

# 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)



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**

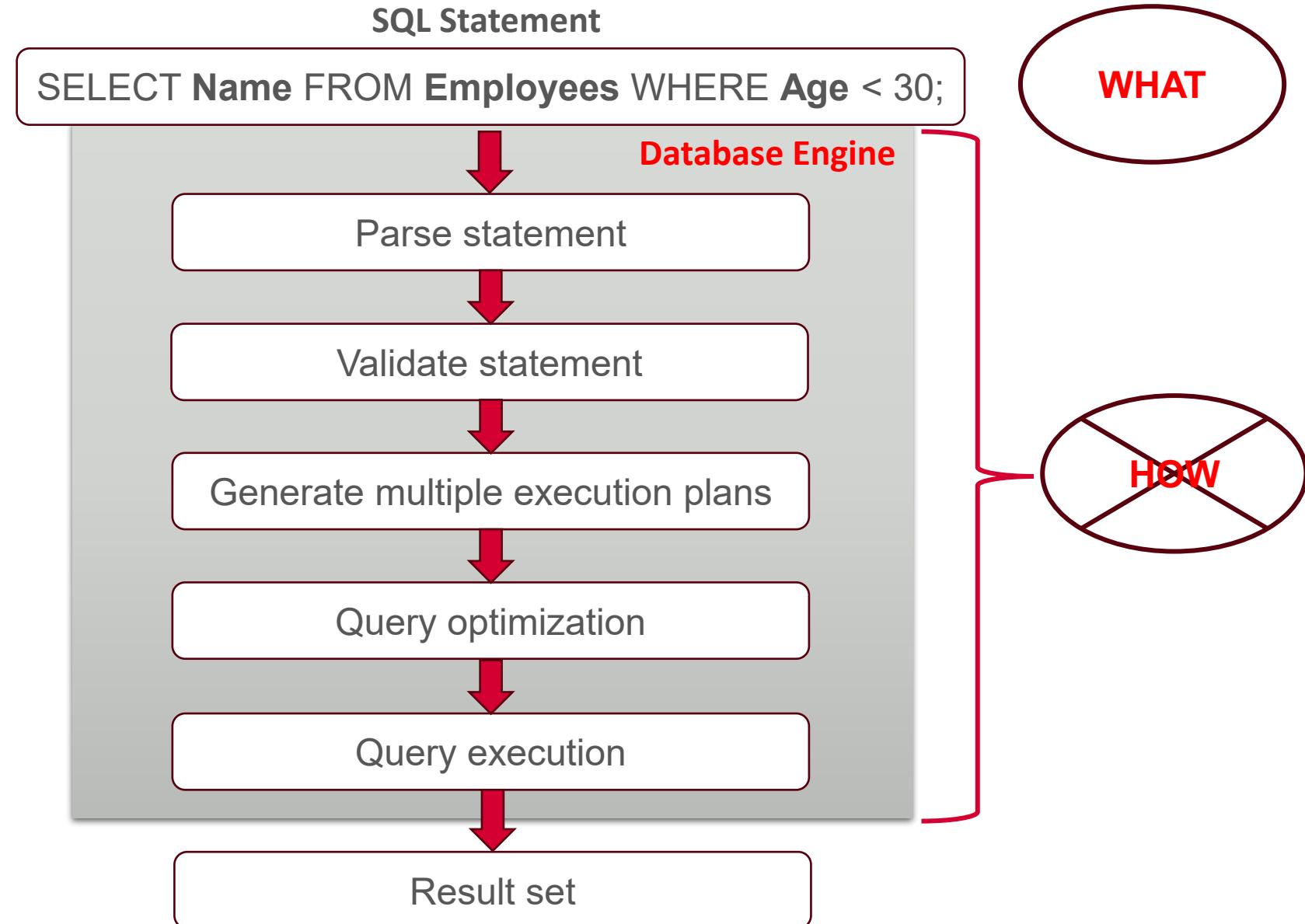
EmpID	Name	Email	Age	ManagerID
0	Alice	alice@example.com	56	Null
1	Bob	bob@example.com	33	0
2	Eve	eve@example.com	26	0
3	Carol	carol@example.com	34	2
4	Dave	dave@example.com	22	3

Find all employees who are less than 30 years old

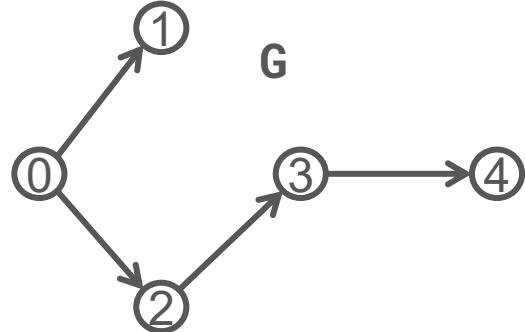


SELECT Name FROM Employees WHERE Age < 30;

# How is the query processed?

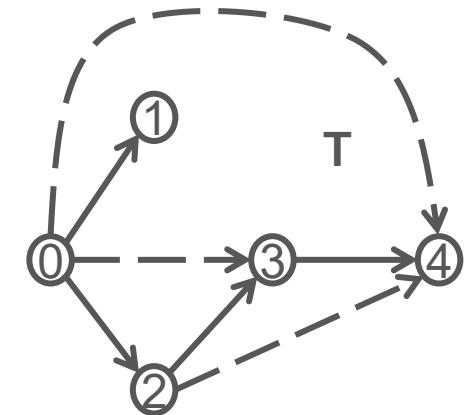


# Recursive Queries using SQL



**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



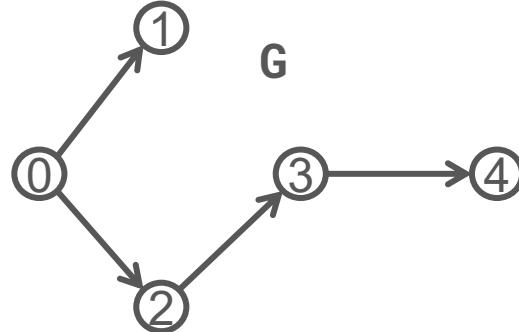
Find all managers, including direct and indirect managers.

```

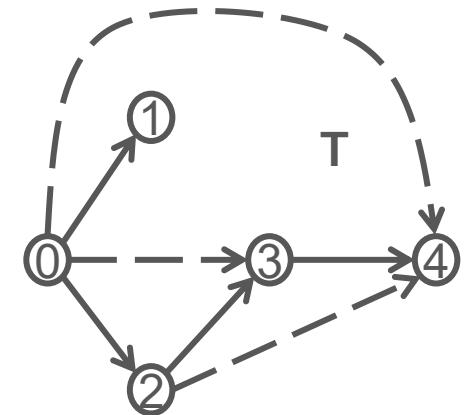
-- Recursive CTE to find all managers (direct and indirect)
WITH RECURSIVE ManagerHierarchy AS (
    -- Base case: direct manager relationship
    SELECT EmpID AS EmployeeID, ManagerID
    FROM Employees WHERE ManagerID IS NOT NULL
    UNION
    -- Recursive case: find manager of managers recursively
    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
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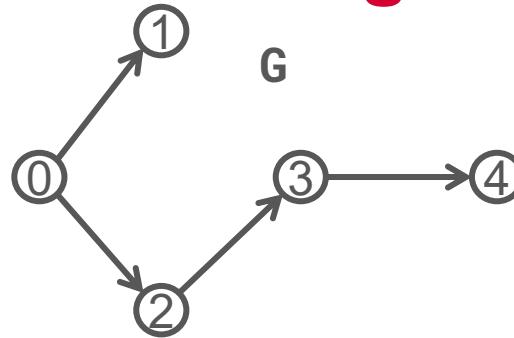
Find all managers, including direct and indirect managers.

## Finding recursive relationships using Datalog

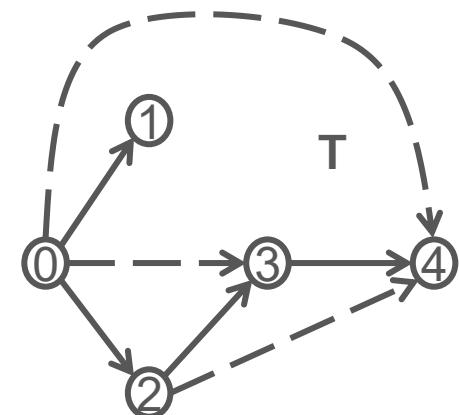
```
// Base case: direct manager manages employee
Manager(mID, eID) :- Employees(eID, mID), mID != NULL.
```

```
// 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: all managers

```

SELECT ManagerID, EmployeeID
FROM Employees
WHERE ManagerID IS NOT NULL
UNION ALL
  
```

Datalog offers greater expressiveness and natural syntax for recursive queries

-- Recursive case: include managers of managers

```

SELECT ManagerID, EmployeeID, ManagerManagerID
FROM ManagerHierarchy
  
```

FROM

JOIN

WH

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

)

```

SELECT DISTINCT ManagerID, EmployeeID FROM ManagerHierarchy
ORDER BY ManagerID;
  
```

# 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

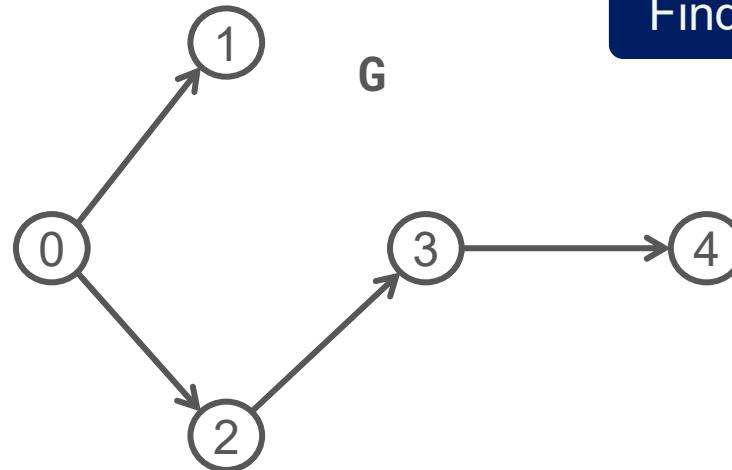
Projection

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.

```

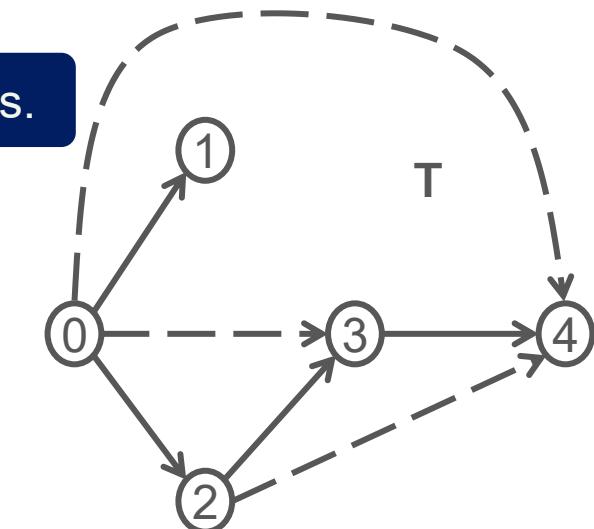
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// Recursive case: indirect managers
Manager(mID, eID) :- Manager(mID, nID), Employees(eID, nID).
  
```

0	1
0	2
2	3
3	4

Transitive closure computation of a graph

```

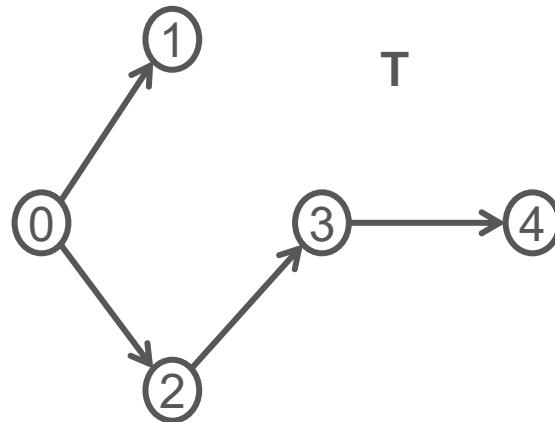
tc(X, Y) :- edges(X, Y).
tc(X, Z) :- tc(X, Y), edges(Y, Z).
  
```



0	1
0	2
2	3
3	4
0	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

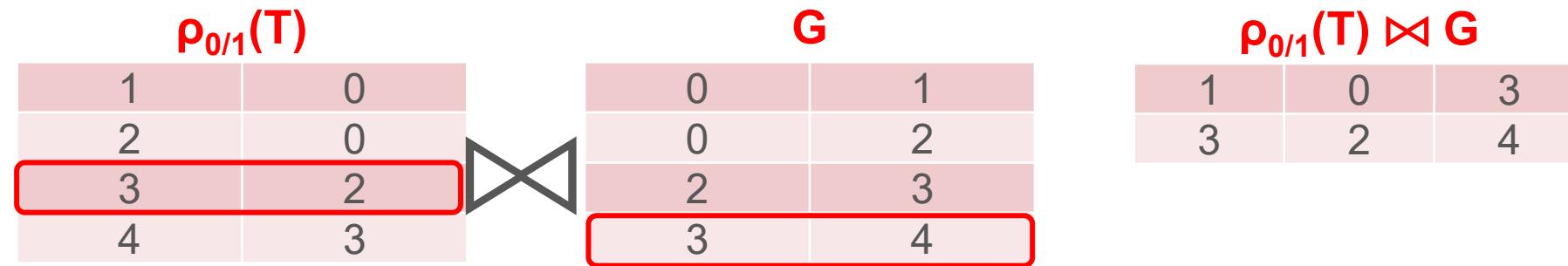
# Transitive Closure: Iterations 1

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



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



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

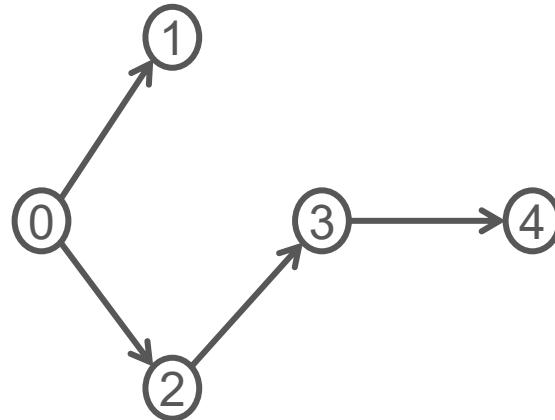
$\rho_{0/1}(T)$		$G$	
1	0	0	1
2	0	0	2
3	2	2	3
4	3	3	4



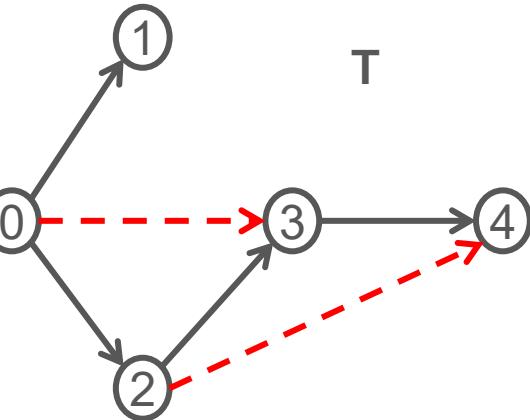
$\rho_{0/1}(T) \bowtie G$			$\Pi_{1,2}(\rho_{0/1}(T) \bowtie G)$	
1	0	3	0	3
3	2	4	2	4

# Transitive Closure: Iterations 1

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



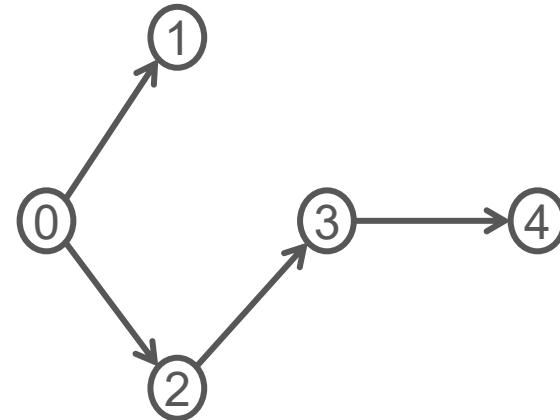
0	1
0	2
2	3
3	4



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

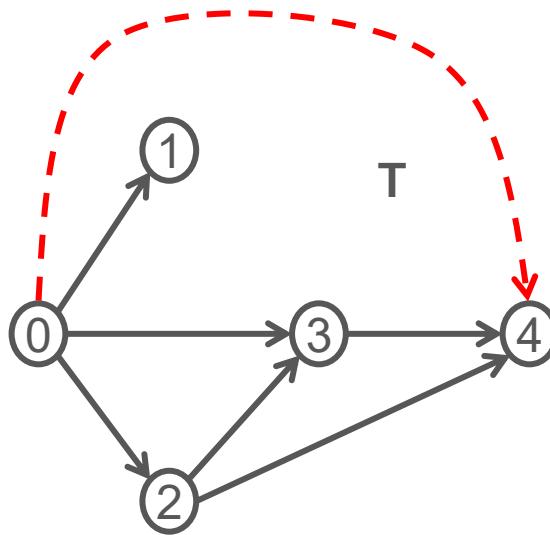
# Transitive Closure: Iterations 2

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



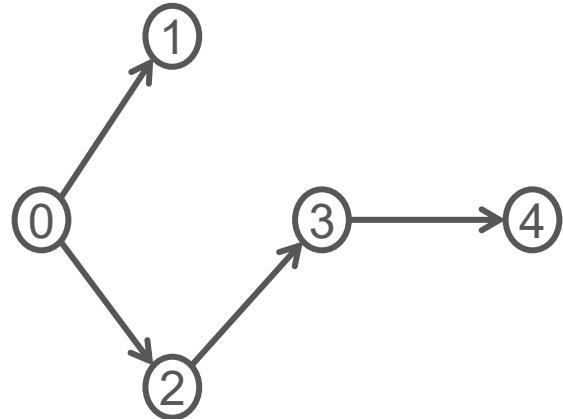
0	1
0	2
2	3
3	4

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

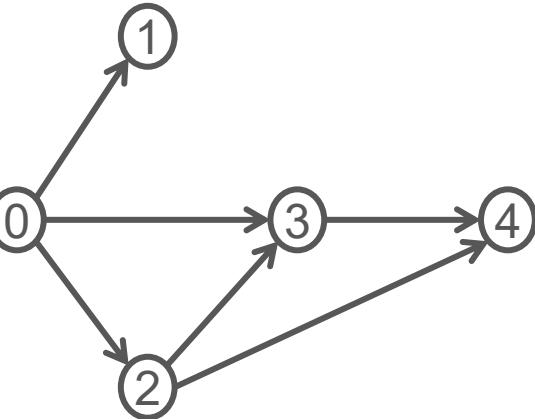


0	1
0	2
2	3
3	4
0	3
2	4
0	4

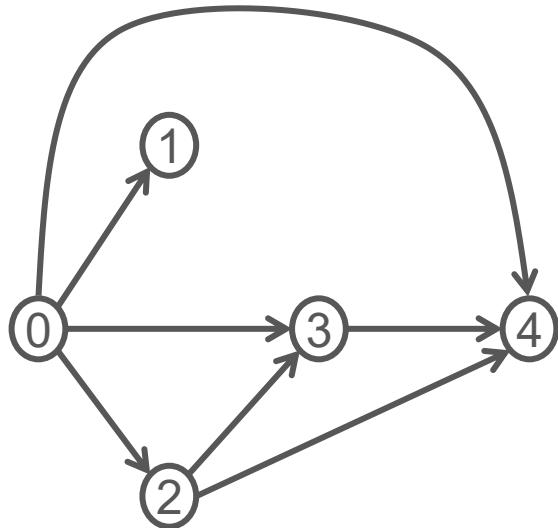
# Transitive Closure: Iterations 3



0	1
0	2
2	3
3	4

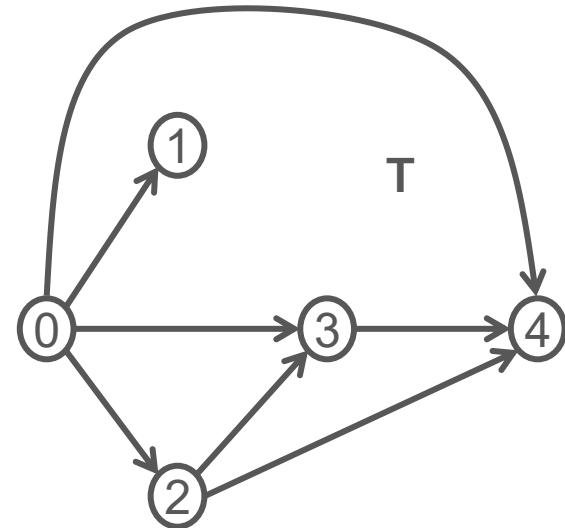


0	1
0	2
2	3
3	4
0	3
2	4



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

Fixed-point reached



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

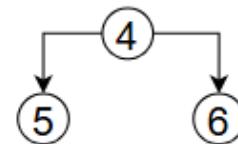
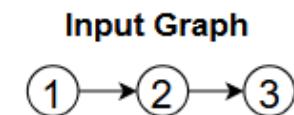
```
cc(X, X) :- edges(X, _).
cc(Y, $MIN(Z)) :- cc(Y, Z), edges(X, Y).
```

