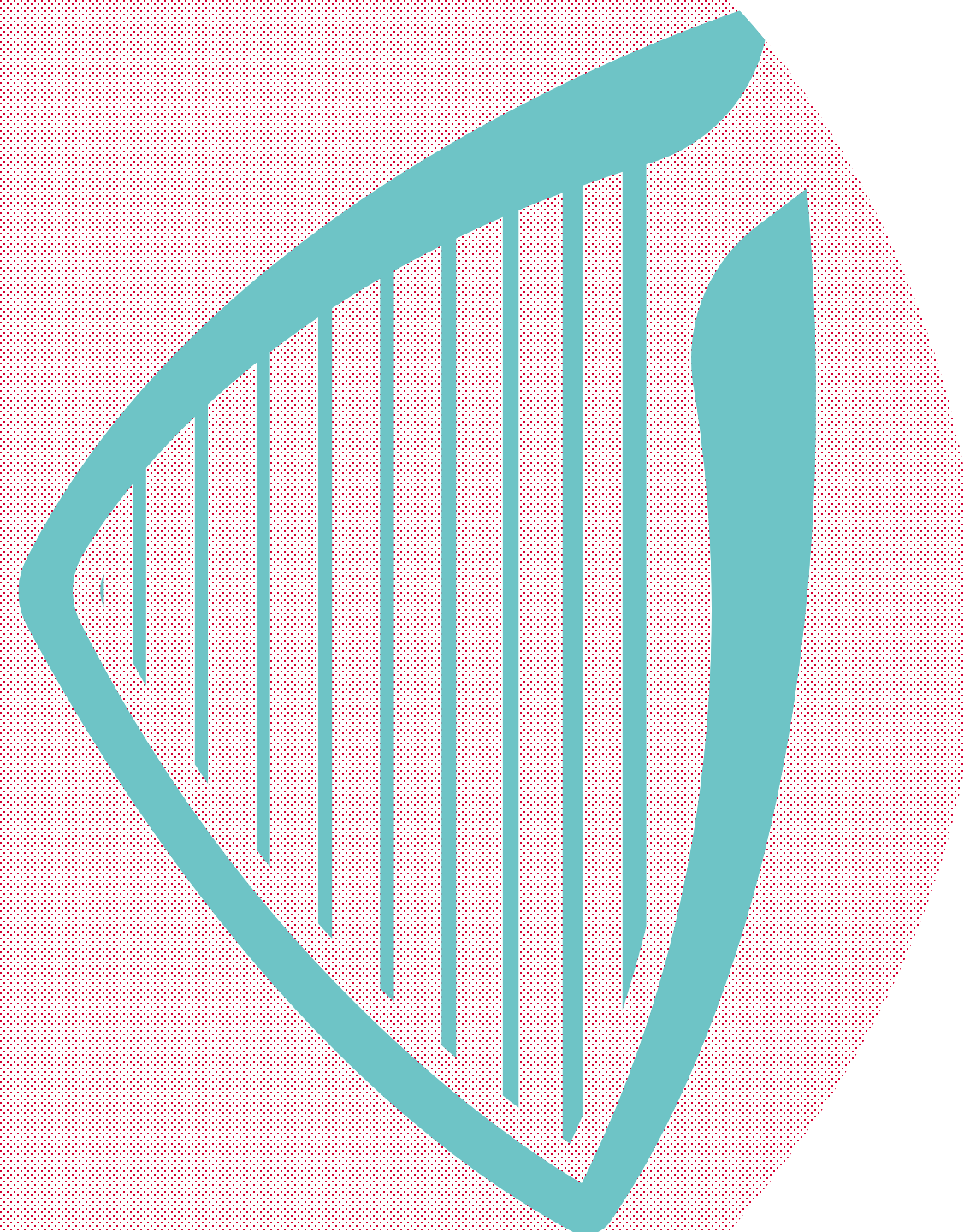




Declarative Analytics on Heterogeneous HPC Systems

Ahmedur Rahman Shovon

Committee:

Dr. Sidharth Kumar, *Chair and Advisor*

Dr. Michael E. Papka

Dr. Zhiling Lan

Dr. Stavros Sintos

Dr. Gopikrishna Deshpande (AU)

Dr. Thomas Gilray (WSU)



UNIVERSITY OF
ILLINOIS CHICAGO

JULY 15, 2025

Roadmap

1. **Introduction and motivation**
2. GPU based Datalog engine development
3. Multi-node multi-GPU Datalog engine development
4. GPU portability of Datalog engines
5. Power analysis of Datalog engines
6. Exploration of high-dimensional scientific analytics
7. Future work
8. Conclusion

Declarative Programming Paradigm

Users expresses **what** to achieve with the data rather than **how** to accomplish it

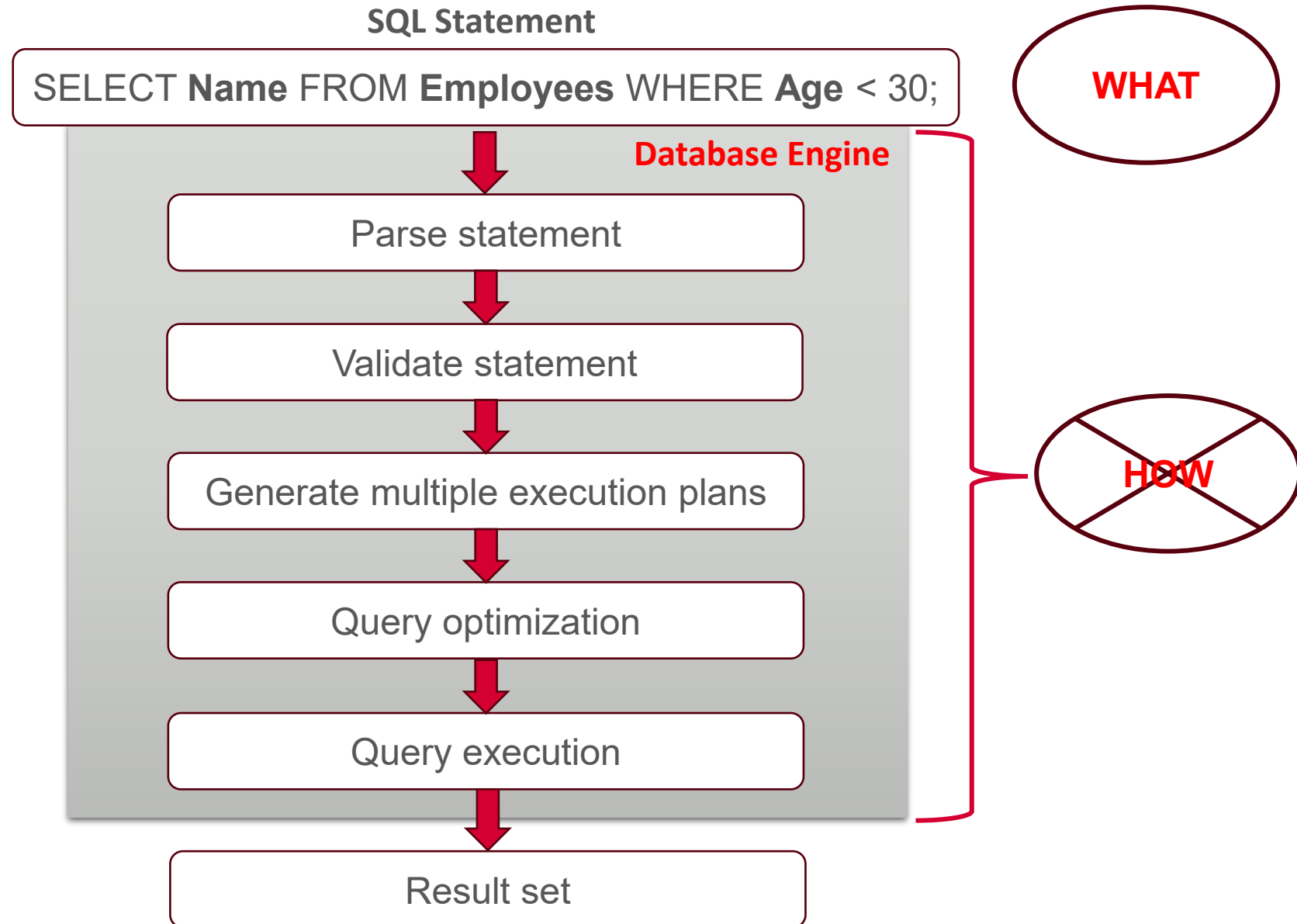
Employees

EmpID	Name	Email	Age	ManagerID
0	Alice	alice@example.com	56	Null
1	Bob	bob@example.com	33	0
2	Eve	eve@example.com	26	0
3	Carol	carol@example.com	34	2
4	Dave	dave@example.com	22	3

Find all employees who are less than 30 years old

SELECT Name FROM Employees WHERE Age < 30;

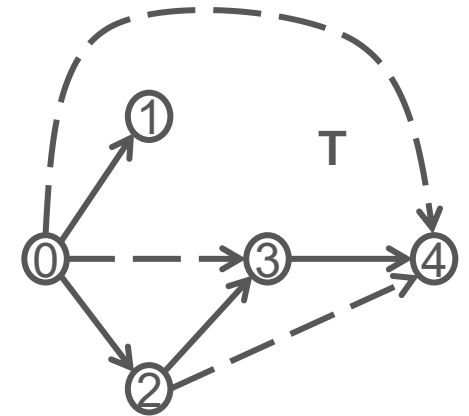
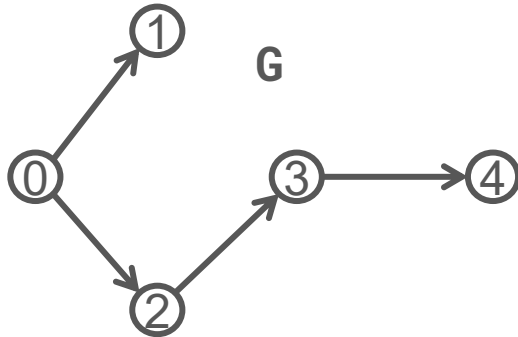
How is the query processed?



Recursive Queries using SQL

Employees

EmpID	Name	Email	Age	ManagerID
0	Alice	alice@example.com	56	Null
1	Bob	bob@example.com	33	0
2	Eve	eve@example.com	26	0
3	Carol	carol@example.com	34	2
4	Dave	dave@example.com	22	3



Find all managers, including direct and indirect managers.

-- Recursive CTE to find all managers (direct and indirect)

WITH RECURSIVE ManagerHierarchy AS (

-- Base case: direct manager relationship

SELECT EmpID AS EmployeeID, ManagerID

FROM Employees WHERE ManagerID IS NOT NULL

UNION

-- Recursive case: find manager of managers recursively

SELECT mh.EmployeeID, e.ManagerID

FROM ManagerHierarchy mh

JOIN Employees e ON mh.ManagerID = e.EmpID WHERE e.ManagerID IS NOT NULL

)

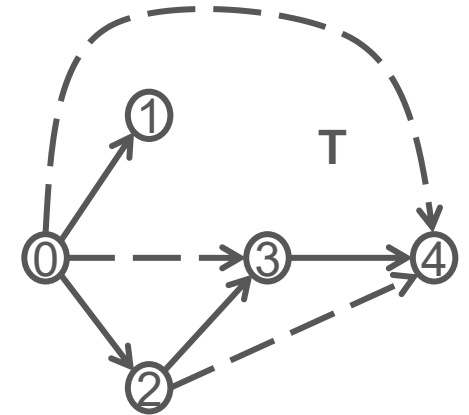
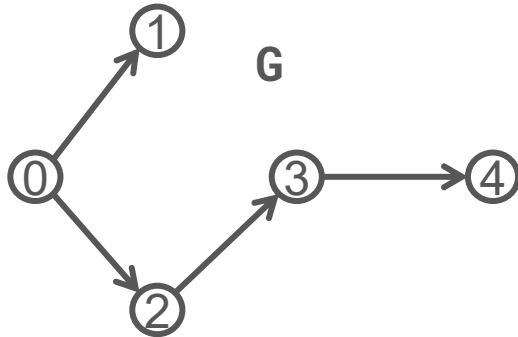
SELECT DISTINCT ManagerID, Employeeid FROM ManagerHierarchy ORDER BY ManagerID;

Advanced approach: Datalog

Recursive Queries using Datalog

Employees

EmpID	Name	Email	Age	ManagerID
0	Alice	alice@example.com	56	Null
1	Bob	bob@example.com	33	0
2	Eve	eve@example.com	26	0
3	Carol	carol@example.com	34	2
4	Dave	dave@example.com	22	3



Find all managers, including direct and indirect managers.

Finding recursive relationships using Datalog

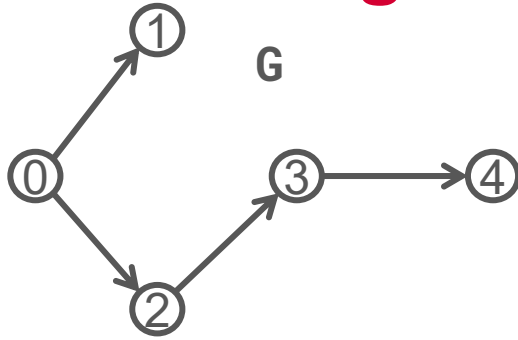
// Base case: direct manager manages employee

Manager(mID, eID) :- Employees(eID, mID), mID != NULL.

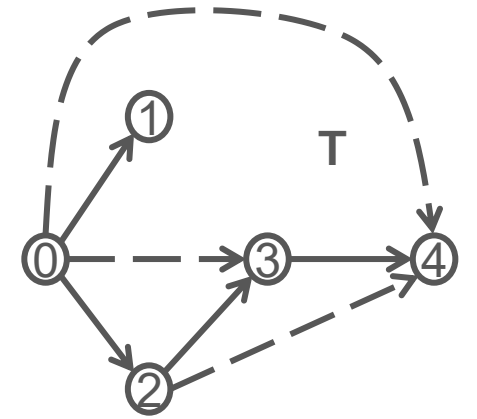
// Recursive case: indirect managers

Manager(mID, eID) :- Manager(mID, nID), Employees(eID, nID).

Advantages of Recursive Queries with Datalog



Find all managers, including direct and indirect managers.



Finding recursive relationships using SQL

-- Recursive CTE to find all managers (direct and indirect)

WITH RECURSIVE ManagerHierarchy AS (

-- Base case: direct managers

SELECT

FROM

WHERE ManagerID IS NOT NULL

UNION

-- Recursive case: indirect managers

SELECT m.ManagerID, e.EmployeeID

FROM

JOIN

WH

)

SELECT DISTINCT ManagerID, EmployeeID FROM ManagerHierarchy

ORDER BY ManagerID;

Finding recursive relationships using Datalog

Datalog offers greater expressiveness and natural syntax for recursive queries

As a logic programming language, it's well-suited for deductive databases

Ideal for declarative analytics to find complex patterns over large datasets

Datalog Rules Execution

Datalog Rules

```
// Base case: direct manager manages employee
Manager(mID, eID) :- Employees(eID, mID), mID != NULL.
// Recursive case: indirect managers
Manager(mID, eID) :- Manager(mID, nID), Employees(eID, nID).
```

Datalog Engine

Parse rules
Generate query plan

Iterative Relational Algebra

$$F_G(T) \triangleq G \cup \Pi_{1,2}(\rho_{0/1}(T) \bowtie_1 G)$$

Union

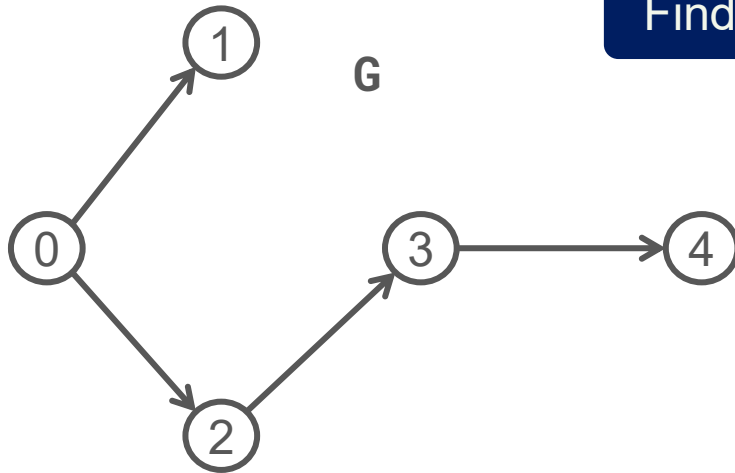
Projection

Join

Relational algebra operations

Loop until $\text{manager}_i = \text{manager}_{i-1}$

Manager Chains Problem as Transitive Closure



Find all managers, including direct and indirect managers.

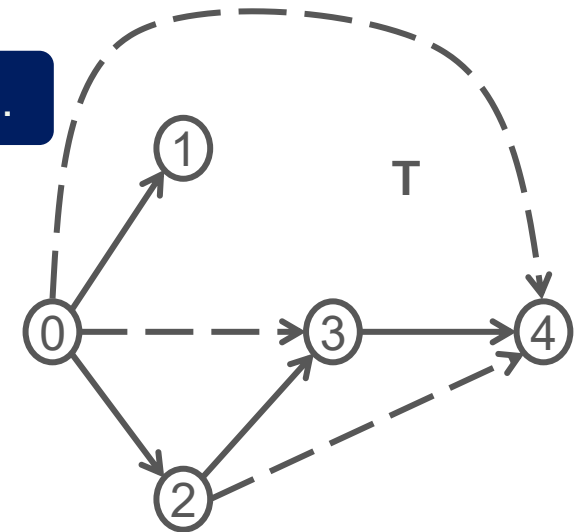
```

// Base case: direct manager manages employee
Manager(mID, eID) :- Employees(eID, mID), mID != NULL.
// Recursive case: indirect managers
Manager(mID, eID) :- Manager(mID, nID), Employees(eID, nID).
  
```

Transitive closure computation of a graph

```

tc(X, Y) :- edges(X, Y).
tc(X, Z) :- tc(X, Y), edges(Y, Z).
  
```

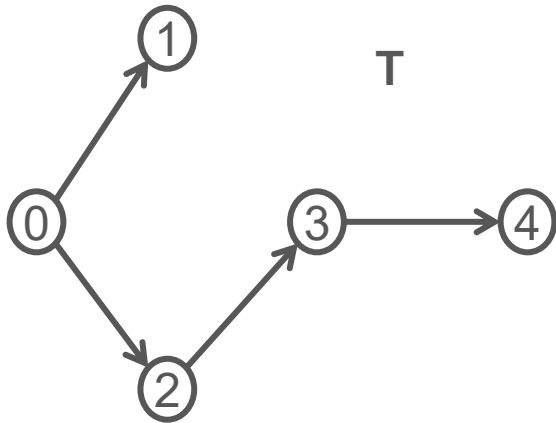


0	1
0	2
2	3
3	4

0	1
0	2
2	3
3	4
0	3
2	4
0	4

Transitive Closure: Iterations 1

$$F_G(T) \triangleq G \cup \Pi_{1,2}(\rho_{0/1}(T) \bowtie_1 G)$$



0	1
0	2
2	3
3	4

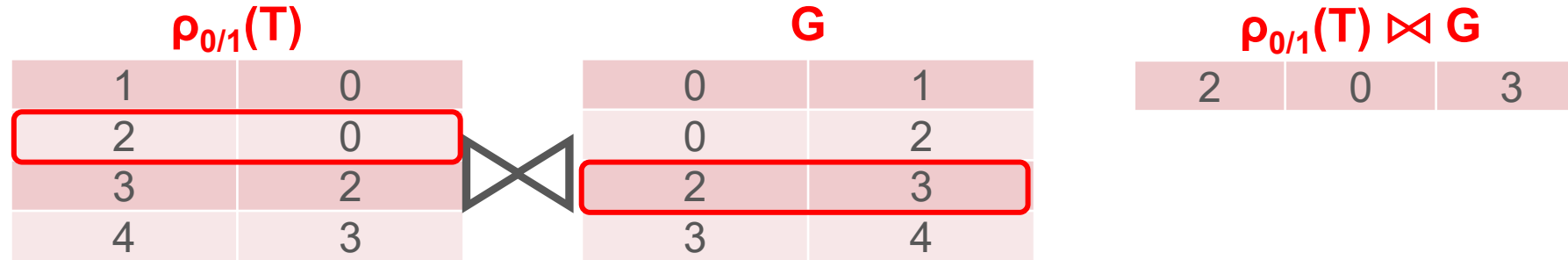
Transitive Closure: Iterations 1

$$F_G(T) \triangleq G \cup \Pi_{1,2}(\rho_{0/1}(T) \bowtie_1 G)$$

$\rho_{0/1}(T)$			G	
1	0		0	1
2	0		0	2
3	2		2	3
4	3		3	4

Transitive Closure: Iterations 1

$$F_G(T) \triangleq G \cup \Pi_{1,2}(\rho_{0/1}(T) \bowtie_1 G)$$



Transitive Closure: Iterations 1

$$F_G(T) \triangleq G \cup \Pi_{1,2}(\rho_{0/1}(T) \bowtie_1 G)$$

$\rho_{0/1}(T)$			G			$\rho_{0/1}(T) \bowtie G$		
1	0		0	1		1	0	3
2	0		0	2		3	2	4
3	2		2	3				
4	3		3	4				

Transitive Closure: Iterations 1

$$F_G(T) \triangleq G \cup \Pi_{1,2}(\rho_{0/1}(T) \bowtie_1 G)$$

$\rho_{0/1}(T)$

1	0
2	0
3	2
4	3



G

0	1
0	2
2	3
3	4

$\rho_{0/1}(T) \bowtie G$

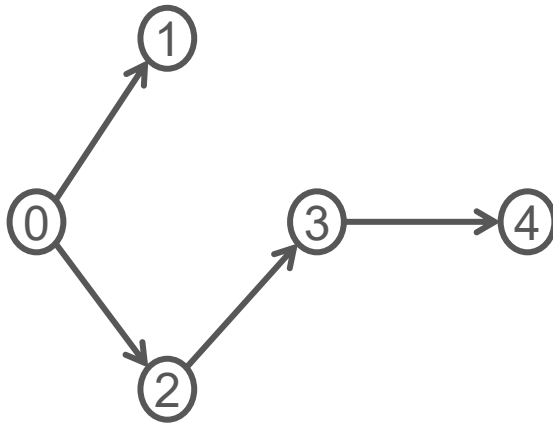
1	0	3
3	2	4

$\Pi_{1,2}(\rho_{0/1}(T) \bowtie G)$

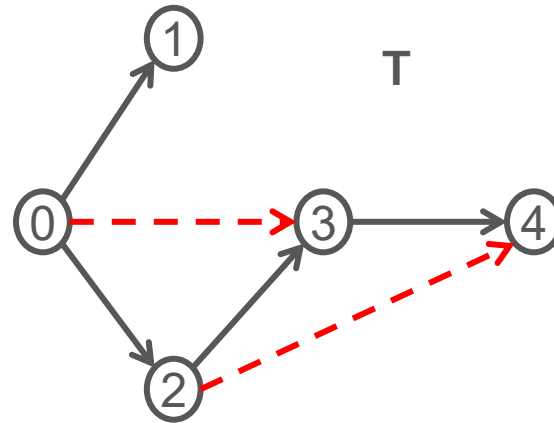
0	3
2	4

Transitive Closure: Iterations 1

$$F_G(T) \triangleq G \cup \Pi_{1,2}(\rho_{0/1}(T) \bowtie_1 G)$$



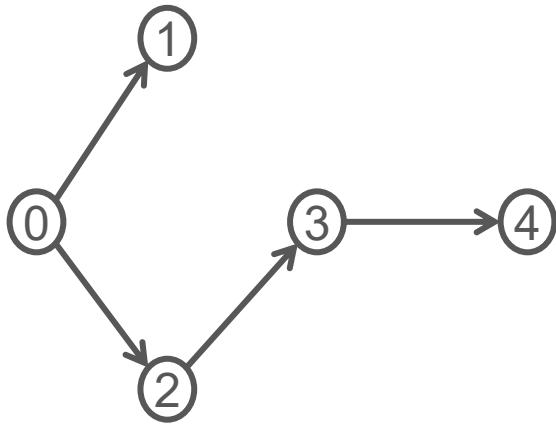
0	1
0	2
2	3
3	4



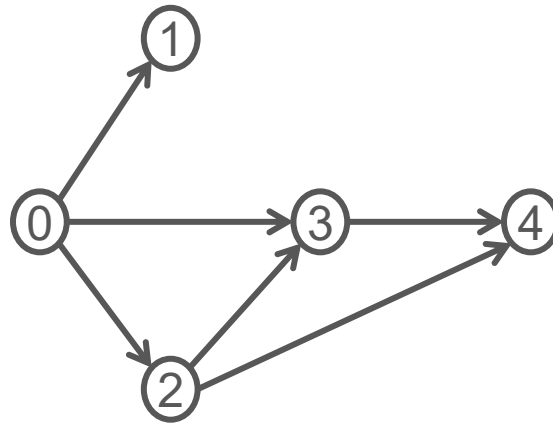
0	1
0	2
2	3
3	4
0	3
2	4

Transitive Closure: Iterations 2

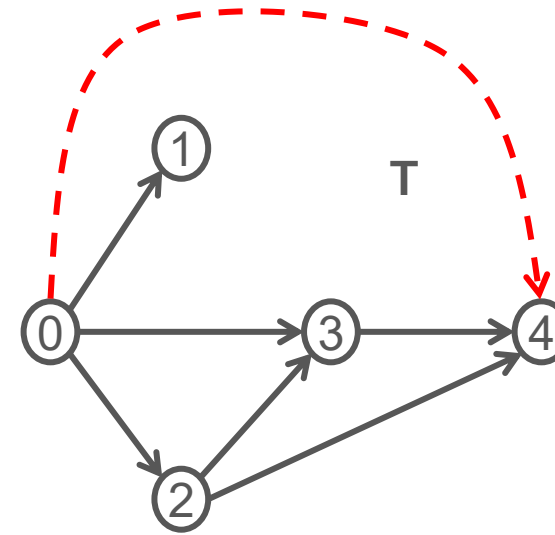
$$F_G(T) \triangleq G \cup \Pi_{1,2}(\rho_{0/1}(T) \bowtie_1 G)$$



