



Scalable Bayesian Network Discovery with Reconfigurable Hardware

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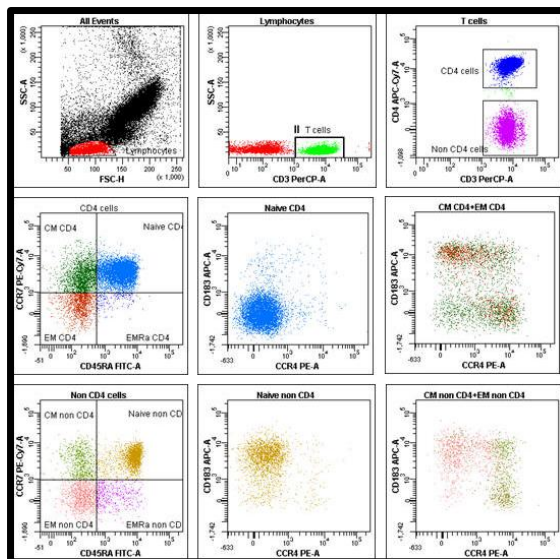
Outline

- Biological Perspective
 - The Motivation: Learning the structure of cell signaling networks
 - The Algorithm: Computational complexity & MCMC
 - Algorithmic approach
- Reconfigurable Computing Perspective
 - Hardware approach
 - FPGA implementation
 - Design scalability
- Results
- Future Work
- Conclusion and Summary

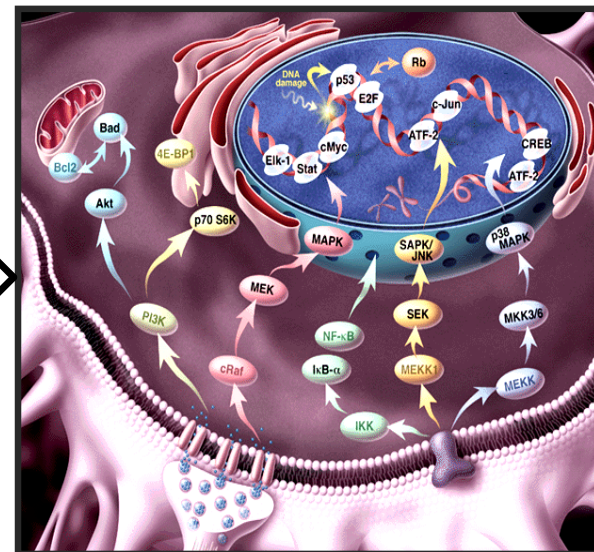
Cell Signaling Networks

Goal: Given flow cytometry data, learn the structure of cell signaling networks

- Flow Cytometry
 - Data in the form of “raw” quantitative observations
 - Measurement of proteins & other components inside cells
- Cell Signaling Networks
 - Structures that model protein signaling pathways
 - Modeling perturbations to a network can help uncover the cause of human disease



This talk

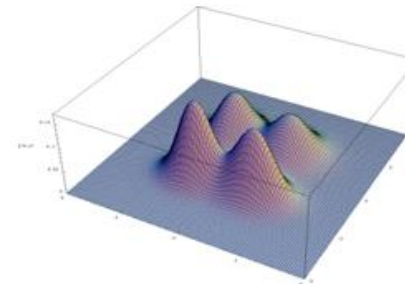


Problem: Kernel is NP-Hard

Goal: Determine which network best explains the data

- Algorithm Bottlenecks
 - Search space grows super-exponentially with the graph's node count
 - Multiple local optima, encoding best-solutions, may exist