### 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).
```

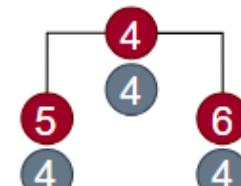
### Same generation

```
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.
```



Edge	CC (Delta, Full)
1	1
2	2
2	3
3	3
4	4
4	5
5	5
5	4
4	6
6	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 Selected Papers. 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 Conference on Management of Data (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

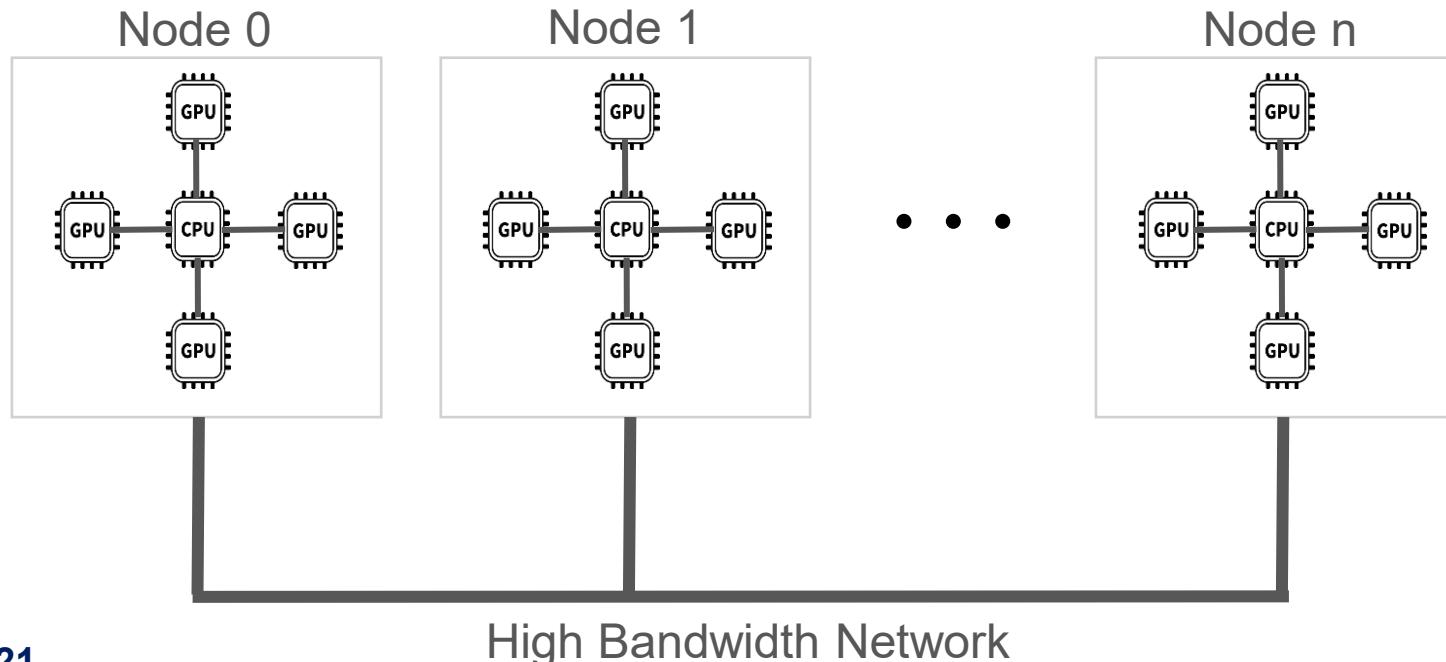
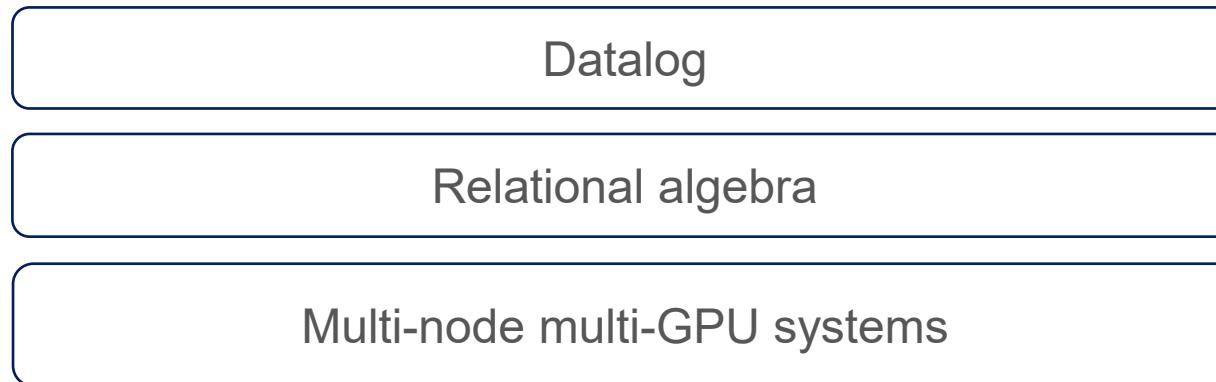
Framework

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



High performance  
implementation for  
highly expressive  
language

# Selected publications

## Conference Proceedings

Towards iterative relational algebra on the {GPU}

Acceptance rate: 19%



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

Acceptance rate: 44%



Optimizing Datalog for the GPU

Acceptance rate: 15%



Multi-node multi-GPU Datalog

Acceptance rate: 23%



## 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



UALCAN: An update to the integrated cancer data analysis platform

Impact factor: 6.3



## Workshop Papers

Accelerating Datalog applications with cuDF

Acceptance rate: 42%



Scalable, interactive and hierarchical visualization of virus taxonomic data



## Submission

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



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# GPU based Datalog engine development

## Off-the-shelf

Started with CPU and GPU based Python libraries

Compare GPU capabilities for iterative join operations

## 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

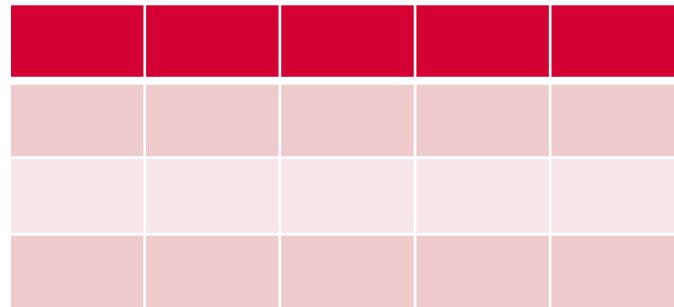
IA<sup>3</sup> 2022



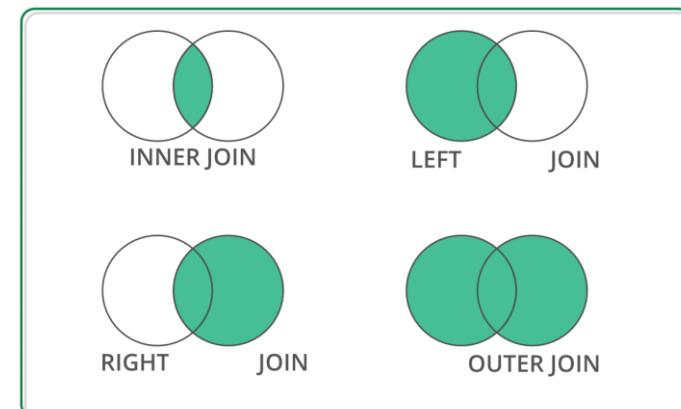
# Off-the-shelf Data Structure for Join Operation



DataFrame: tabular data structure



DataFrame has RA primitives APIs

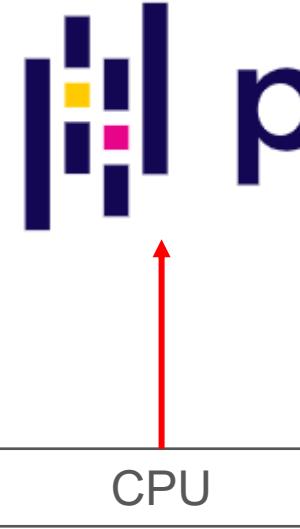


• Reback, J., McKinney, W., Van Den Bossche, J., Augspurger, T., Cloud, P., Klein, A., ... & Seabold, S. (2020). pandas-dev/pandas: Pandas 1.0. 5. Zenodo.

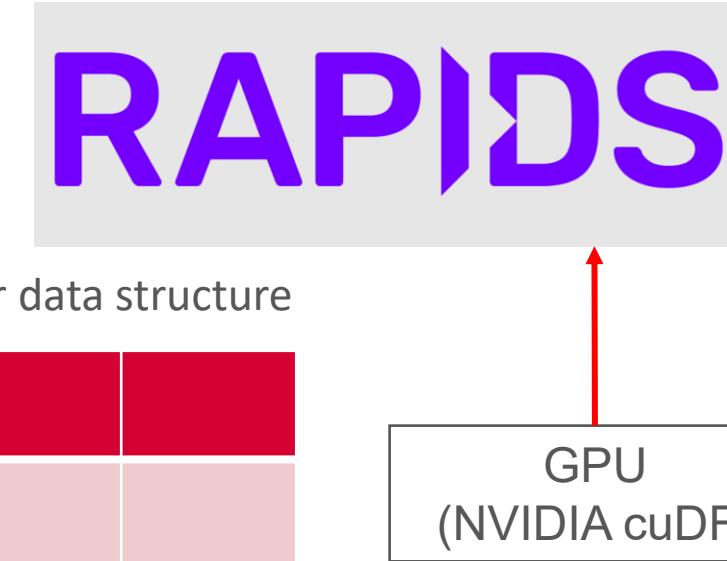
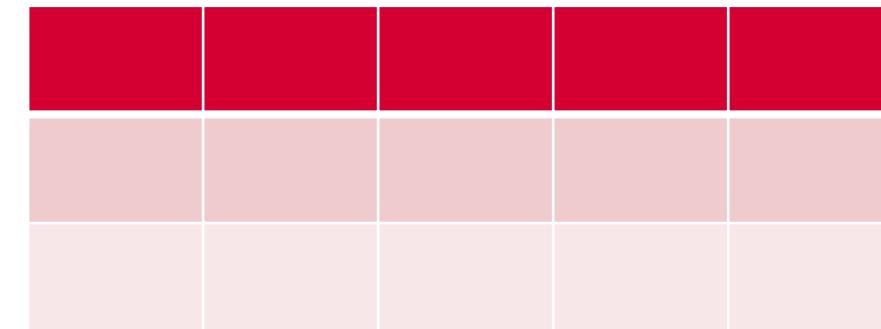
• Chen, D. Y. (2017). Pandas for everyone: Python data analysis. Addison-Wesley Professional.

• Singh, R. (2020, July 1). Merging DataFrames with Pandas: Pd.merge(). Medium. Retrieved April 8, 2023, from <https://medium.com/swlh/merging-dataframes-with-pandas-pd-merge-7764c7e2d46d>

# Off-the-shelf Python Libraries



DataFrame: 2D labeled tabular data structure



*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
import cudf

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

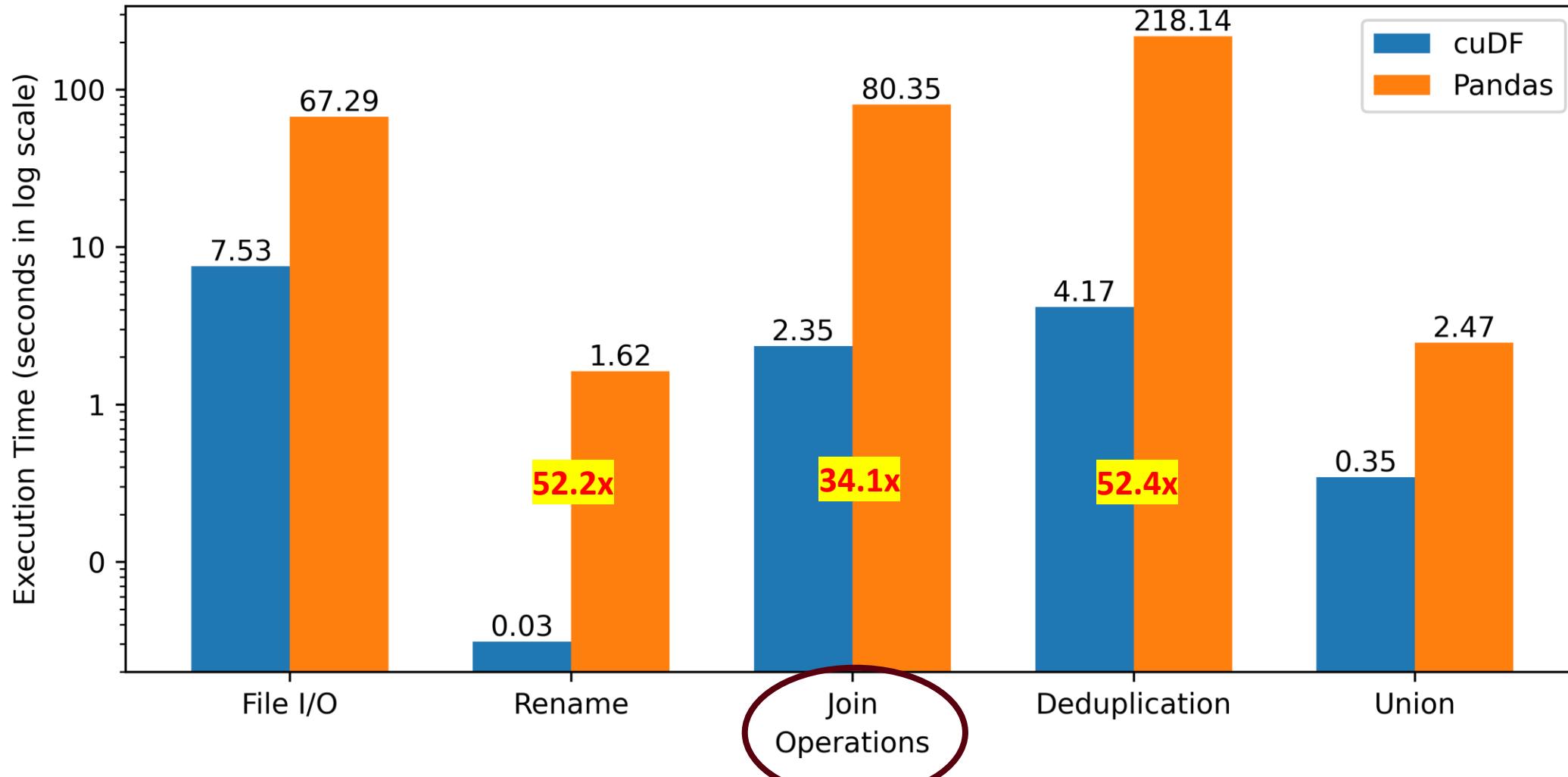


**Apps:** Transitive Closure, Triangle Counting

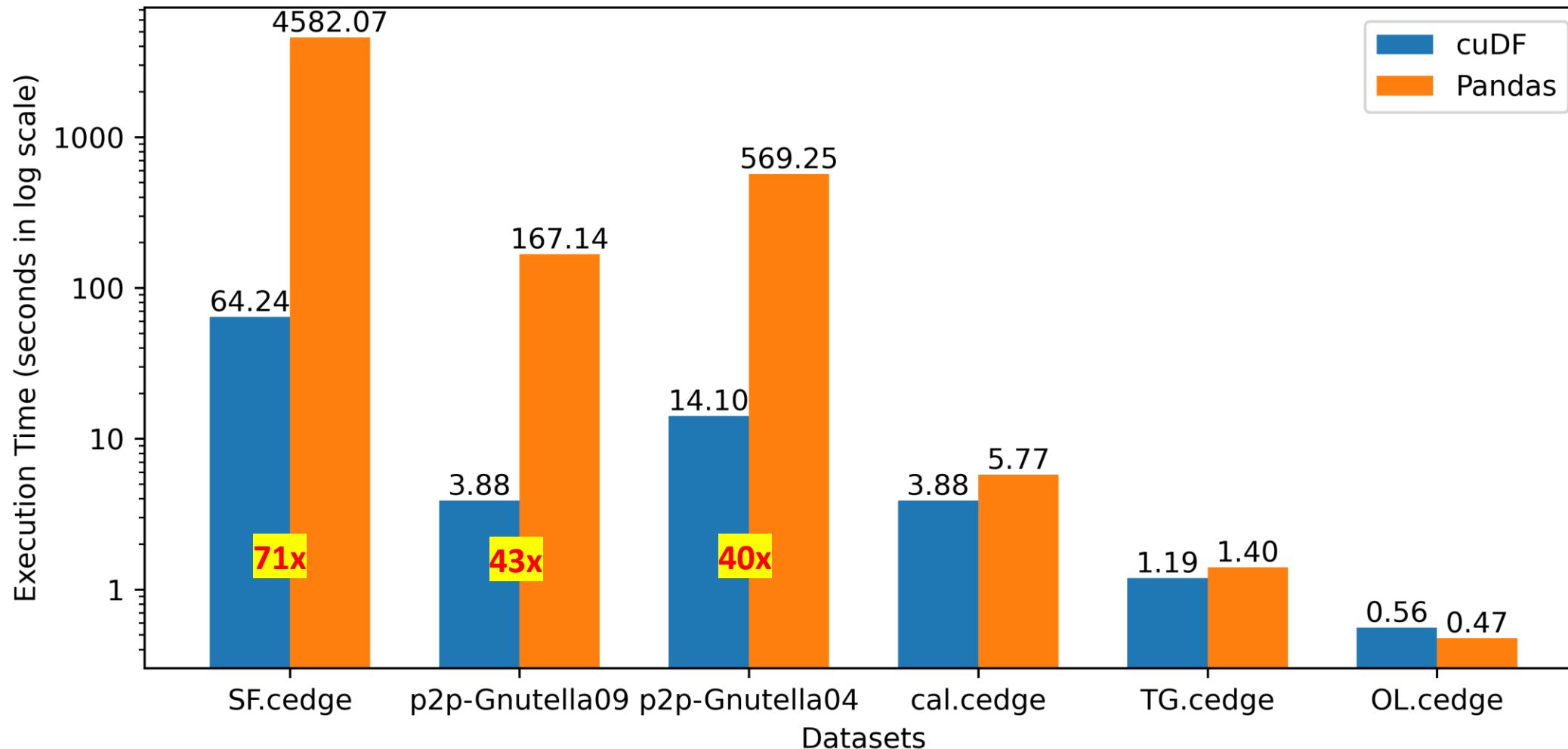
**Datasets:** Stanford large network, SuiteSparse, Road network

- 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

- A. R. Shovon, L. R. Dyken, O. Green, T. Gilray and S. Kumar, "Accelerating Datalog applications with cuDF," 2022 IEEE/ACM Workshop on Irregular Applications: Architectures and Algorithms (IA3), Dallas, TX, USA, 2022, pp. 41-45
- 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.

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Compare GPU capabilities  
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GPUJoin

Relational Algebra (RA)  
operations using CUDA

High performance GPU  
hashtable for iterative RA