0	1
0	2
2	3
3	4



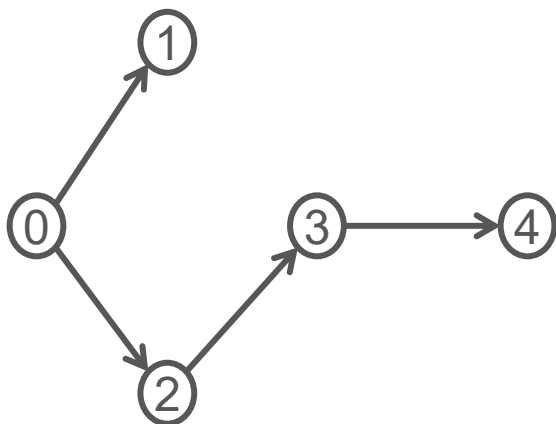
0	1
0	2
2	3
3	4
0	3
2	4



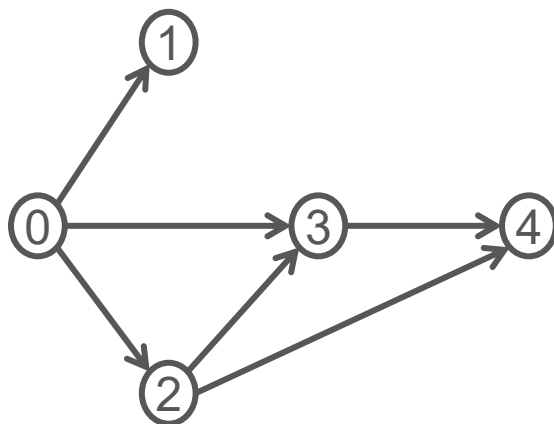
0	1
0	2
2	3
3	4
0	3
2	4
0	4

Transitive Closure: Iterations 3

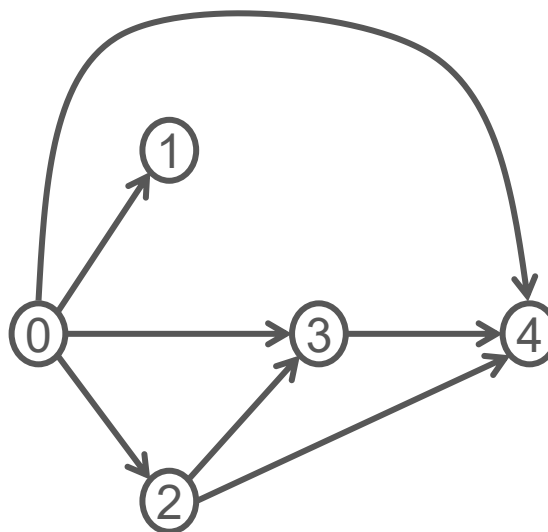
Fixed-point reached



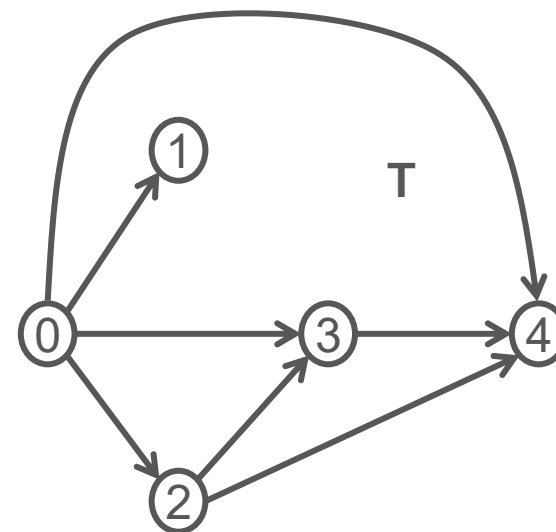
0	1
0	2
2	3
3	4



0	1
0	2
2	3
3	4
0	3
2	4



0	1
0	2
2	3
3	4
0	3
2	4
0	4



0	1
0	2
2	3
3	4
0	3
2	4
0	4

Graph Mining Applications

Applications

Transitive closure

```
tc(X, Y) :- edges(X, Y).
tc(X, Z) :- tc(X, Y), edges(Y, Z).
```

Connected components

```
cc(X, X) :- edges(X, _).
cc(Y, $MIN(Z)) :- cc(Y, Z), edges(X, Y).
```

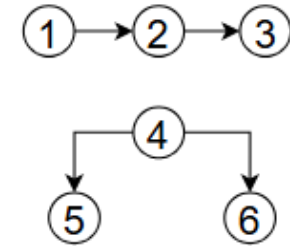
Triangle counting

```
2cl(X, Y) :- edges(X, Y), X < Y.
2cl(X, Y) :- edges(Y, X), X < Y.
triangles(X, Y, Z) :- 2cl(X, Y), 2cl(Y, Z), 2cl(X, Z).
```

Same generation

```
sg(x, y) :- edges(p, x), edges(p, y), x ≠ y.
sg(x, y) :- edges(a, x), sg(a, b), edges(b, y), x ≠ y.
```

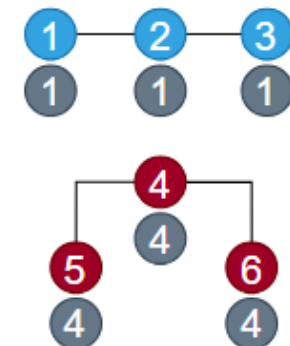
Input Graph



Initialization

Edge		CC (Delta, Full)	
1	2	1	1
2	1	2	2
2	3	3	3
3	2	4	4
4	5	5	5
5	4	6	6
4	6		
6	4		

WCC



- De Moor, O., Gottlob, G., Furche, T., & Sellers, A. (Eds.). (2012). Datalog Reloaded: First International Workshop, Datalog 2010, Oxford, UK, March 16-19, 2010. Revised Se Springer.
- Huang, S. S., Green, T. J., & Loo, B. T. (2011, June). Datalog and emerging applications: an interactive tutorial. In Proceedings of the 2011 ACM SIGMOD International Conf (pp. 1213-1216).
- Gilray, T., & Kumar, S. (2017). Toward parallel cfa with datalog, mpi, and cuda. In Scheme and Functional Programming Workshop.
- Zomorodian, A. (2012). Topological data analysis. Advances in applied and computational topology, 70, 1-39.

Datalog's Applications

Graph
mining

Static
analysis

Deductive
database

Machine
learning

Datalog

Iterative Relational Algebra

Limitations of Current Datalog Engines

Multi-threaded

Distributed
(Apache Spark)Multi-node
Multi-threaded

Single-GPU

Multi-node
Multi-GPU

Soufflé

RDFox

SLOG

GPUJoin

LogicBlox

Radlog

PRAM

GPULog

Nemo

BigDatalog

GPUDatalog

Scalability

Framework

CPU-only

VRAM

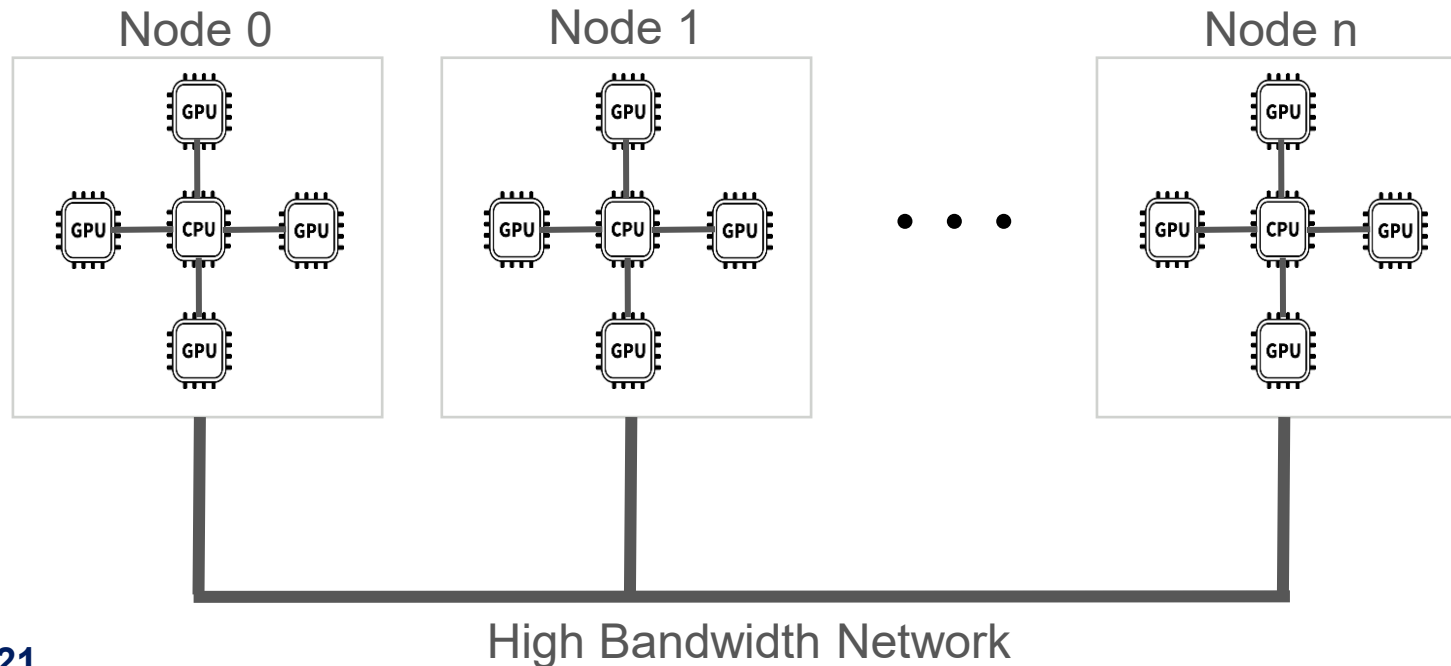
- Herbert Jordan, Bernhard Scholz, and Pavle Subotić. 2016. Soufflé: On synthesis of program analyzers. In Computer Aided Verification: 28th International Conference, CAV 2016, Toronto, ON, Canada, July 17-23, 2016, Proceedings, Part II 28, Swarat Chaudhuri and Azadeh Farzan (Eds.). Springer, Springer International Publishing, Cham, 422–430.
- Boris Motik, Yavor Nenov, Robert Piro, Ian Horrocks, and Dan Olteanu. 2014. Parallel materialisation of datalog programs in centralised, main-memory RDF systems. In Proceedings of the AAAI Conference on Artificial Intelligence, Vol. 28.
- Shkapsky, A., Yang, M., Interlandi, M., Chiu, H., Condie, T., & Zaniolo, C. (2016, June). Big data analytics with datalog queries on spark. In Proceedings of the 2016 International Conference on Management of Data (pp. 1135-1149).
- Yavor Nenov, Robert Piro, Boris Motik, Ian Horrocks, Zhe Wu, and Jay Banerjee. 2015. RDFox: A Highly-Scalable RDF Store. In The Semantic Web - ISWC 2015, Marcelo Arenas, Oscar Corcho, Elen Simperl, Markus Strohmaier, Mathieu d'Aquin, Kavitha Srinivas, Paul Groth, Michel Dumontier, Jeff Heflin, Krishnaprasad Thirunarayan, and Steffen Staab (Eds.). Springer International Publishing, Cham, 3–20.
- Thomas Gilray, Arash Sahebolamri, Yihao Sun, Sowmith Kunapaneni, Sidharth Kumar, and Kristopher Micinski. 2024. Datalog with First-Class Facts. Proc. VLDB Endow. 18, 3 (November 2024), 651–665.

Motivation: Datalog on Multi-Node Multi-GPU Systems

Datalog

Relational algebra

Multi-node multi-GPU systems



High performance
implementation for
highly expressive
language

Selected publications

Conference Proceedings

Towards iterative relational algebra on the {GPU}

Acceptance rate: 19%

**USENIX
ATC '23**

Bruck Algorithm Performance Analysis for Multi-GPU All-to-All Communication

Acceptance rate: 44%

**HPC
Asia
2024**

Optimizing Datalog for the GPU

Acceptance rate: 15%

 **ASPLOS 2025**

Multi-node multi-GPU Datalog

Acceptance rate: 23%

ACM ICS '25

Journal Articles

The robustness of persistent homology of brain networks to data acquisition-related non-neural variability in resting state fMRI

Impact factor: 5.0

HUMAN BRAIN MAPPING

UALCAN: An update to the integrated cancer data analysis platform

Impact factor: 6.3

NEOPLASIA

Workshop Papers

Accelerating Datalog applications with cuDF

Acceptance rate: 42%

IA³ 2022

Scalable, interactive and hierarchical visualization of virus taxonomic data

VAHC 2023

Submission

Topology assisted clustering of temporal fMRI brain networks with use-case in mitigating non-neural multi-site variability

IEEE Access

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IEEE Access

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5. Power analysis of Datalog engines
6. Exploration of high-dimensional scientific analytics
7. Future work
8. Conclusion

GPU based Datalog engine development

Off-the-shelf

Started with CPU and GPU
based Python libraries

Compare GPU capabilities
for iterative join operations

IA³ 2022

GPUJoin

Relational Algebra (RA)
operations using CUDA

High performance GPU
hashtable for iterative RA

GPULog

CUDA based SIMD API for
deductive analytics

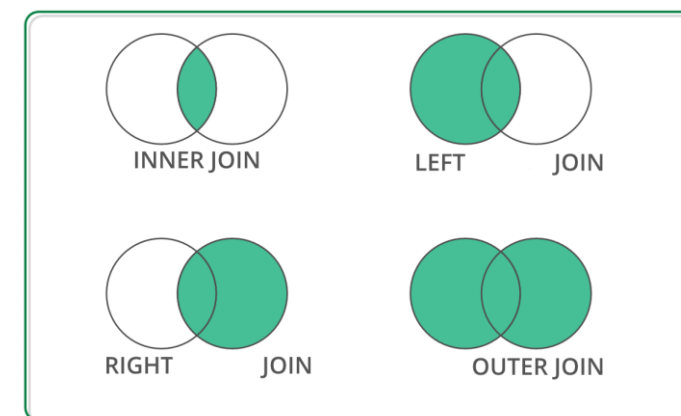
Novel Hash-Indexed Sorted
Array data structure

Off-the-shelf Data Structure for Join Operation

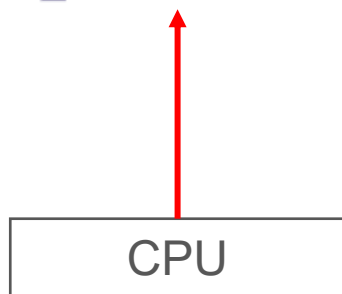


DataFrame: tabular data structure

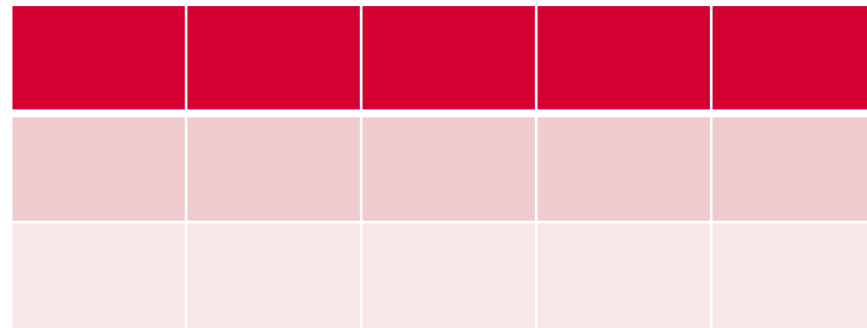
DataFrame has RA primitives APIs



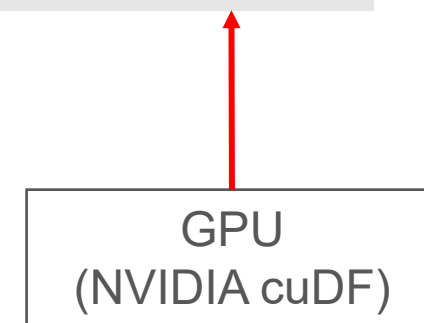
Off-the-shelf Python Libraries



DataFrame: 2D labeled tabular data structure



RAPIDS



Both supports join operation with similar APIs

- Reback, J., McKinney, W., Van Den Bossche, J., Augspurger, T., Cloud, P., Klein, A., ... & Seabold, S. (2020). pandas-dev/pandas: Pandas 1.0. 5. Zenodo.
- Chen, D. Y. (2017). Pandas for everyone: Python data analysis. Addison-Wesley Professional.
- Green, O., Du, Z., Patel, S., Xie, Z., Liu, H., & Bader, D. A. (2021, December). Anti-Section Transitive Closure. In 2021 IEEE 28th International Conference on High Performance Computing, Data, and Analytics (HiPC) (pp. 192-201). IEEE.
- Fender, A., Rees, B., & Eaton, J. RAPIDS cuGraph. In Massive Graph Analytics (pp. 483-493). Chapman and Hall/CRC.

CPU (Pandas) and GPU (cuDF)

```
import pandas as pd
```

CPU Environment

```
import cudf
```

GPU Environment

```
def get_read_csv(filename, method='cudf', n):  
    column_names = ['column 1', 'column 2']  
    if method == 'df':  
        return pd.read_csv(filename, sep='\t', header=None,  
                           names=column_names, nrows=n)  
    return cudf.read_csv(filename, sep='\t', header=None,  
                        names=column_names, nrows=n)  
  
def get_join(relation_1, relation_2):  
    column_names = ['column 1', 'column 2']  
    return relation_1.merge(relation_2, on=column_names[0],  
                           how="inner",  
                           suffixes=('_relation_1', '_relation_2'))
```

Evaluation environment, applications, datasets

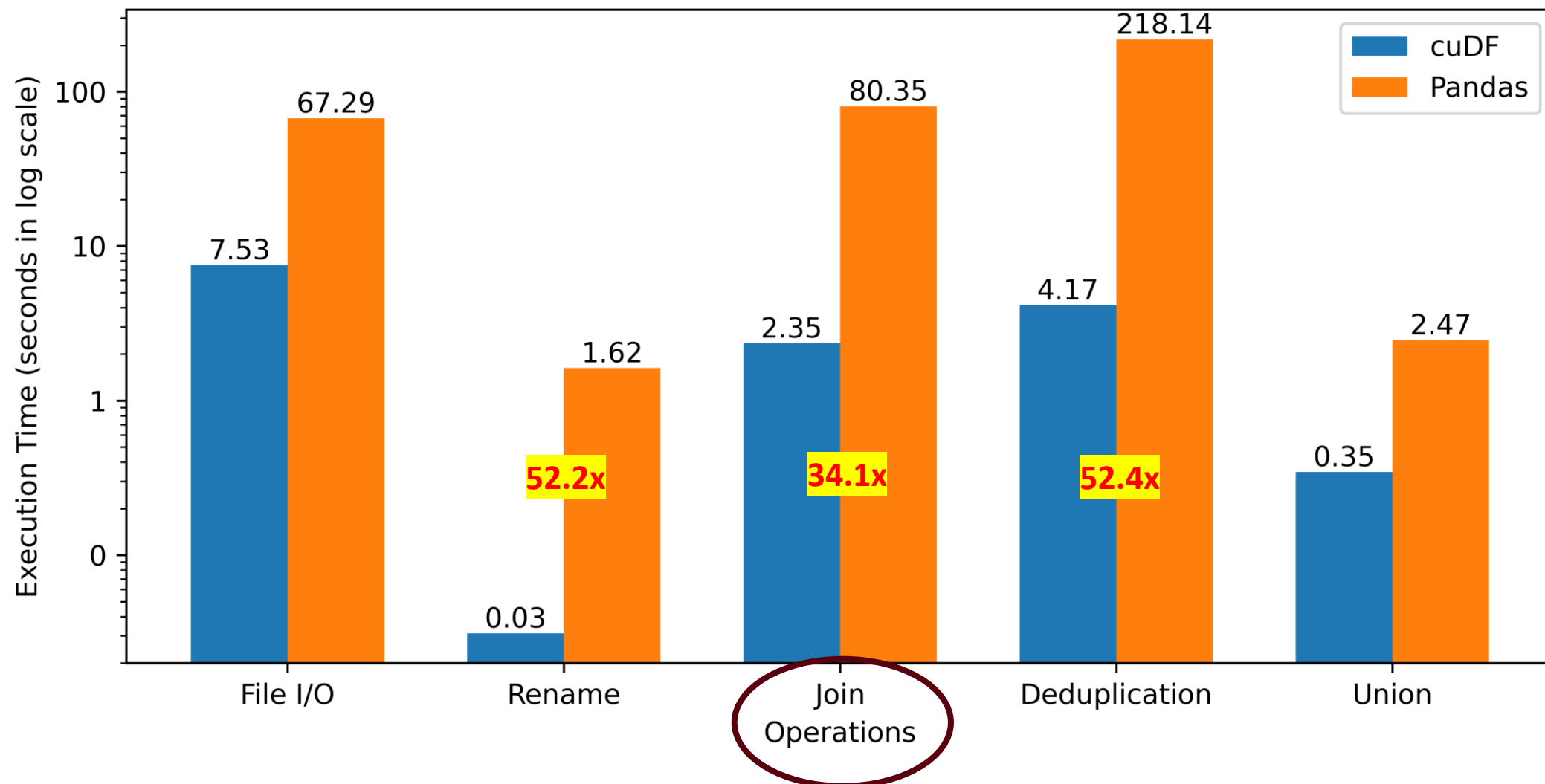


Apps: Transitive Closure, Triangle Counting

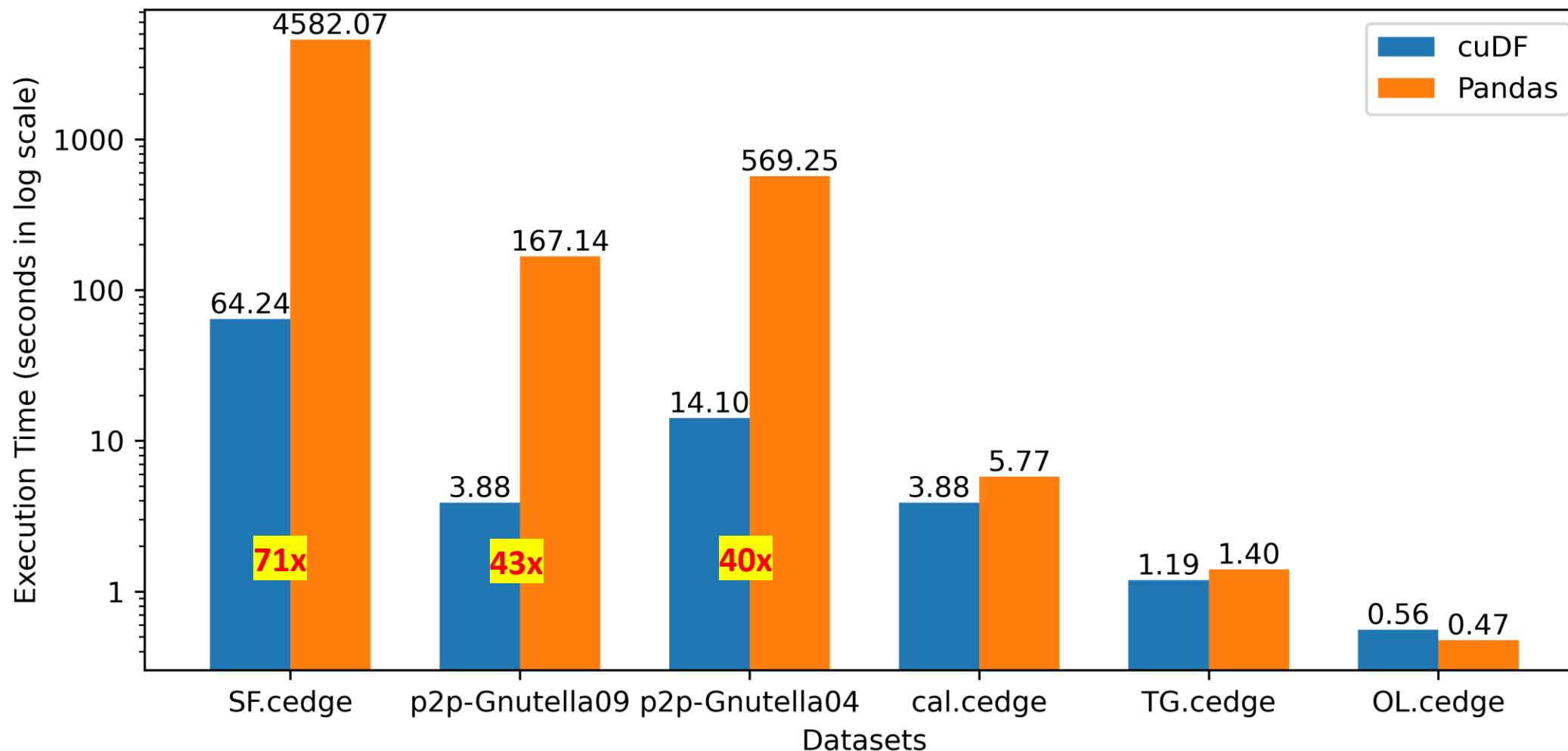
Datasets: Stanford large network, SuiteSparse, Road network