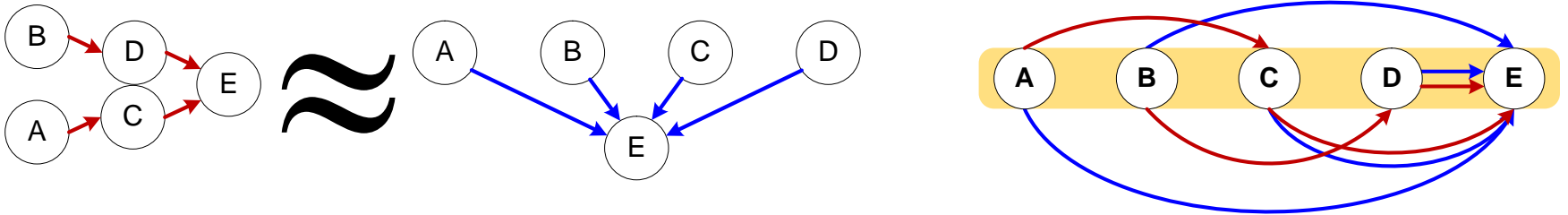
Nodes	Graphs
4	453
5	29281
10	4.7×10^{17}
20	2.34×10^{72}



- Alternative Approach: “MCMC Sampling”
 - Markov Chain Monte Carlo
 - Slower than search methods
 - More reliable and less prone to get stuck in local optima (higher “QoR”)

Algorithmic Approach

- Graph vs. Order Space
 - The “order space” is much smaller than the “graph space”
 - Swapping nodes in the order space results in a larger move



- Computational Strategy
 - (1) Calculate local scores per parent set
 - (2) “Order Sampler”: Determine the likely orders (**algorithm kernel**)