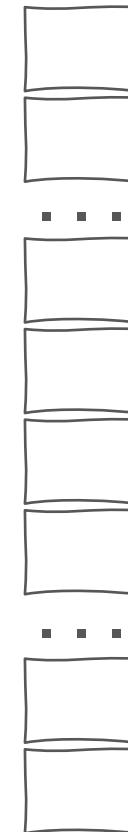
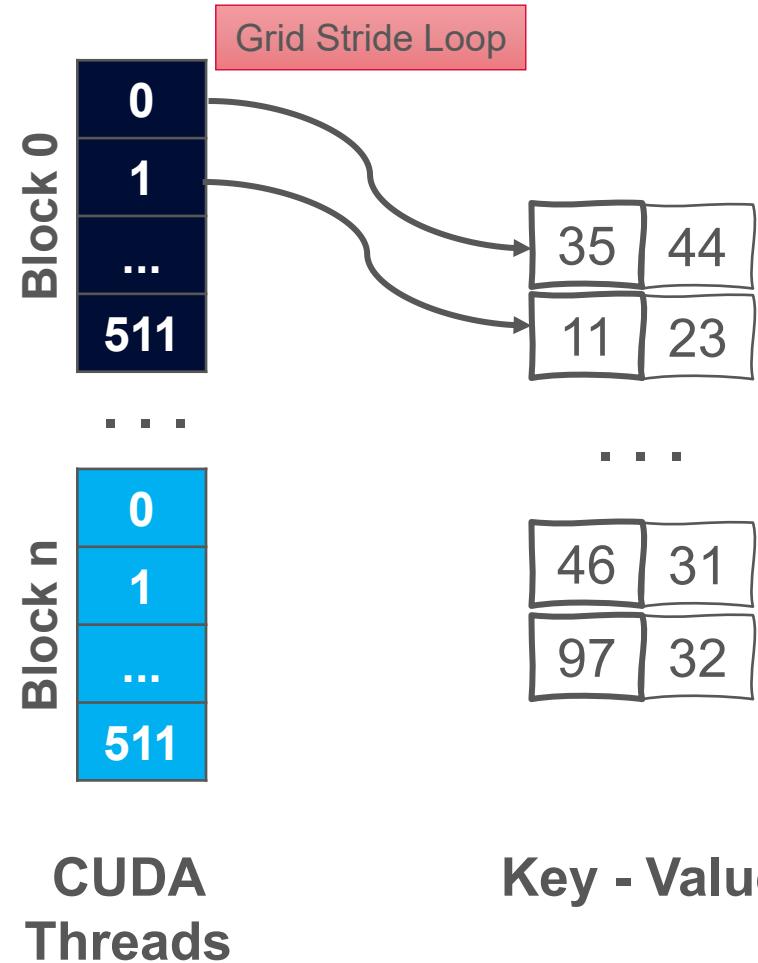
GPULog

CUDA based SIMD API for  
deductive analytics

Novel Hash-Indexed Sorted  
Array data structure



# Hash Table (Open Addressing, Linear Probing)

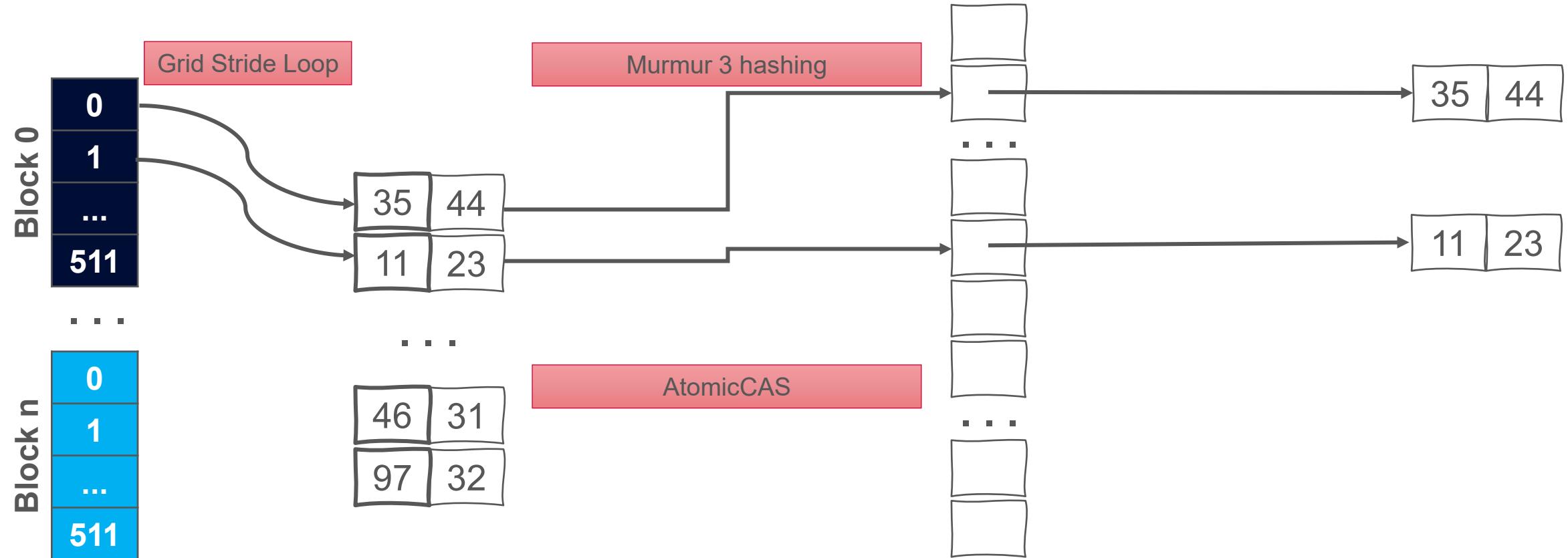


Hash Table

Key-Value Pair



# Hash Table (Open Addressing, Linear Probing)



CUDA  
Threads

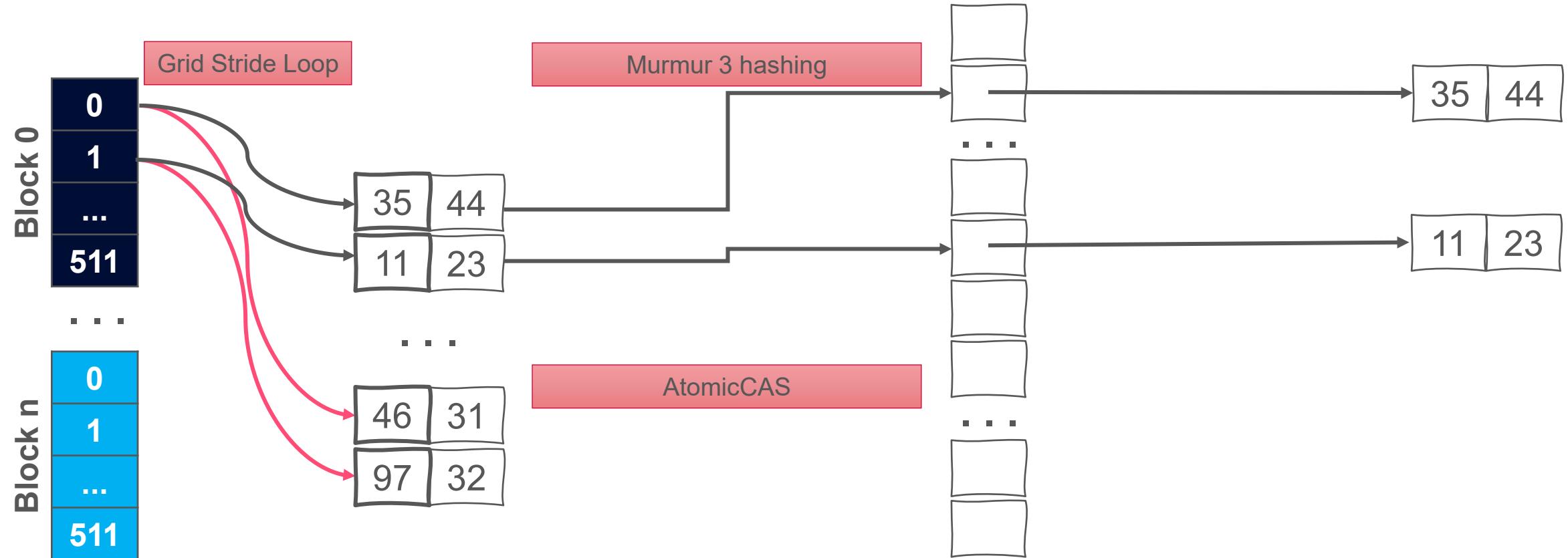
Key - Value

Hash Table

Key-Value Pair



# Hash Table (Open Addressing, Linear Probing)



CUDA  
Threads

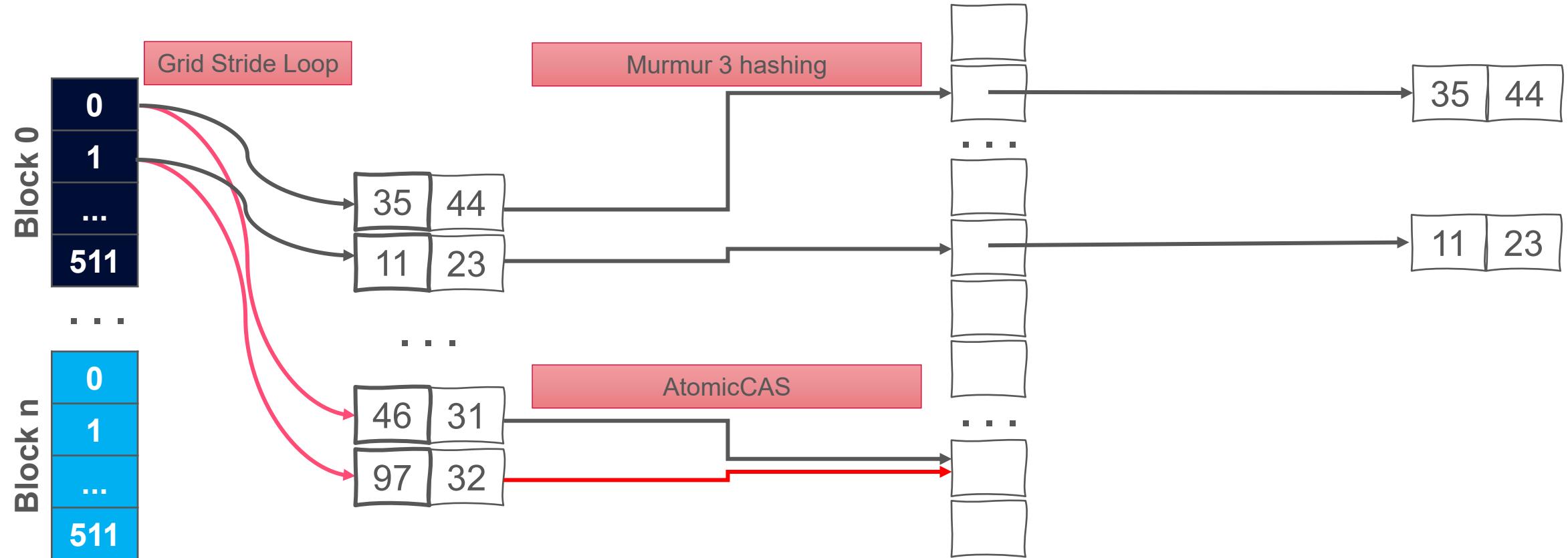
Key - Value

Hash Table

Key-Value Pair



# Hash Table (Open Addressing, Linear Probing)



CUDA  
Threads

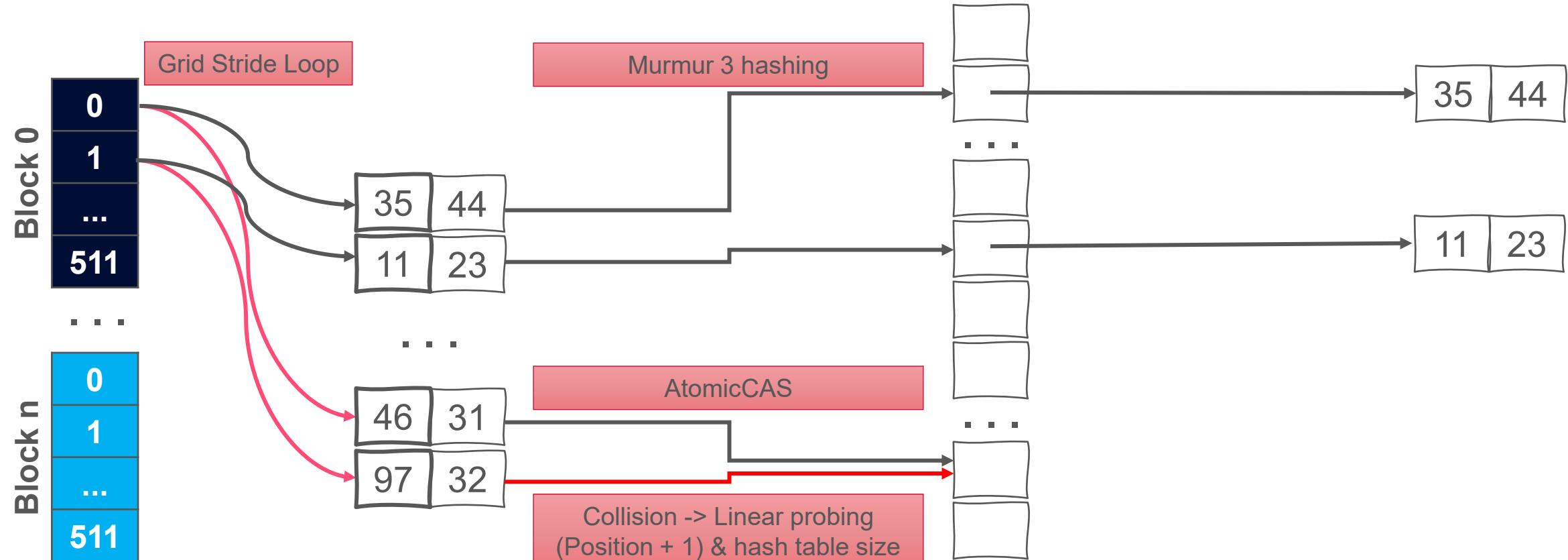
Key - Value

Hash Table

Key-Value Pair



# Hash Table (Open Addressing, Linear Probing)



CUDA  
Threads

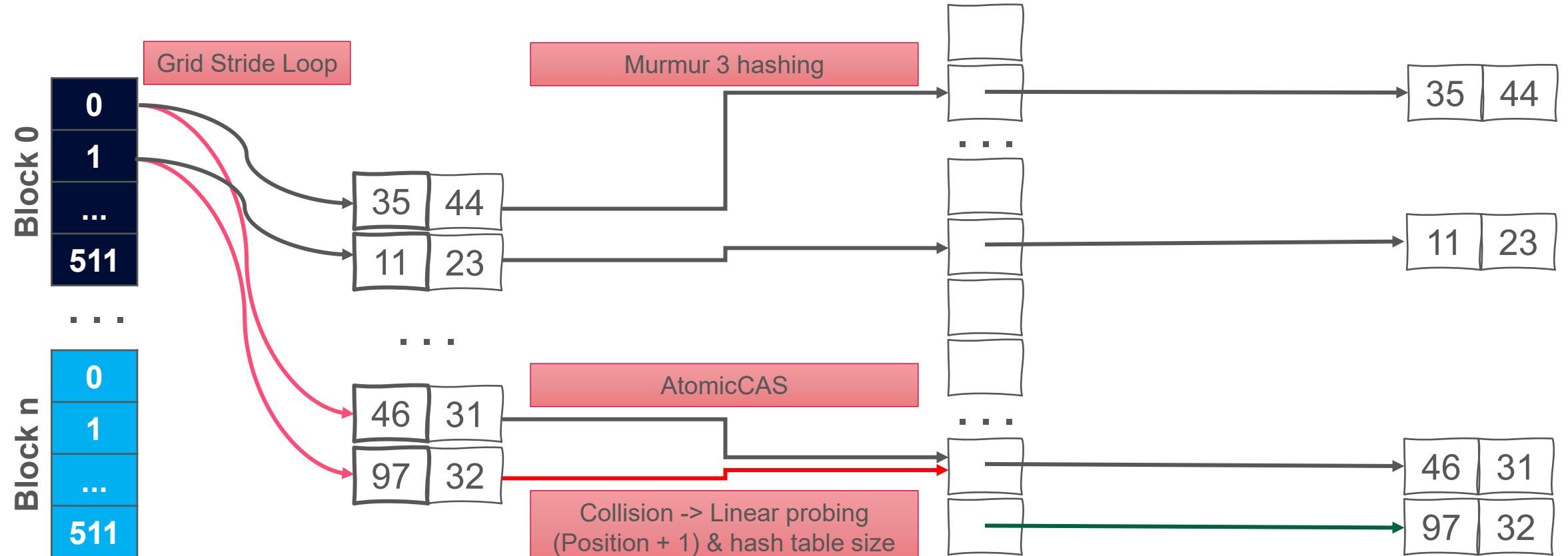
Key - Value

Hash Table

Key-Value Pair



# Hash Table (Open Addressing, Linear Probing)



CUDA  
Threads

Key - Value

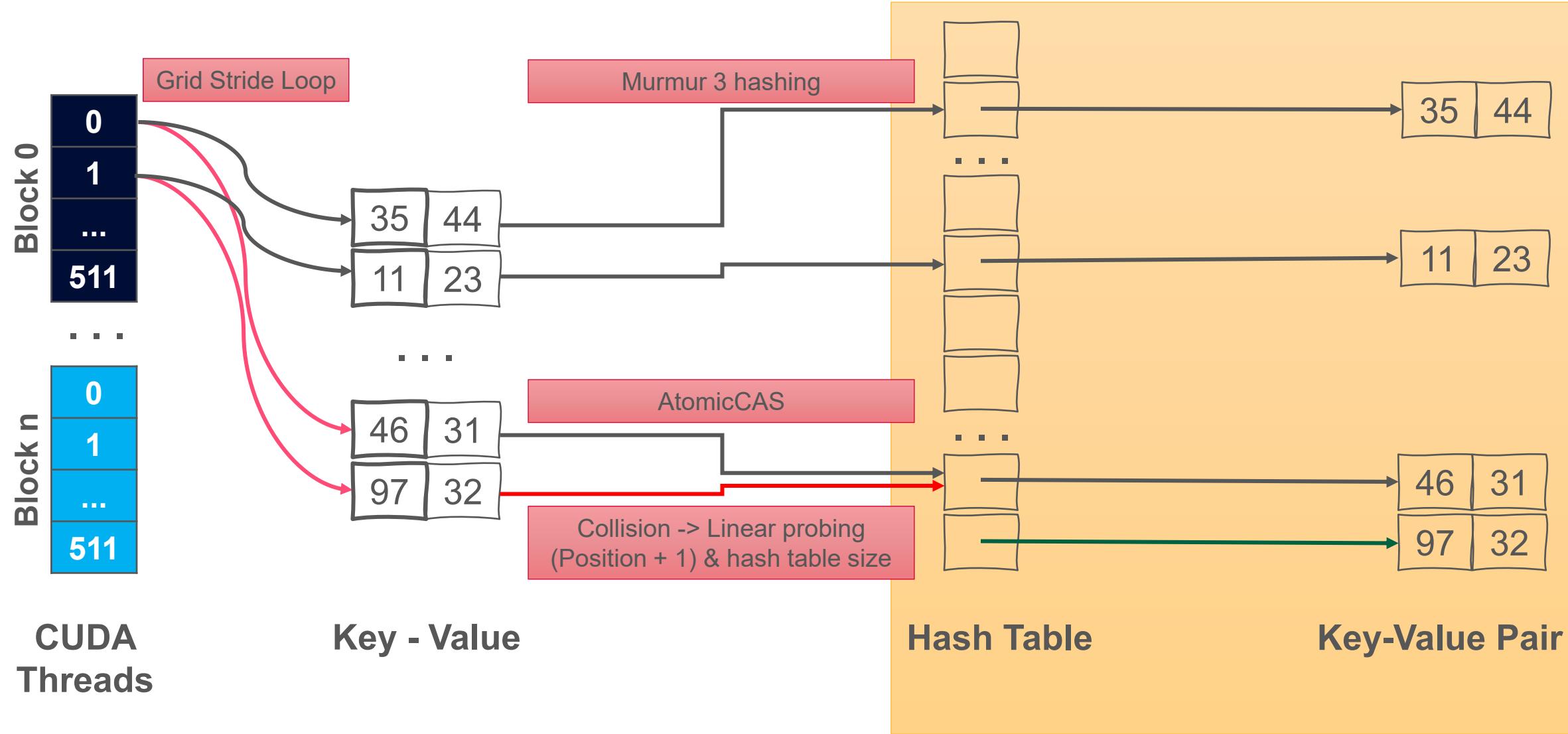
Hash Table

Key-Value Pair



# Hash Table (Open Addressing, Linear Probing)

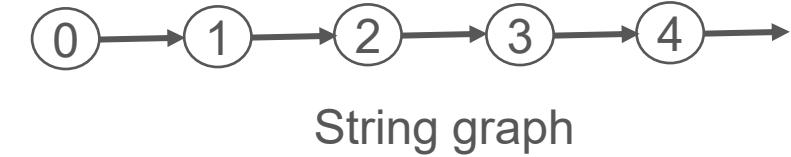
Stores key-value pairs in sparse space  
Uses extra memory



# 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



Key	Value

First pass

Calculate join size

# Performing Hash Join on GPU

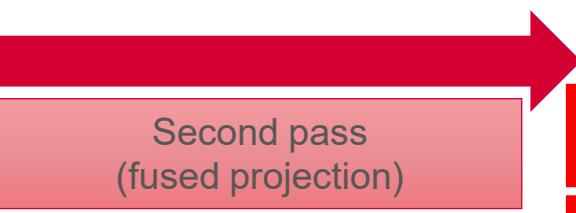
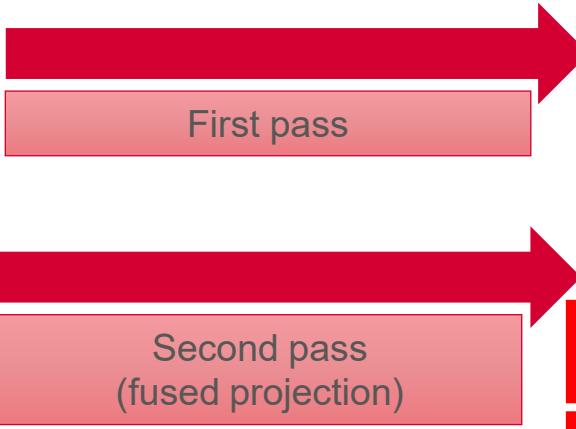
$$F_G(T) \triangleq G \cup \Pi_{1,2}(\rho_{0/1}(T) \bowtie_1 G)$$

**Static Hash Table**

Key	Value

**Inner Relation**

Key	Value



**Calculate join size**

Prefix sum

**Join Result**

Join key	Value 1	Value 2
X		

# Performing Hash Join on GPU

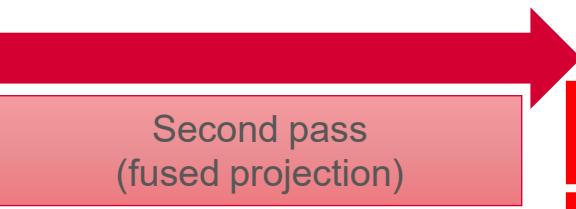
$$F_G(T) \triangleq G \cup \Pi_{1,2}(\rho_{0/1}(T) \bowtie_1 G)$$

**Static Hash Table**

Key	Value

**Inner Relation**

Key	Value



**Calculate join size**

Prefix sum

**Join Result**

Join key	Value 1	Value 2
X		
X		
X		
X		
X		
X		
X		

Sort and Unique

**Deduplicated Join Result**

Key	Value

# Evaluation environment, applications, datasets

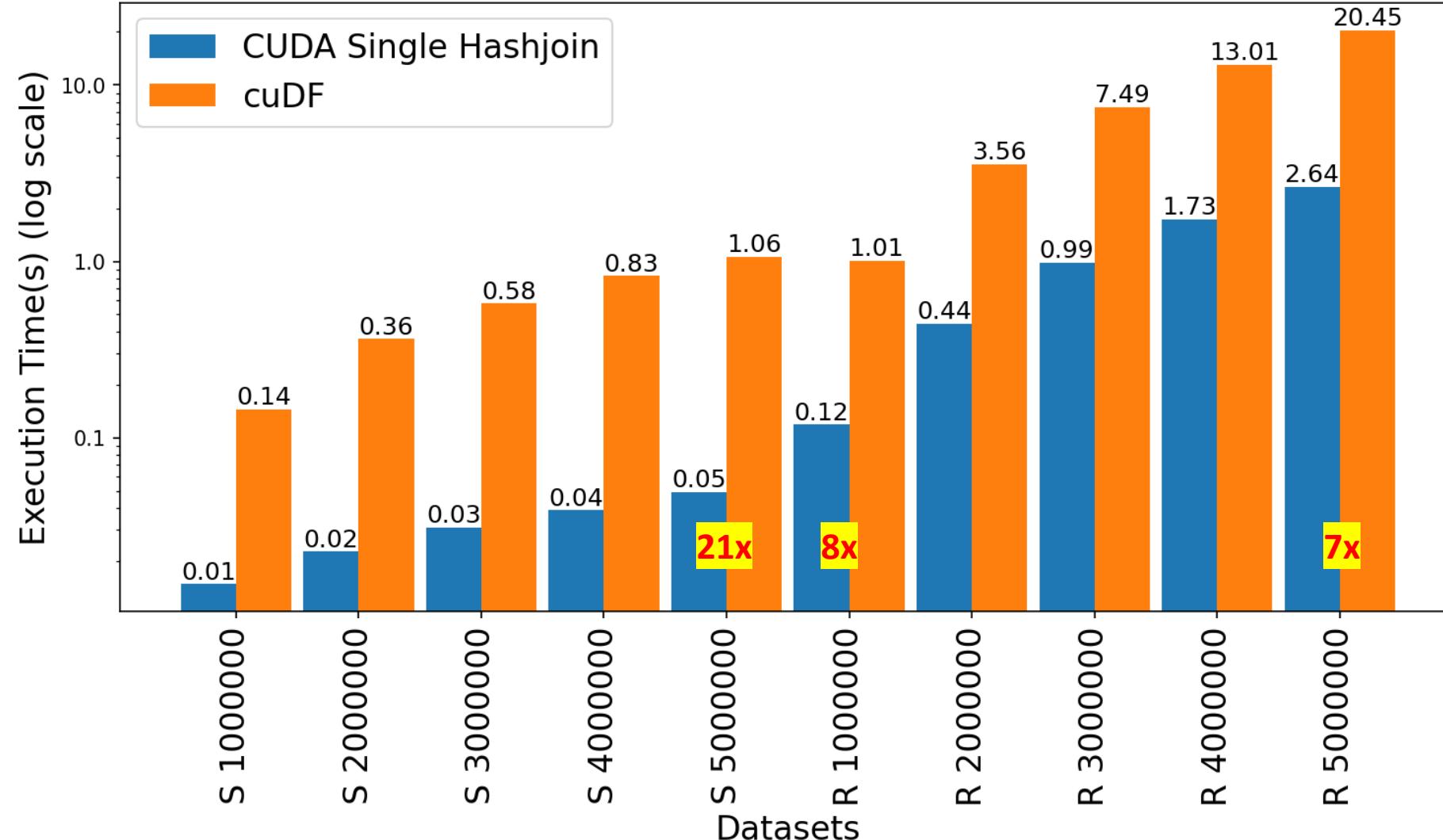


**Apps:** Transitive Closure, Iterated Hash Join

**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
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
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

## Off-the-shelf

Started with CPU and GPU based Python libraries

Compare GPU capabilities for iterative join operations

## GPUJoin

Relational Algebra (RA) operations using CUDA

High performance GPU hashtable for iterative RA

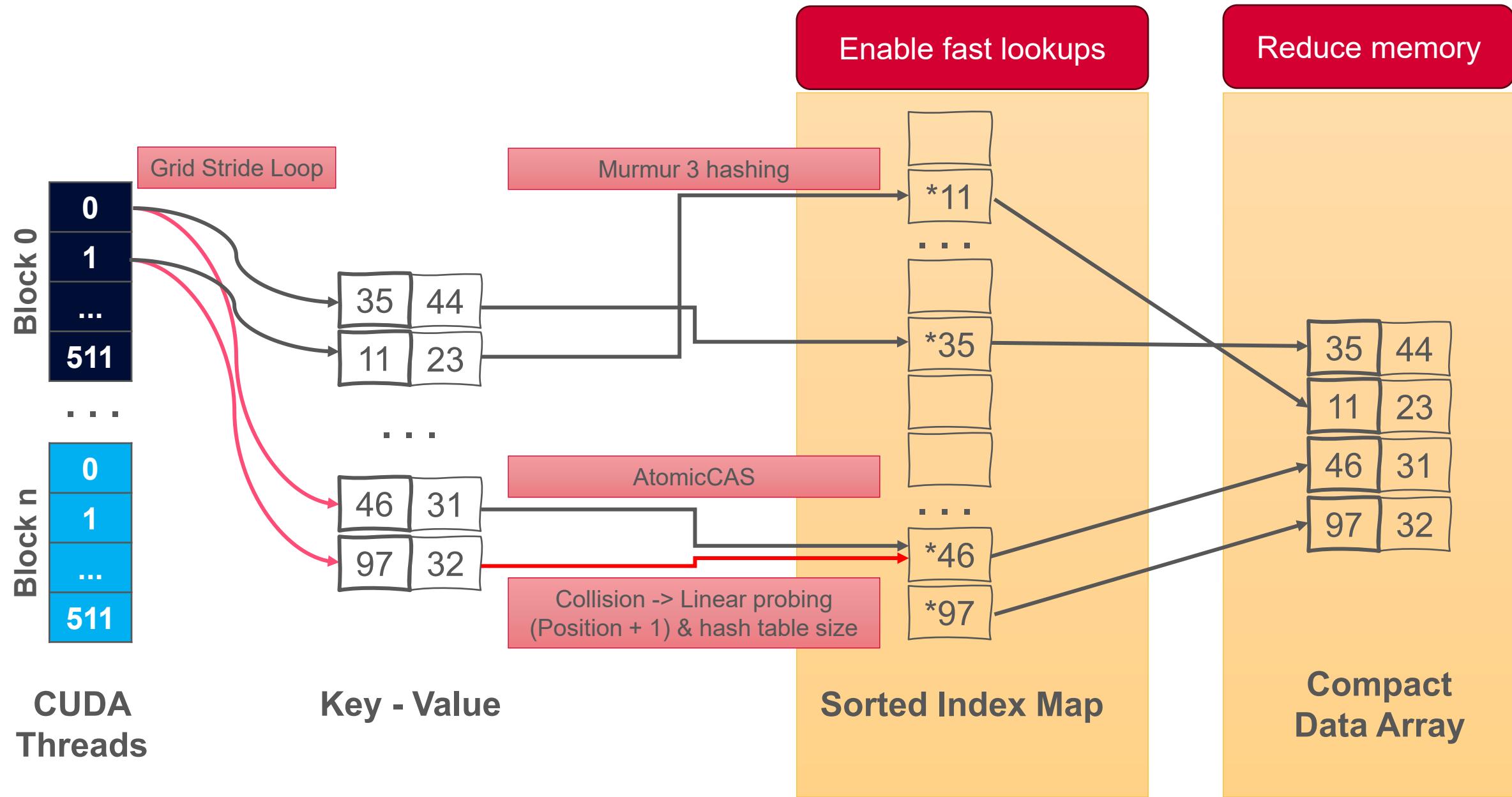
## GPULog

CUDA based SIMD API for deductive analytics

Novel Hash-Indexed Sorted Array data structure



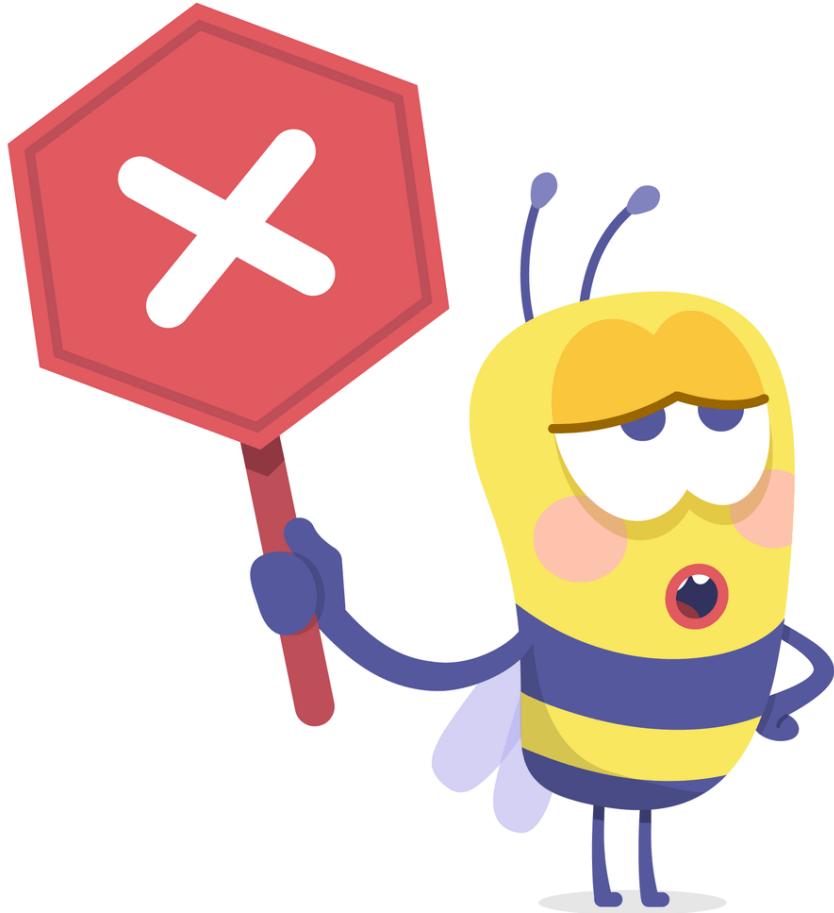
# Hash Indexed Sorted Array (HISA)



# Performance Enhancement (Transitive closure)

Dataset name	<i>Reach edges</i>	GPULOG	Soufflé	GPUJoin	cuDF
com dblp	1.91B	<b>14.30</b>	232.99	OOM	OOM
fe ocean	1.67B	<b>23.36</b>	292.15	100.30	OOM
vsp finan	910M	<b>21.91</b>	239.33	125.94	OOM
Gnutella31	884M	<b>5.58</b>	96.82	OOM	OOM
fe body	156M	<b>3.76</b>	23.40	22.35	OOM
SF cedge	80M	<b>1.63</b>	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

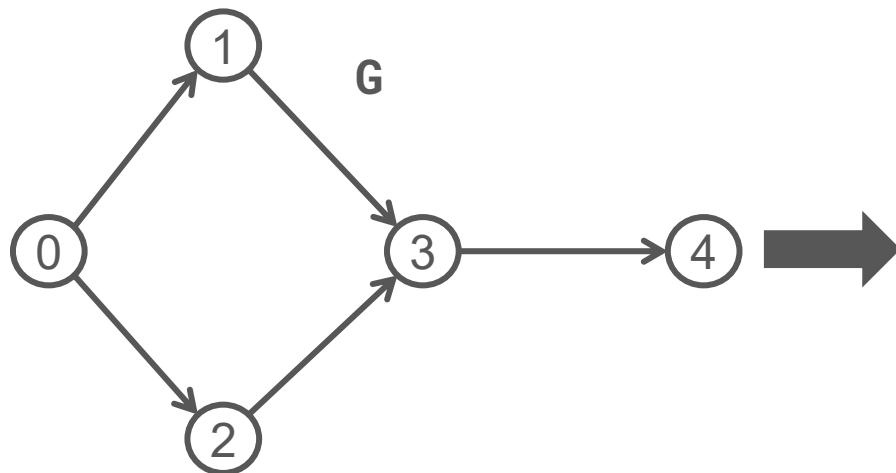


MPI + X (Cuda) design

3<sup>rd</sup> Best Poster Award at Computational Research Symposium 2025

ACM ICS '25

# Requirements for Multi-node Multi-GPU Datalog

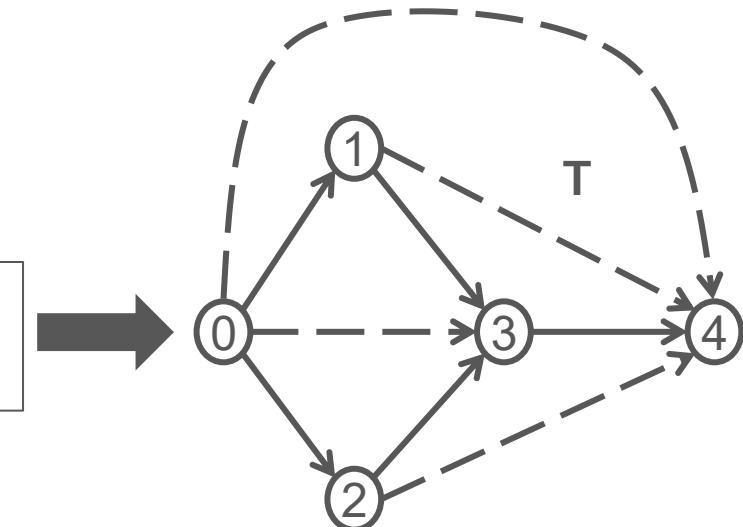