- Argonne Leadership Computing Facility. 2022. Theta. <https://www.alcf.anl.gov/alcf-resources/theta>
- Jure Leskovec and Andrej Krevl. 2014. SNAP Datasets: Stanford Large Network Dataset Collection. <http://snap.stanford.edu/data>
- Timothy A. Davis and Yifan Hu. 2011. The University of Florida Sparse Matrix Collection. ACM Trans. Math. Softw. 38, 1, Article 1 (dec 2011), 25 pages. doi:10.1145/2049662.2049663

Performance Improvement for cuDF over Pandas (Standalone Operations)

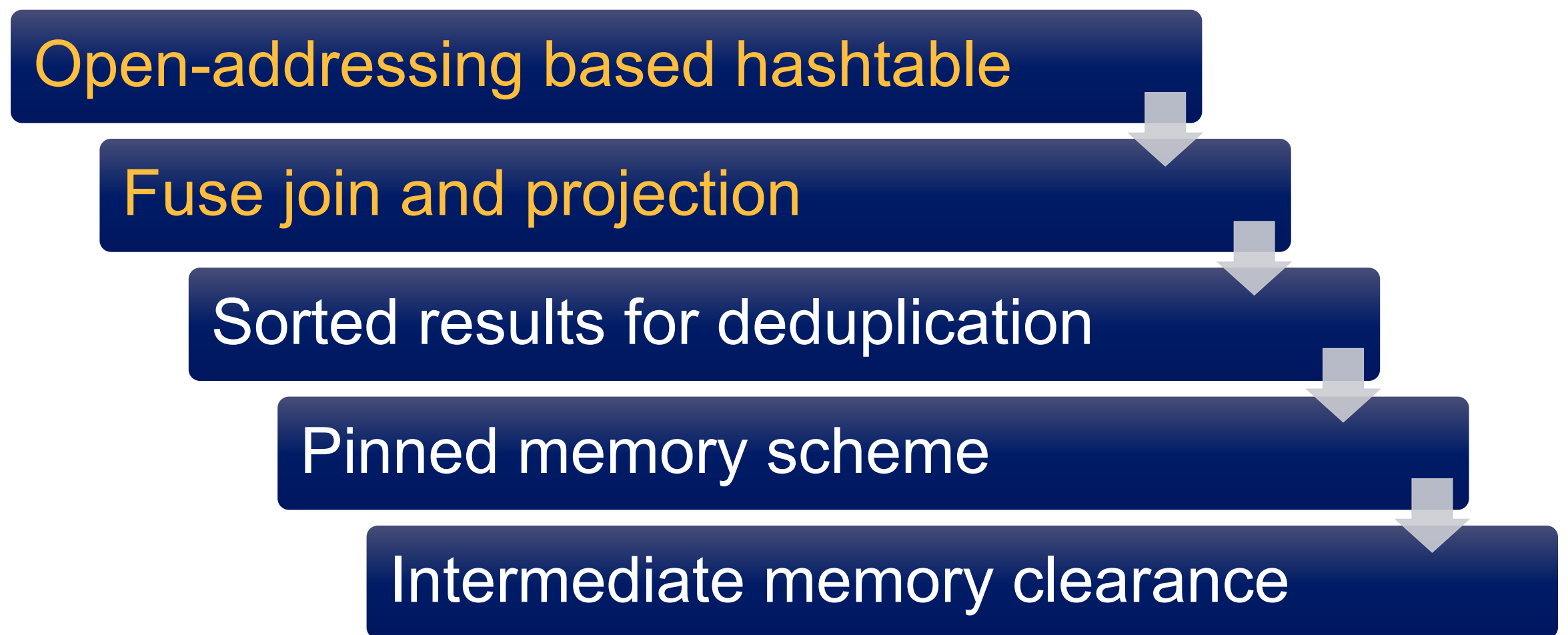


Performance Improvement for cuDF over Pandas (Transitive Closure)



Improvement Opportunity

Open-addressing based hashtable



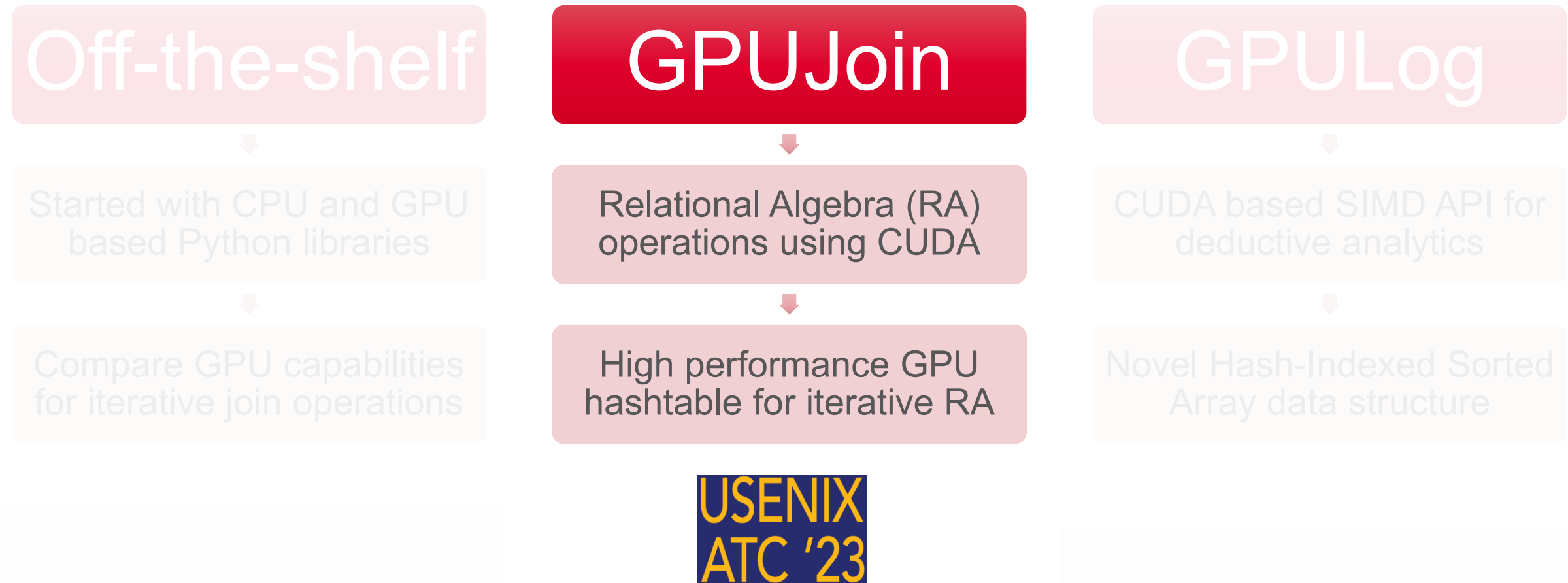
Fuse join and projection

Sorted results for deduplication

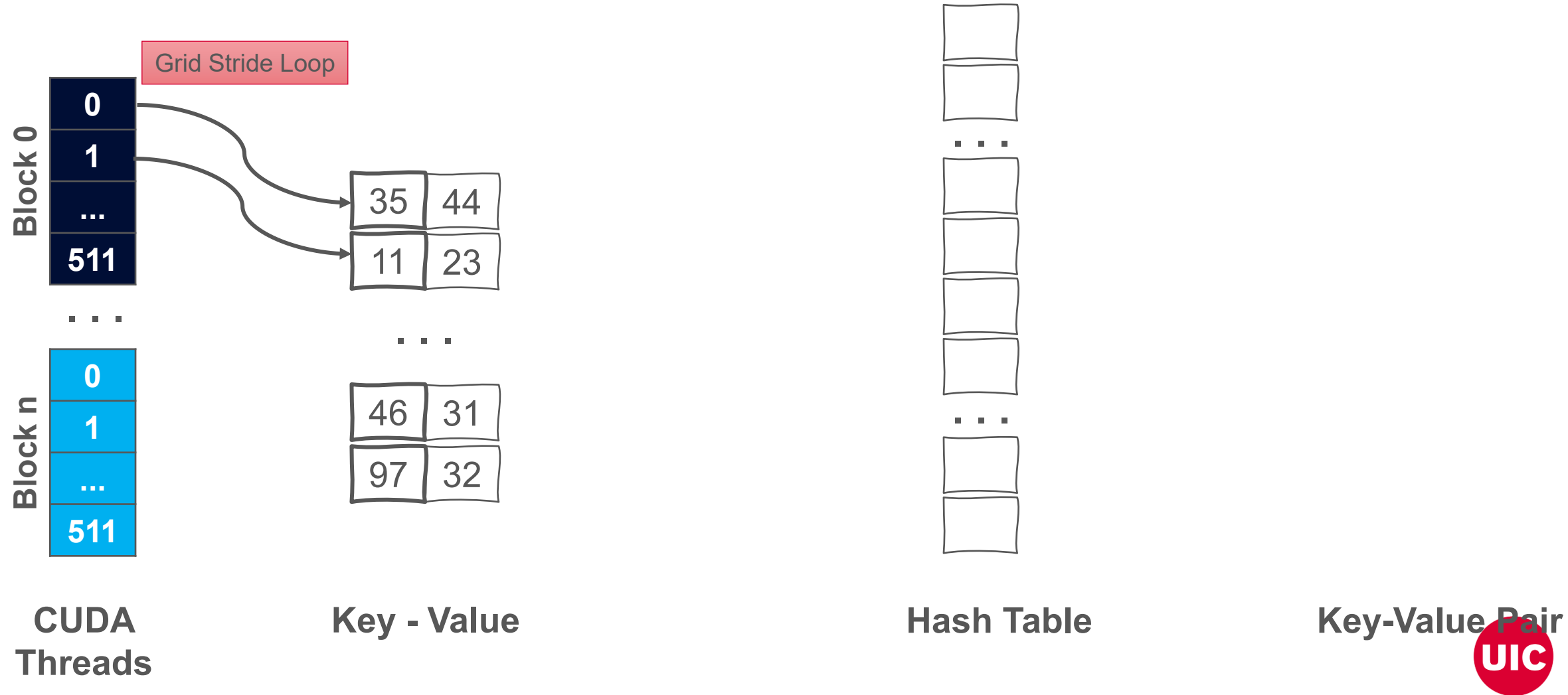
Pinned memory scheme

Intermediate memory clearance

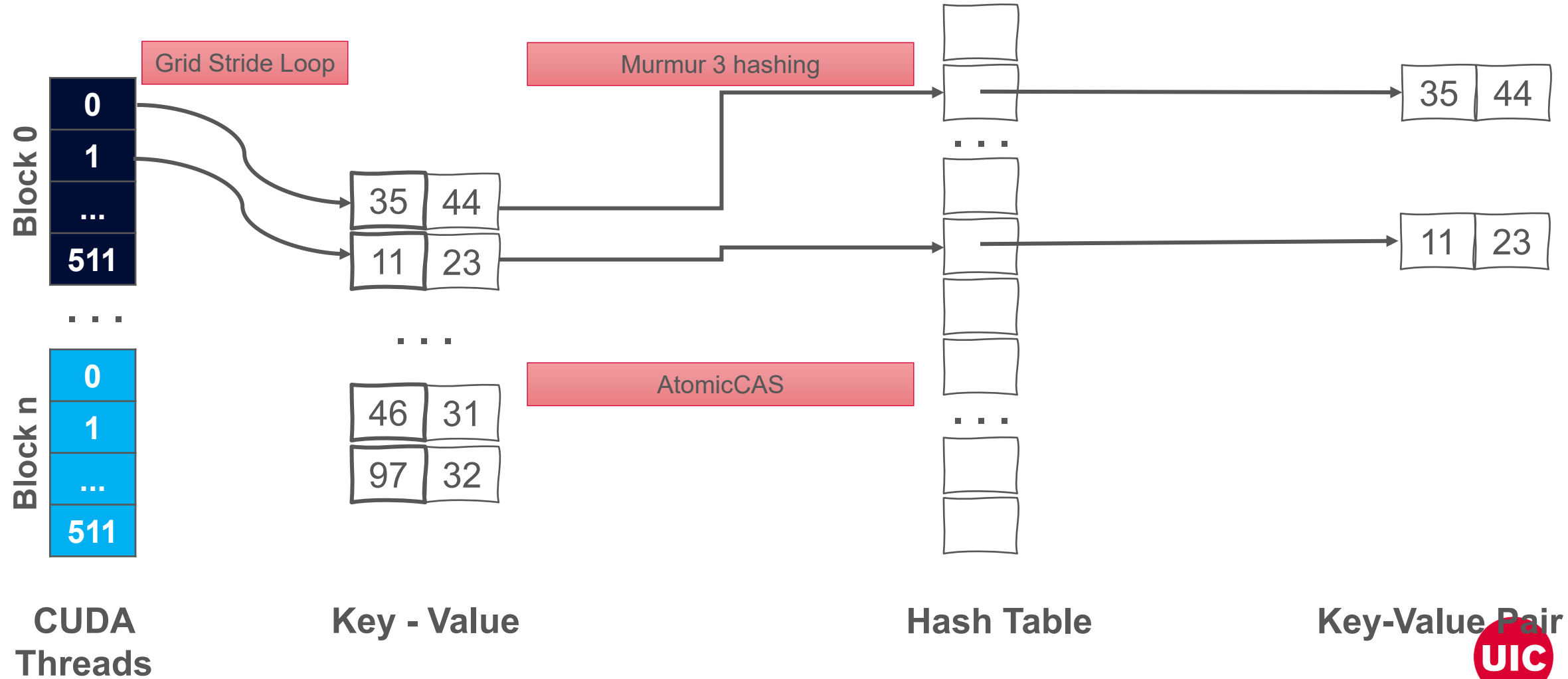
GPU based Datalog engine development



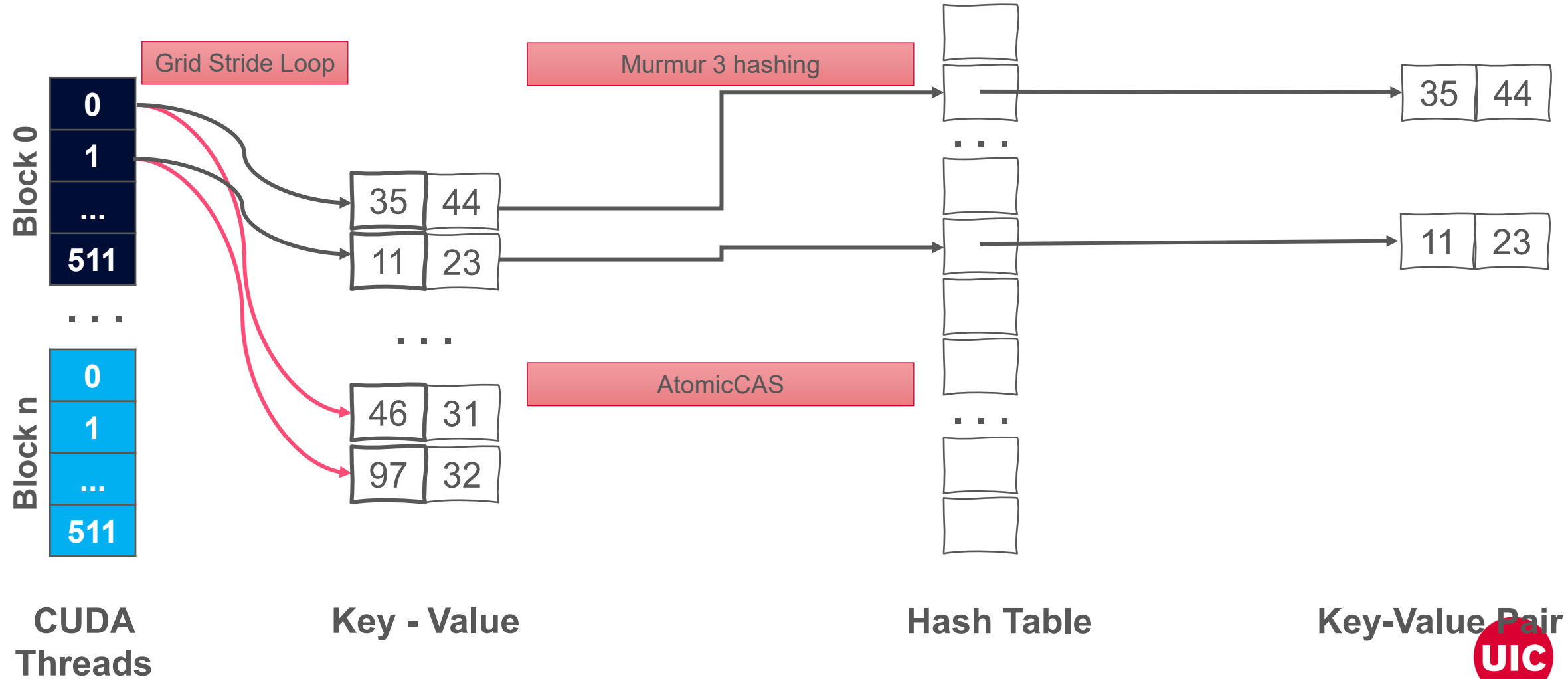
Hash Table (Open Addressing, Linear Probing)



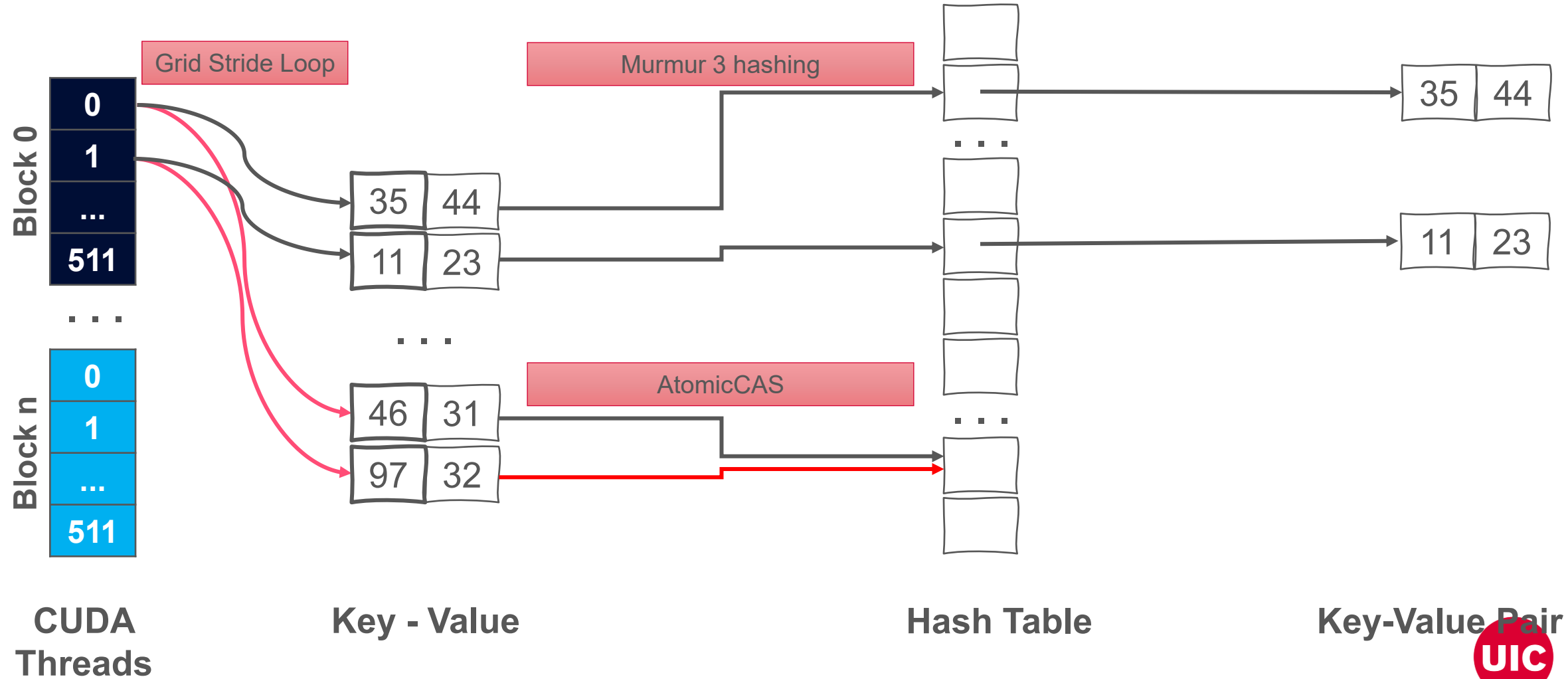
Hash Table (Open Addressing, Linear Probing)



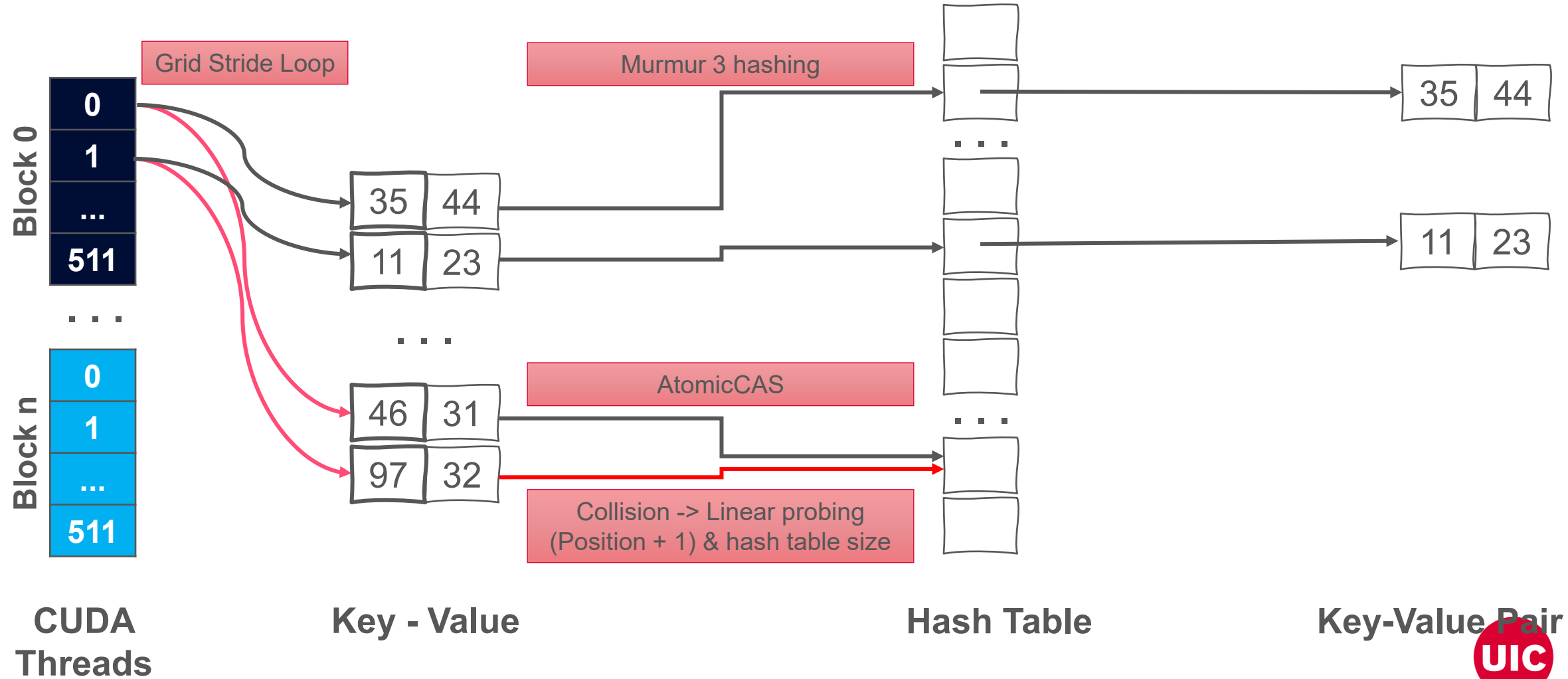
Hash Table (Open Addressing, Linear Probing)