 - (3) “Graph Sampler”: Extract graphs from probable orders
- Idea: Implement the Order Sampler in Hardware
 - Minimize the time it takes to score an **order**
 - Reduce the computational complexity to score an order

Hardware Approach

- Scoring an order is “embarrassingly” parallel
 - Divide computation by node

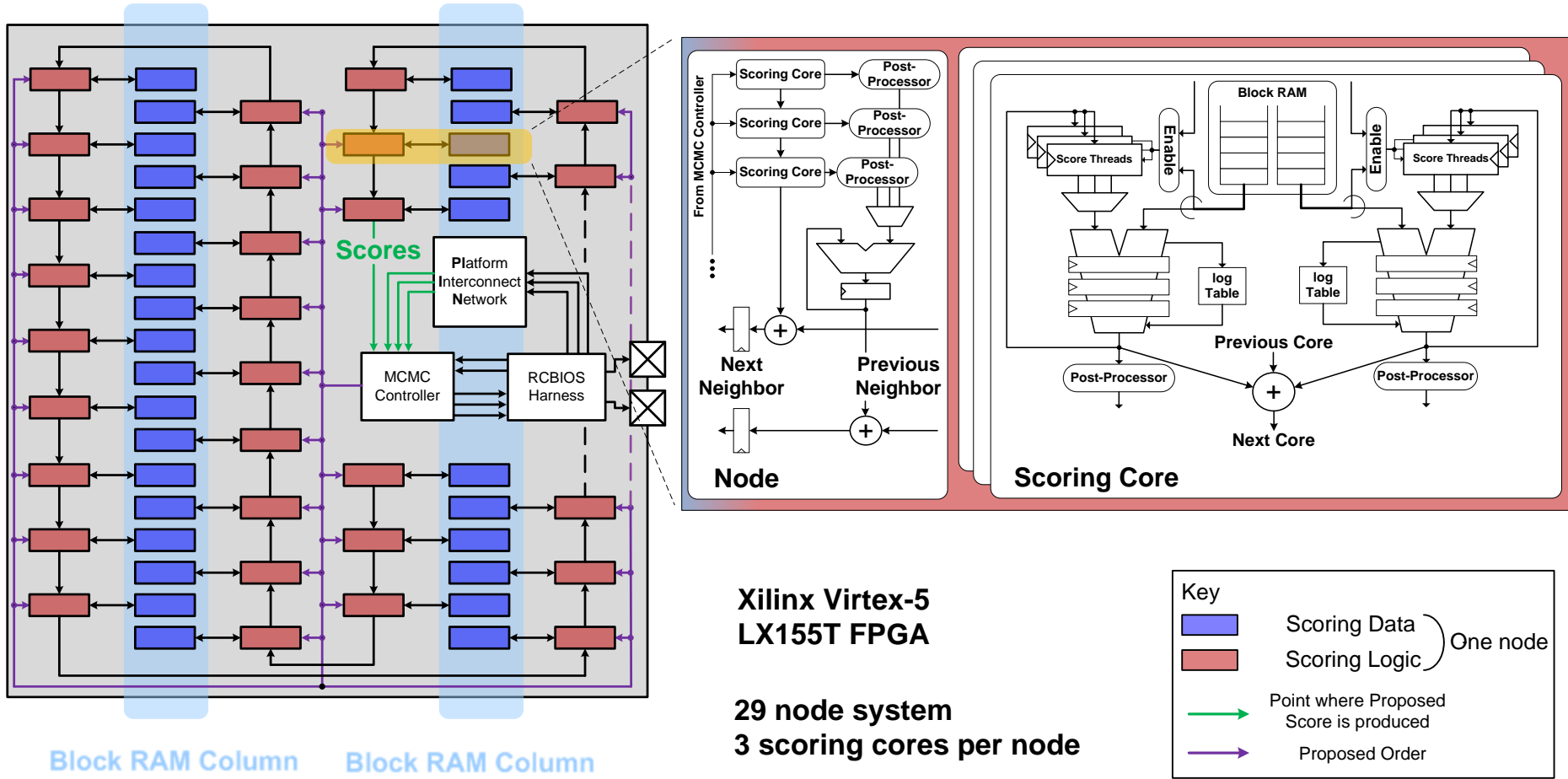
$$\begin{aligned} \text{Score for Order } \prec &= \sum_{i=1}^N \text{Local score of node } V_i \\ \text{score}(\prec) &= \sum_{i=1}^N \text{score}(V_i) \\ &= \text{score}(V_1) + \text{score}(V_2) + \dots + \text{score}(V_N) \end{aligned}$$