```
tc(X, Y) :- edges(X, Y).
tc(X, Z) :- tc(X, Y), edges(Y, Z).
```

2 Data representation

0	1
0	2
1	3
2	3
3	4

1 Workload partitioning



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

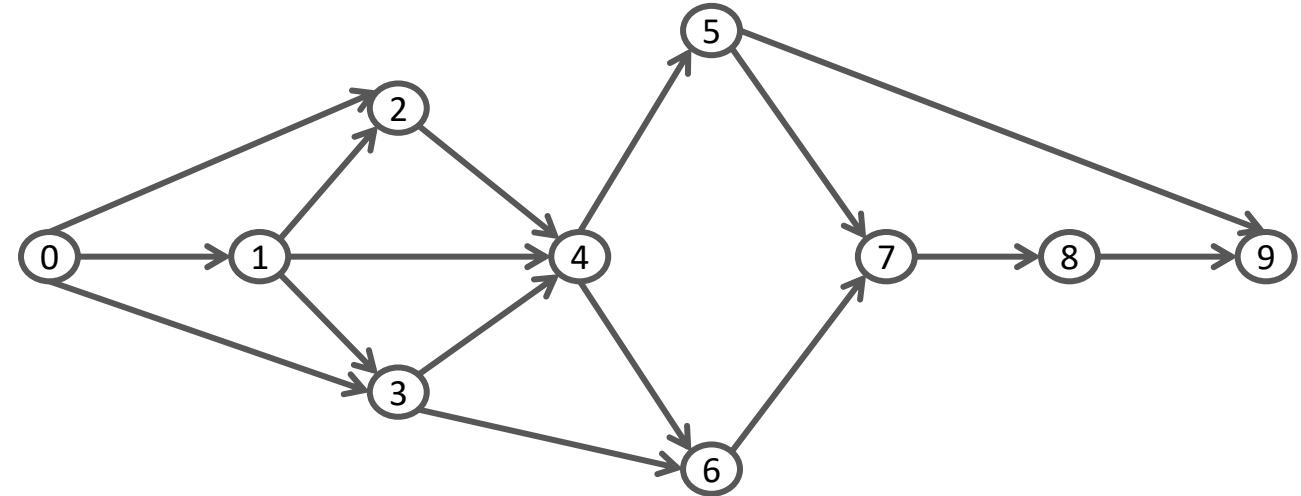
Communication

(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)



(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(k) % 3

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

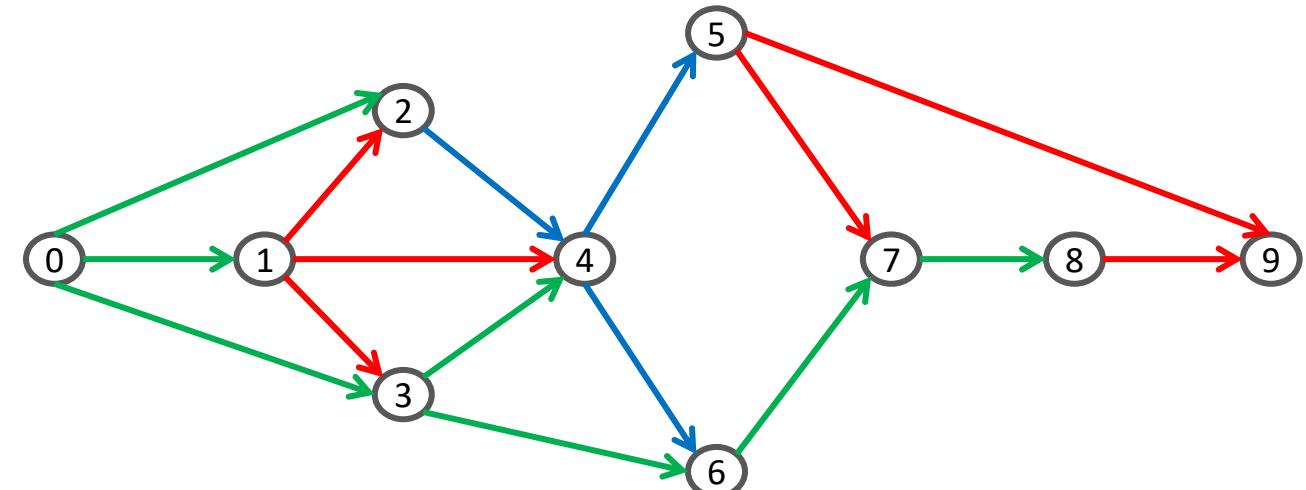
0	1
0	2
0	3
3	4
3	6
7	6
7	8
5	7
5	9
8	9
1	2
1	3
1	4
2	4
4	5
4	6
1	2
1	3
2	4
4	5
4	6
4	6
7	8
8	9

0

1

2

The join column decides  
the GPU (bucket-id)



GPU (ID)

# MNMGDatalog: Data representation

Hash-based data distribution

GPU-optimized data representation

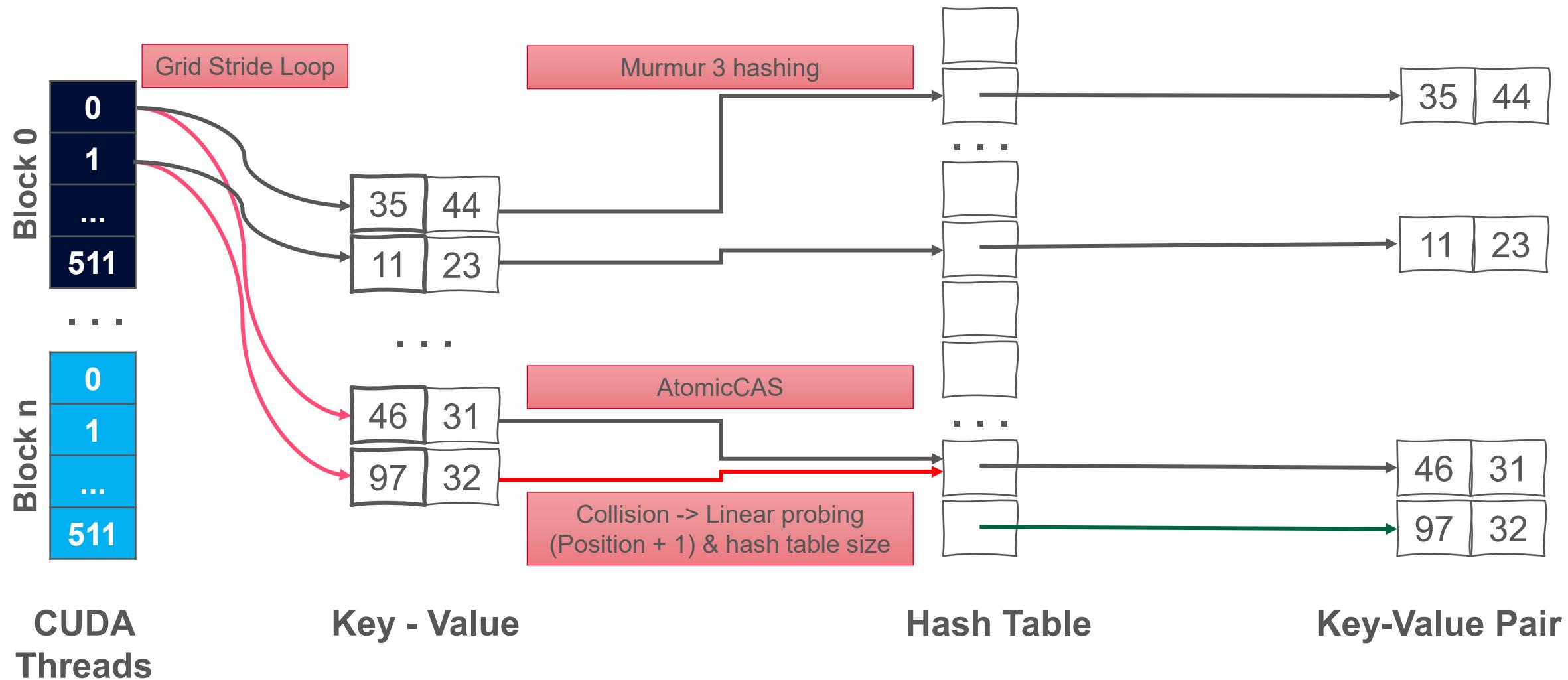
GPU-Aware all-to-all communication

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

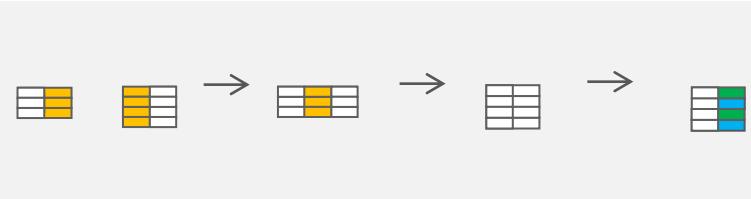
3 Communication

## Relational Algebra Kernel

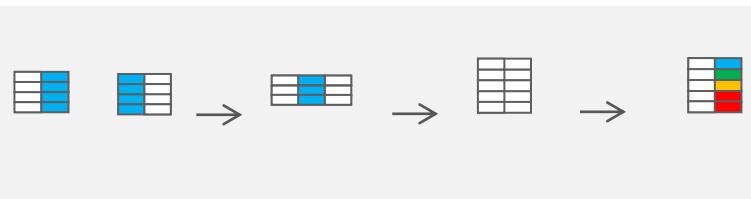
All to all  
comm

Join      Project      Hash

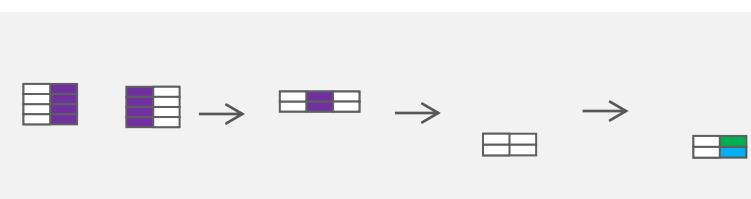
**GPU 0**



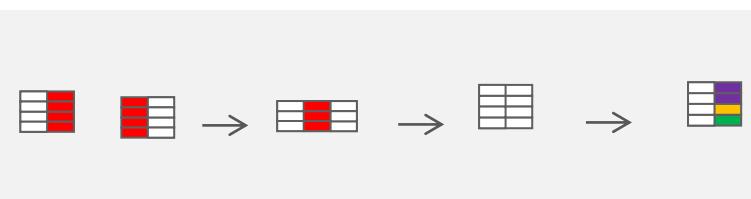
**GPU 1**



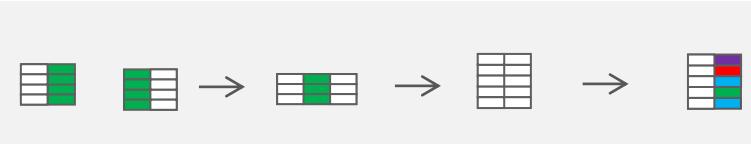
**GPU 2**



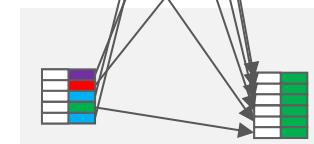
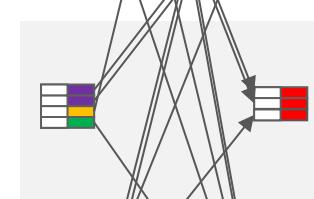
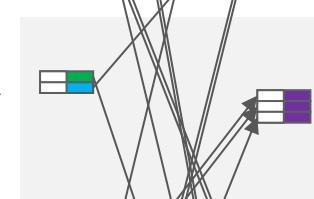
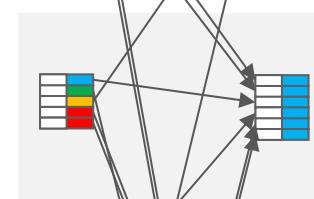
**GPU 3**

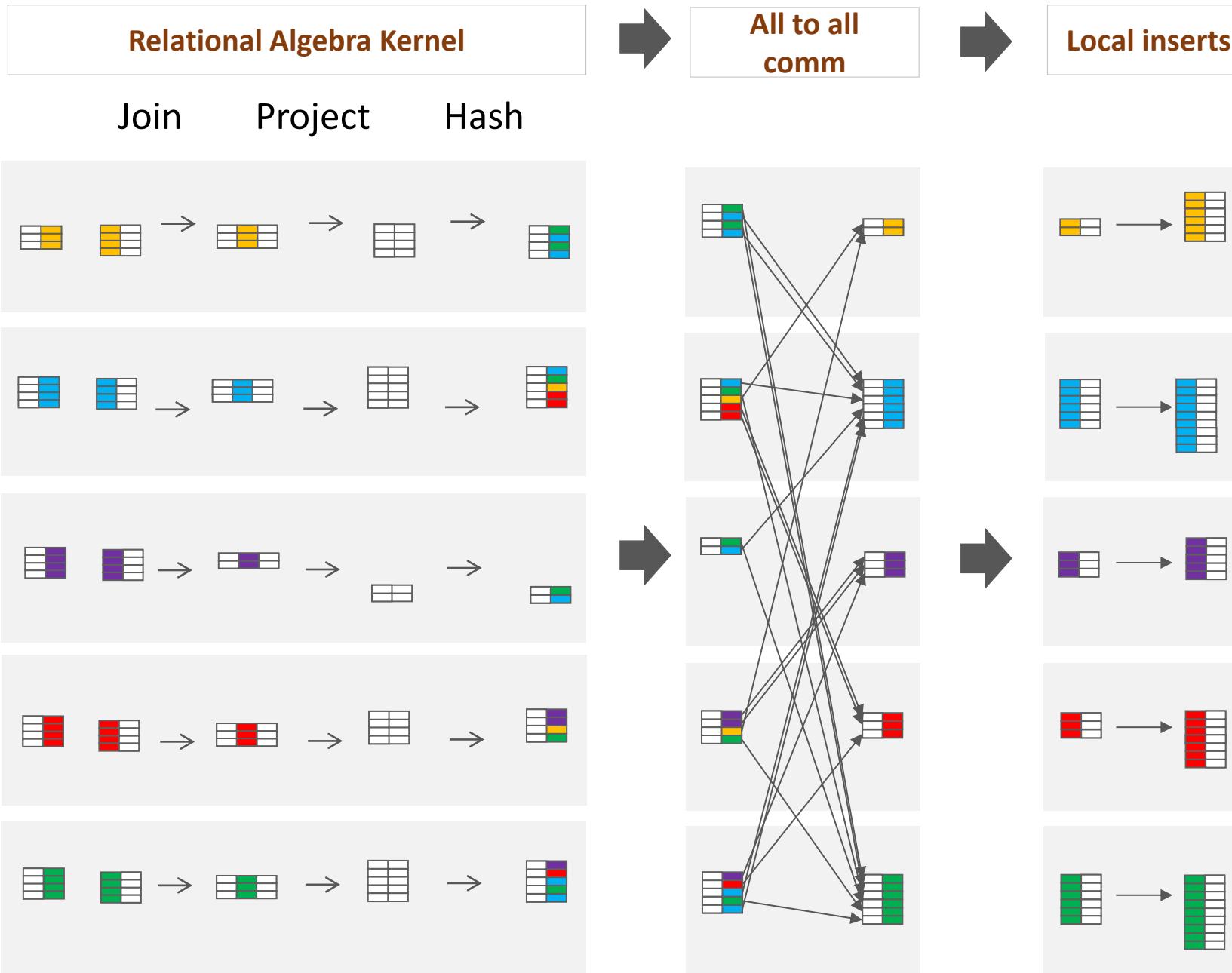


**GPU 4**



All to all  
comm





**Relational Algebra Kernel**

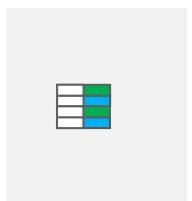
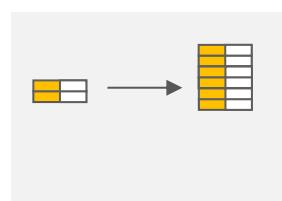
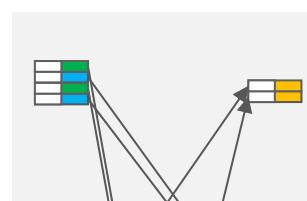
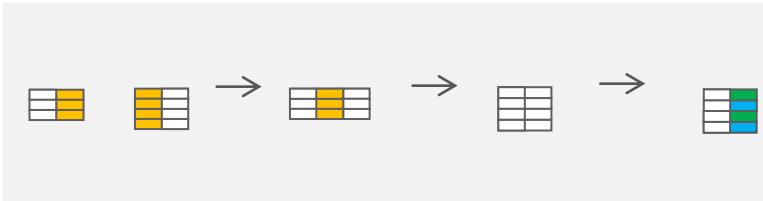
Join      Project      Hash

All to all  
comm

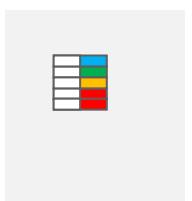
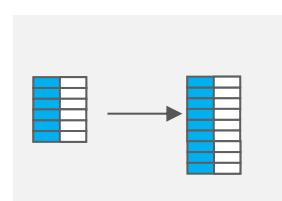
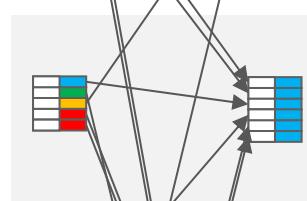
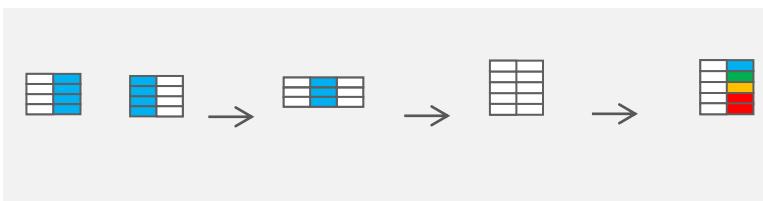
Local inserts

Fixed  
point  
?

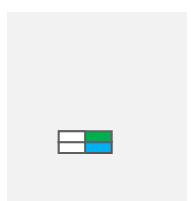
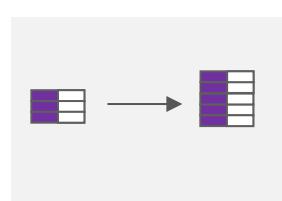
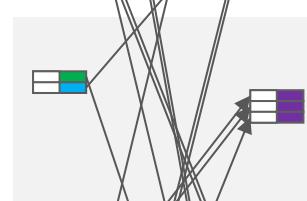
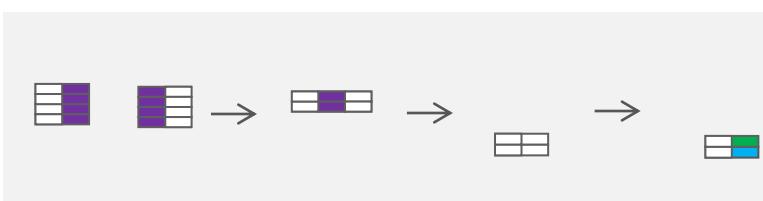
**GPU 0**



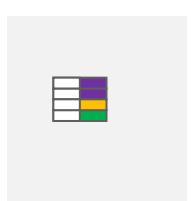
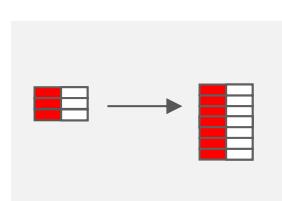
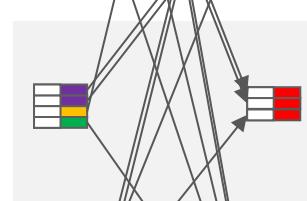
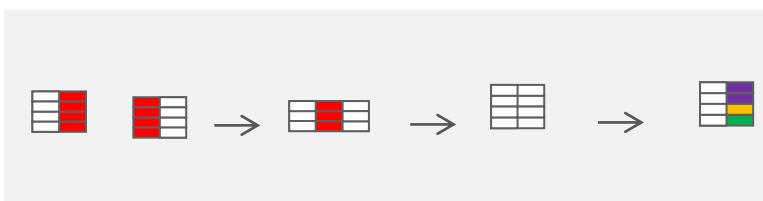
**GPU 1**



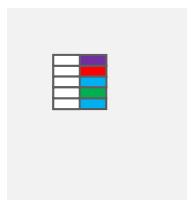
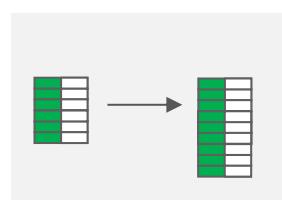
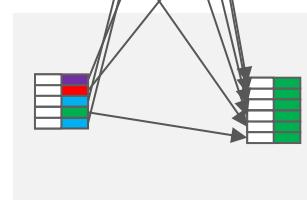
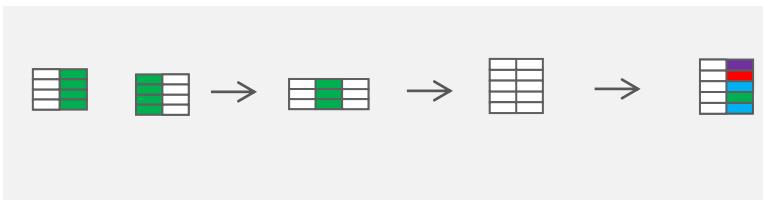
**GPU 2**

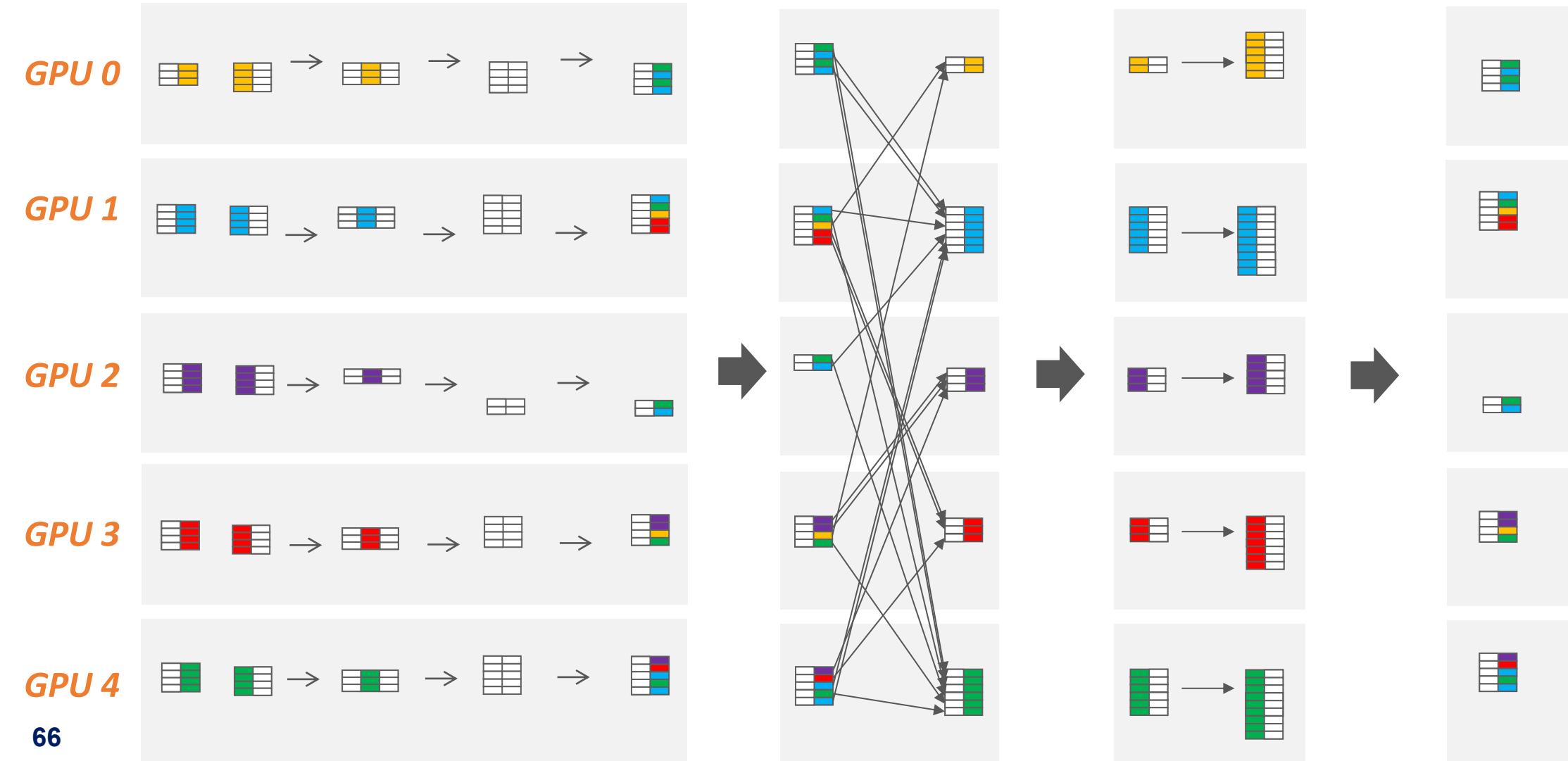