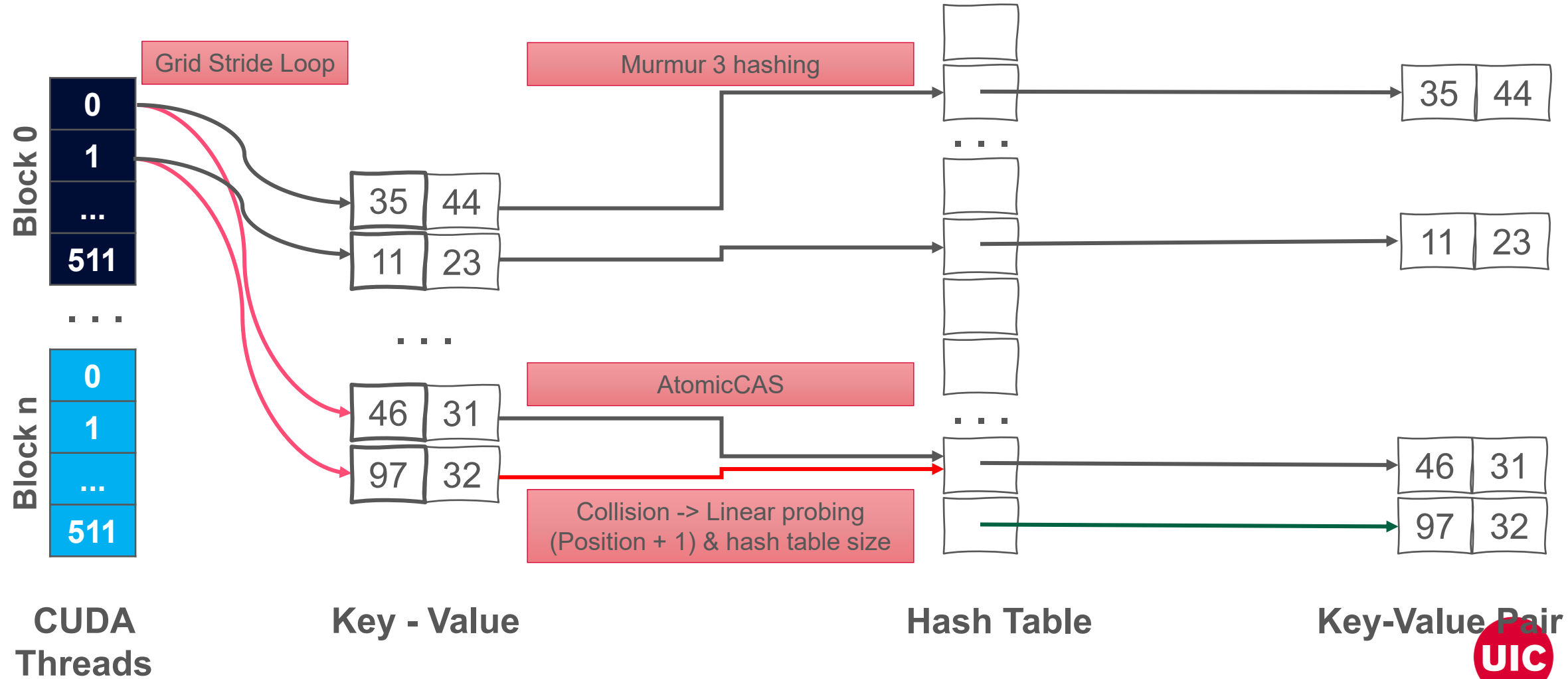
Hash Table (Open Addressing, Linear Probing)



Hash Table (Open Addressing, Linear Probing)

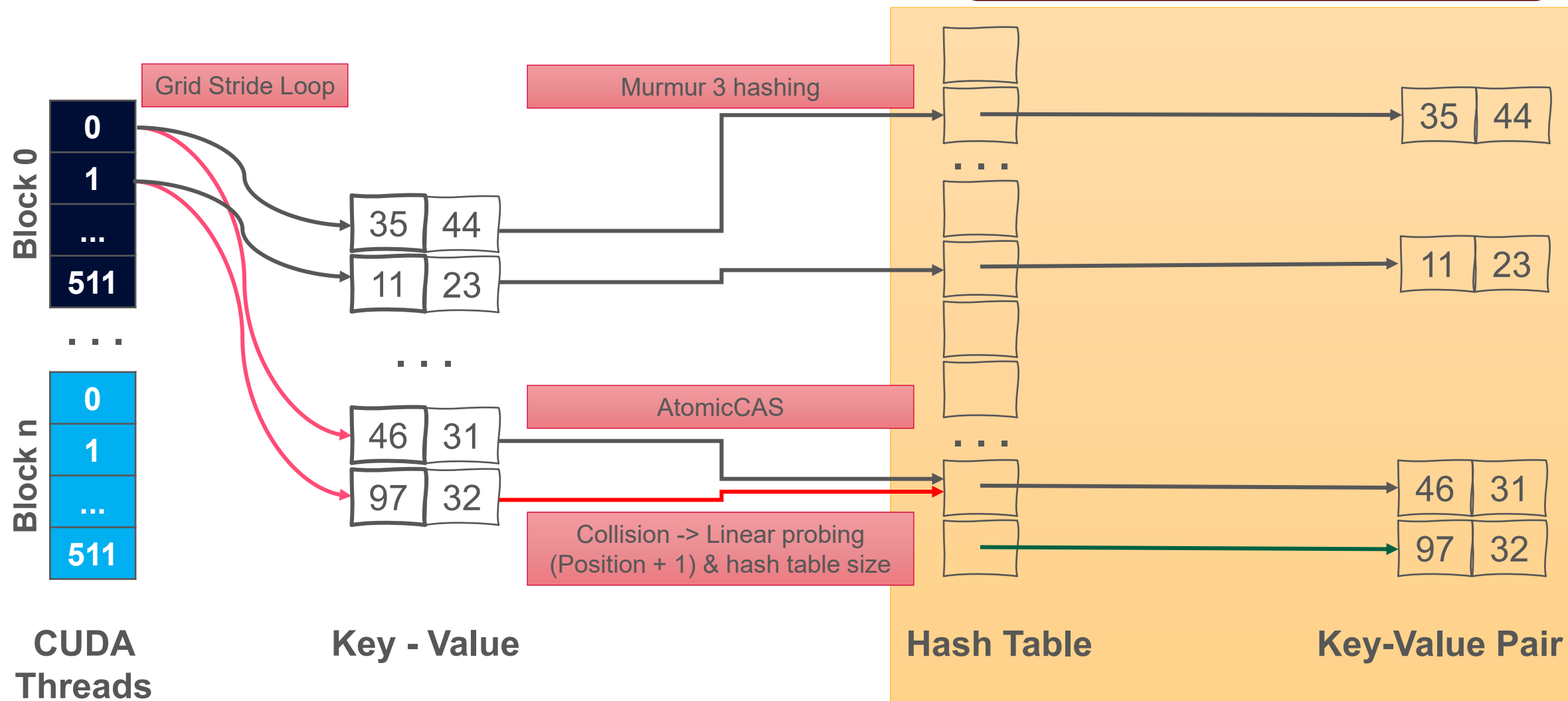


Hash Table (Open Addressing, Linear Probing)



Hash Table (Open Addressing, Linear Probing)

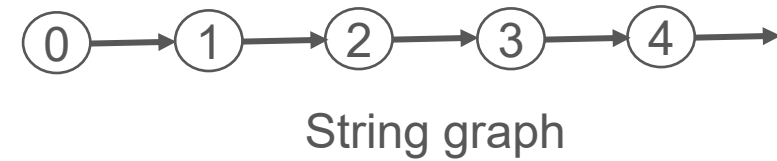
Stores key-value pairs in sparse space
Uses extra memory



Hash Table Performance

Build rate:

- Random synthetic graph: **400 million keys/second**
- String graph: **4 billion keys/second**



Performing Hash Join on GPU

$$F_G(T) \triangleq G \cup \Pi_{1,2}(\rho_{0/1}(T) \bowtie_1 G)$$

Static Hash Table

Key	Value

Inner Relation

Key	Value



First pass

Calculate join size

Performing Hash Join on GPU

$$F_G(T) \triangleq G \cup \Pi_{1,2}(\rho_{0/1}(T) \bowtie_1 G)$$

Static Hash Table

Key	Value

Inner Relation

Key	Value



First pass

Second pass
(fused projection)

Calculate join size

Prefix sum

Join Result

Join K	Value 1	Value 2

Performing Hash Join on GPU

$$F_G(T) \triangleq G \cup \Pi_{1,2}(\rho_{0/1}(T) \bowtie_1 G)$$

Static Hash Table

Key	Value

Inner Relation

Key	Value



First pass

Second pass
(fused projection)

Calculate join size

Prefix sum

Join Result

Join Key	Value 1	Value 2

Sort and Unique

Deduplicated Join Result

Key	Value

Evaluation environment, applications, datasets

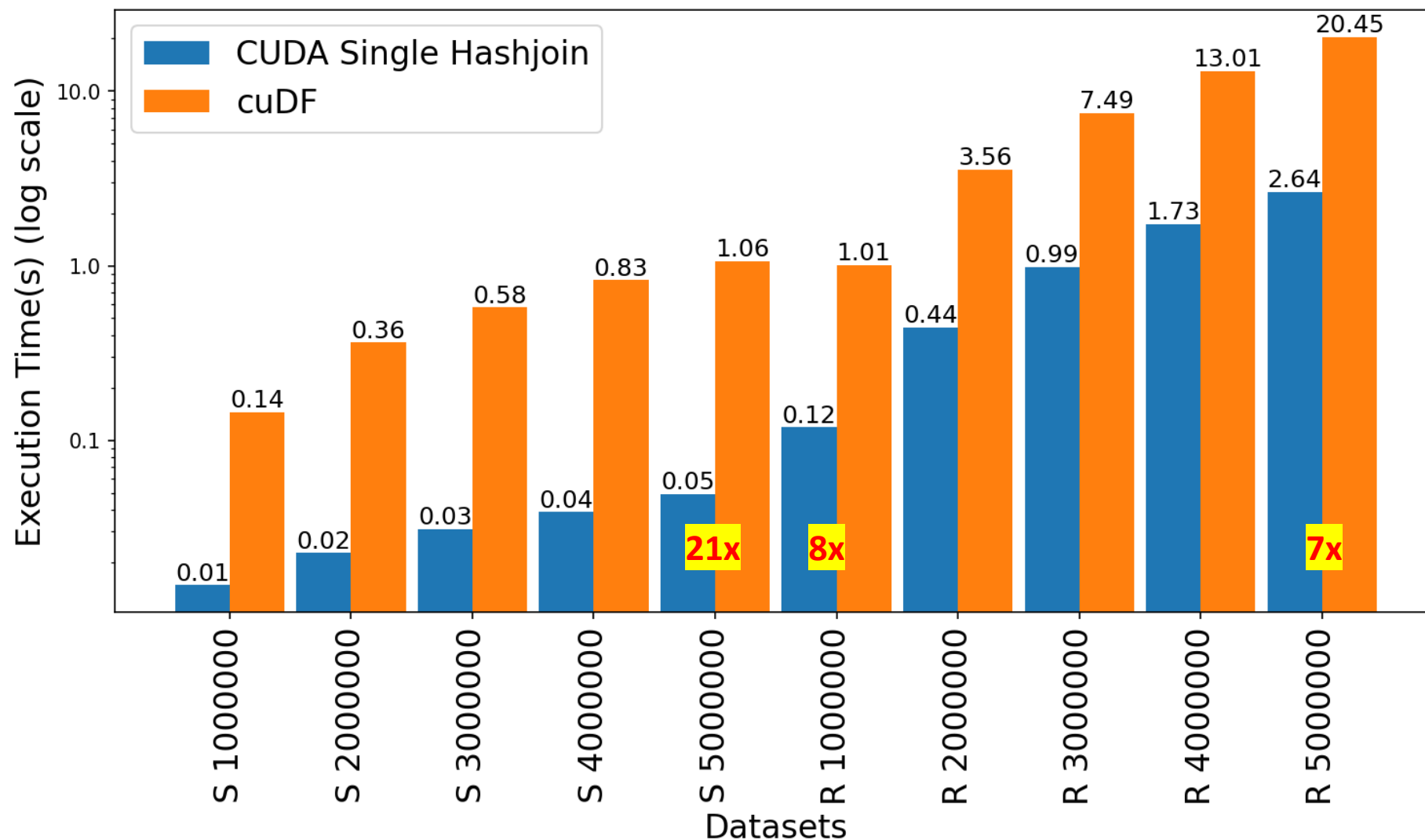


Apps: Transitive Closure, Iterated Hash Join

Datasets: Stanford large network, SuiteSparse, Road network

- Argonne Leadership Computing Facility. 2022. Theta. <https://www.alcf.anl.gov/alcf-resources/theta>
- Jure Leskovec and Andrej Krevl. 2014. SNAP Datasets: Stanford Large Network Dataset Collection. <http://snap.stanford.edu/data>
- Timothy A. Davis and Yifan Hu. 2011. The University of Florida Sparse Matrix Collection. ACM Trans. Math. Softw. 38, 1, Article 1 (dec 2011), 25 pages. doi:10.1145/2049662.2049663

Join Performance Comparison: GPUJoin vs cuDF



Transitive closure performance evaluation

Dataset	Type	Rows	TC size	Iterations	GPUJoin(s)	Soufflé(s)	cuDF(s)
fe_ocean	U	409,593	1,669,750,513	247	138.237 3.9x	536.233	OOM
p2p-Gnutella31	D	147,892	884,179,859	31	OOM	128.917	OOM
usroads	U	165,435	871,365,688	606	364.554	222.761	OOM
fe_body	U	163,734	156,120,489	188	47.758	29.070	OOM
loc-Brightkite	U	214,078	138,269,412	24	15.880	29.184	OOM
SF.cedge	U	223,001	80,498,014	287	11.274	17.073	64.417
fe_sphere	U	49,152	78,557,912	188	13.159	20.008	80.077
CA-HepTh	D	51,971	74,619,885	18	4.318	15.206	26.115
p2p-Gnutella04	D	39,994	47,059,527	26	2.092	7.537	14.005
p2p-Gnutella09	D	26,013	21,402,960	20	0.720	3.094	3.906
wiki-Vote	D	103,689	11,947,132	10	1.137	3.172	6.841
cti	U	48,232	6,859,653	53	0.295	1.496 10x	3.181
delaunay_n16	U	196,575	6,137,959	101	1.137	1.612	5.596
luxembourg_osm	U	119,666	5,022,084	426	1.322	2.548	8.194
ego-Facebook	U	88,234	2,508,102	17	0.544	0.606	3.719
cal.cedge	U	21,693	501,755	195	0.489	0.455	2.756
TG.cedge	U	23,874	481,121	58	0.198	0.219	0.857
wing	U	121,544	329,438	11	0.085	0.193	0.905
OL.cedge	U	7,035	146,120	64	0.148	0.181	0.523

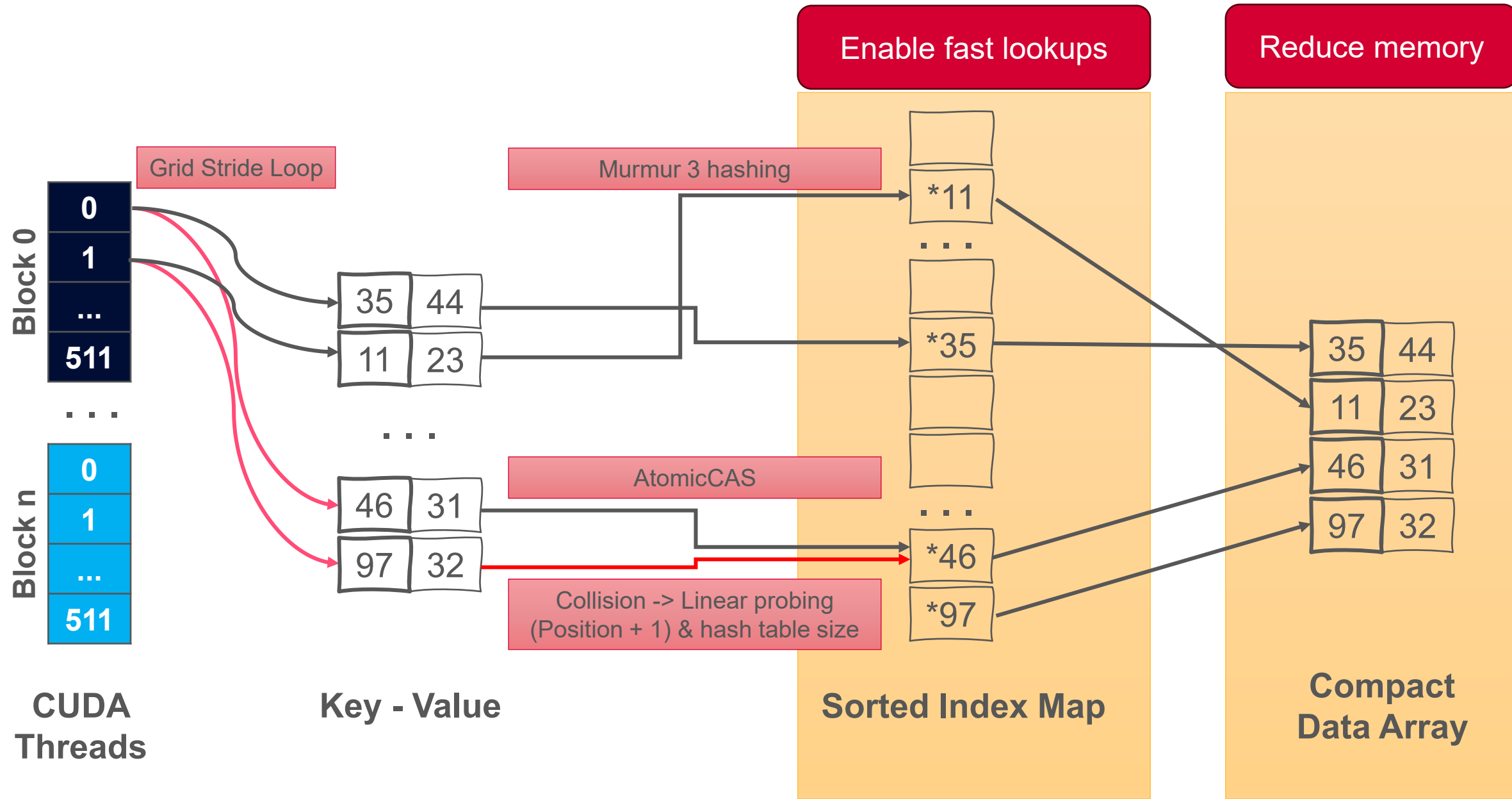


Fallback: Memory usage

GPU based Datalog engine development



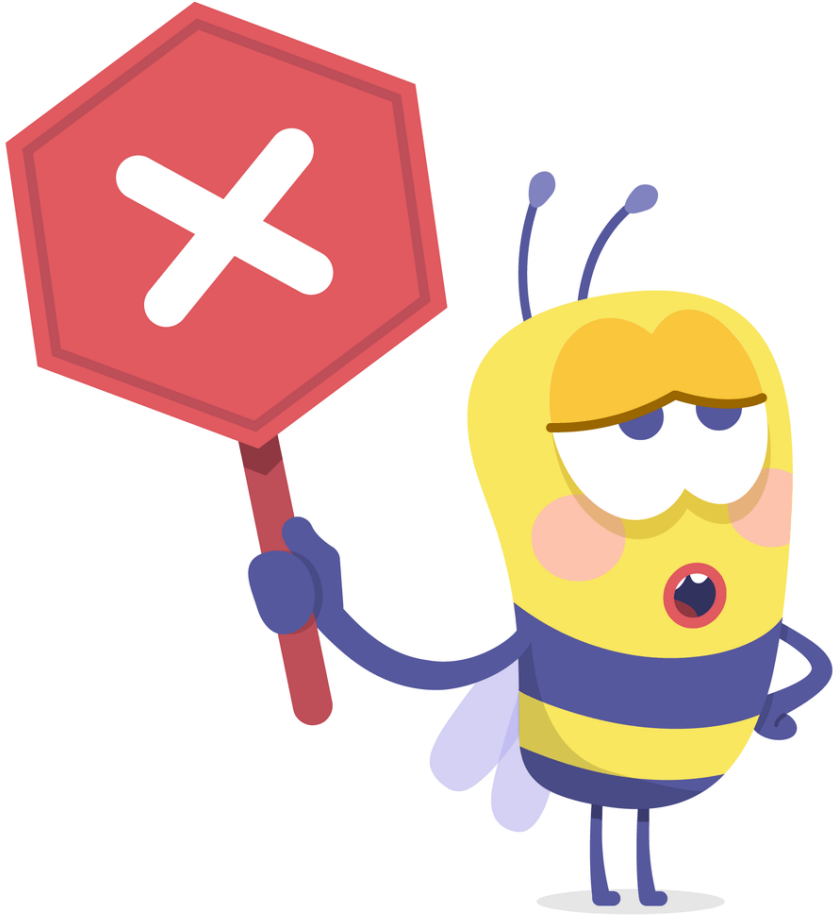
Hash Indexed Sorted Array (HISA)



Performance Enhancement (Transitive closure)

Dataset name	<i>Reach edges</i>	GPULOG	12x to 20x	3x to 6x	39x
			Soufflé	GPUJoin	cuDF
com-dblp	1.91B	14.30	232.99	OOM	OOM
fe_ocean	1.67B	23.36	292.15	100.30	OOM
vsp_finan	910M	21.91	239.33	125.94	OOM
Gnutella31	884M	5.58	96.82	OOM	OOM
fe_body	156M	3.76	23.40	22.35	OOM
SF.cedge	80M	1.63	33.27	3.76	64.29

Limitations



Limited to a single GPU

Memory overflow error for larger graphs

Roadmap

1. Introduction and motivation
2. GPU based Datalog engine development
- 3. Multi-node multi-GPU Datalog engine development**
4. GPU portability of Datalog engines
5. Power analysis of Datalog engines
6. Exploration of high-dimensional scientific analytics
7. Future work
8. Conclusion