Comparisons, Multiplications, and Subtractions

Built as separate parallel units in hardware

- Partition parent sets into block RAMs
- Perform (3) – the “Graph Sampler” step – alongside the Order Sampler
- Map computations to **log** space
 - Bulk of computations are on probabilities (small values)
 - Multiplications \rightarrow Additions

FPGA Implementation

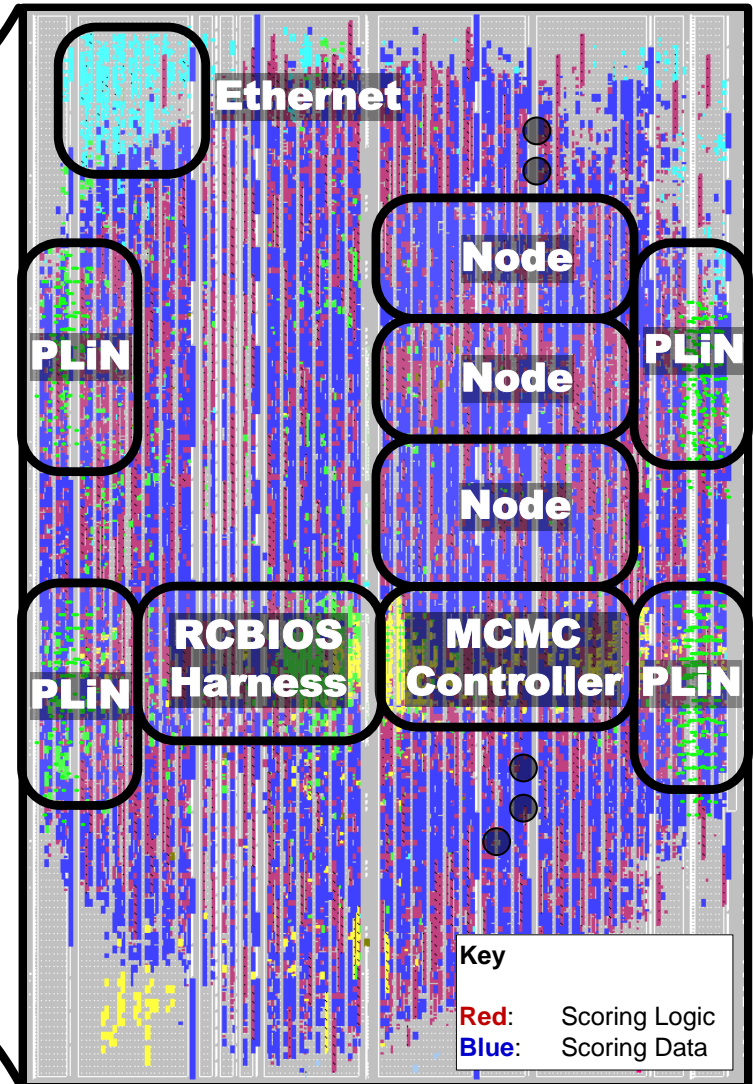
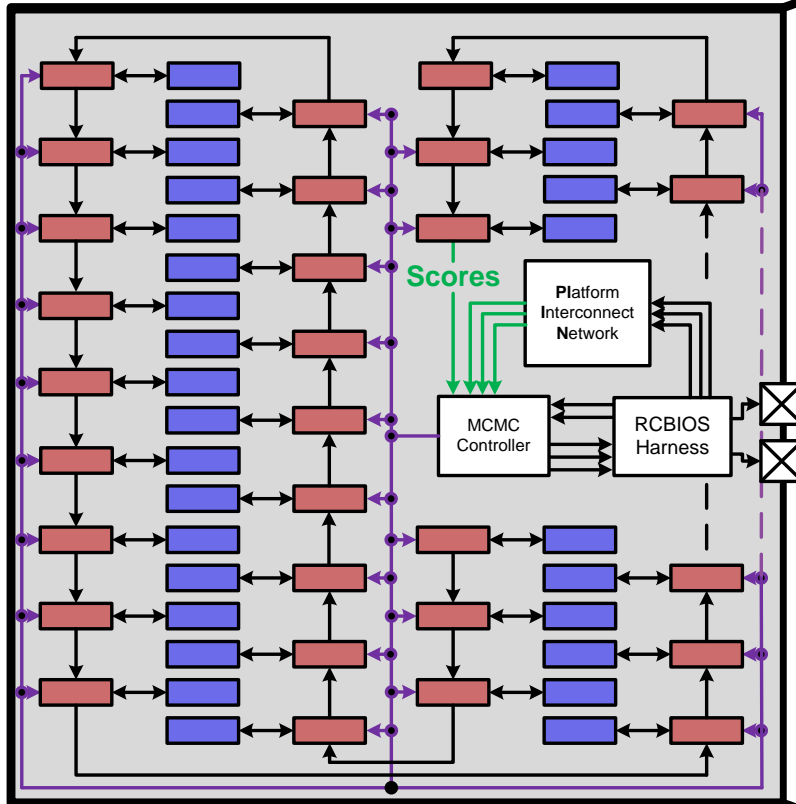


