can help improve the efficiency
- We compared FJ-DRA against FJ-AMVA and found that it performs better in most of our tests

### Questions?

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