Multi-node multi-GPU Datalog engine development

MNMGDatalog



Multi-node multi-GPU Datalog
engine

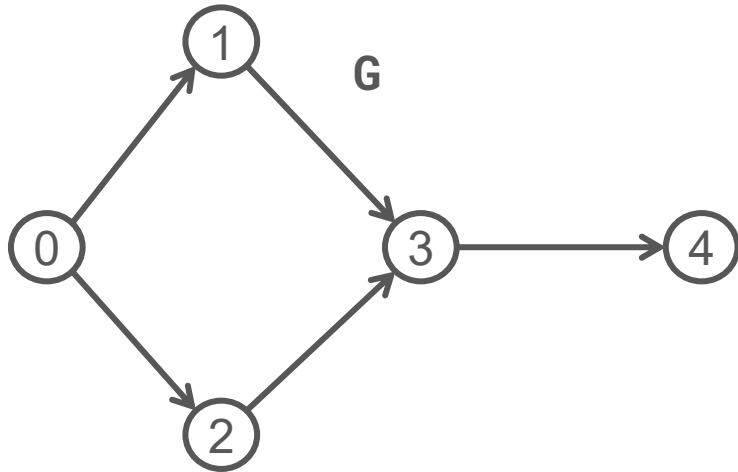


MPI + X (Cuda) design

3rd Best Poster Award at Computational Research Symposium 2025

ACM ICS '25

Requirements for Multi-node Multi-GPU DataLog



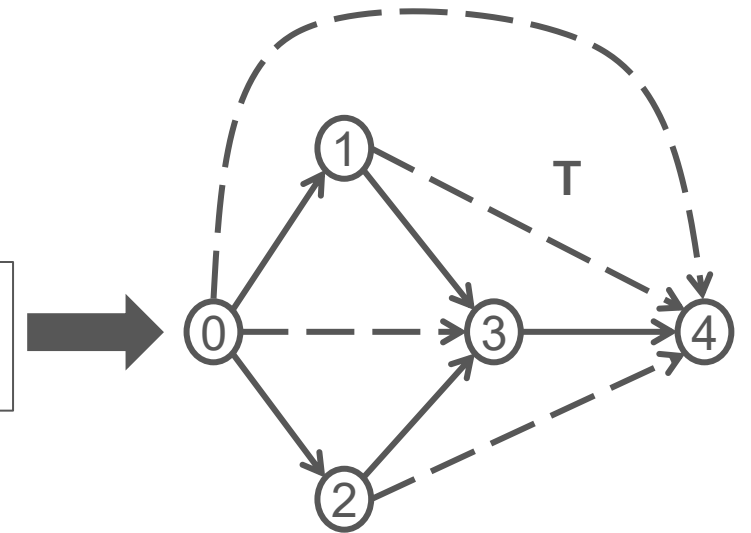
0	1
0	2
1	3
2	3
3	4

1 Workload partitioning

```

tc(X, Y) :- edges(X, Y).
tc(X, Z) :- tc(X, Y), edges(Y, Z).
  
```

2 Data representation



0	1
0	2
1	3
2	3
3	4
0	3
1	4
2	4
0	4

3 Communication

MNMGDataLog Implementation

Hash-based data
distribution

GPU-optimized data
representation

GPU-Aware all-to-all
communication

1

Workload partitioning

2

Data representation

3

Communication

MNMGDataLog: Radix-hash-based partitioning

Hash-based data
distribution

GPU-optimized data
representation

GPU-Aware all-to-all
communication

1 Workload partitioning

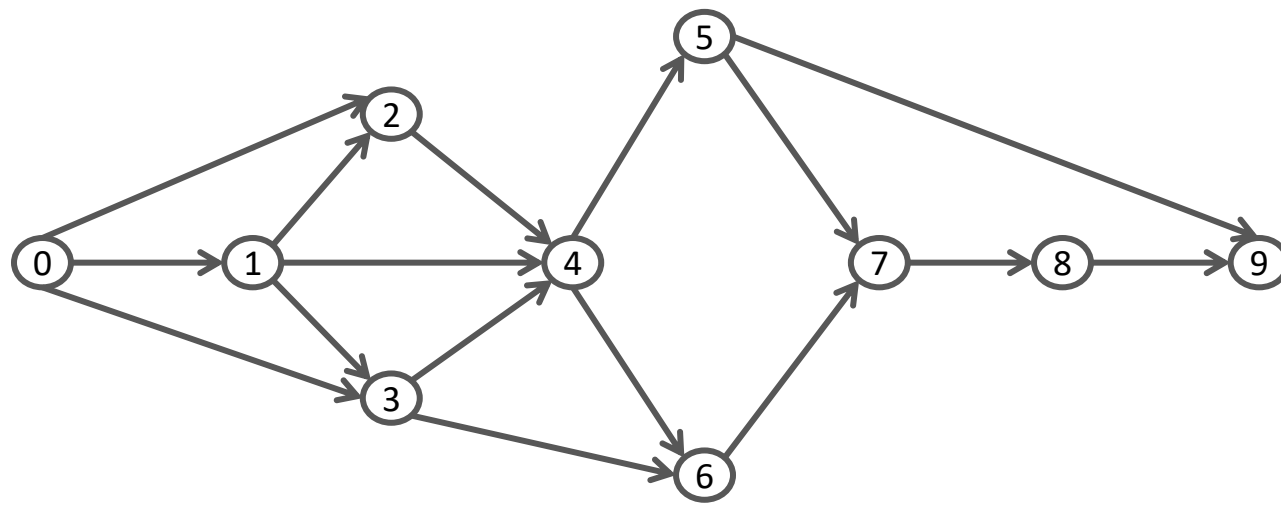
2 Data representation

3 Communication

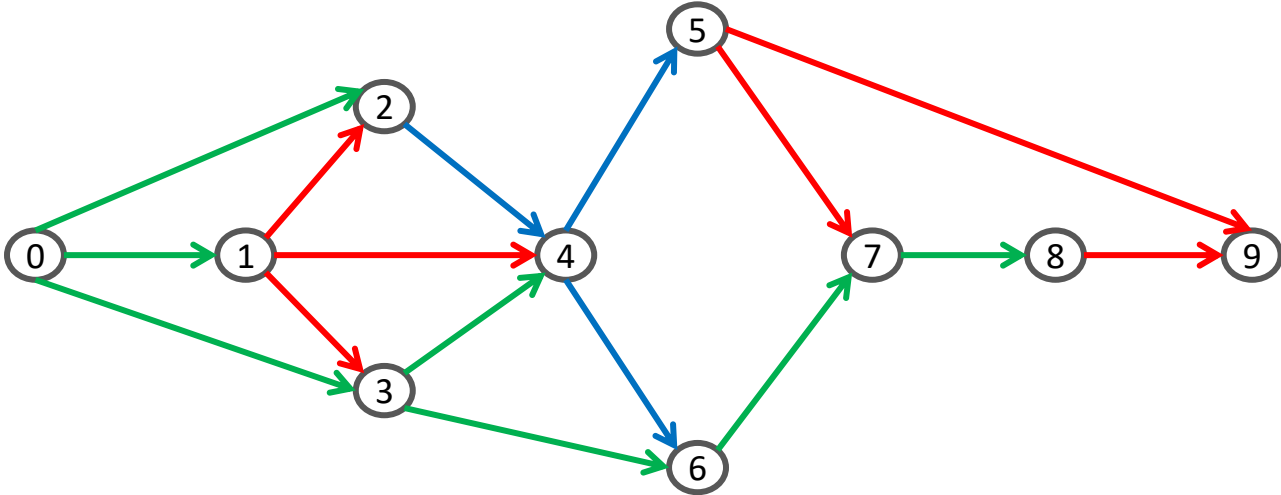
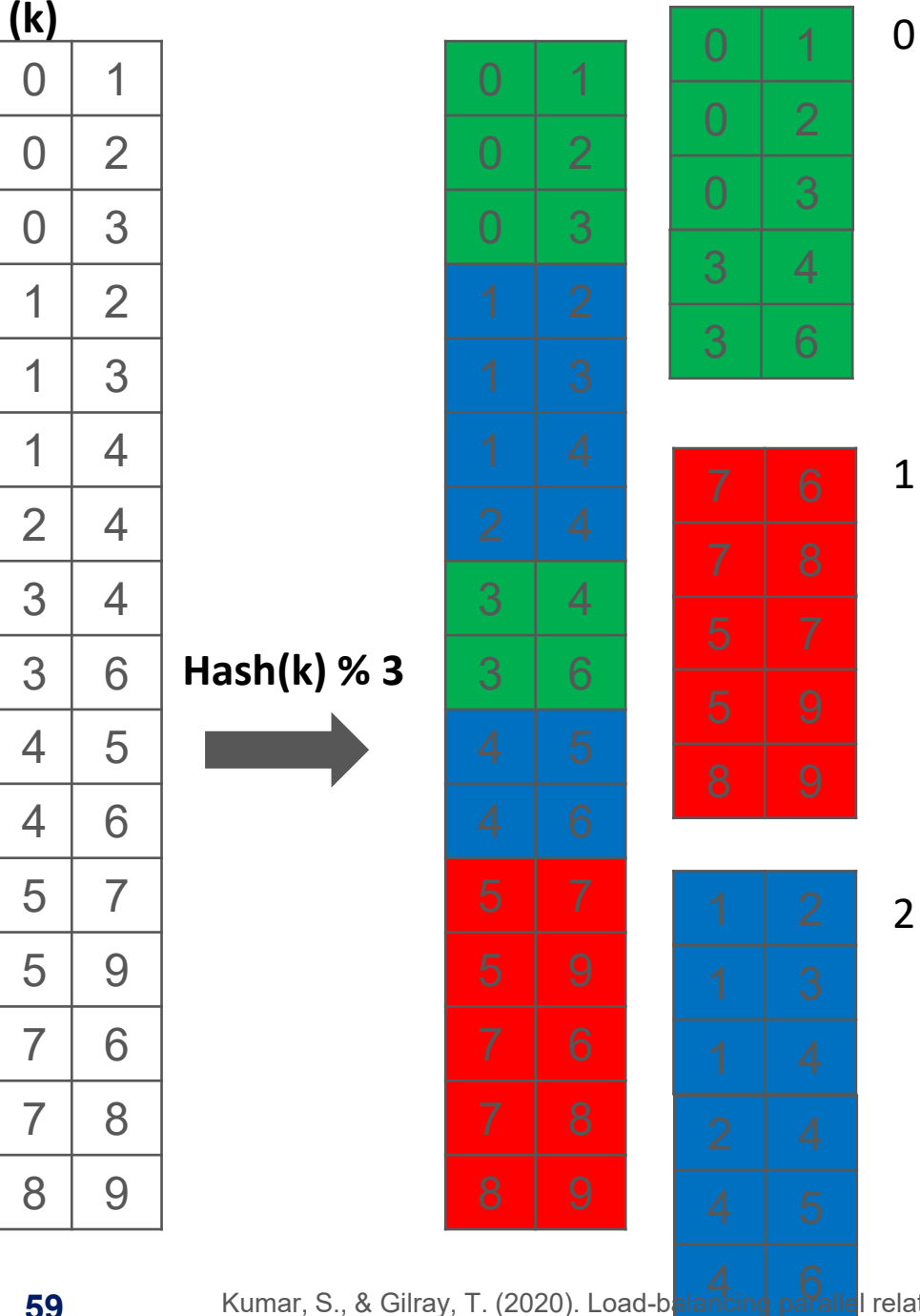
The join column decides the GPU (bucket-id)

(k)	
0	1
0	2
0	3
1	2
1	3
1	4
2	4
3	4
3	6
4	5
4	6
5	7
5	9
7	6
7	8
8	9

Hash-partitioning based on the Join Column



The join column decides the GPU (bucket-id)



MNMGDataLog: Data representation

Hash-based data
distribution

GPU-optimized data
representation

GPU-Aware all-to-all
communication

1

Workload partitioning

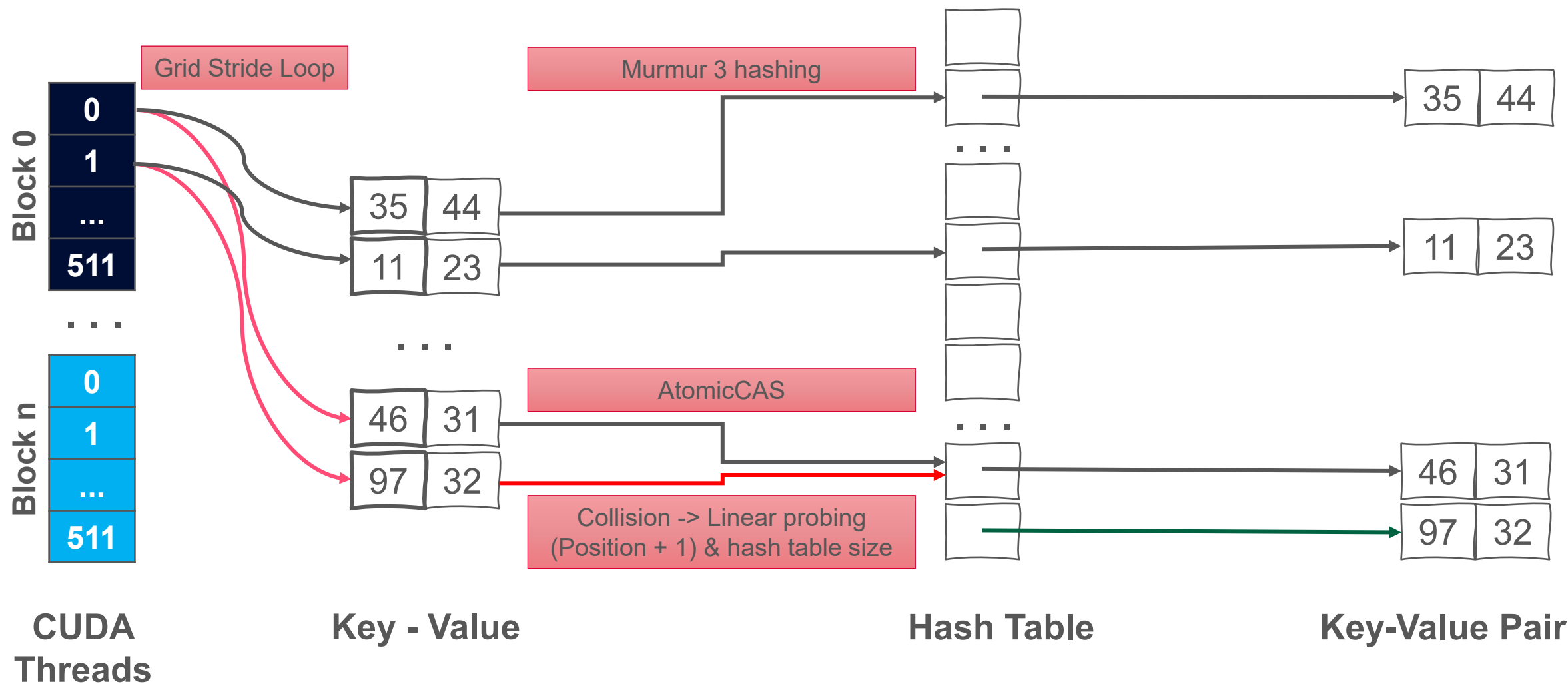
2

Data representation

3

Communication

GPU Hash Table (Open Addressing, Linear Probing)



MNMGDataLog: Communication for Iterative RA

Hash-based data
distribution

GPU-optimized data
representation

GPU-Aware all-to-all
communication

1

Workload partitioning

2

Data representation

3

Communication

Relational Algebra Kernel

Join

Project

Hash

All to all
comm

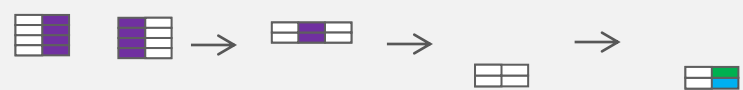
GPU 0



GPU 1



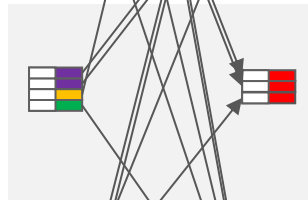
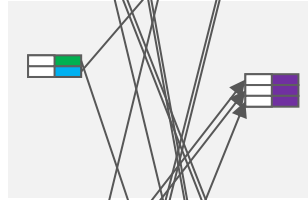
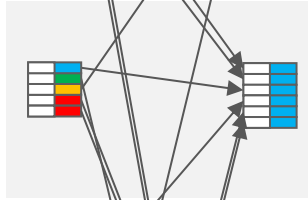
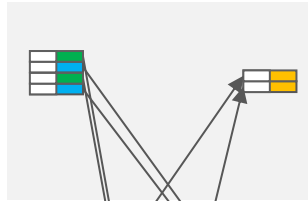
GPU 2



GPU 3



GPU 4



Relational Algebra Kernel

Join

Project

Hash

All to all
comm

Local inserts

GPU 0



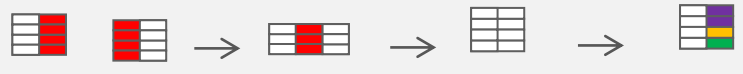
GPU 1



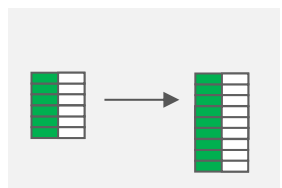
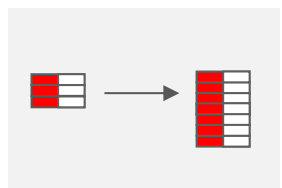
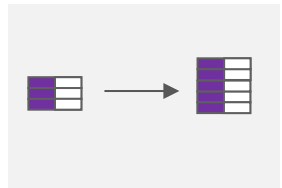
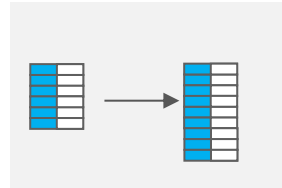
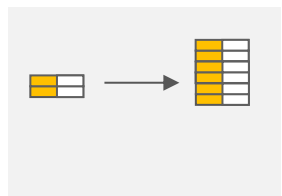
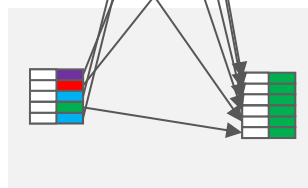
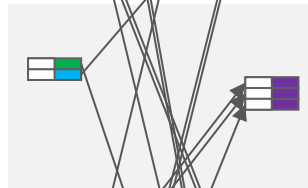
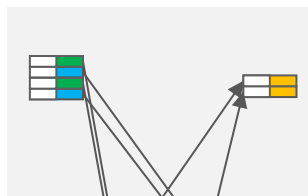
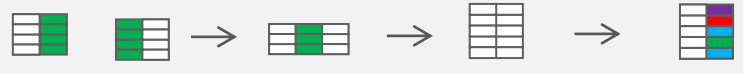
GPU 2

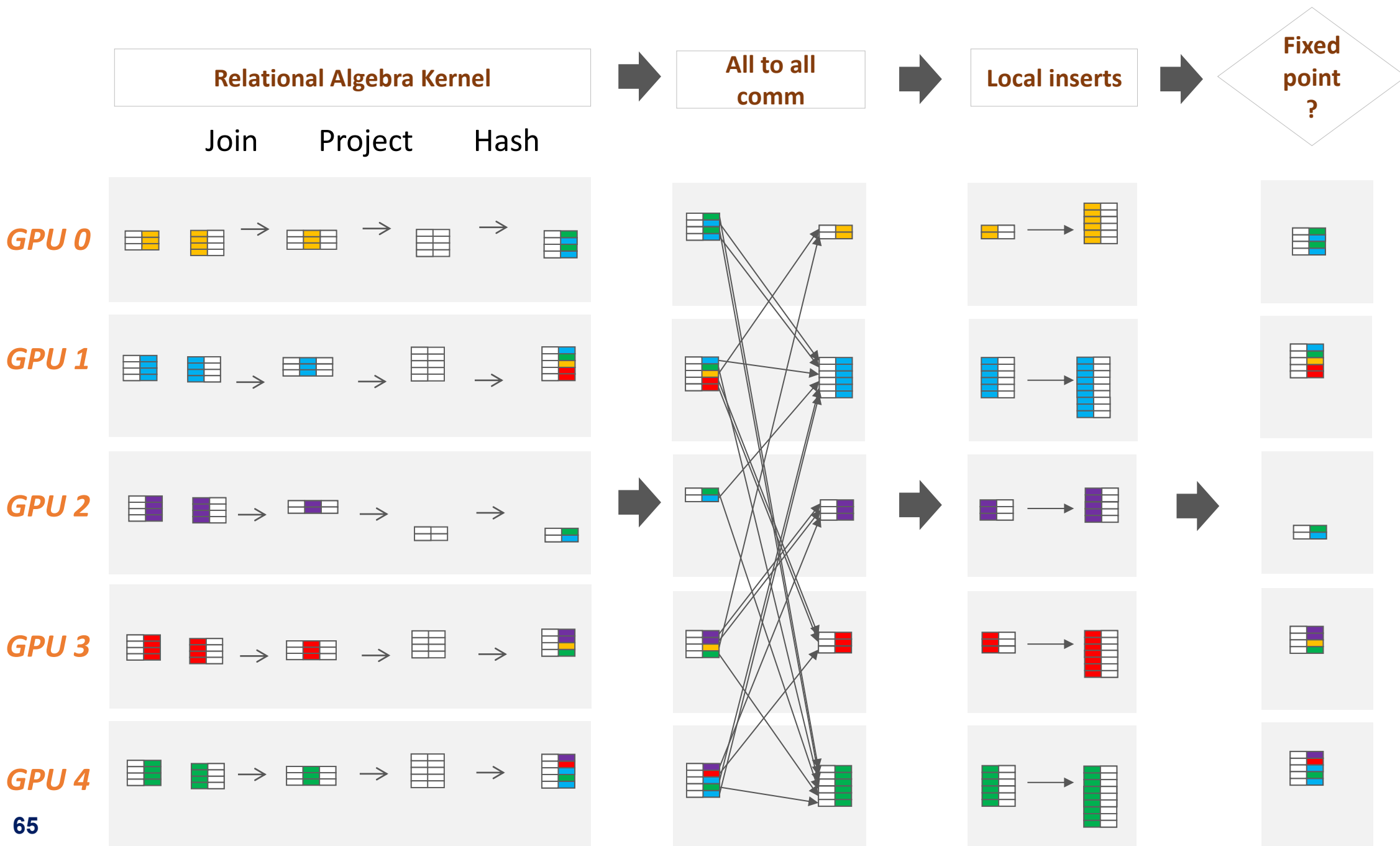


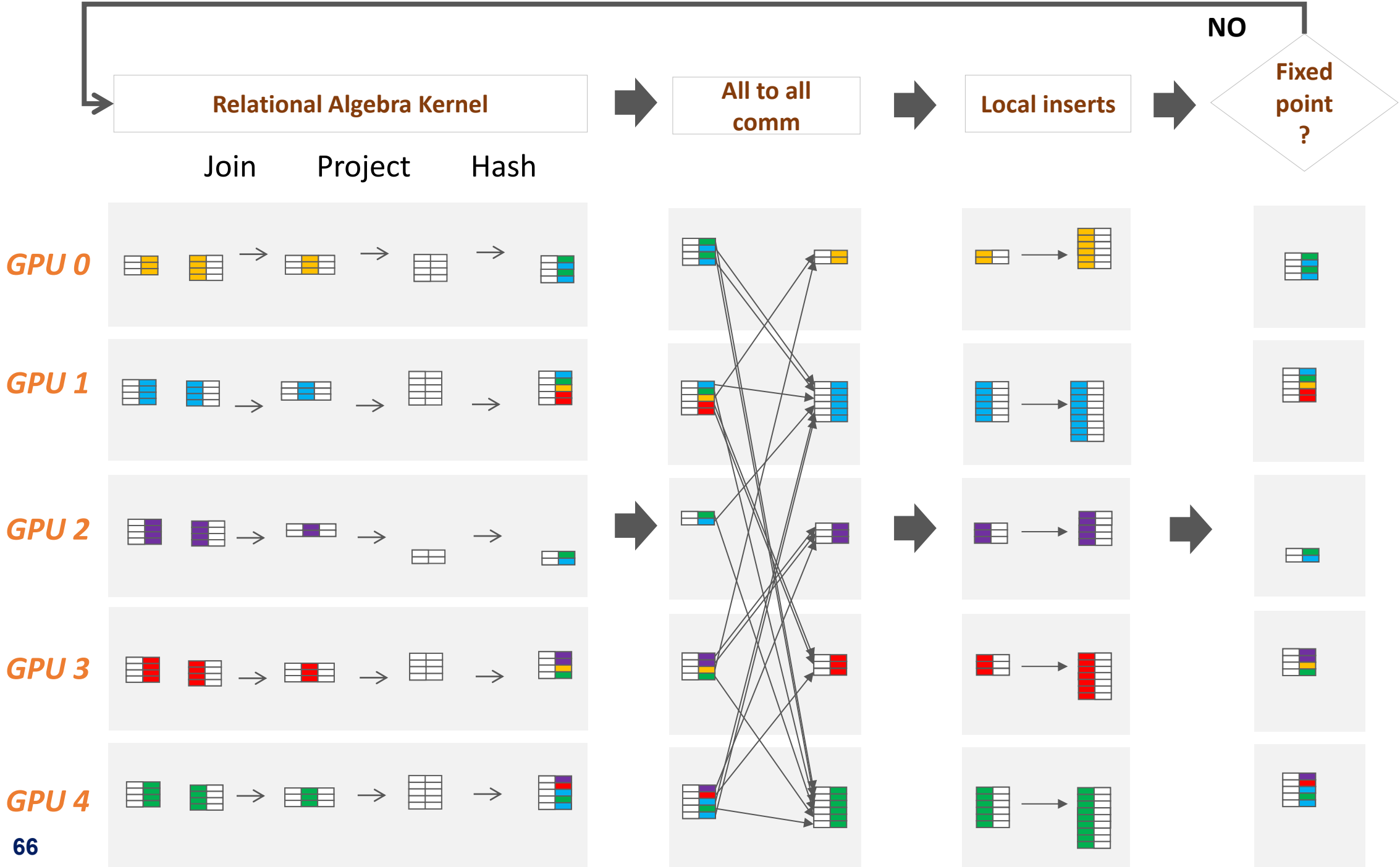
GPU 3

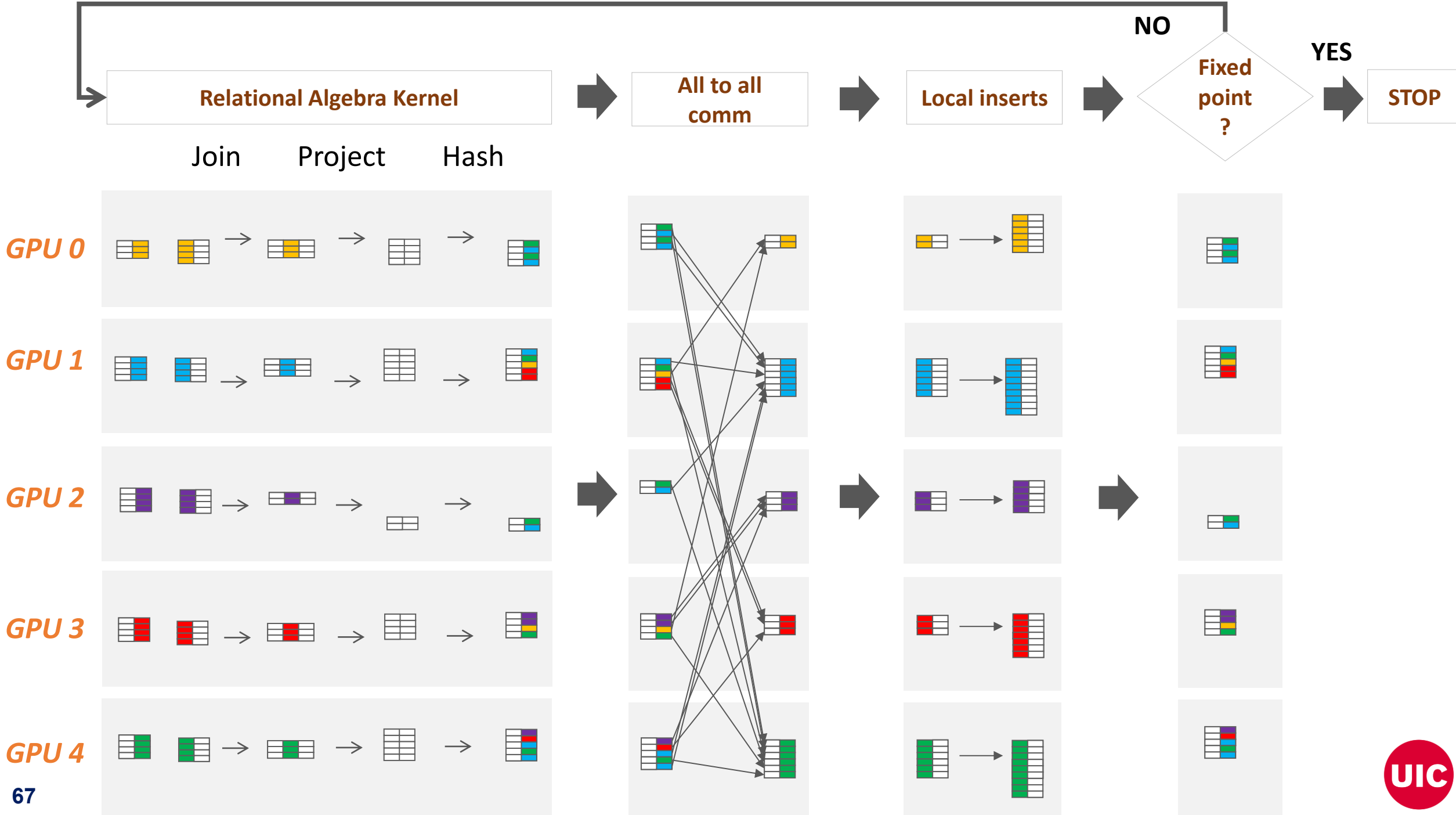


GPU 4









MNMGDataLog Evaluation



Polaris (Argonne National Lab)