**Xilinx Virtex-5
LX155T FPGA**

**29 node system
3 scoring cores per node**

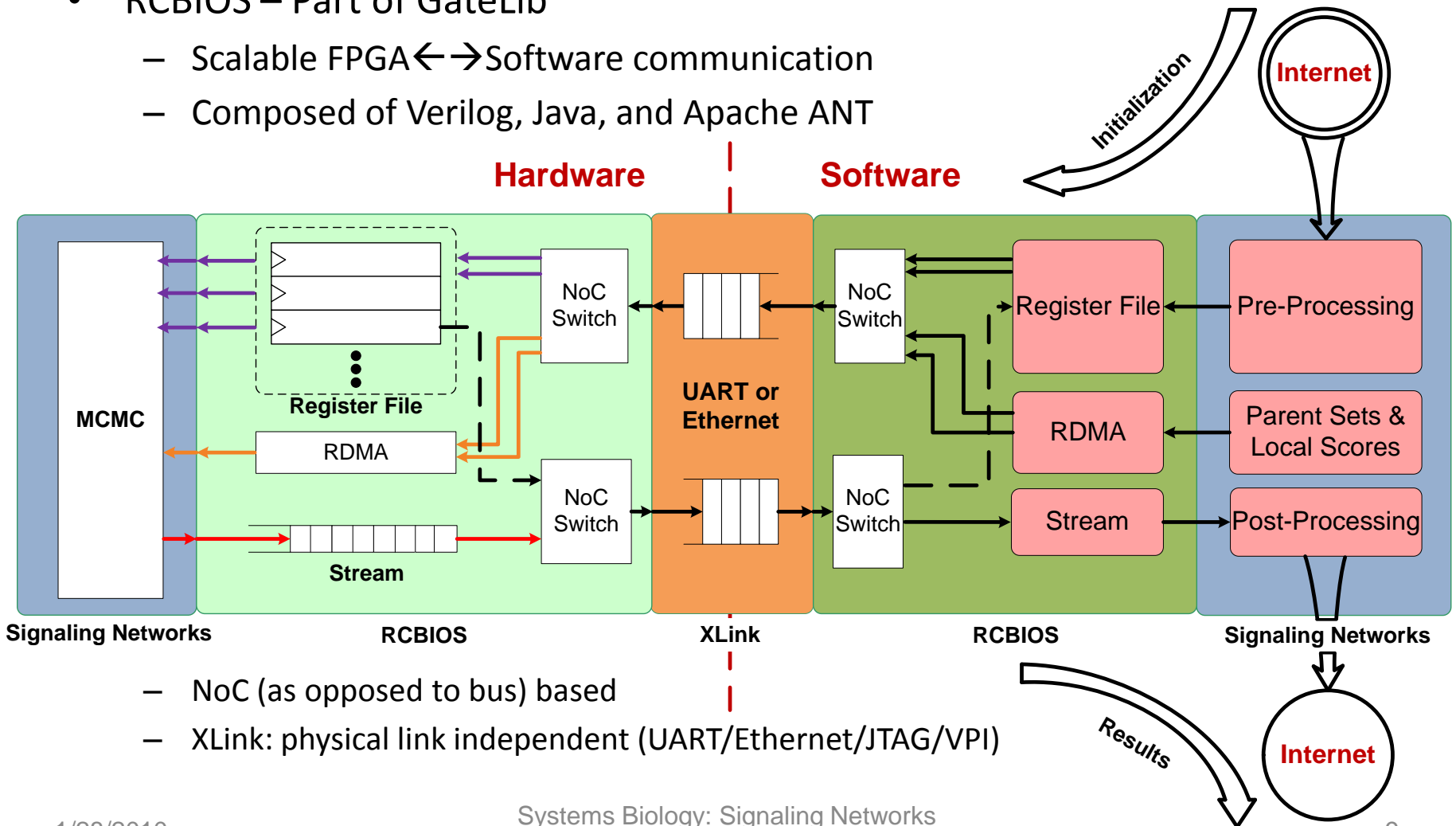
FPGA Floorplanner (LX155T)

Abstract View vs. Actual Implementation



FPGA Infrastructure

- RCBIOS – Part of GateLib
 - Scalable FPGA \leftrightarrow Software communication
 - Composed of Verilog, Java, and Apache ANT



- NoC (as opposed to bus) based
- XLink: physical link independent (UART/Ethernet/JTAG/VPI)

Design Scalability

“MCMC Mesh”