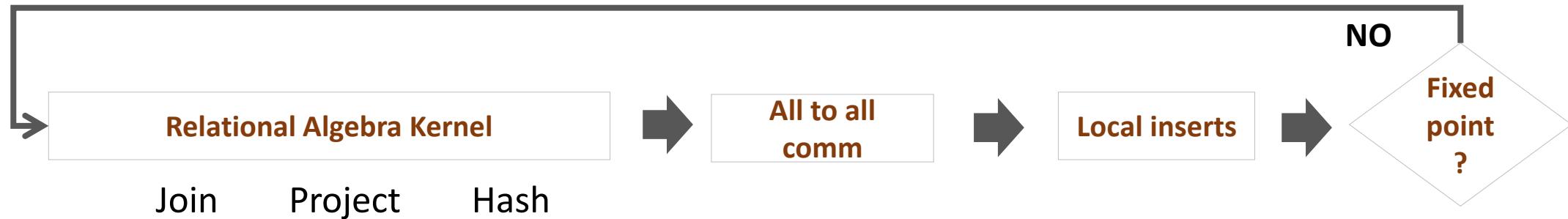


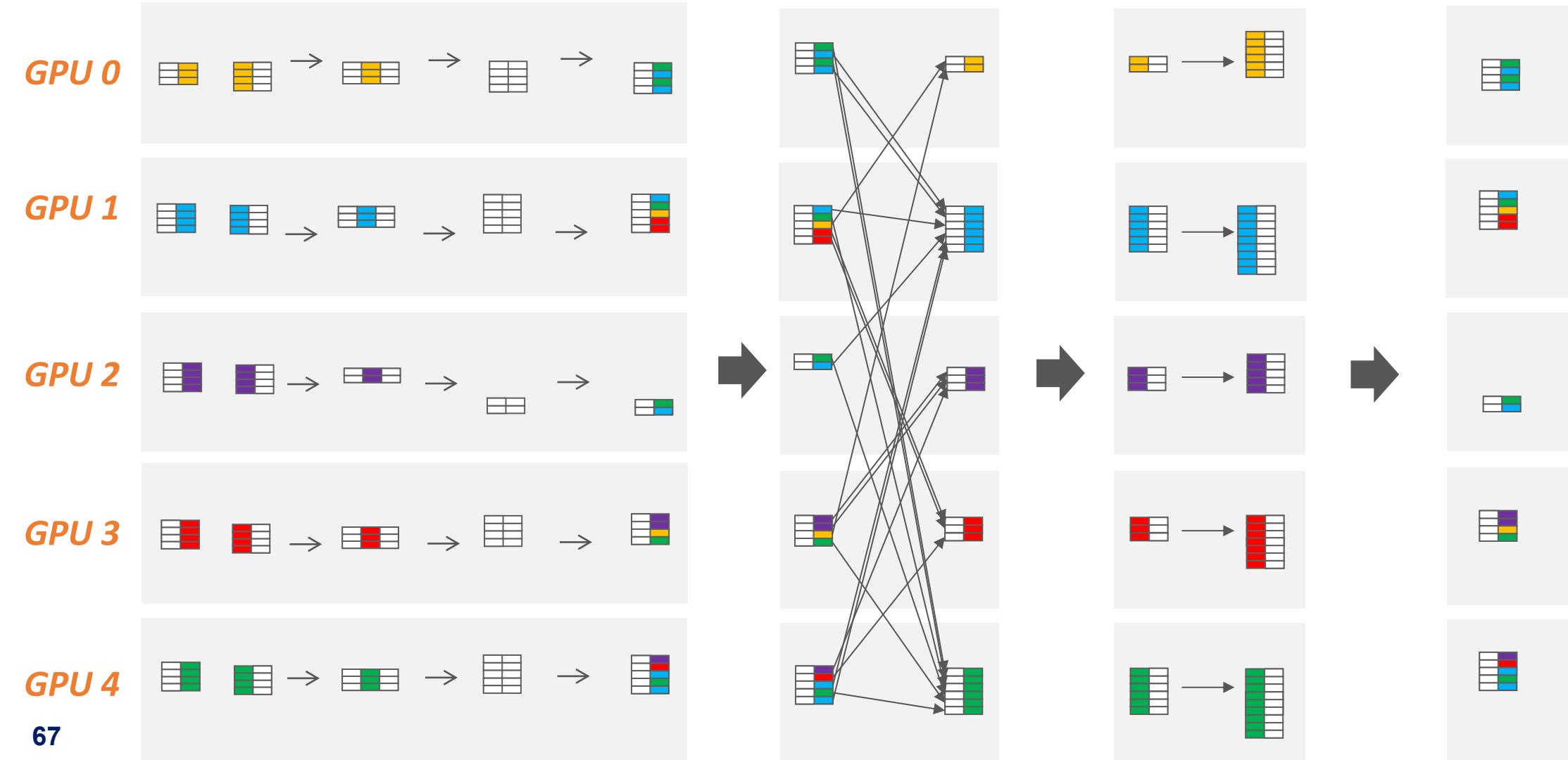
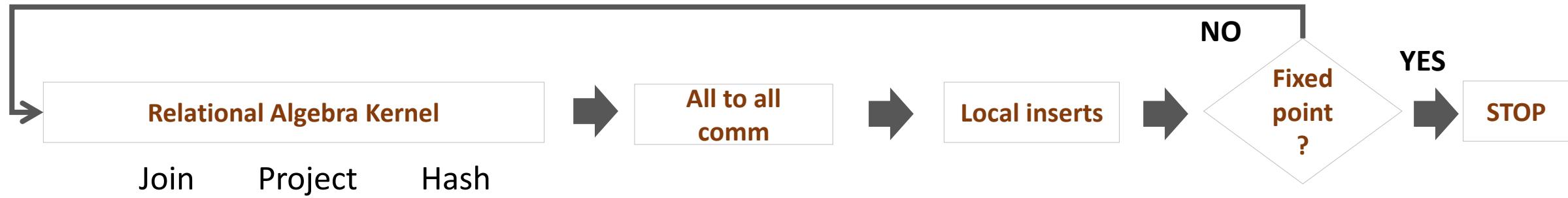
**GPU 3**



**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	<b>13.58</b>	26.95	232.99	OOM
fe ocean	1.67B	<b>66.34</b>	72.74	292.15	100.30
usroads	871M	<b>75.07</b>	78.08	222.76	364.55
vsp finan	910M	<b>81.14</b>	82.75	239.33	125.94

Transitive closure benchmark on single-GPU

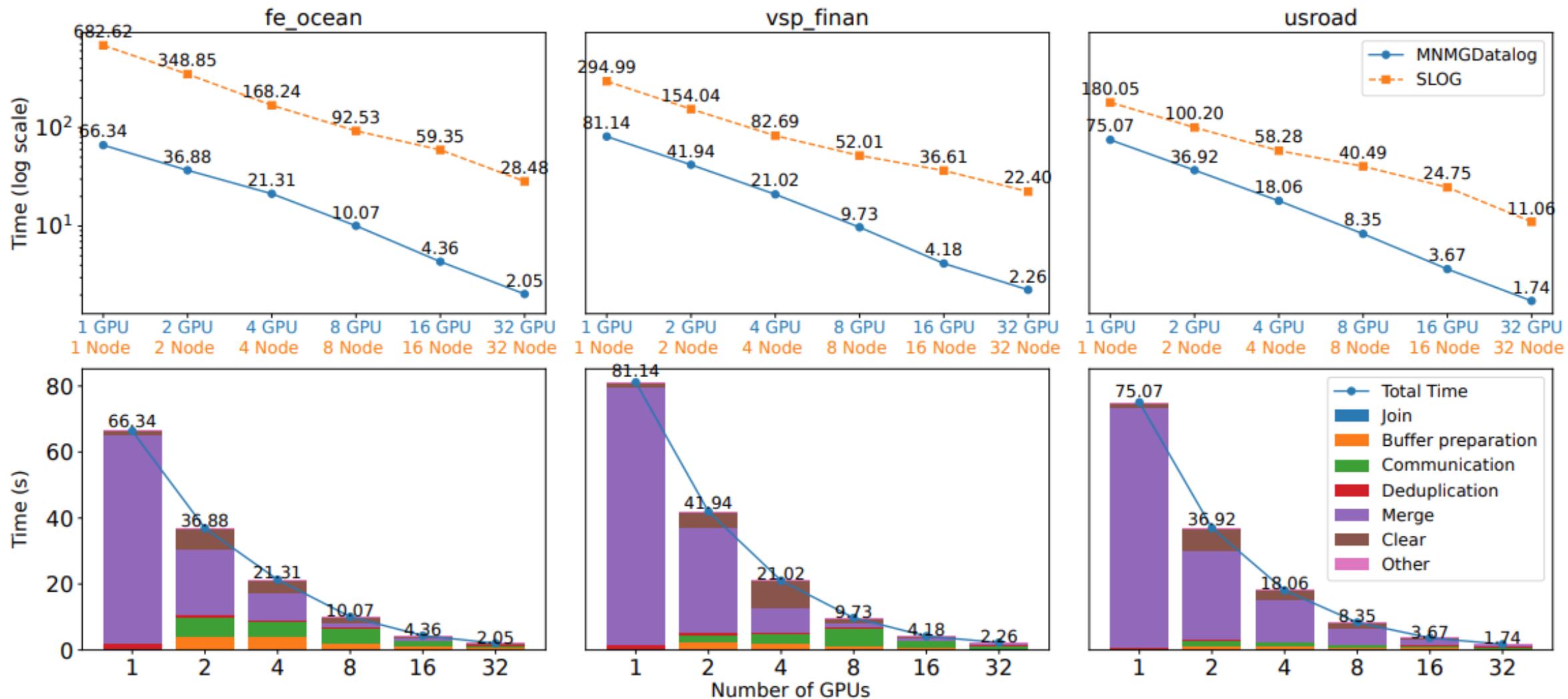
**Speedup up to 1.9x over GPULog, 20x over Souffle,  
4.8x over GPUJoin**

Dataset name	SG size	Time (s)			
		MNMGDATALOG	GPULOG	Soufflé	cuDF
fe_body	408M	<b>9.08</b>	18.41	74.26	OOM
loc-Brightkite	92.3M	<b>1.66</b>	11.67	48.18	OOM
fe_sphere	205M	<b>3.55</b>	7.88	48.12	OOM
CA-HepTH	74M	<b>0.60</b>	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

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

# Parallel Programming Models



## W.J. Cody Associates

Rosetta parallel programming models benchmarking framework

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

**Rosetta Benchmark**

Benchmark Report									Code
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) ≈ 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

# Roadmap

1. Introduction and motivation
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- 5. Power analysis of Datalog engines**
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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



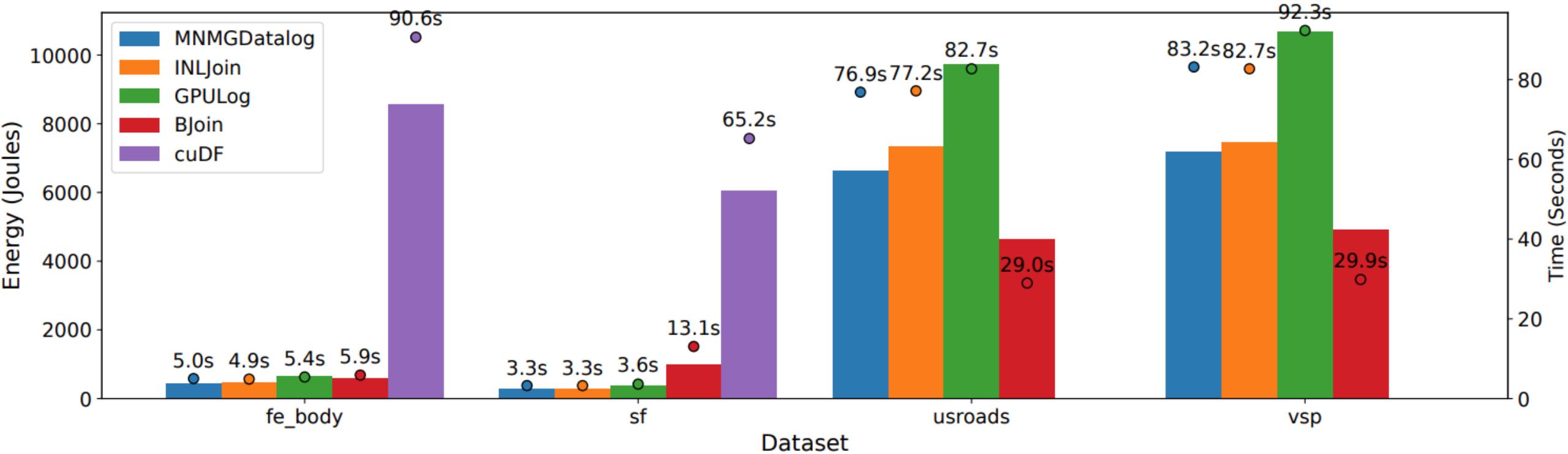
Polaris (Argonne National Lab)

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

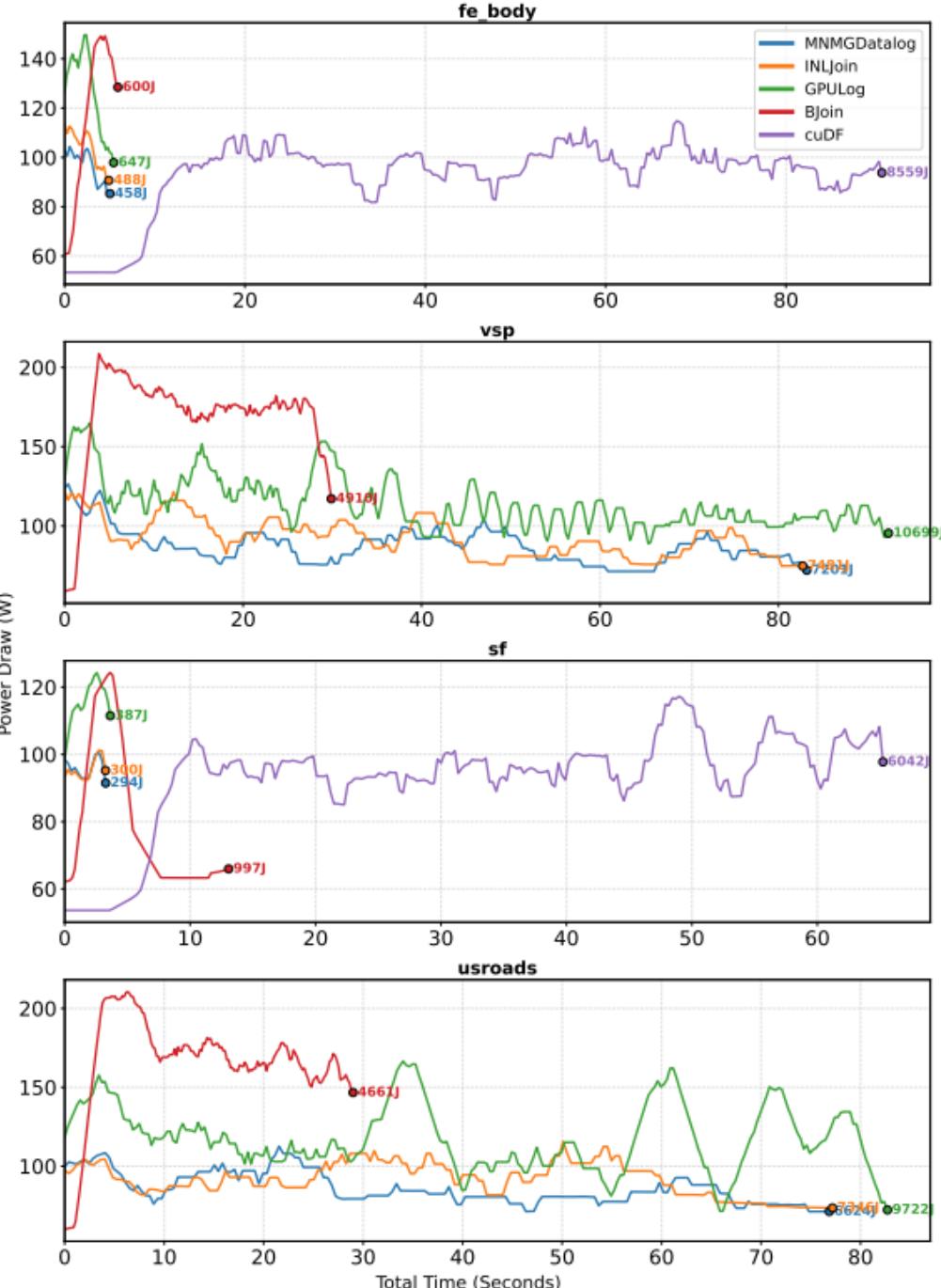
# Energy profile: TC energy consumption (single-GPU)

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

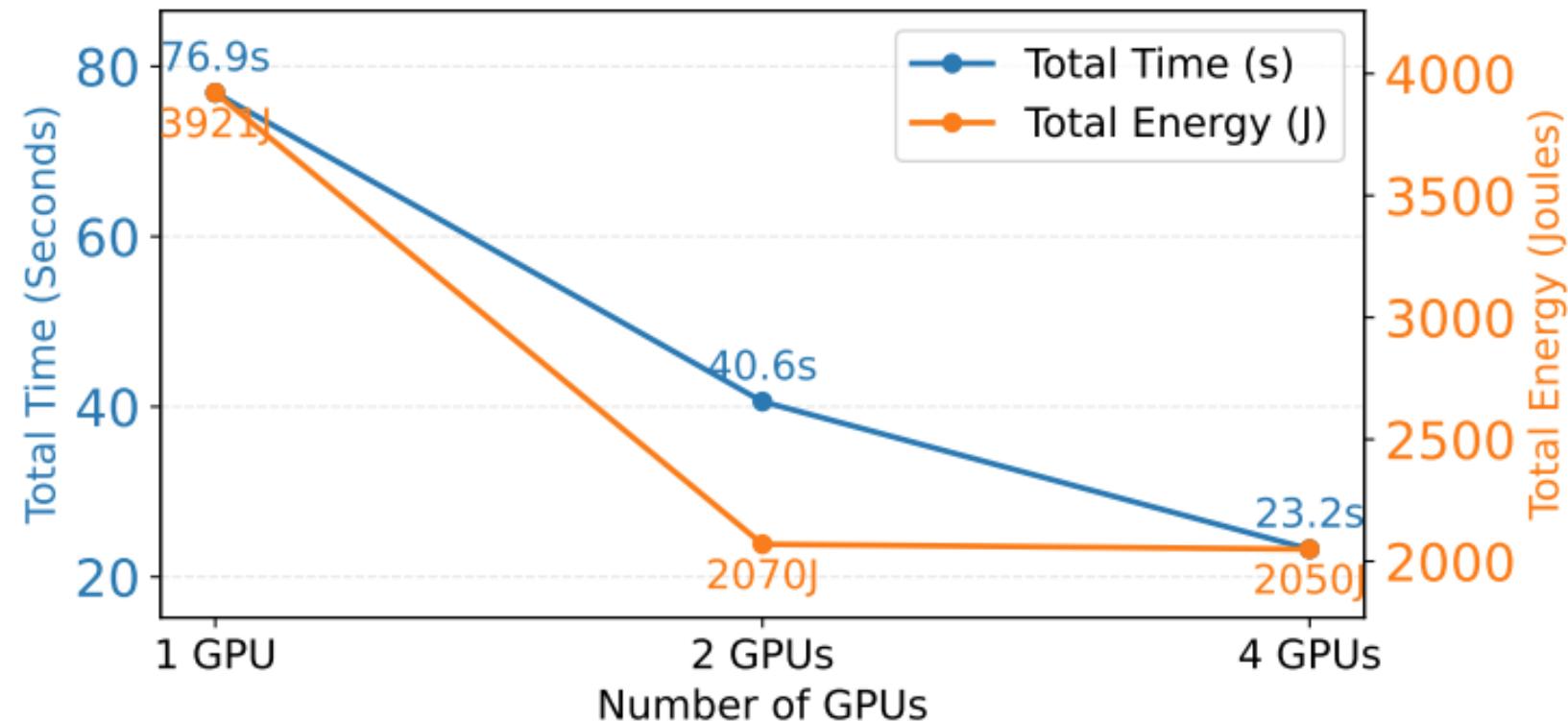
- N is the total number of sampling intervals
- $P_i$  is the power draw (in watts) measured at interval i
- $\Delta t_i$  is the time (in seconds) between samples i and i - 1



- 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

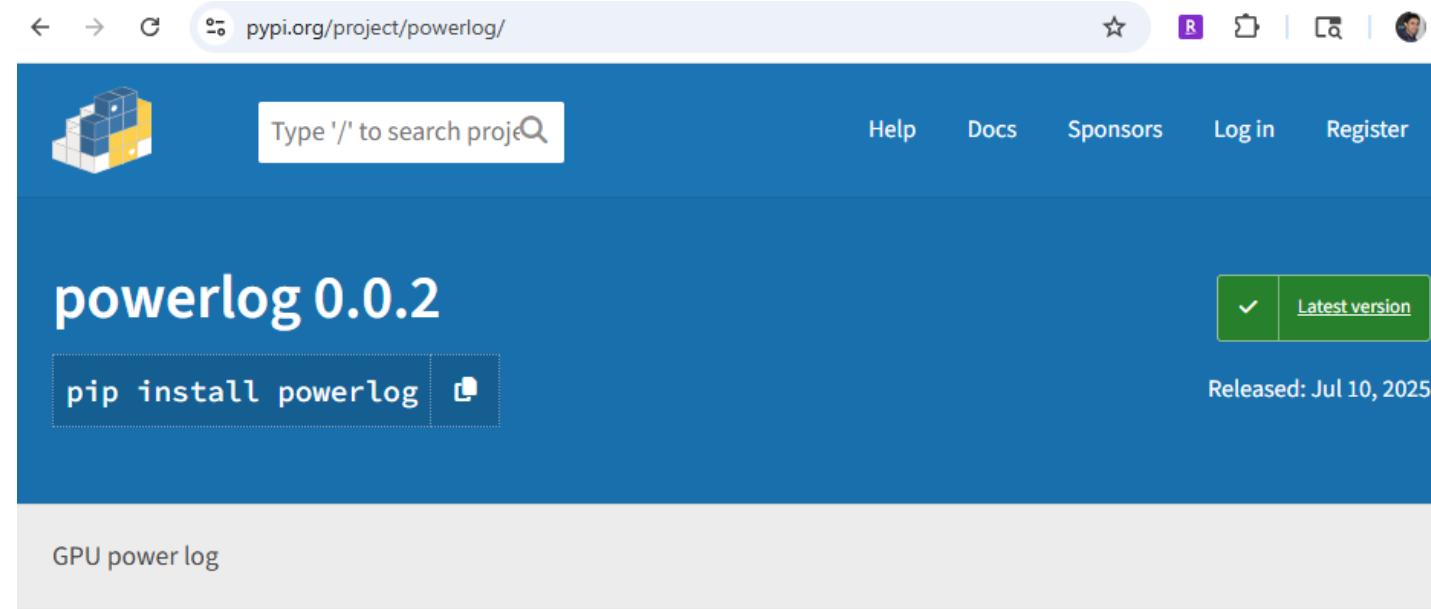


# Energy profile: TC energy consumption (multi-GPUs)



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

# Powerlog CLI Tool for Energy Profiling GPU Applications



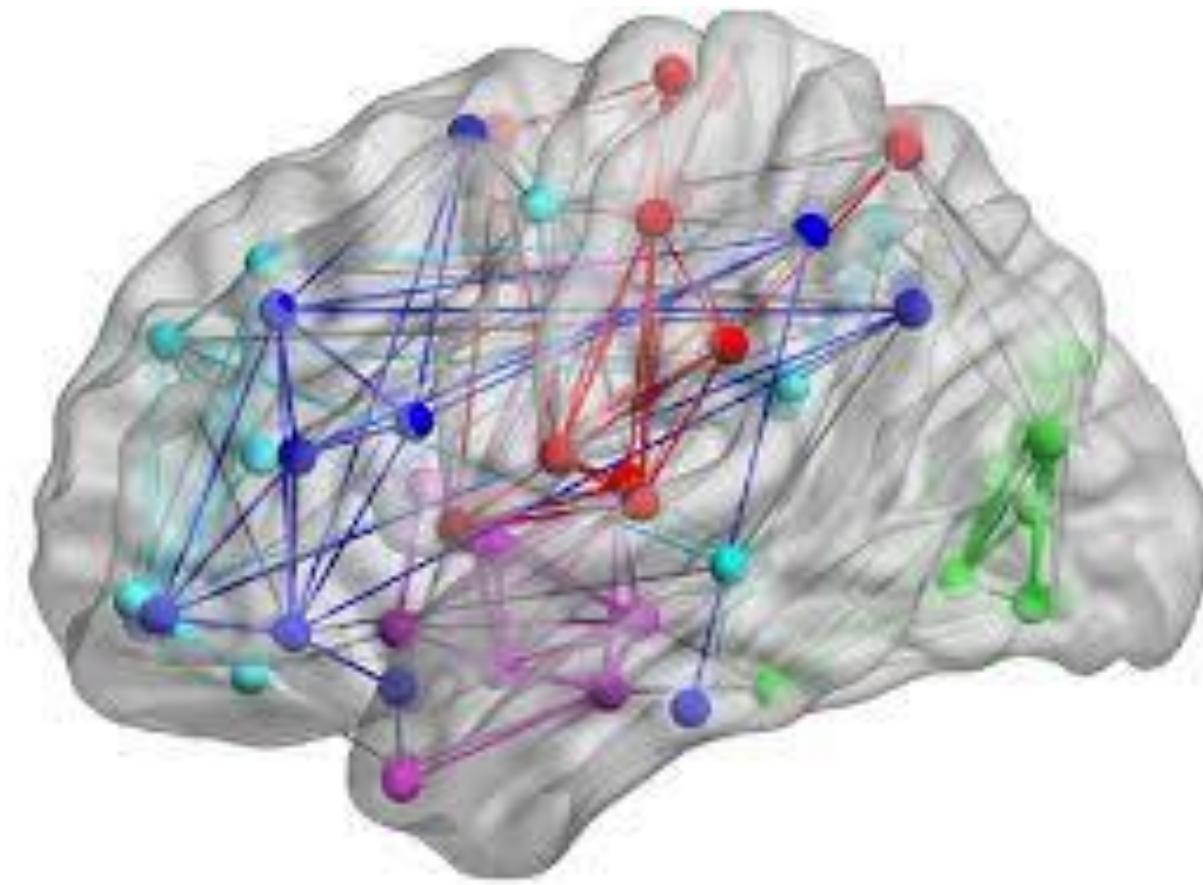