Multi-threaded

Multi-node
Multi-threaded

Single-GPU

Soufflé

SLOG

GPUJoin

GPULog

cuDF

- Argonne Leadership Computing Facility. 2022. Polaris. <https://www.alcf.anl.gov/polaris>
- Jure Leskovec and Andrej Krevl. 2014. SNAP Datasets: Stanford Large Network Dataset Collection. <http://snap.stanford.edu/data>
- Timothy A. Davis and Yifan Hu. 2011. The University of Florida Sparse Matrix Collection. ACM Trans. Math. Softw. 38, 1, Article 1 (dec 2011), 25 pages. doi:10.1145/2049662.2049663

Single GPU benchmark

Dataset name	TC edges	Time (s)			
		MNMGDataLog	GPULOG	Soufflé	GPUJoin
com-dblp	1.91B	13.58	26.95	232.99	OOM
fe_ocean	1.67B	66.34	72.74	292.15	100.30
usroads	871M	75.07	78.08	222.76	364.55
vsp_finan	910M	81.14	82.75	239.33	125.94

Transitive closure benchmark on single-GPU

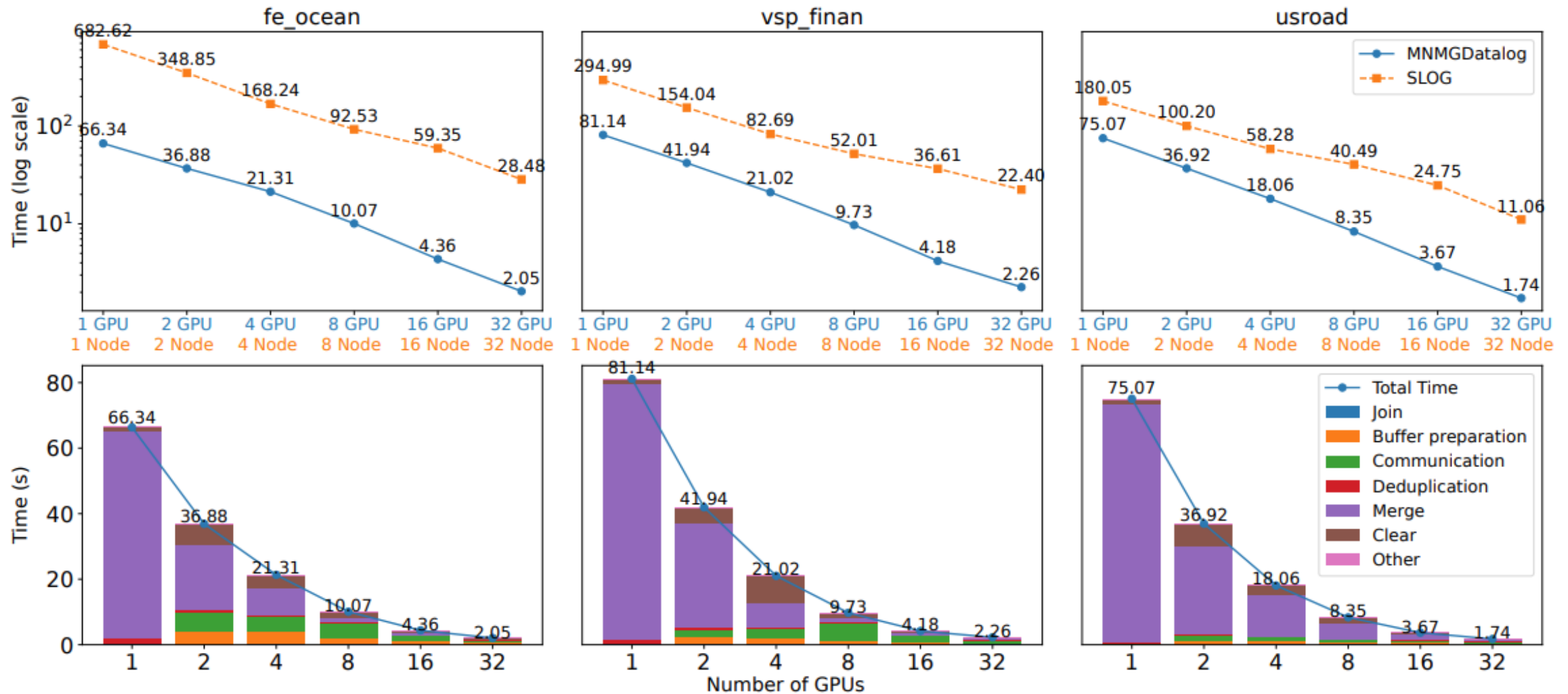
**Speedup up to 1.9x over GPULog, 20x over Souffle,
4.8x over GPUJoin**

Dataset name	SG size	Time (s)			
		MNMGDataLog	GPULOG	Soufflé	cuDF
fe_body	408M	9.08	18.41	74.26	OOM
loc-Brightkite	92.3M	1.66	11.67	48.18	OOM
fe_sphere	205M	3.55	7.88	48.12	OOM
CA-HepTH	74M	0.60	4.79	20.12	21.24

Same generation benchmark on single-GPU

**Speedup up to 7x over GPULog, 33.5x over Souffle,
35.4x over cuDF**

Multi-node multi-GPU benchmark



Transitive closure benchmark and breakdown up to 32 GPUs

Speedup up to 32x

Roadmap

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7. Future work
8. Conclusion

Parallel Programming Models



W.J. Cody Associates

Rosetta parallel programming models benchmarking framework

- Sequential
- OpenMP
- OpenMP-target
- CUDA
- HIP
- SYCL

Rosetta Benchmark		Code							
Sections		Benchmark Report							
- All Results (Table) - Walltime Plot - Speedup Plot: openmp-parallel - Speedup Plot: openmp-task - Speedup Plot: serial		All Results (Table)							
Benchmark	PPM	# Samples	Wall	User	Kernel	ompwtime	Max RSS	Peak Allocation	
idioms.assign	serial	54	1.57µs	1.63µs	None	N/A	N/A	78.12KiB	
idioms.assign	openmp-parallel	37925	50.58µs	2.14ms	45.35µs	50.43µs	22.17MiB	78.12KiB	
idioms.assign	openmp-task	11644	757.55µs	42.62ms	192.99µs	757.34µs	17.44MiB	78.12KiB	
idioms.pointwise	serial	103	1.59µs	1.57µs	None	N/A	N/A	78.12KiB	
idioms.pointwise	openmp-parallel	38113	56.17µs	2.48ms	55.12µs	56.35µs	22.23MiB	78.12KiB	
idioms.pointwise	openmp-task	10837	786.66µs	45.08ms	176.15µs	786.75µs	14.20MiB	78.12KiB	
idioms.reduction	serial	9	8.29µs	8.22µs	None	N/A	N/A	8.00B	
idioms.reduction	openmp-parallel	32145	68.30µs	2.93ms	53.20µs	68.10µs	17.71MiB	8.00B	
idioms.reduction	openmp-task	11618	760.66µs	43.64ms	172.65µs	760.60µs	17.00MiB	8.00B	
suites.polybench.2mm	serial	4	7.26ms	7.25ms	7.00µs	N/A	N/A	1.19MiB	
suites.polybench.2mm	openmp-parallel	13124	638.60µs	37.28ms	187.96µs	638.60µs	13.33MiB	1.19MiB	
suites.polybench.3mm	serial	4	11.72ms	11.72ms	None	N/A	N/A	1.72MiB	
suites.polybench.3mm	openmp-parallel	15288	503.06µs	27.71ms	265.82µs	503.43µs	15.01MiB	1.72MiB	
suites.polybench.adi	serial	4	71.42ms	71.42ms	None	N/A	N/A	1.22MiB	
suites.polybench.adi	openmp-parallel	566	17.36ms	1.05	5.26ms	17.36ms	6.20MiB	1.22MiB	
suites.polybench.atax	serial	4	148.06µs	142.25µs	5.75µs	N/A	N/A	1.16MiB	
suites.polybench.atax	openmp-parallel	69372	56.16µs	2.92ms	40.66µs	56.45µs	36.61MiB	1.16MiB	

GPULog-HIP

- Ported GPULog to GPULog-HIP
- AMD MI250 (dual chiplet) $\approx$ Nvidia A100
- **GPULog-HIP uses only one MI250 chiplet**
- **GPULog benefits from Nvidia RMM library**

Query	Dataset	GPULog (NVIDIA)		GPULog (AMD)	
		H100	A100	MI250	MI50
SG	fe_body	5.05	8.61	19.57	41.99
	BrightKite	3.42	6.79	14.00	30.05
	fe_sphere	2.36	4.64	8.48	19.426
CSPA	httpd	1.33	2.73	6.75	15.27
	linux	0.39	0.77	1.39	3.32
	postgres	1.27	2.68	6.79	14.55

GPULog and GPULog-HIP comparison on Nvidia and AMD GPUs

GPUJoin-SYCL

oneAPI Hackathon: CUDA to SYCL Migration

Hosted by Intel



- Ported GPUJoin to SYCL using SYCLomatic
- **Required bug fixing of the generated code**
- Scratch implementation of TC using SYCL
- Won Intel® Arc A770 Limited Edition GPU



Clean	Clean CUDA Project
Install	Install SYCLomatic
Convert	Convert CUDA code to SYCL
Check	Check the SYCL code and modify if necessary
Execute	Run in Intel Dev Cloud

CUDA-based GPUJoin to SYCL using SYCLomatic

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Compare energy profiles across Datalog engines

Engine	Join strategy	Data structure
MNMGDatalog	Radix hash join	GPU Hash table
INLJoin	Index nested loop join	GPU struct array
GPULog	Sorted hash join	Hash-Indexed Sorted Array
BJoin	Batch join with memory pooling	Hash-Indexed Sorted Array
cuDF	Dataframe's merge	Python's Dataframe



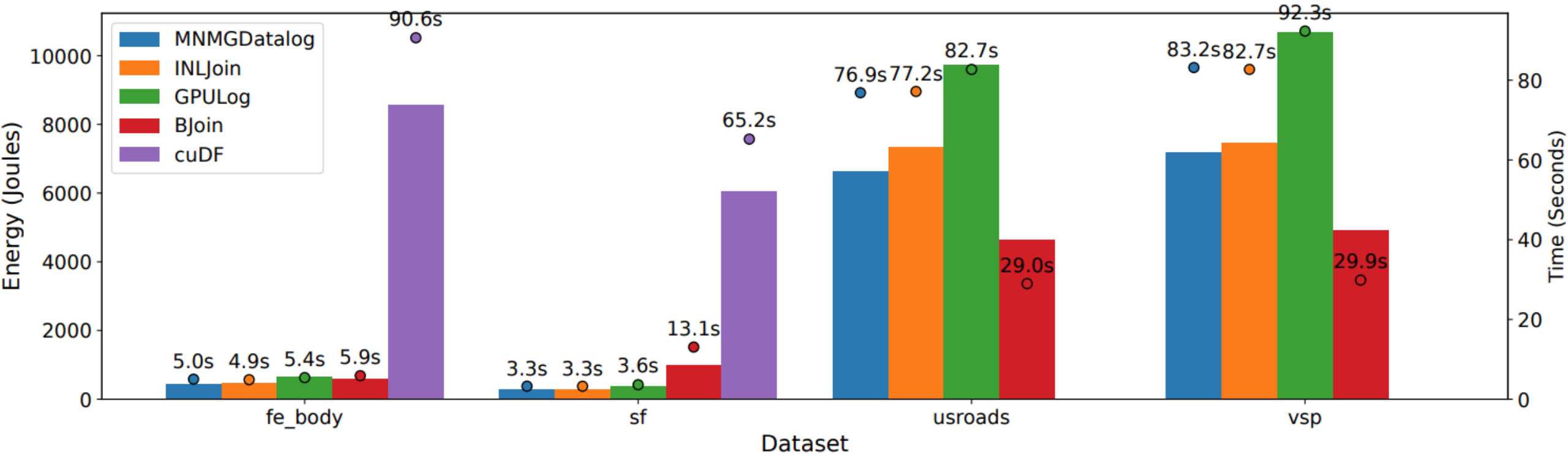
Polaris (Argonne National Lab)

Tools: nvidia-smi, nvml, NVIDIA Nsight systems, NVIDIA Nsight compute

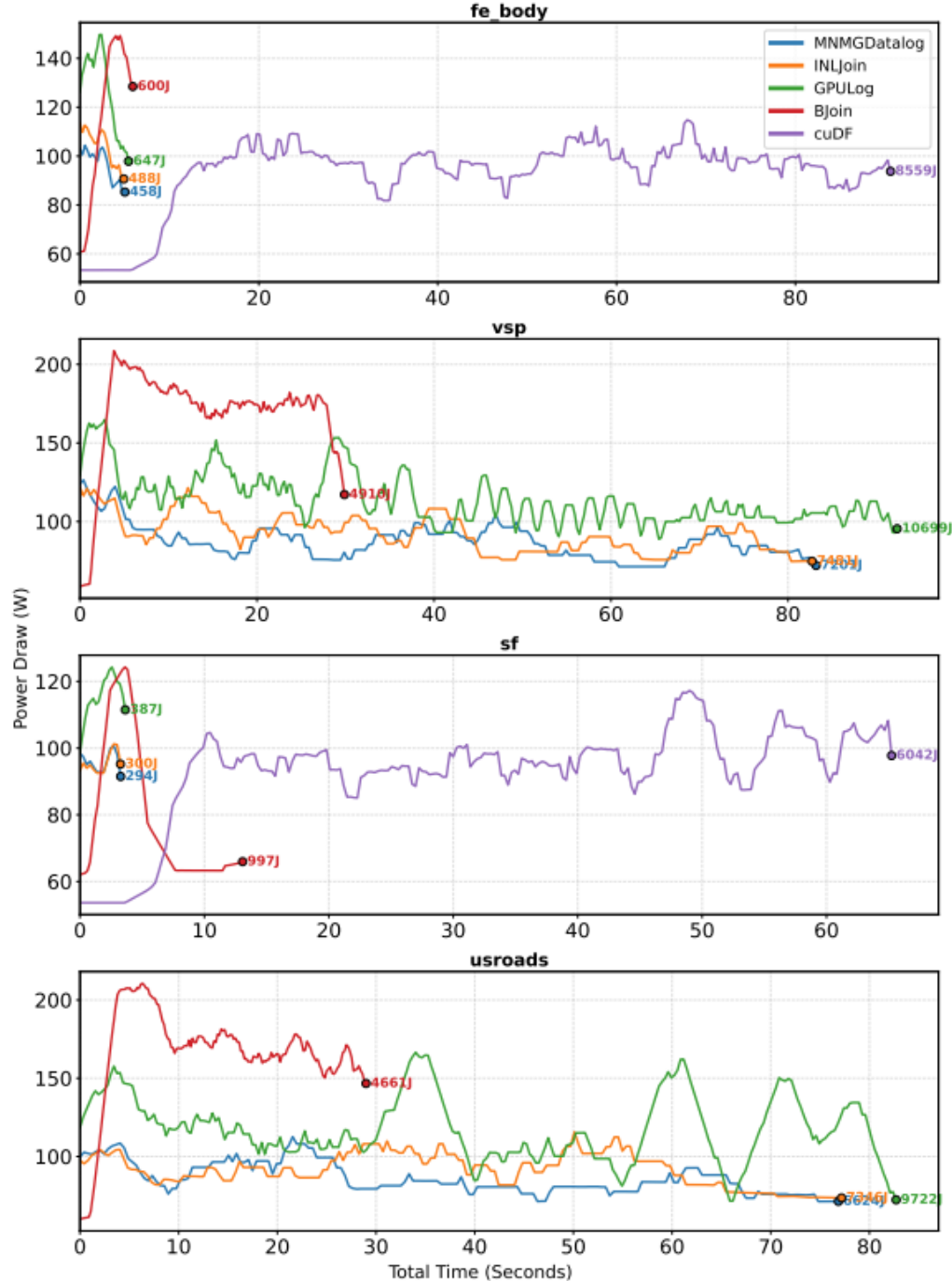
Energy profile: TC energy consumption (single-GPU)

$$E = \sum_{i=1}^N P_i \cdot \Delta t_i$$

- N is the total number of sampling intervals
- P_i is the power draw (in watts) measured at interval i
- Δt_i is the time (in seconds) between samples i and i - 1

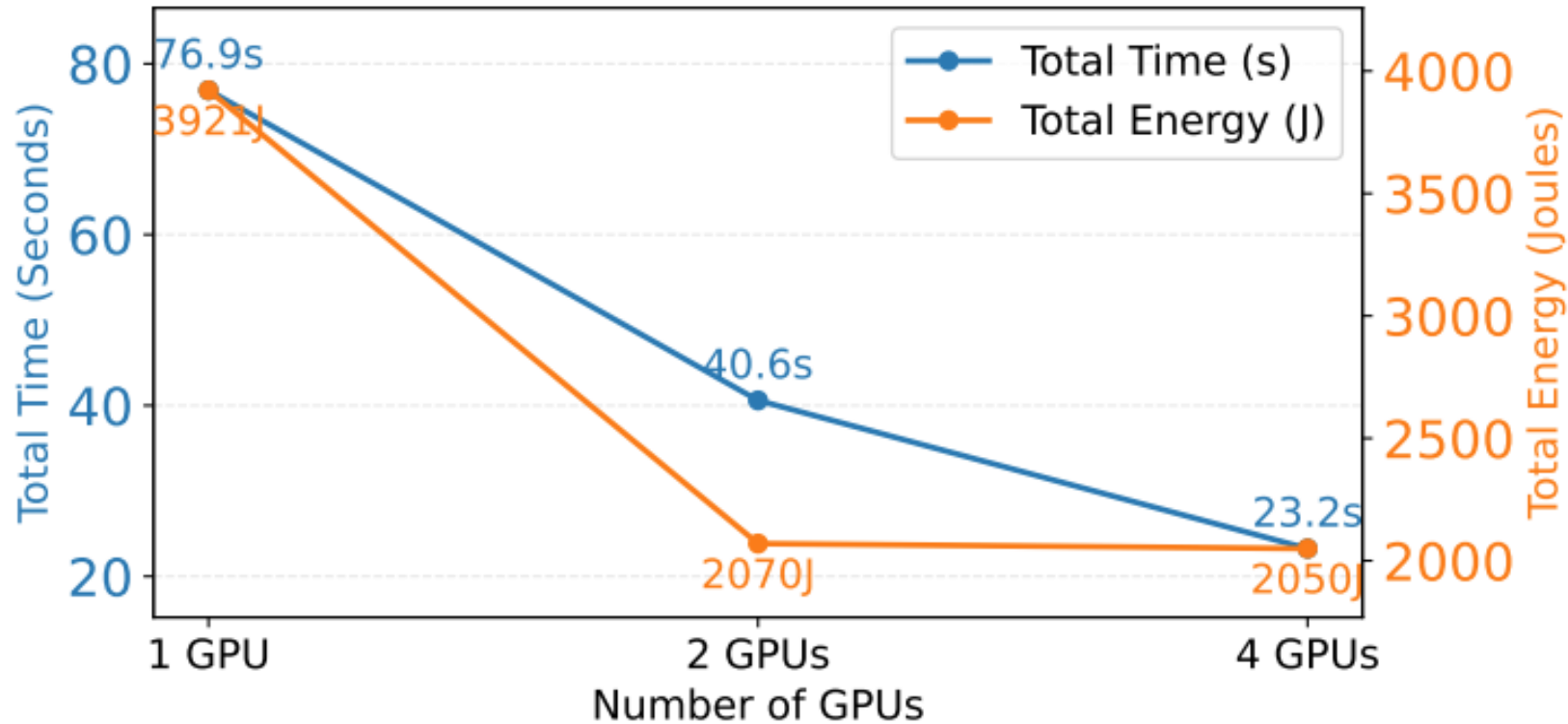


- BJoin's memory pooling draws more power, runs faster for larger iterations
- HISA-based joins (BJoin, GPULog) use more energy
- Simpler GPU friendly data structure (MNMGDatalog and INLJoin) draws less power but could run longer