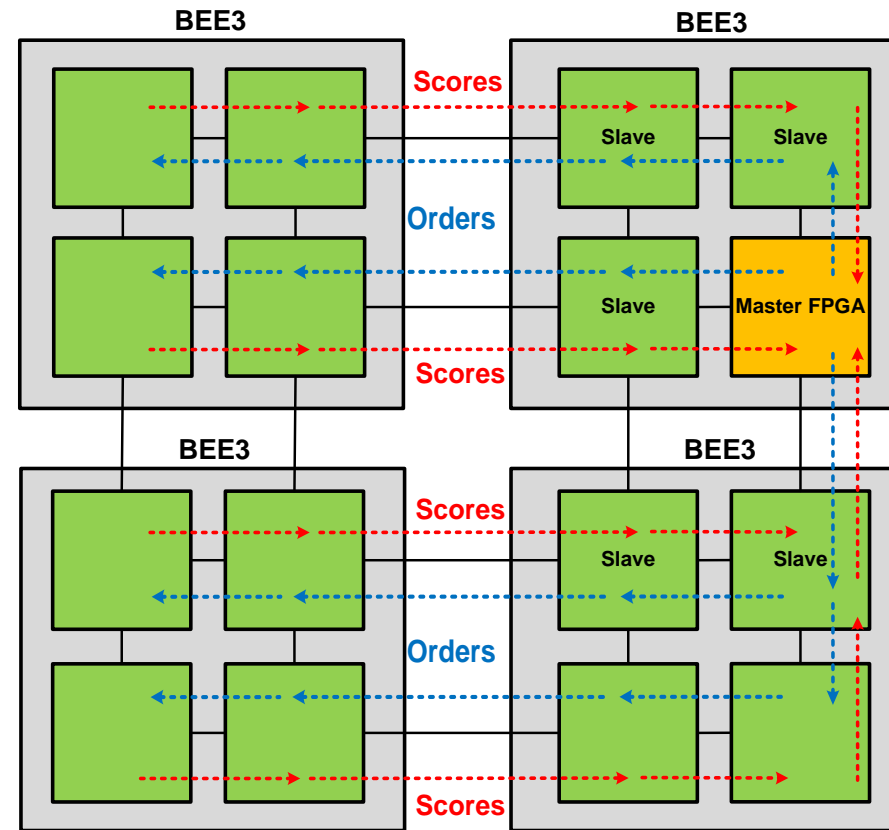
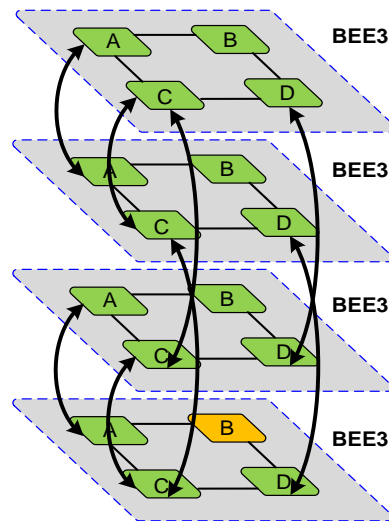
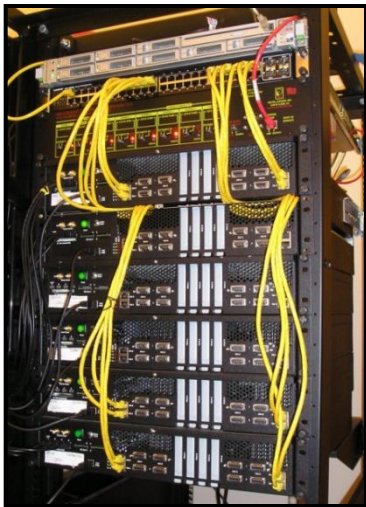
Idea: Split larger problems across multiple FPGAs

* While maintaining base design

Additional Infrastructure

- (1) Inter-chip ring connections
- (2) Inter-board Aurora high-speed links
- (3) **Platform Interconnect Network (PLiN)**

built on (1) and (2)



Results

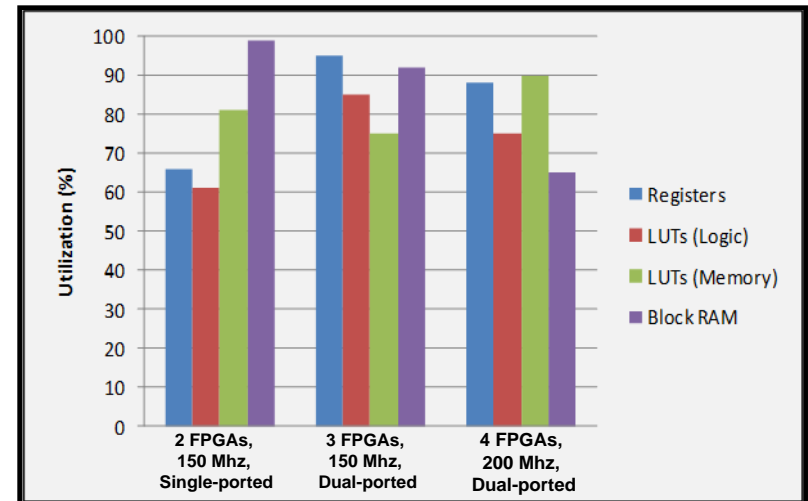
Problem Specification

22 nodes
7547 parent sets per node
100 random restarts
10,000 iterations per restart