The screenshot shows the PyPI project page for the `powerlog` package. At the top, there's a navigation bar with links for Help, Docs, Sponsors, Log in, and Register. Below the header, the package name `powerlog 0.0.2` is displayed, along with a green "Latest version" button. A "pip install powerlog" button is also present. To the right, the release date "Released: Jul 10, 2025" is shown. The main content area has a light gray background and contains sections for "Navigation" (with "Project description" highlighted), "Project description" (with a "Powerlog" section), and "Verified details" (with a green checkmark icon). The "Project description" section includes a "pypi package 0.0.2" badge and a detailed description of the tool's purpose: "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." A note below states, "These details have been verified by PyPI".

Powerlog Python package (<https://pypi.org/project/powerlog/>)

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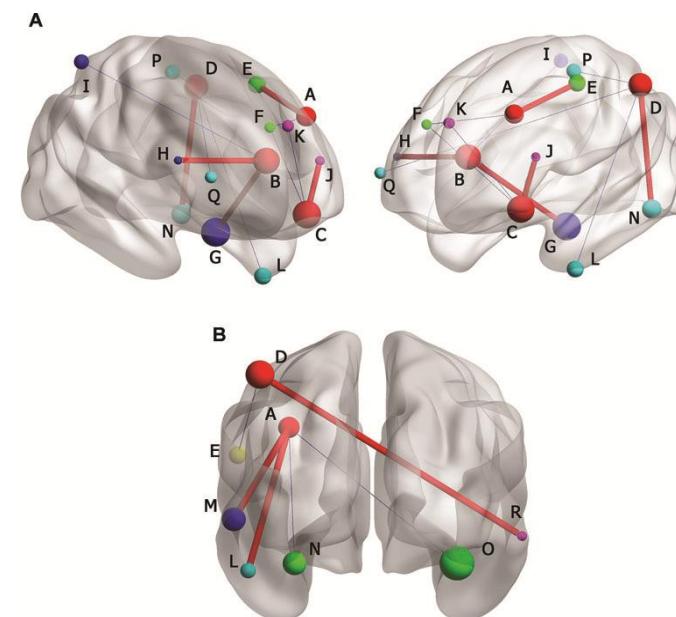
# What is a brain network?



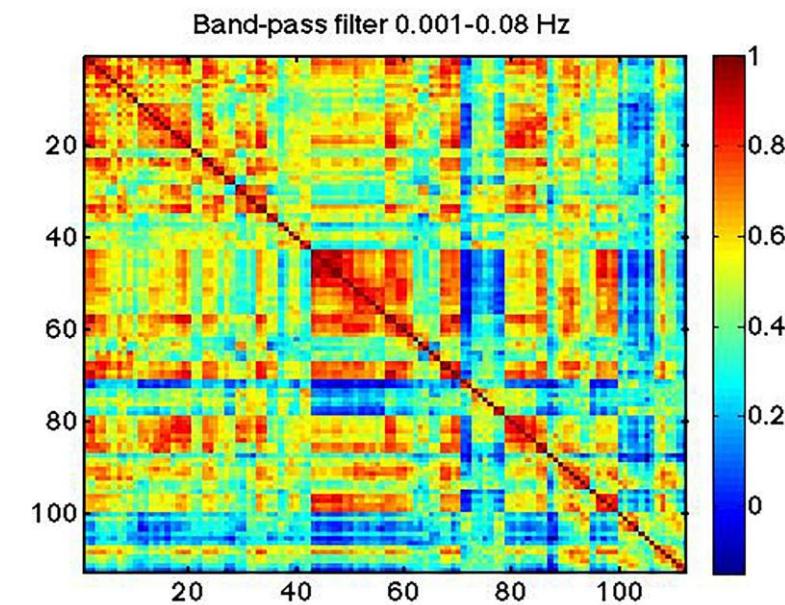
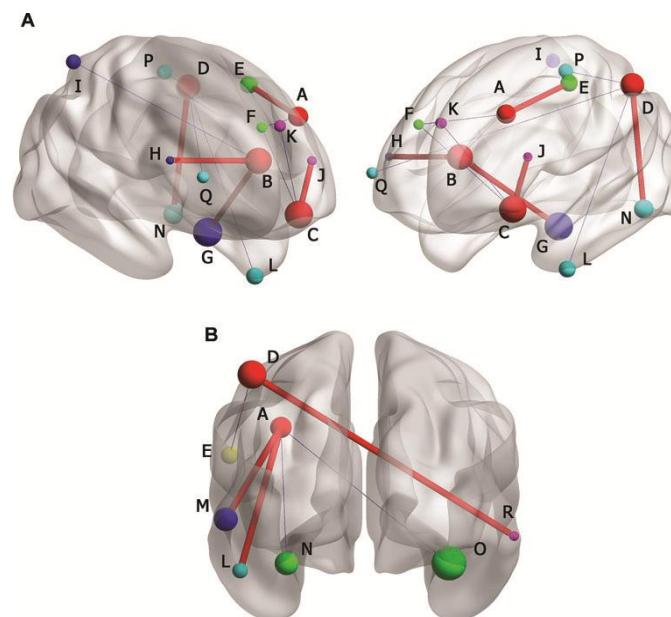
# How to construct brain network?



# How to construct brain network?



# How to construct brain network?



# Multi-site brain image analysis



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

- B. B. Biswal, M. Mennes, X.-N. Zuo, S. Gohel, C. Kelly, S. M. Smith, C. F. Beckmann, J. S. Adelstein, R. L. Buckner, S. Colcombe et al., "Toward discovery science of human brain function," *Proceedings of the national academy of sciences*, vol. 107, no. 10, pp. 4734–4739, 2010.

# Multi-site brain image analysis (continue)



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SOUTH ALABAMA

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

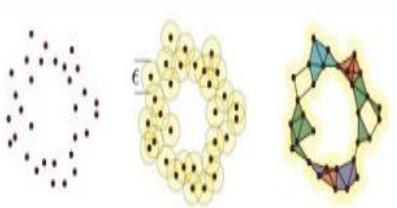
Convert fMRI  
image to FCN



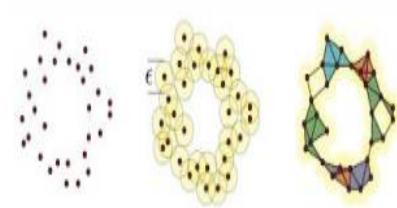
Apply Topological  
Data Analysis on  
FCN



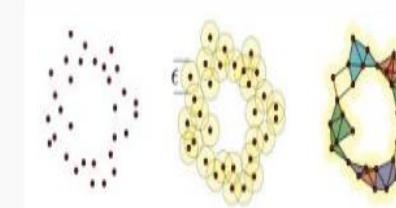
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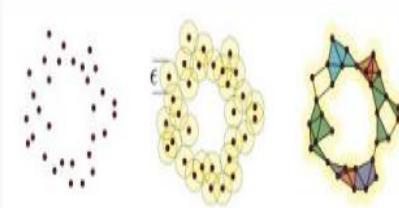
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**UIC**

# Topological Data Analysis (TDA)



Studies shape of data like connected components, loops, voids



Main tool: persistent homology, adapts homology to point cloud

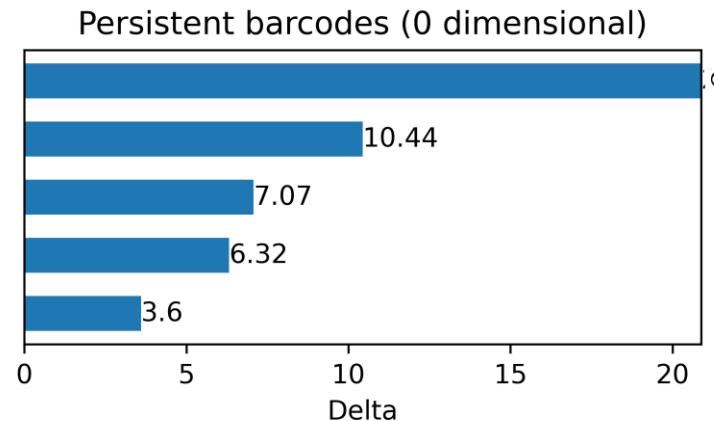


Analyzes social networks, brain connectivity, internet

# Persistent Barcodes using Connected Component

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

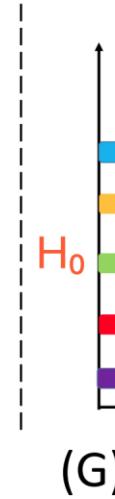
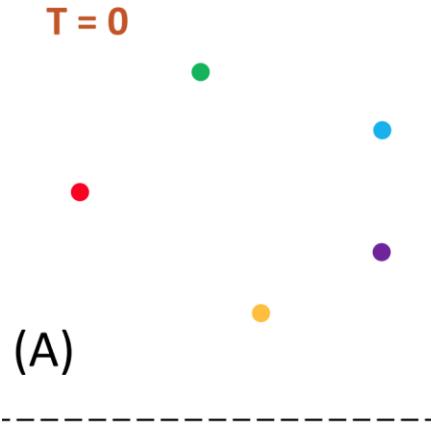
**0-dim barcodes:** each bar shows when each features (**connected component**) born and die



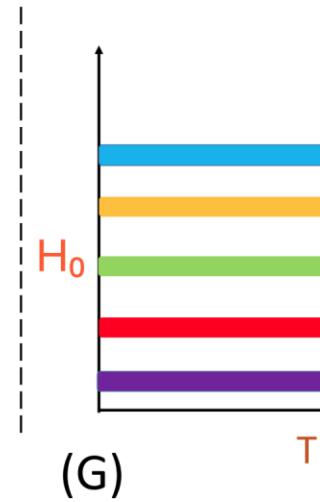