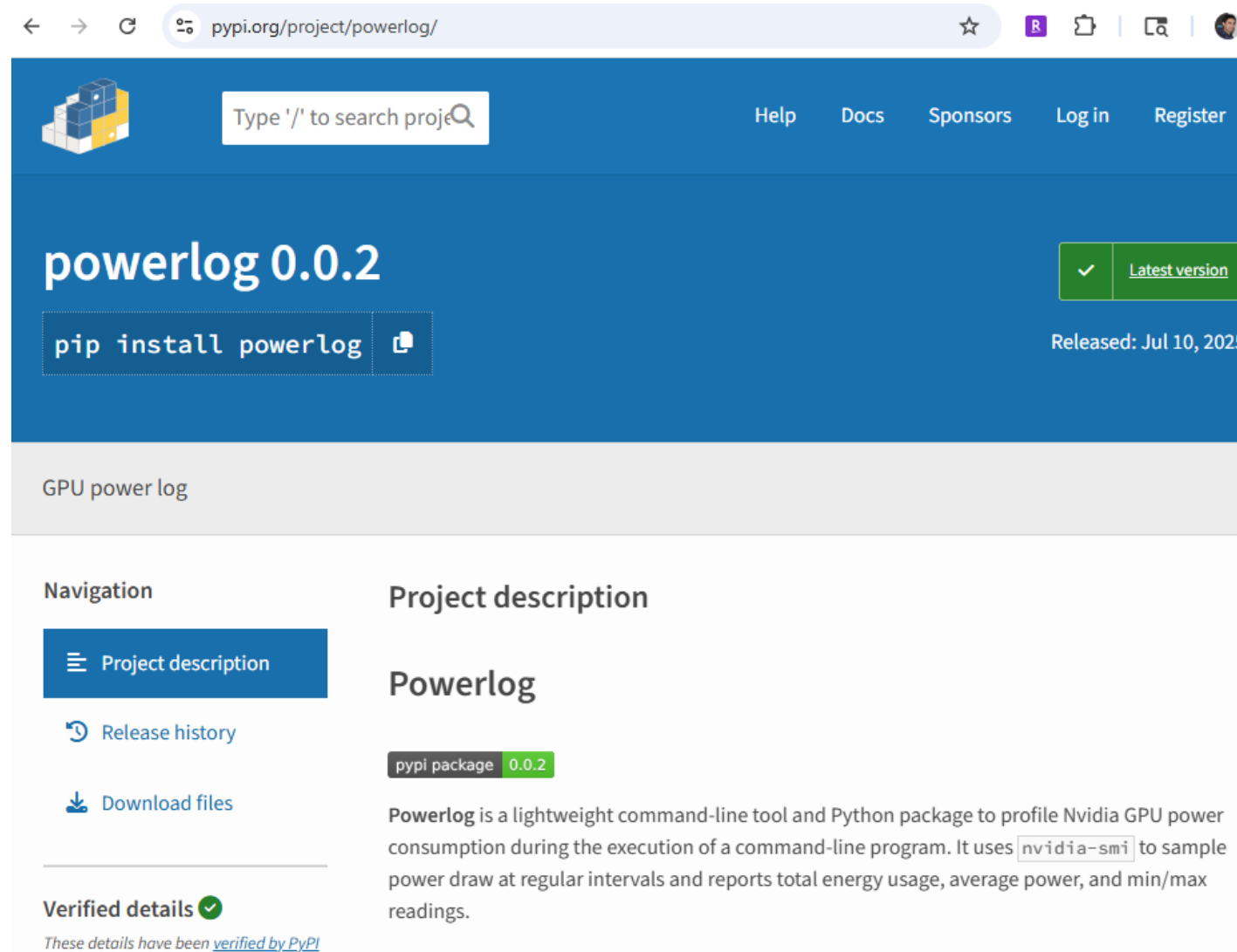
Power draw (TC)

Energy profile: TC energy consumption (multi-GPUs)



TC energy consumption (1 - 4 GPUs) for usroads dataset using MNMGDatalog

Powerlog CLI Tool for Energy Profiling GPU Applications

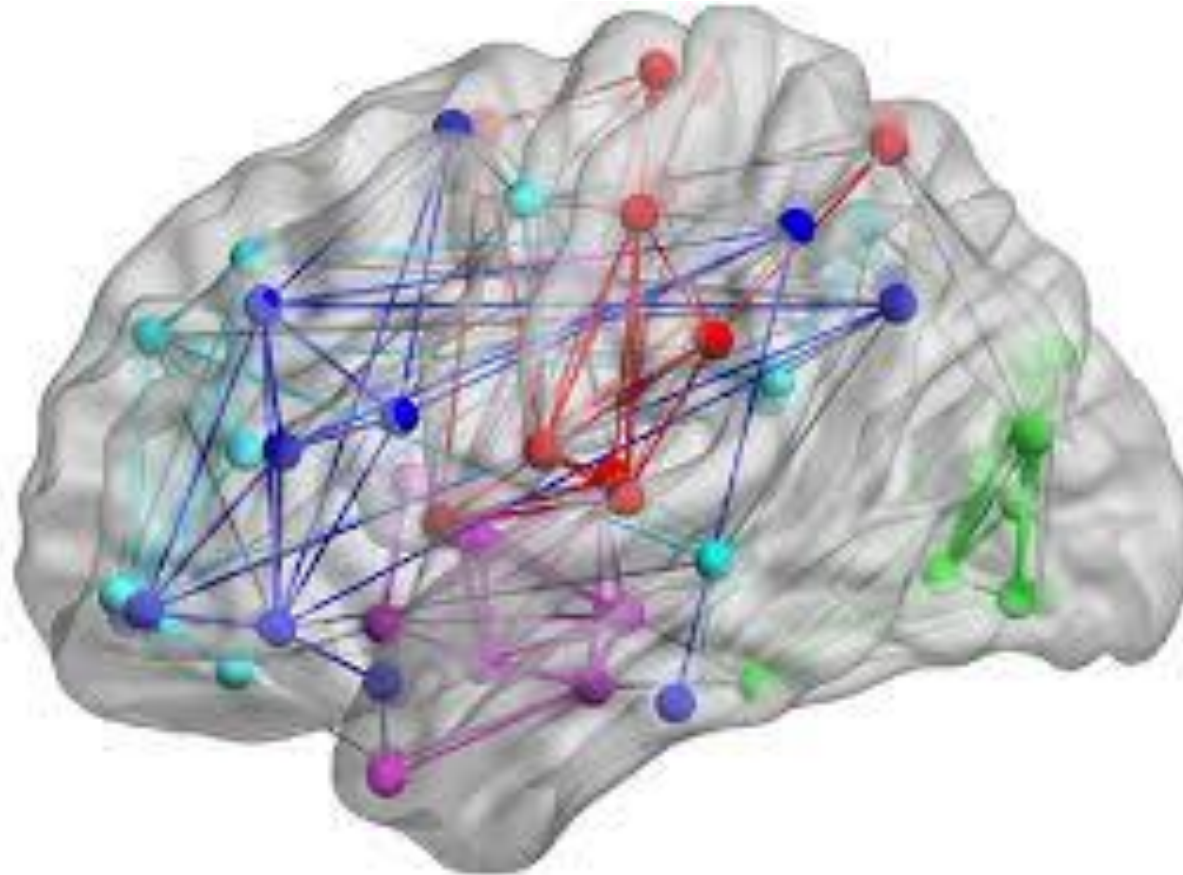


The screenshot shows the PyPI project page for **powerlog 0.0.2**. The page has a blue header with navigation links: Help, Docs, Sponsors, Log in, and Register. Below the header, the project name **powerlog 0.0.2** is displayed in large white text on a blue background. To the right of the project name is a green checkmark icon and a link to the **Latest version**. Below the project name is a code block containing the command `pip install powerlog` and a copy icon. To the right of the code block is the release date: **Released: Jul 10, 2025**. Below the blue header is a light gray section with the text **GPU power log**. The main content area is divided into two columns. The left column is titled **Navigation** and contains three links: **Project description** (highlighted with a blue background), **Release history**, and **Download files**. Below these links is a section titled **Verified details** with a green checkmark icon and the text *These details have been [verified by PyPI](#)*. The right column is titled **Project description** and contains the heading **Powerlog**. Below the heading is a badge that says **pypi package 0.0.2**. The description text reads: **Powerlog** is a lightweight command-line tool and Python package to profile Nvidia GPU power consumption during the execution of a command-line program. It uses `nvidia-smi` to sample power draw at regular intervals and reports total energy usage, average power, and min/max readings.

Roadmap

1. Introduction and motivation
2. GPU based Datalog engine development
3. Multi-node multi-GPU Datalog engine development
4. GPU portability of Datalog engines
5. Power analysis of Datalog engines
- 6. Exploration of high-dimensional scientific analytics**
7. Future work
8. Conclusion

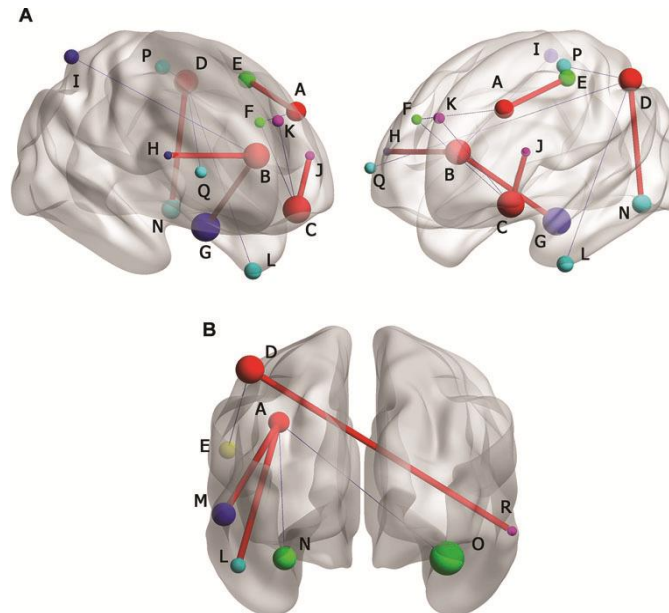
What is a brain network?



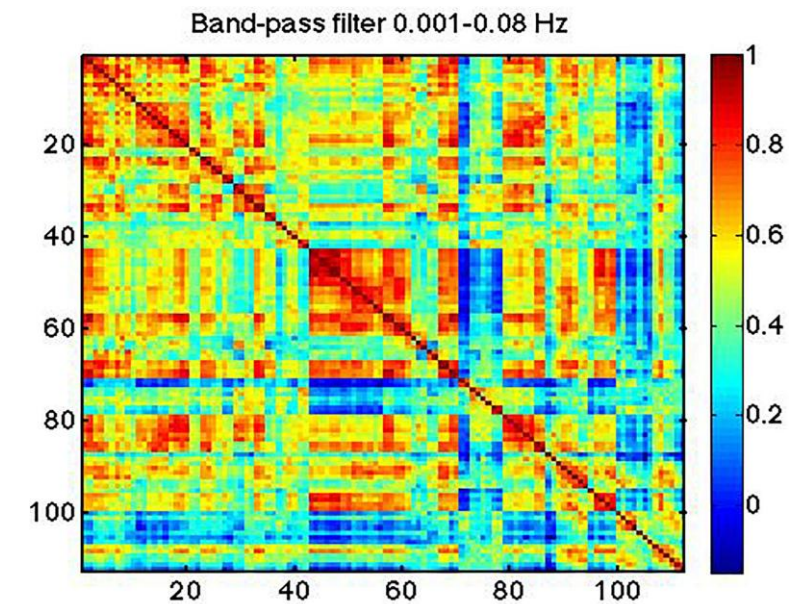
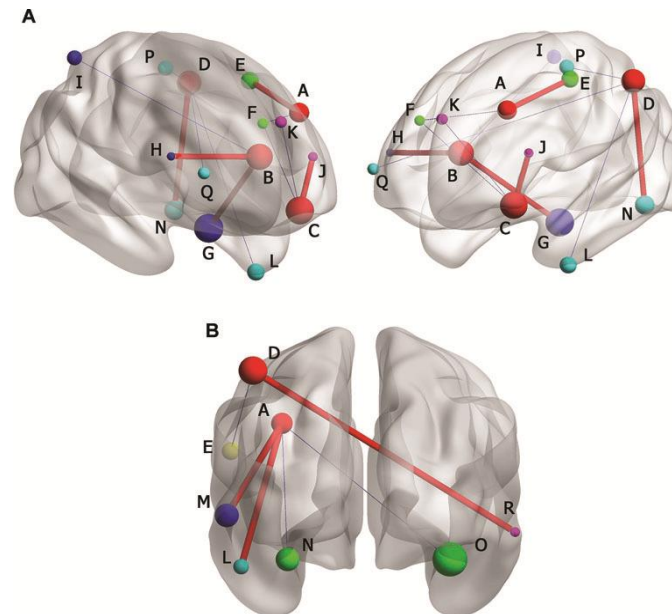
How to construct brain network?



How to construct brain network?



How to construct brain network?



Multi-site brain image analysis



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Multi-site rs-fMRI studies allow gathering
larger sample sizes

Multi-site brain image analysis (continue)



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But different **scanners** and **sampling periods**
negates the advantages of larger sample sizes

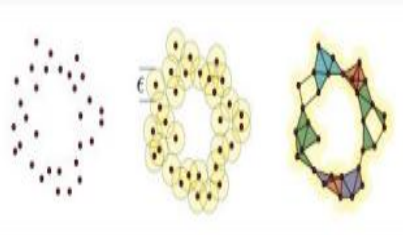
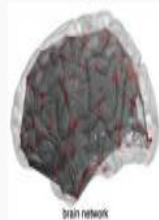
Convert fMRI
image to FCN



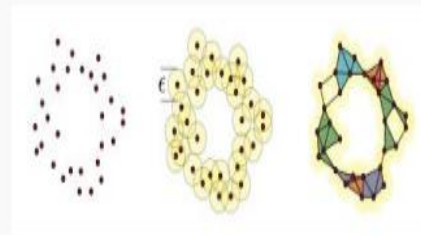
Apply Topological
Data Analysis on
FCN



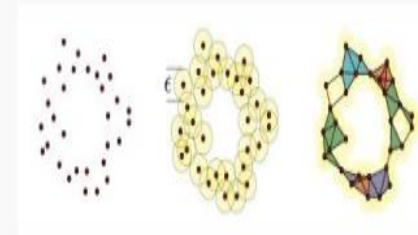
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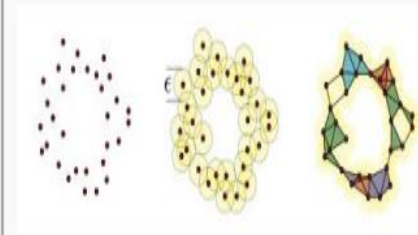
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Topological Data Analysis (TDA)

8
9

Studies shape of data like connected components, loops, voids



Main tool: persistent homology, adapts homology to point cloud



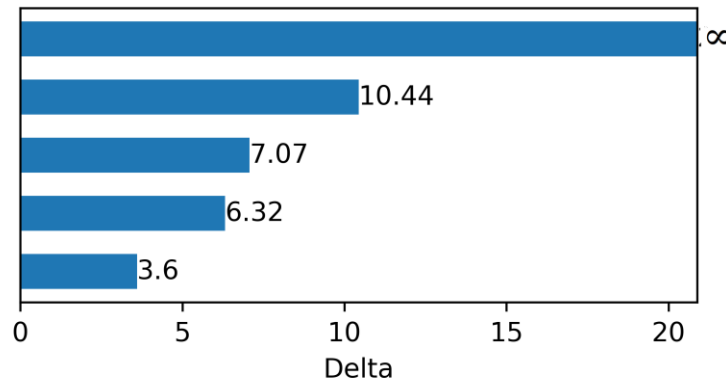
Analyzes social networks, brain connectivity, internet

Persistent Barcodes using Connected Component

Persistent Homology tracks **birth and death of topological features** in the graph

0-dim barcodes: each bar shows when each features (**connected component**) born and die

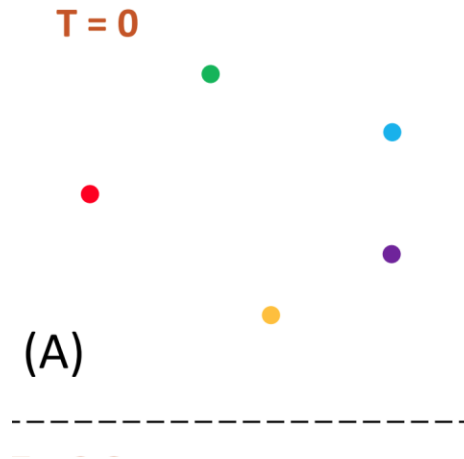
Persistent barcodes (0 dimensional)



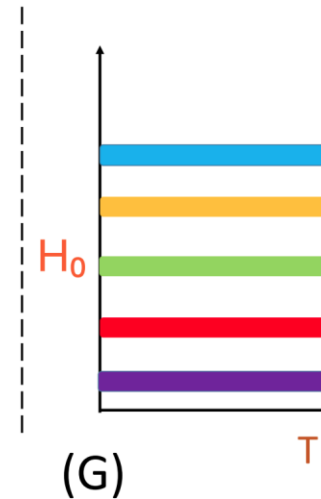
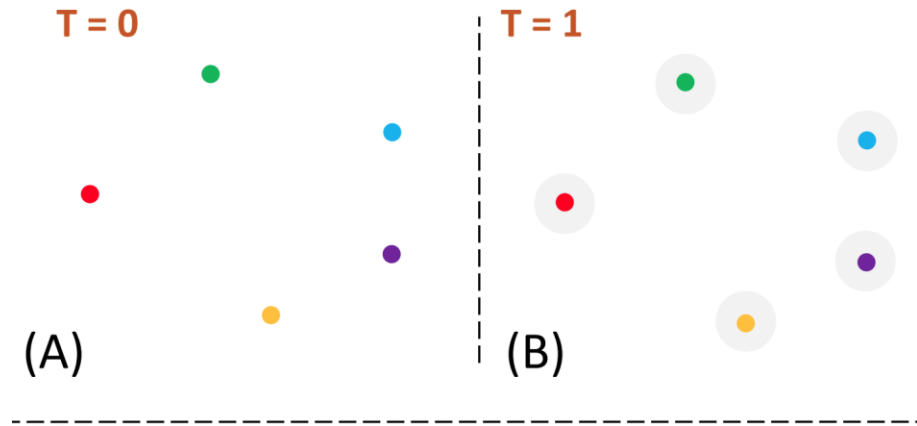
Adjacency matrix

0.00	8.54	12.36	7.07	10.44
8.54	0.00	20.88	3.60	18.00
12.36	20.88	0.00	18.78	6.32
7.07	3.60	18.78	0.00	15.13
10.44	18.00	6.32	15.13	0.00

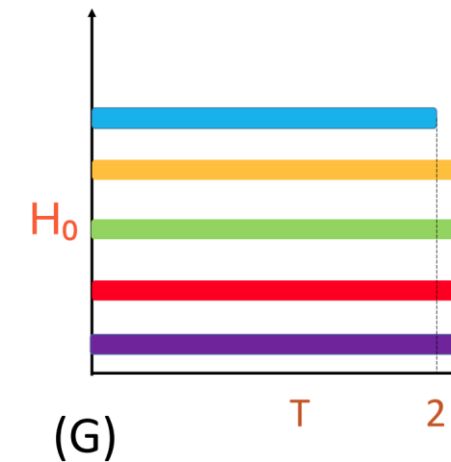
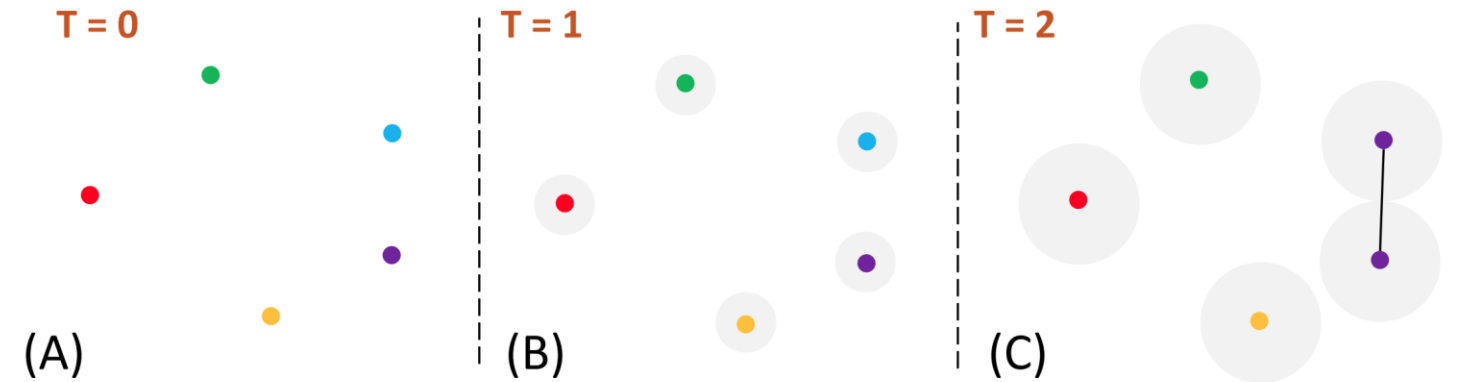
Computing Persistent Homology (Point Cloud)

9
1

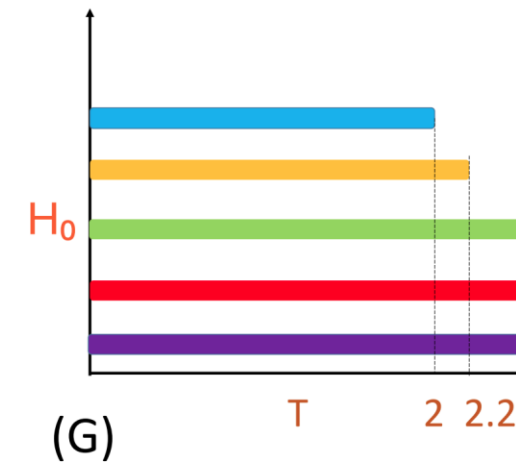
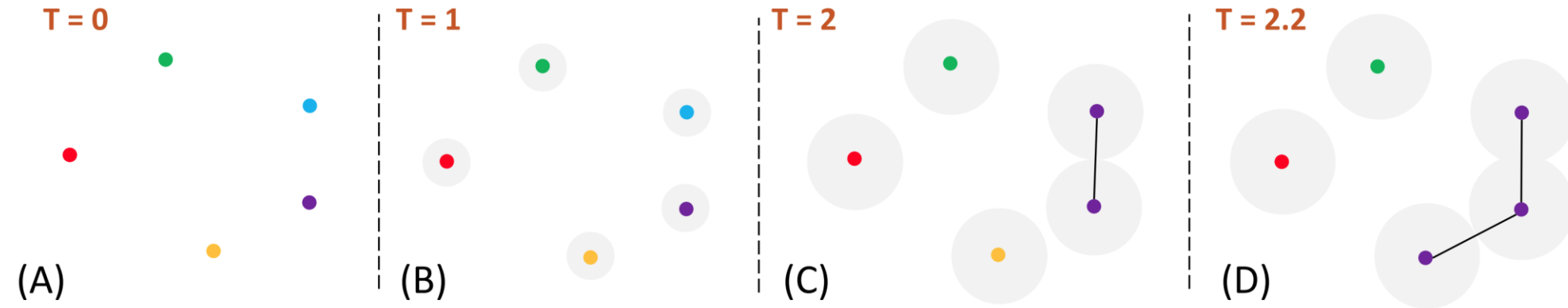
Computing Persistent Homology (Point Cloud)

9
2

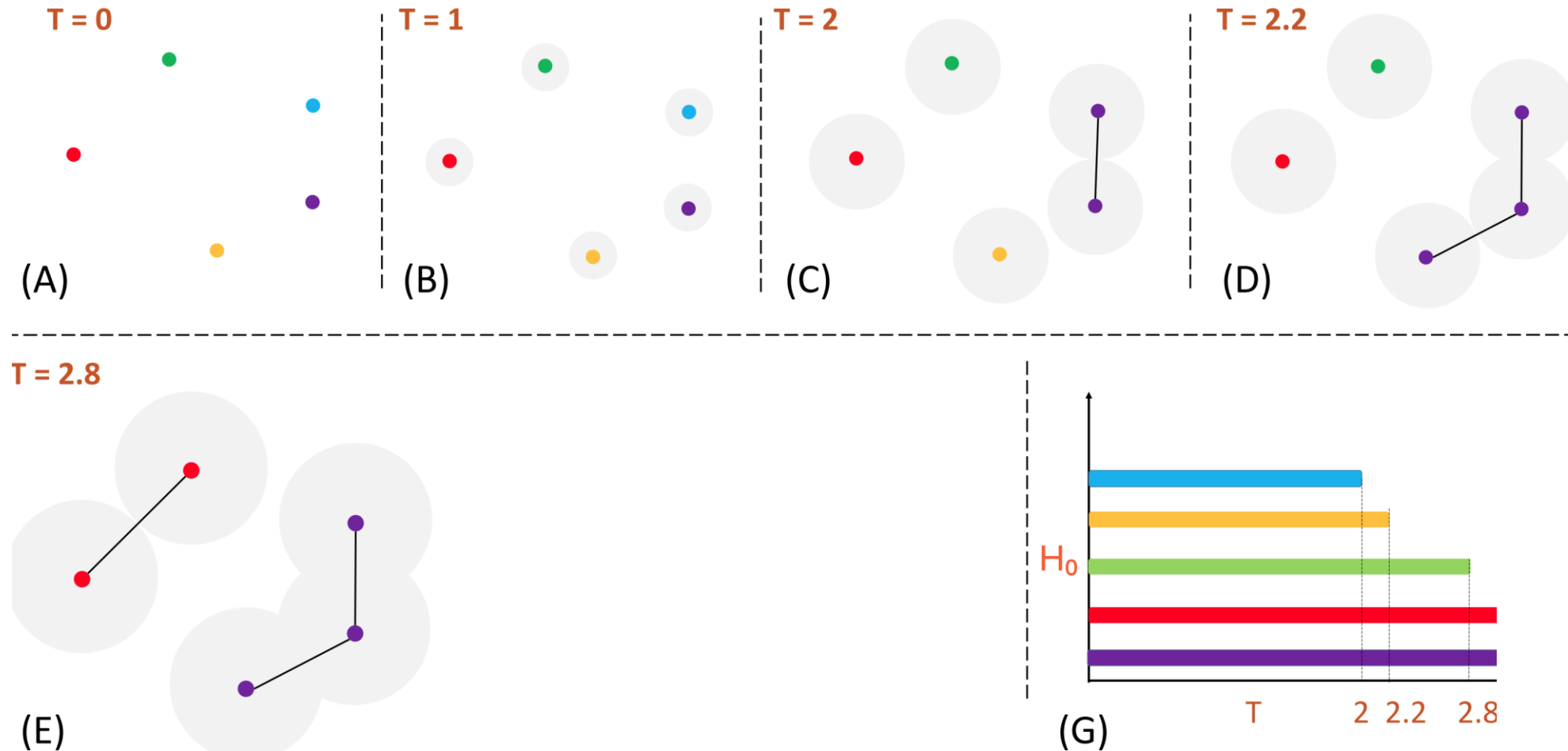
Computing Persistent Homology (Point Cloud)