Questions

- What's the deal with the 1x vs. 4x GPP?
- What is the "Caching Algorithm"?

Times (s):	Order		Graph
1x GPP*:	62.33	+	12.67
4x GPP*:	343.62	+	12.67
GPU:	98.42	+	12.67
2x FPGA:	8.13	+	0
3x FPGA:	5.11	+	0
4x FPGA:	4.42	+	0



GPP: 4-core Intel Xeon 3.00GHz (PowerEdge 1850), 7.71 GB RAM, 10.00 GB swap (Caching algorithm)
GPU: 1.3 GHz NVIDIA Tesla c1060 (Caching algorithm)
FPGA: Xilinx Virtex-5 LX155T (-2)

Future Work

- Order caching

Insight: A given order will always produce the same score

- Optimization used by both GPU & GPP implementations
- Can be made at an order or “local order” granularity

- Pre-processing on FPGA

- (1) “Pre-processing” has become new bottleneck
- Map “Local score” generation to each FPGA in network
- Transport “observations” data to FPGA

Insight: Observation files are small, score files are large

- Map Kernel to OpenRCL platform

Conclusion and Summary

This work coordinates clusters of FPGA accelerators
In order to learn protein network structure

- Reconfigurable Computing gives us the ability to...
 - Build each accelerator to best-fit different problems
 - Provide arbitrary design scaling with low overhead

Acknowledgements

For making this work possible, a special thanks to:

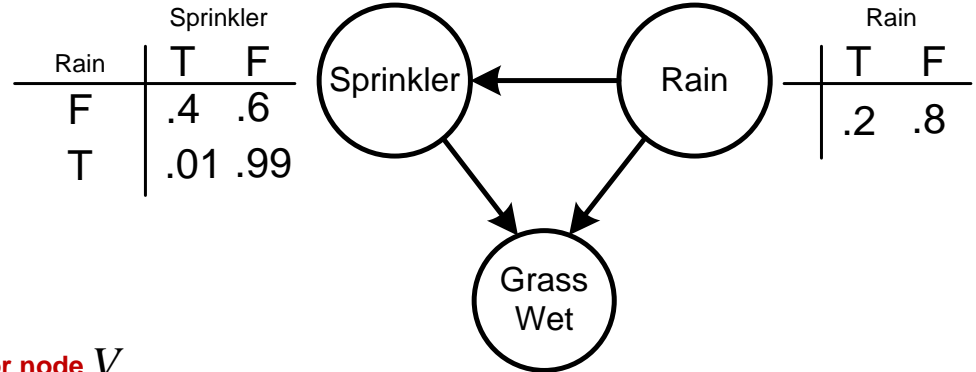
- Ilia Lebedev & Mingjie Lin for the **Platform Interconnect Network**
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BACKUP SLIDES

Bayesian Networks

- “Belief Network”
 - Directed acyclic graph
 - Structure encodes...

- Conditional independence
- Causal relationships



Rain	Sprinkler	
	T	F
F	.4	.6
T	.01	.99

		Rain	
		T	F
.2	.8		

		Grass Wet	
		T	F
F	F	0	1
		.8	.2
T	F	.9	.1
		.99	.01

Courtesy of Tom Griffiths (U.C. Berkeley)

- Bayesian Score
 - A basis for comparing Bayesian Structures
 - Based on prior belief and observations

Experimental data

$$P(\underbrace{D, G}_{\text{Graph}}) = \underbrace{P(G)}_{\text{Prior probability}} P(D | G)$$