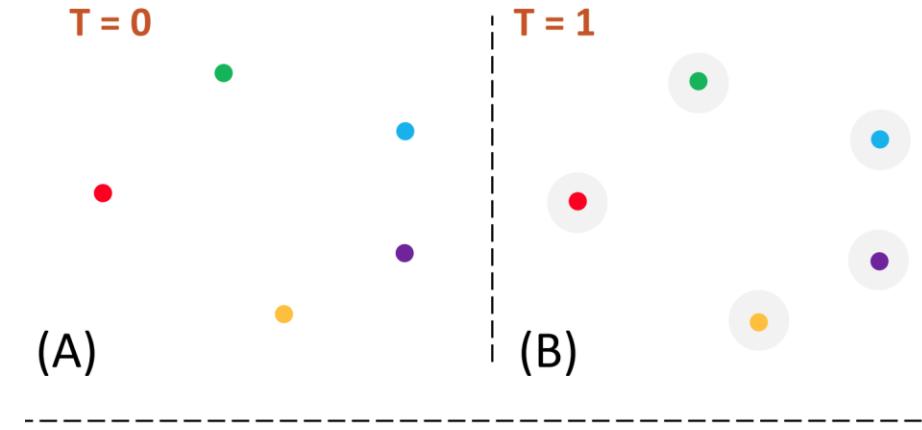
Adjacency matrix

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

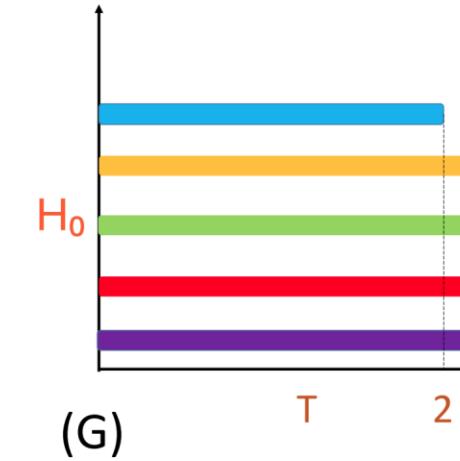
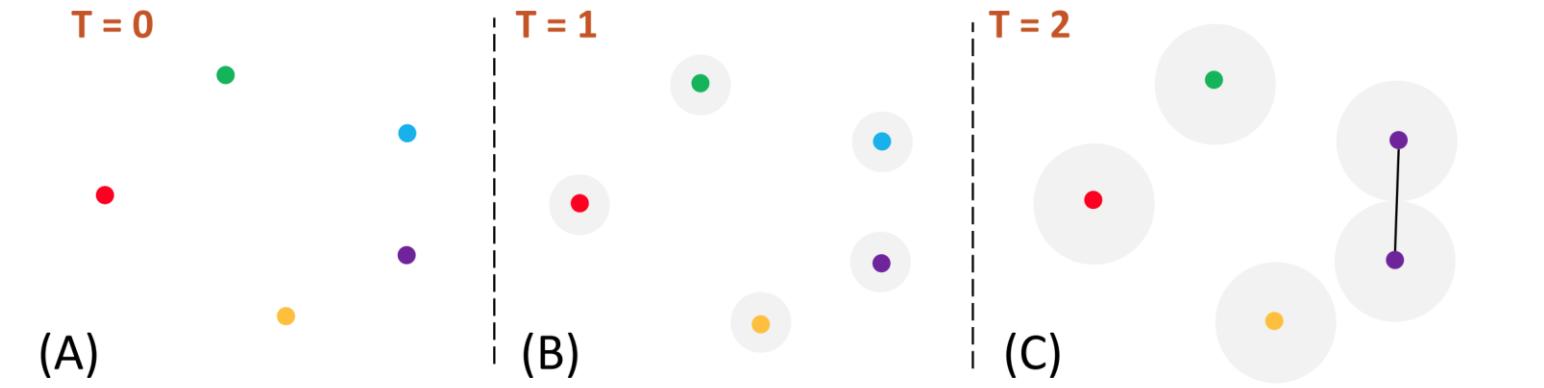
# Computing Persistent Homology (Point Cloud)

9  
1

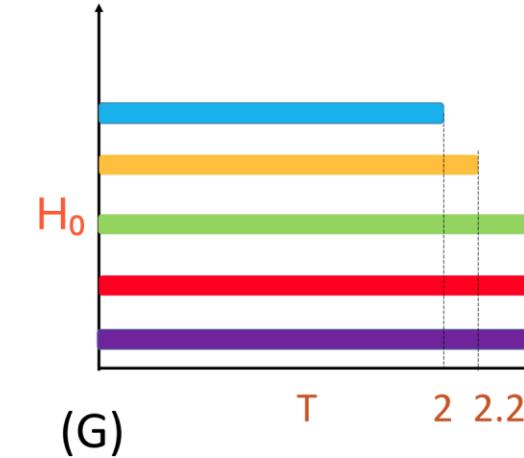
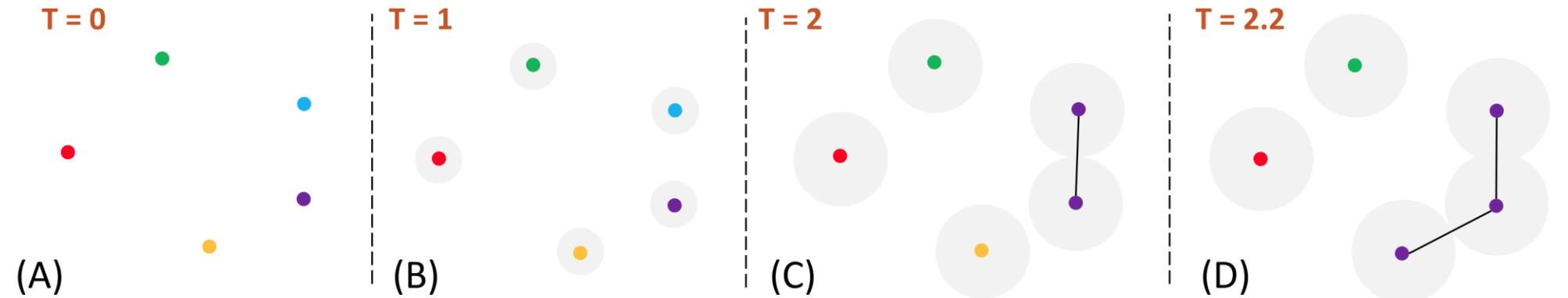
# Computing Persistent Homology (Point Cloud)



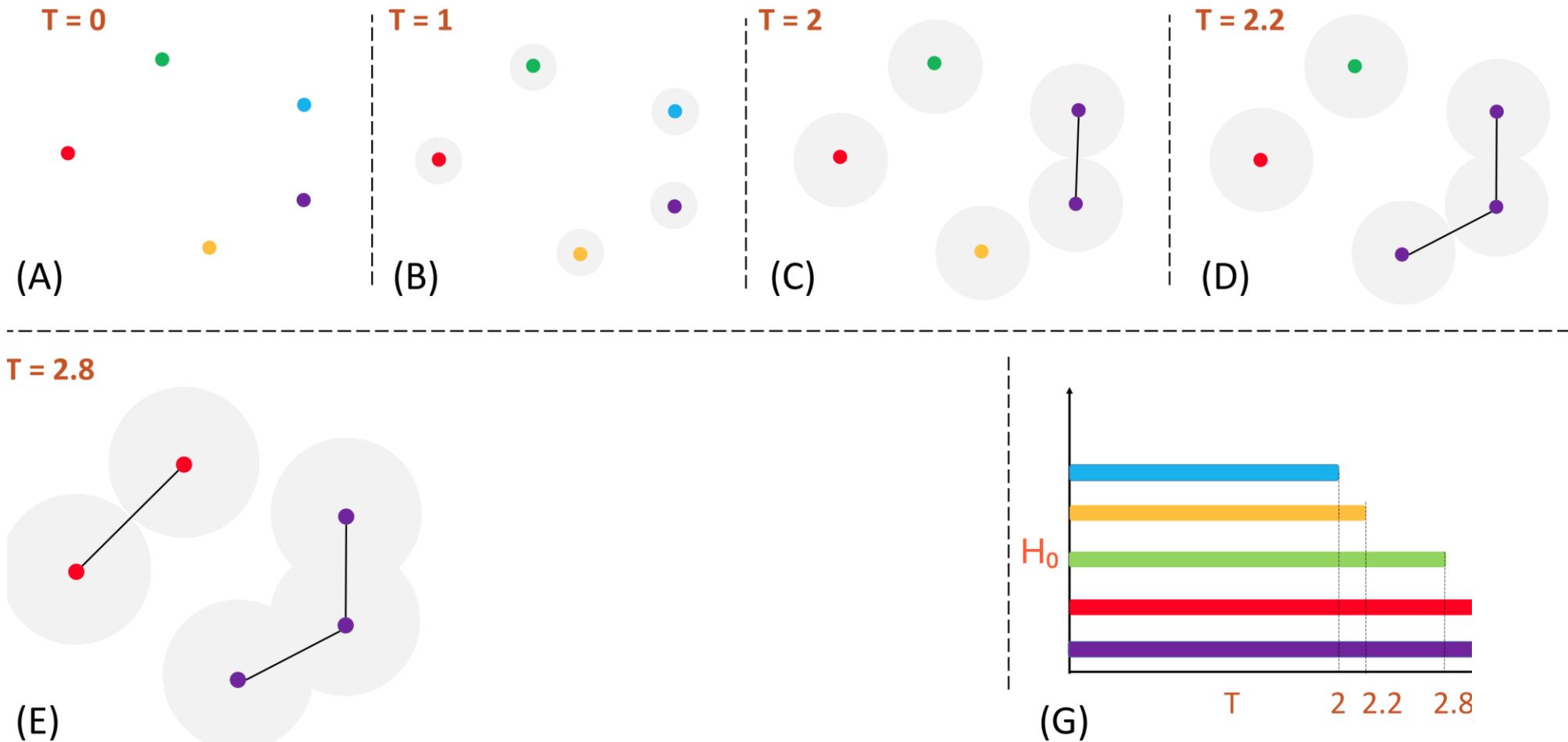
# Computing Persistent Homology (Point Cloud)



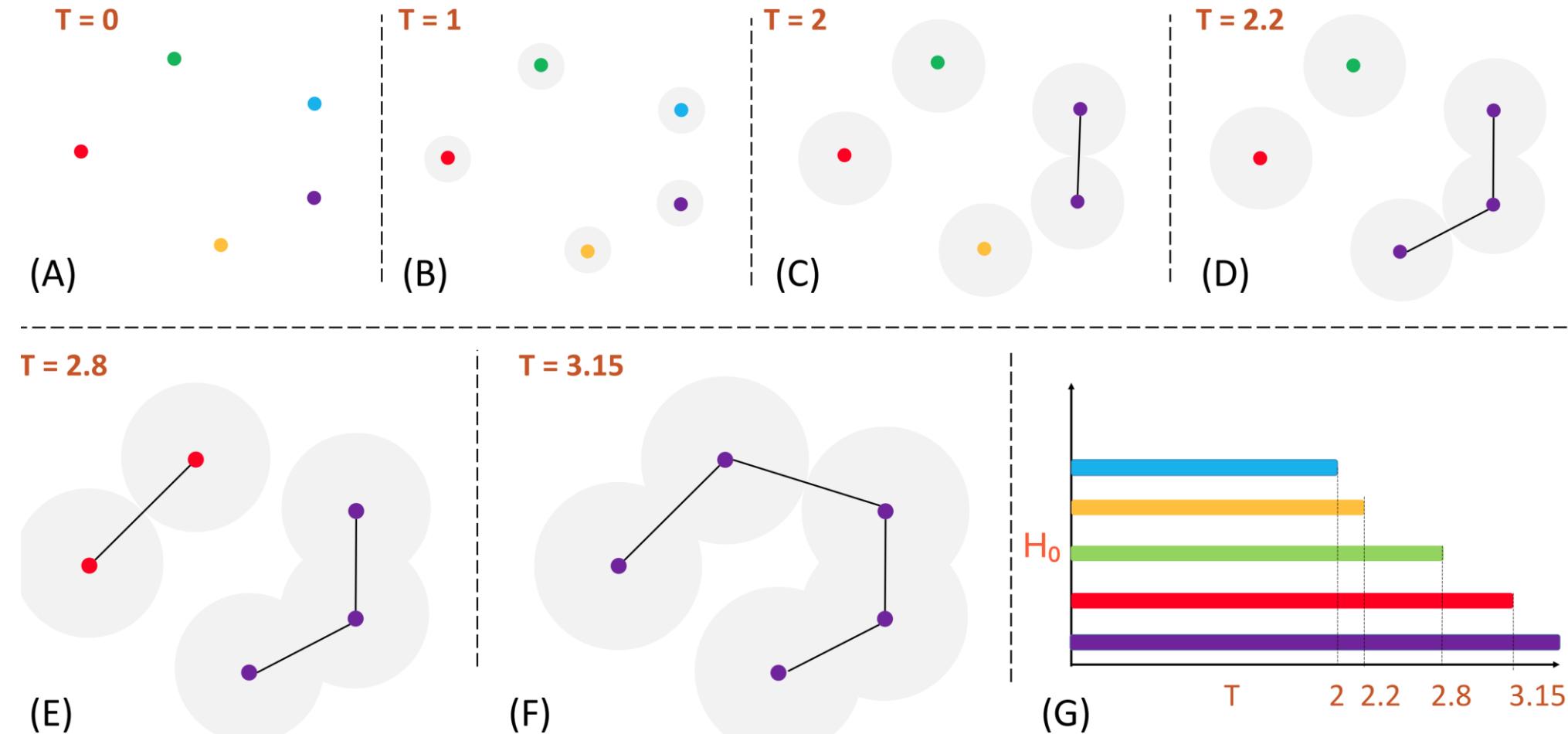
# Computing Persistent Homology (Point Cloud)

9  
4

# Computing Persistent Homology (Point Cloud)

9  
5

# Computing Persistent Homology (Point Cloud)



# PH for High-Dimensional Data Analysis

Verify

PH of rs-fMRI can represent topological features of FCNs

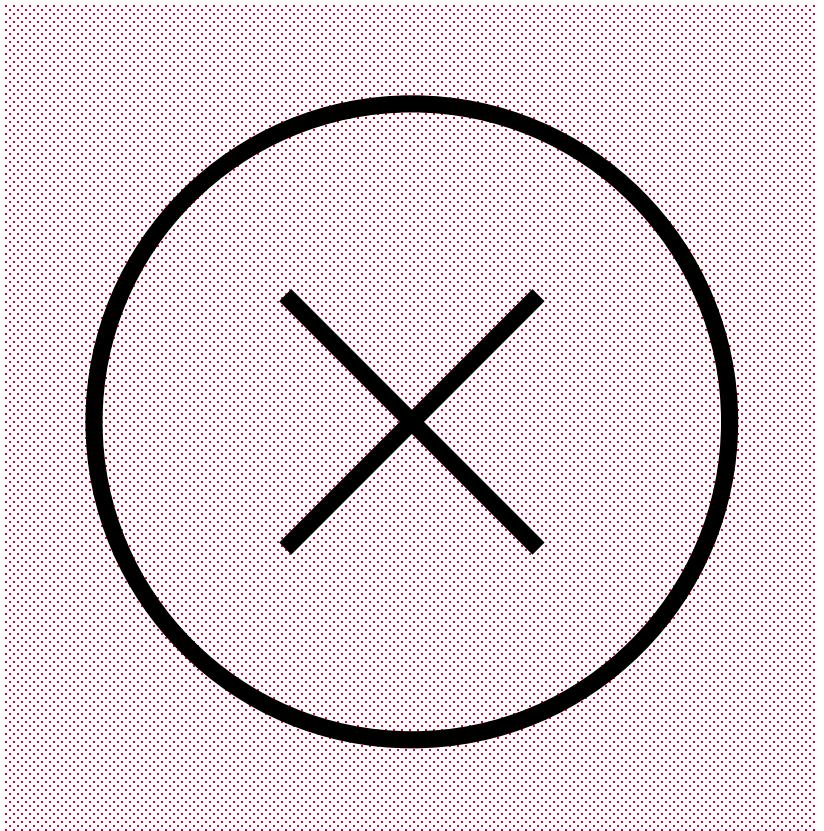
Demonstrate

FCN networks are statistically similar despite variations in TRs

Remove

Non-neural variability in multi-site brain networks

- Ashley Suh, Mustafa Hajij, Bei Wang, Carlos Scheidegger, and Paul Rosen. Persistent homology guided force-directed graph layouts. *IEEE Transactions on Visualization and Computer Graphics*, 26(1):697–707, January 2020
- Tananun Songdechakraiwit, Li Shen, and Moo Chung. Topological learning and its application to multimodal brain network integration. In *Medical Image Computing and Computer Assisted Intervention—MICCAI 2021: 24th International Conference, Strasbourg, France, September 27–October 1, 2021, Proceedings, Part II* 24, pages 166–176. Springer, 2021.



## Limitations

Imperative and complex high-dimensional  
data analysis pipelines

# Declarative Topological Data Analysis

1

Use Datalog to simplify persistent barcodes computations

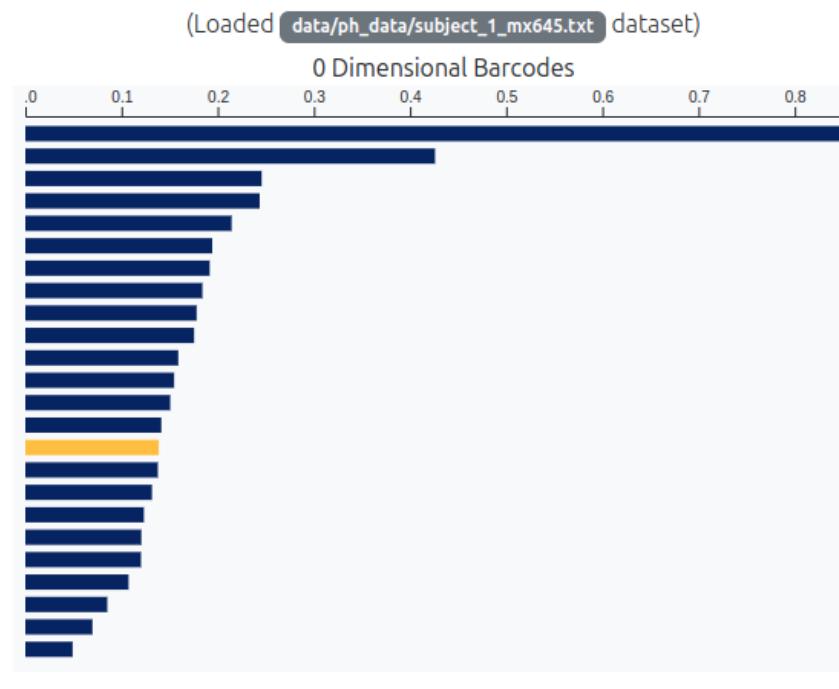
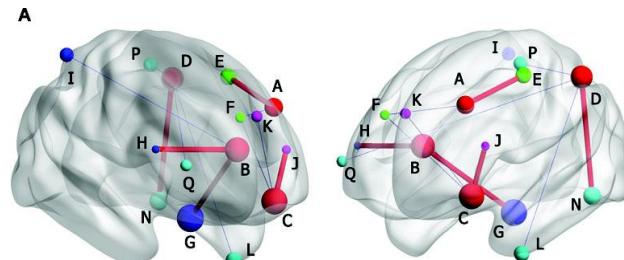
2

Explore potential applications of using declarative TDA

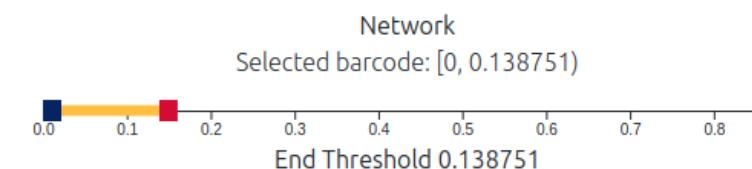
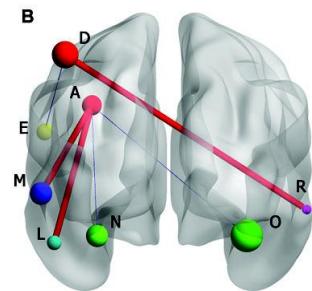
3

Scale TDA methods to large datasets in HPC environment

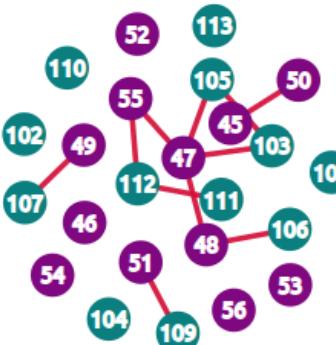
# Persistent Barcodes using Declarative Analytics



Construct a filtration



Compute persistence of connected components at each threshold

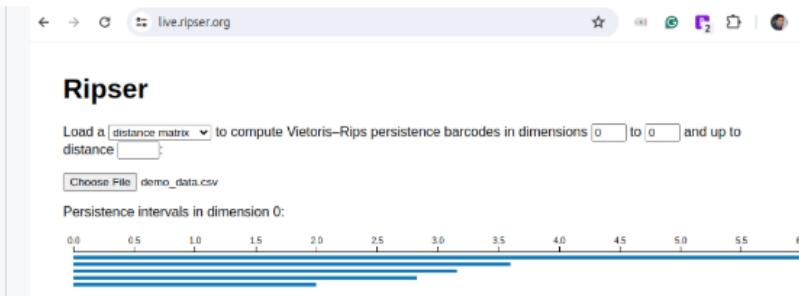


Build graph based on the selected threshold

# 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 2.830000000000001 1  
0 3.160000000000001 1  
0 3.600000000000001 1  
0 6.000000000000001 1
```