9
3

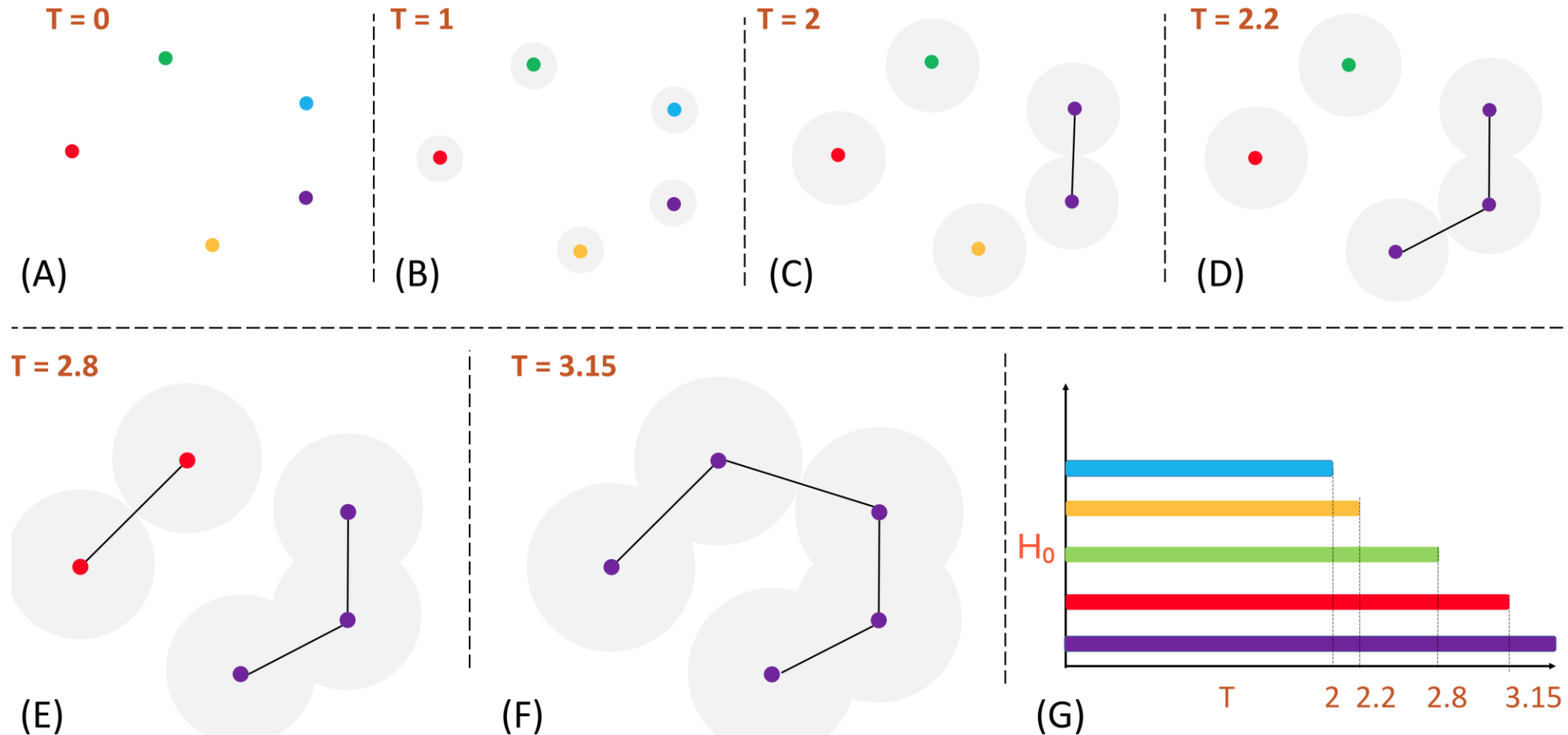
Computing Persistent Homology (Point Cloud)

9
4

Computing Persistent Homology (Point Cloud)

9
5

Computing Persistent Homology (Point Cloud)

9
6

PH for High-Dimensional Data Analysis

9
7

```
graph TD; A[Verify] --> B[Demonstrate]; B --> C[Remove];
```

Verify

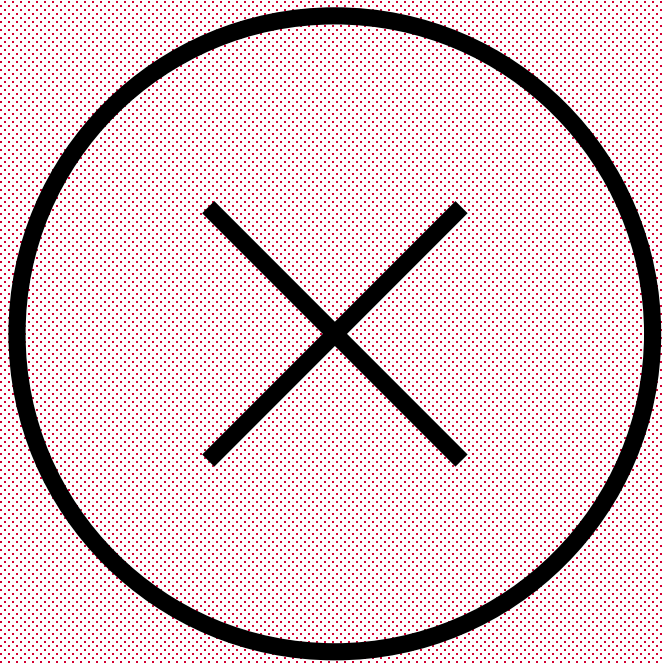
PH of rs-fMRI can represent topological features of FCNs

Demonstrate

FCN networks are statistically similar despite variations in TRs

Remove

Non-neural variability in multi-site brain networks



Limitations

Imperative and complex high-dimensional data analysis pipelines

Declarative Topological Data Analysis

1

Use Datalog to simplify persistent barcodes computations

2

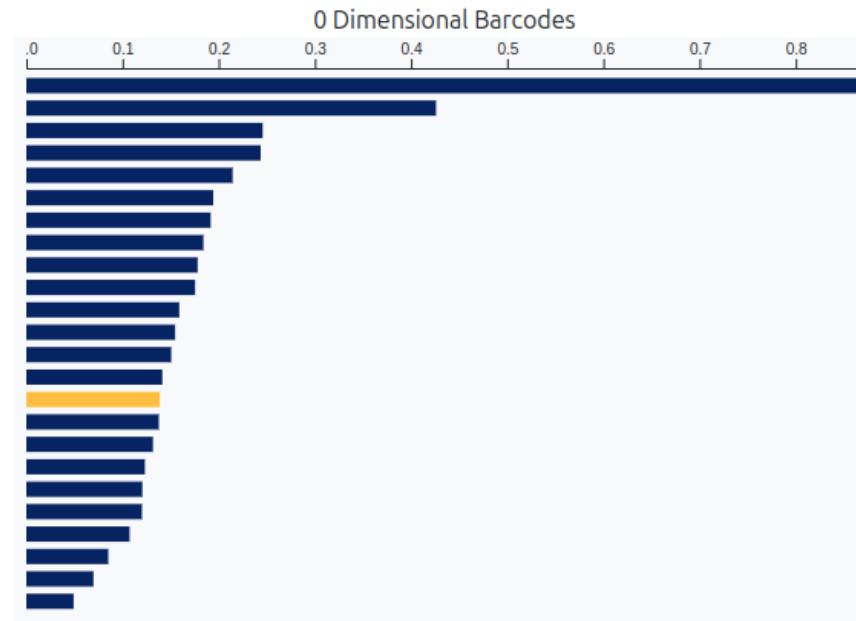
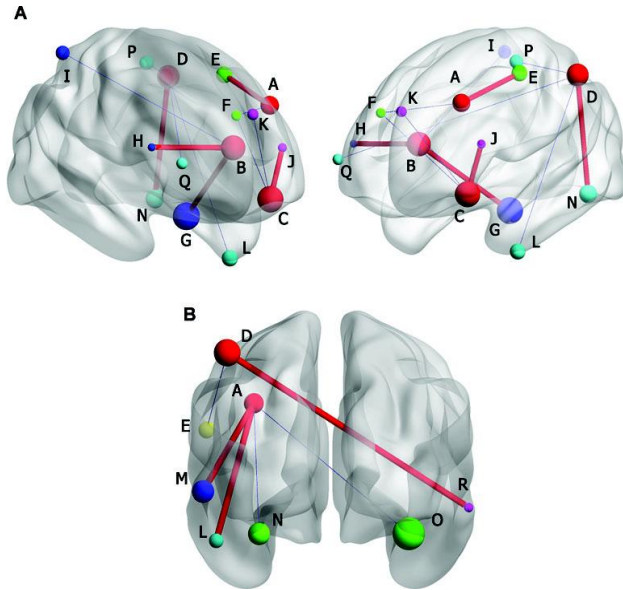
Explore potential applications of using declarative TDA

3

Scale TDA methods to large datasets in HPC environment

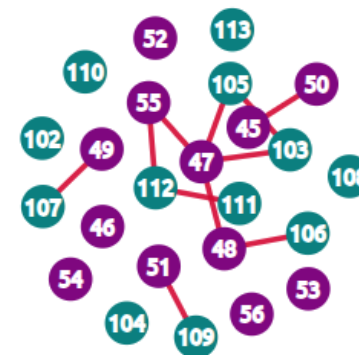
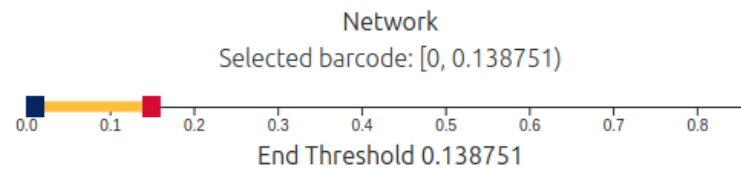
Persistent Barcodes using Declarative Analytics

(Loaded `data/ph_data/subject_1_mx645.txt` dataset)



← Construct a filtration

← Compute persistence of connected components at each threshold

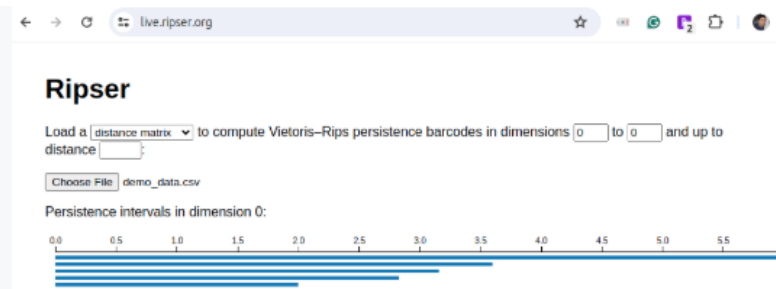


← Build graph based on the selected threshold

Persistent Barcodes using Declarative Analytics

- Use Datalog for efficient connected components calculation
- Successfully implemented 0-dimensional barcodes using Soufflé
- Extend barcodes computation to the **SLOG** engine for scalable multi-node HPC environments

```
root@c4a52625bf20:/workspace# souffle bars.dl -F./data
-----
barcodes
-----
0      2      1
0      2.8300000000000001      1
0      3.1600000000000001      1
0      3.6000000000000001      1
0      6.0000000000000001      1
-----
```



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Future/ongoing work

Enhance
MNMGDatalog
implementation



Extend
declarative
applications



Balance energy
and
performance

Enhance MNMGDatalog Implementation

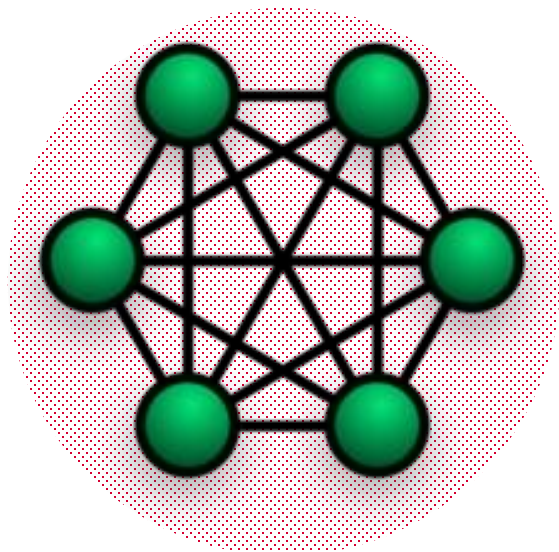
Add per-iteration checkpoint/restart capability



Develop HIP and SYCL version for MNMGDatalog

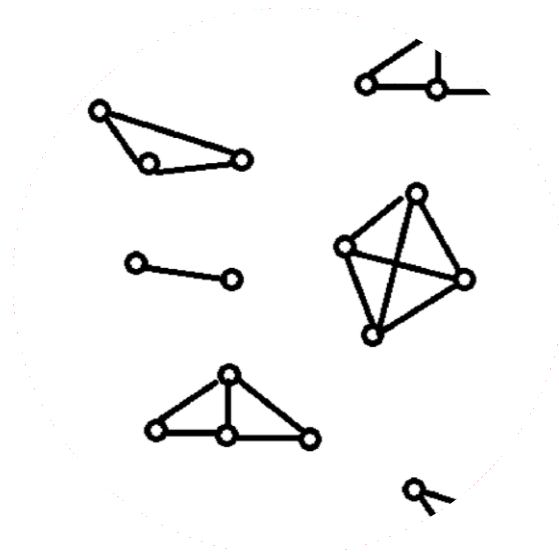


MNMGDataLog for High-dimensional Brain Imaging



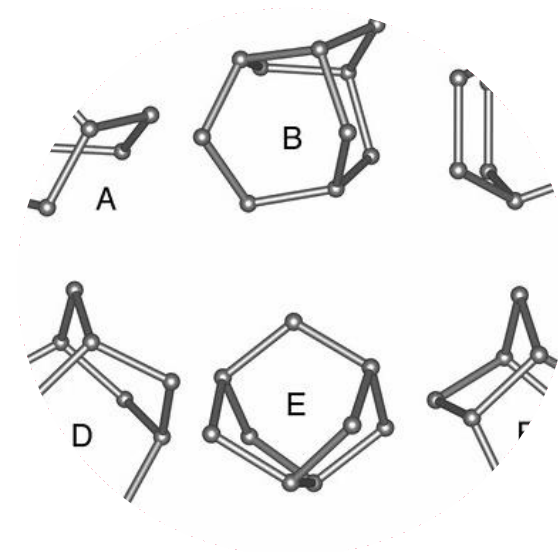
MNMGDataLog for fully connected network

Use high-level expressive query to handle dense connectivity graphs from brain imaging data



Declarative modeling of topological structure

Encode high-dimensional relationships using recursive rules over pairwise similarities

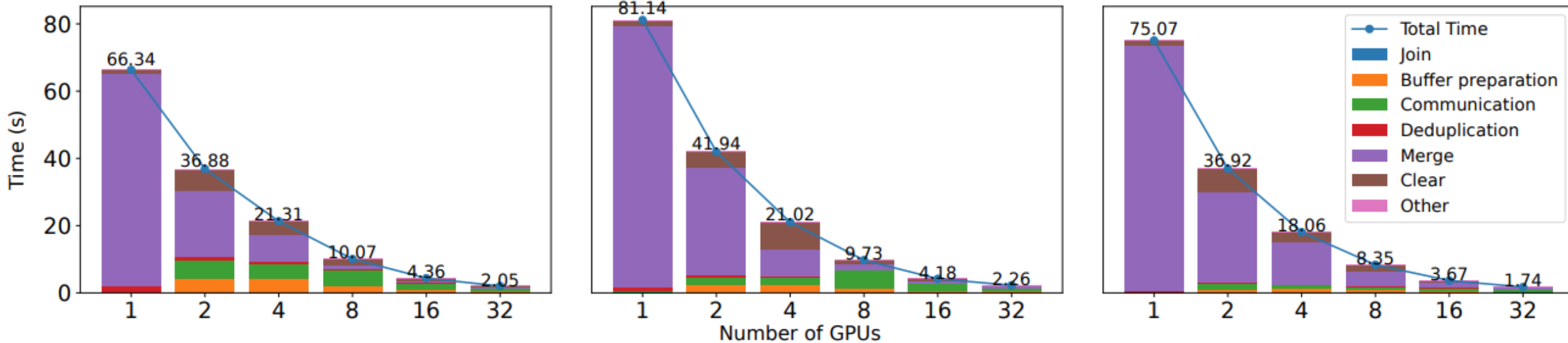


Topological pattern matching

Discover persistent structural motifs across subjects using distributed recursive evaluation

Power-aware Declarative Analytics

Compare energy profiles across Datalog engines



Design runtime tools to correlate iterative operations with power draw

Develop AI-driven models to optimize query plans for energy efficiency

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Conclusion

Declarative analytics at scale



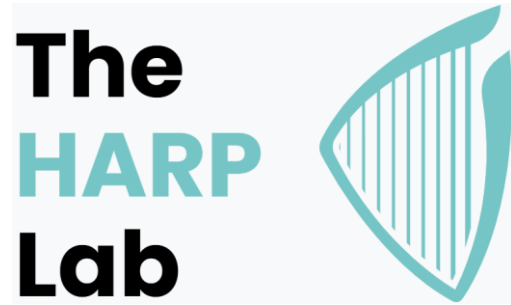
GPU Portability and power analysis



Scientific analytics



Acknowledgements





Thank You

ashov@uic.edu

<https://arshovon.com/>



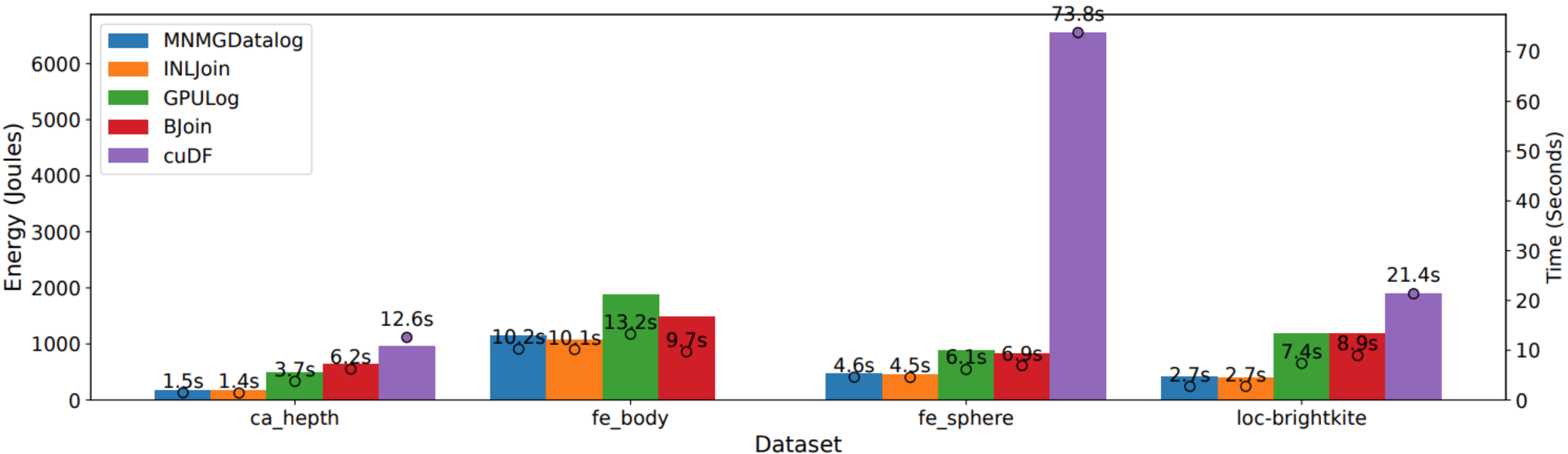
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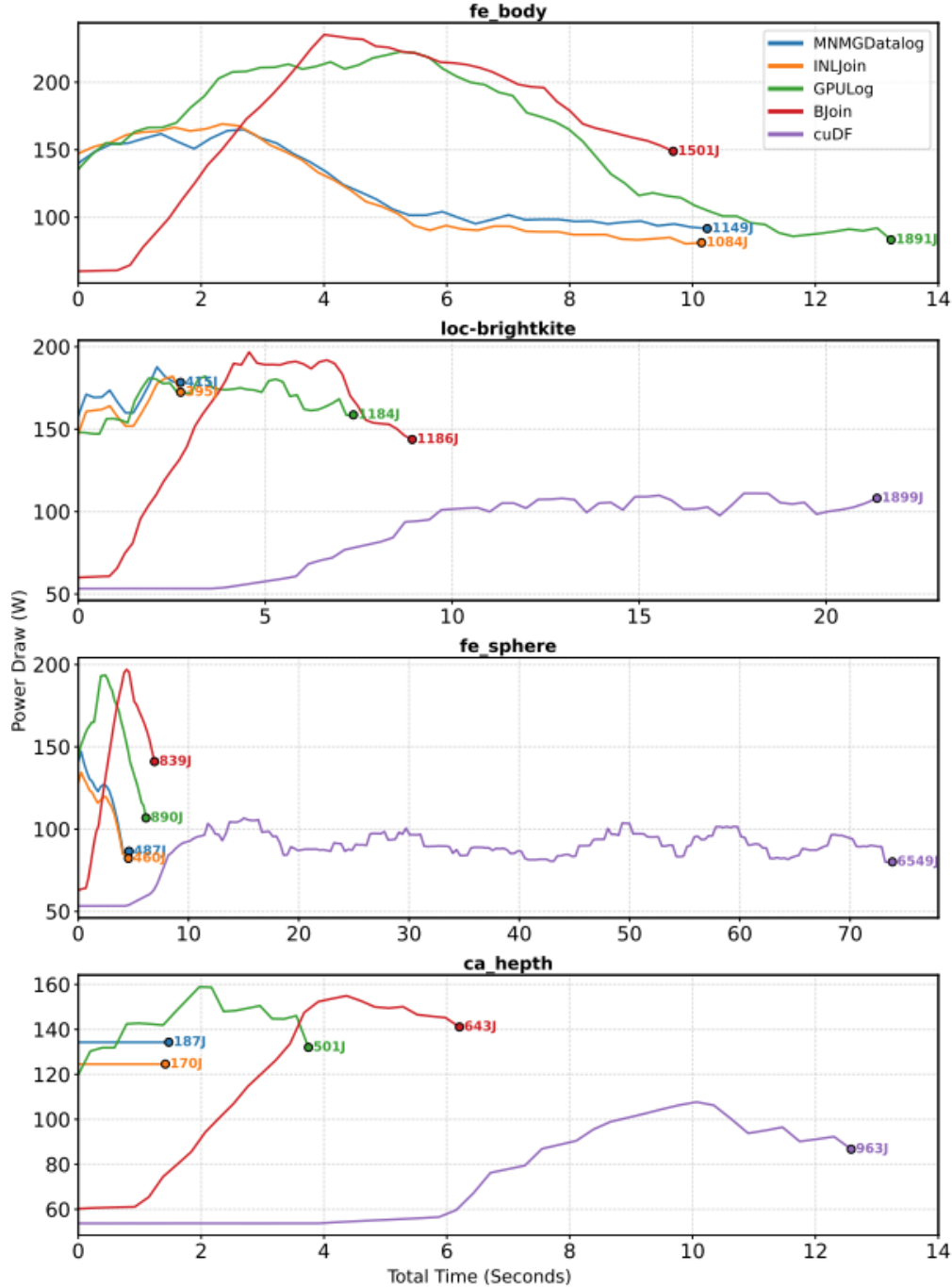
Appendix

Energy profile: SG energy consumption (single-GPU)

$$E = \sum_{i=1}^N P_i \cdot \Delta t_i$$

- N is the total number of sampling intervals
- P_i is the power draw (in watts) measured at interval i
- Δt_i is the time (in seconds) between samples i and i - 1





Power draw (SG)