# Roadmap

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# Future/ongoing work

Enhance  
MNMGDatalog  
implementation

Extend  
declarative  
applications

Balance energy  
and  
performance

# Enhance MNMGDatalog Implementation

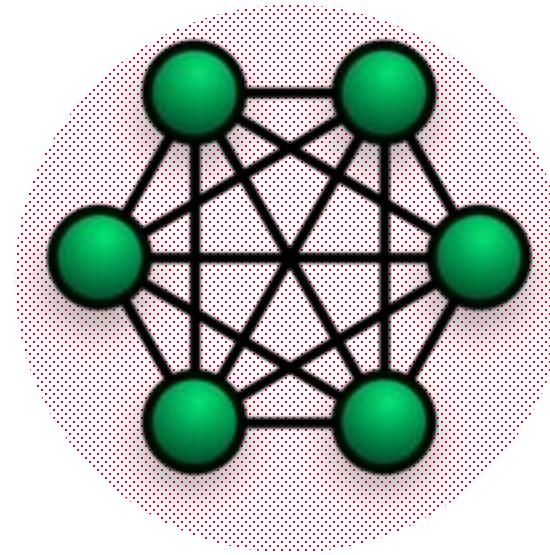
Add per-iteration checkpoint/restart capability



Develop HIP and SYCL version for MNMGDatalog

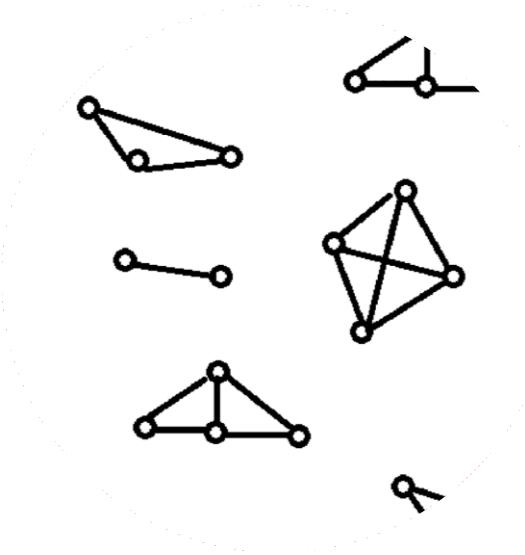


# MNMGDatalog for High-dimensional Brain Imaging



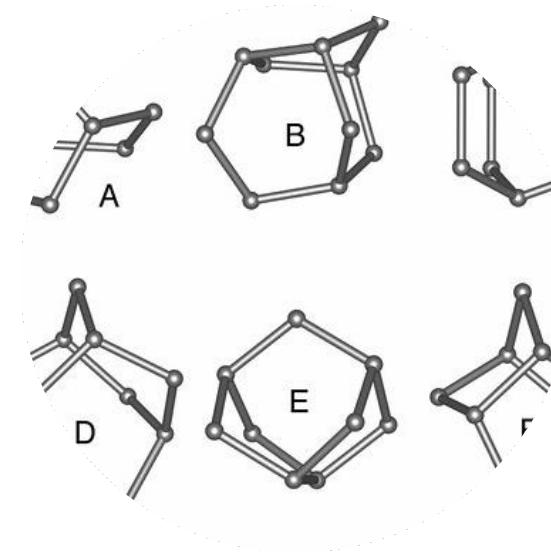
## MNMGDatalog for fully connected network

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

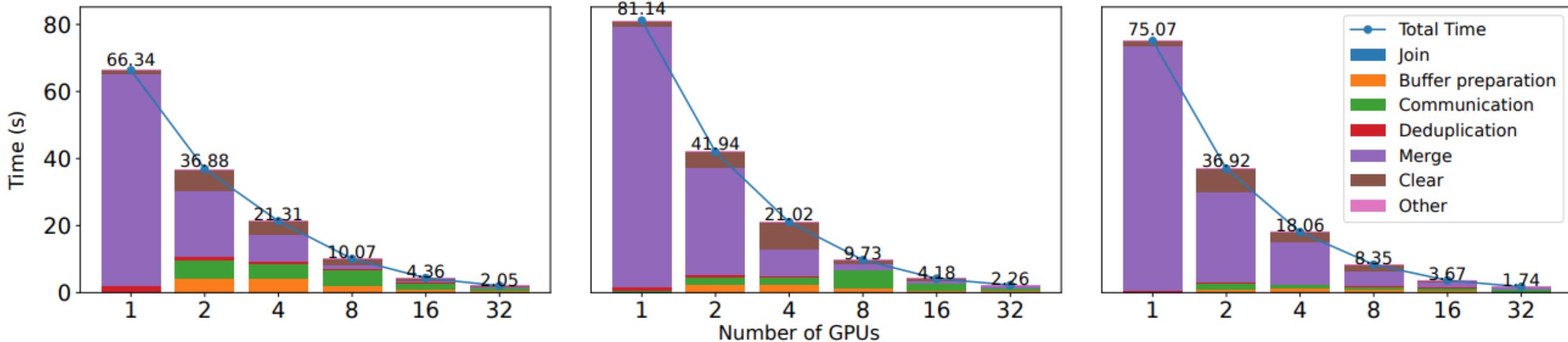


## Topological pattern matching

Discover persistent structural motifs across subjects using distributed recursive evaluation

# Power-aware Declarative Analytics

Compare energy profiles across Datalog engines



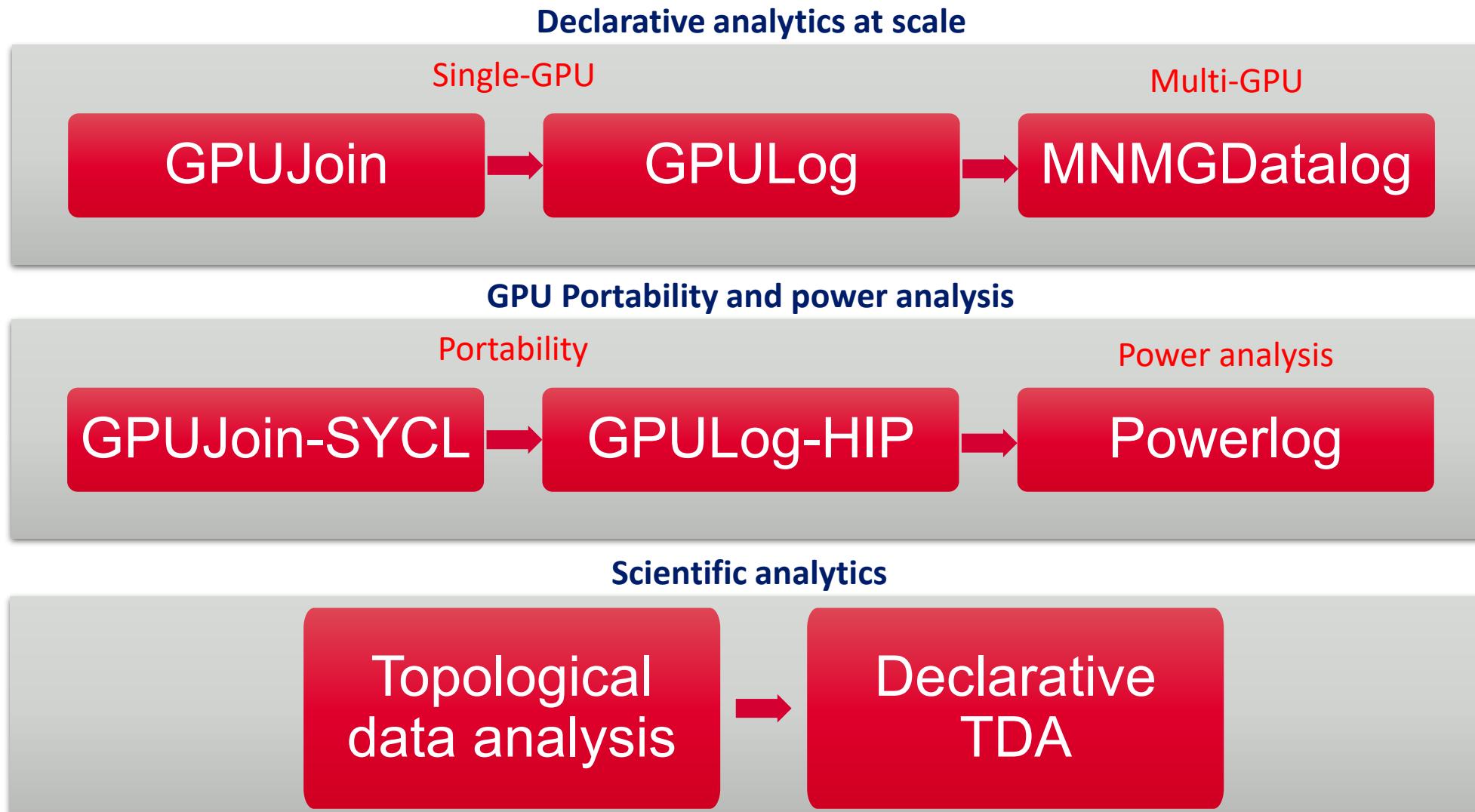
Design runtime tools to correlate iterative operations with power draw

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

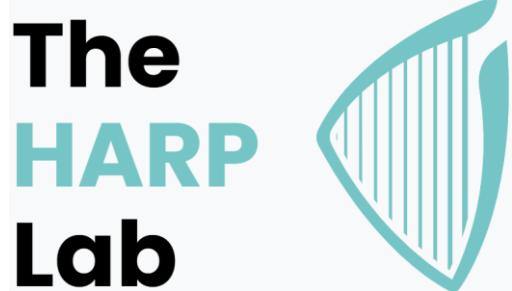
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# Conclusion



# Acknowledgements



**UAB** The University of Alabama at Birmingham.





# Thank You

[ashov@uic.edu](mailto:ashov@uic.edu)  
<https://arshovon.com/>



# 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
- $\Delta t_i$  is the time (in seconds) between samples i and i - 1

