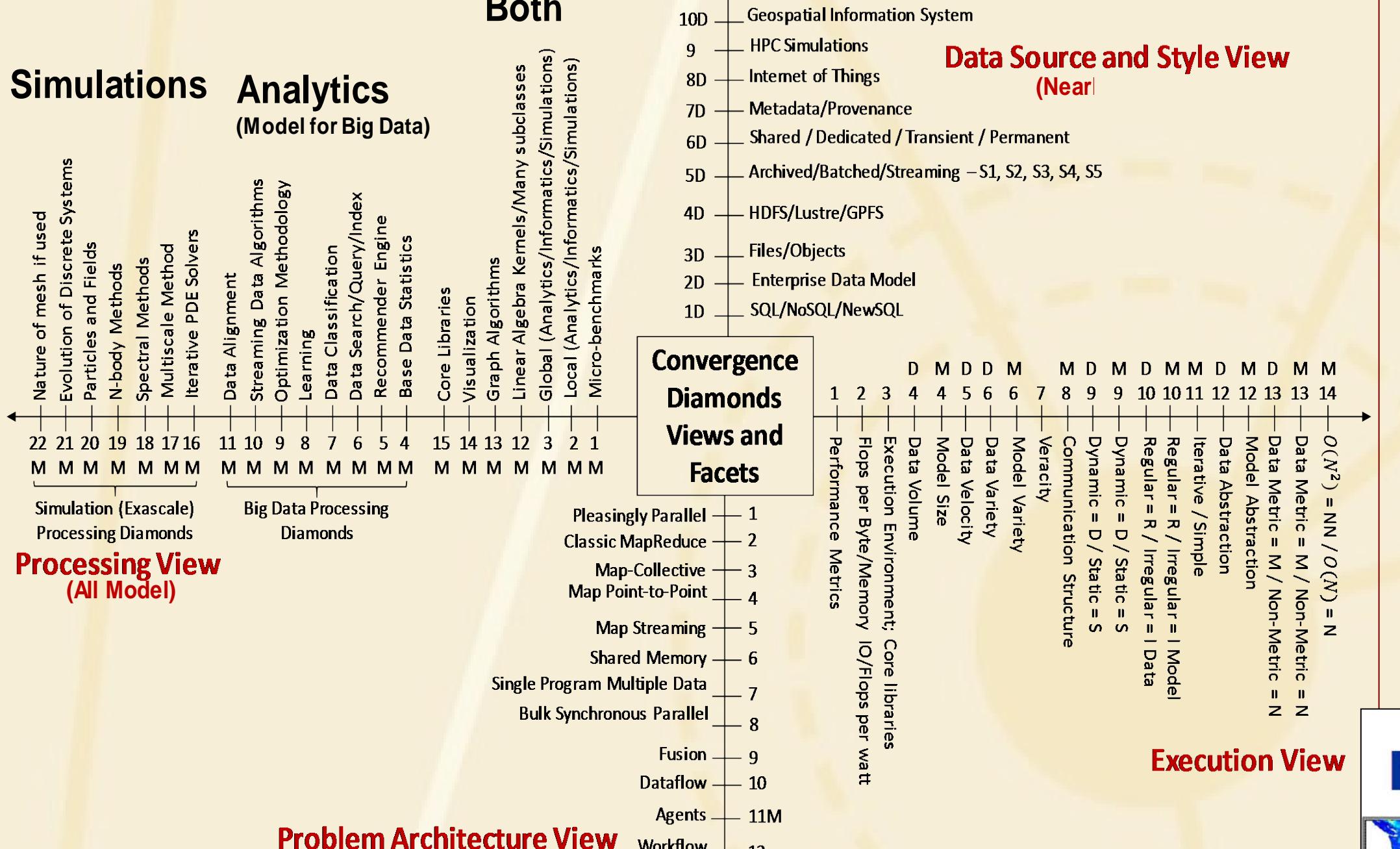


## Components of NSF 144305

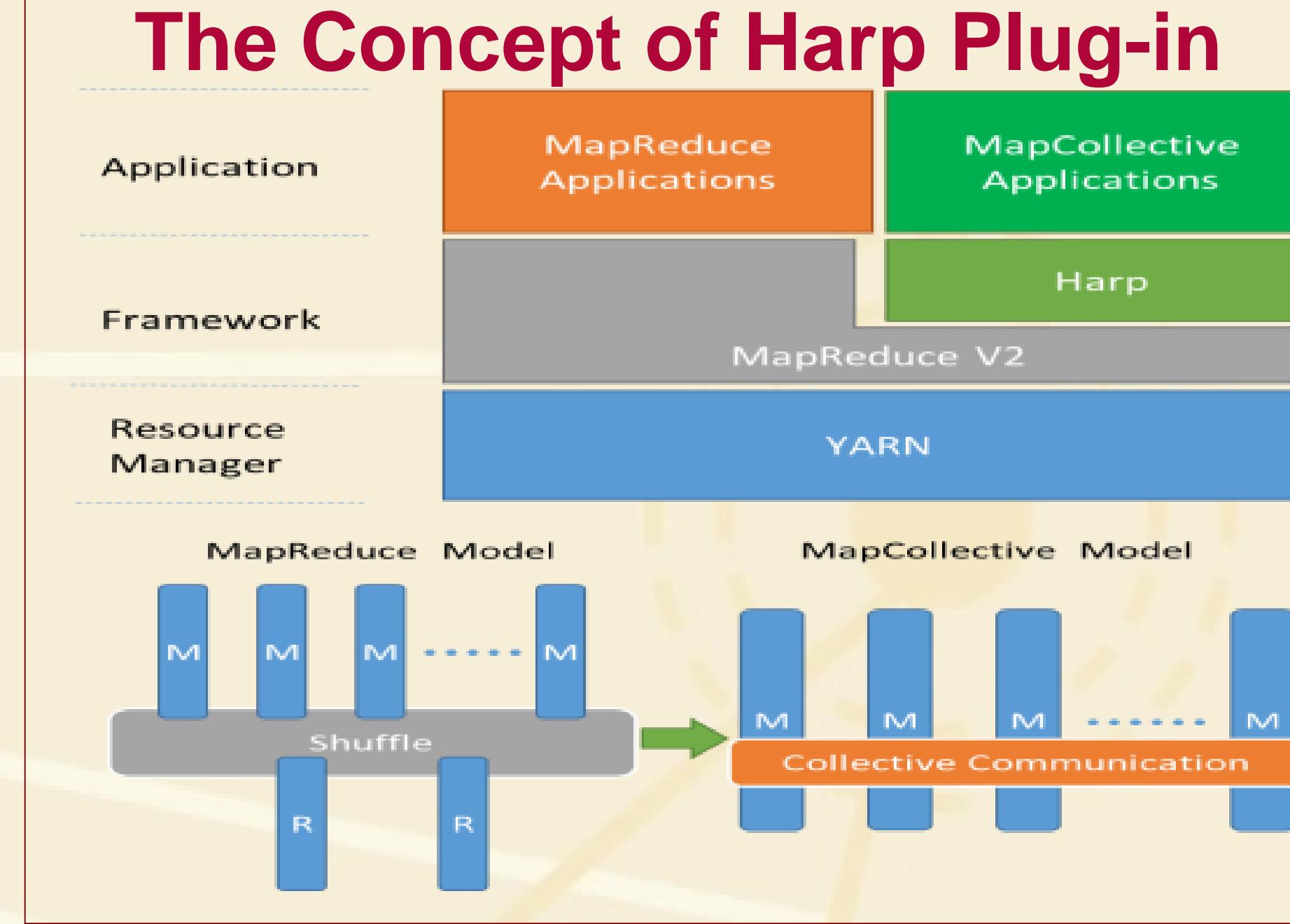
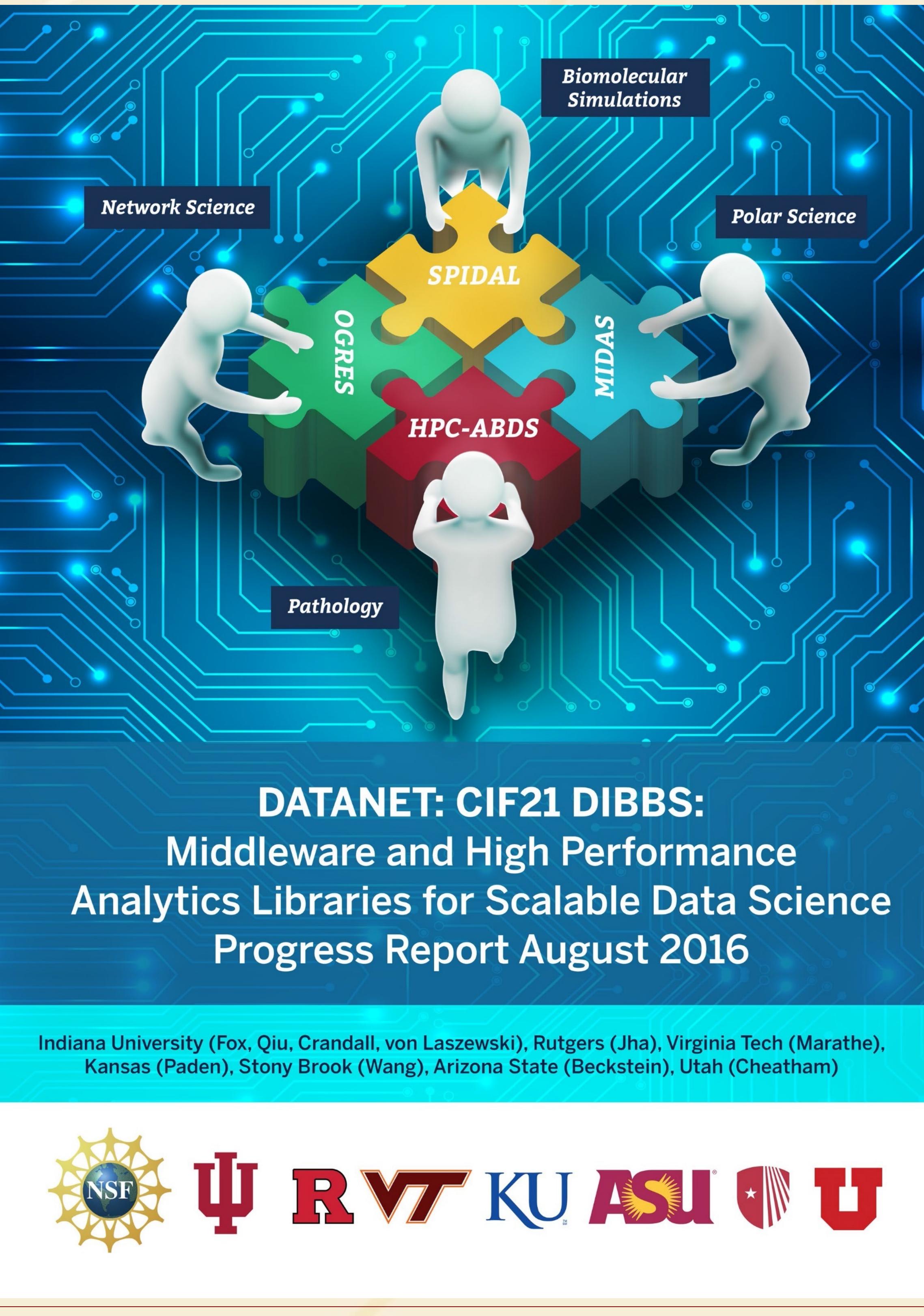
- Design and build **SPIDAL** – a Scalable Parallel Interoperable Data Analytics Library
- Domain specific libraries – mainly from project
- Core Machine Learning Libraries
- High Performance for Java, Libraries and MIDAS
- **NIST Big Data Application Analysis:** Features of data intensive Applications deriving 50 Ogres and 64 Convergence Diamonds
- HPC-ABDS: Cloud-HPC Interoperable software with performance of HPC and rich functionality of commodity Apache Stack
- Implementations: HPC and Clouds with DevOps
- Applications: Biomolecular Simulations, Network Science, Epidemiology, Computer Vision, Spatial Geographical Information Systems, Remote Sensing for Polar Science and Pathology Informatics.

## Building Blocks of Proposal

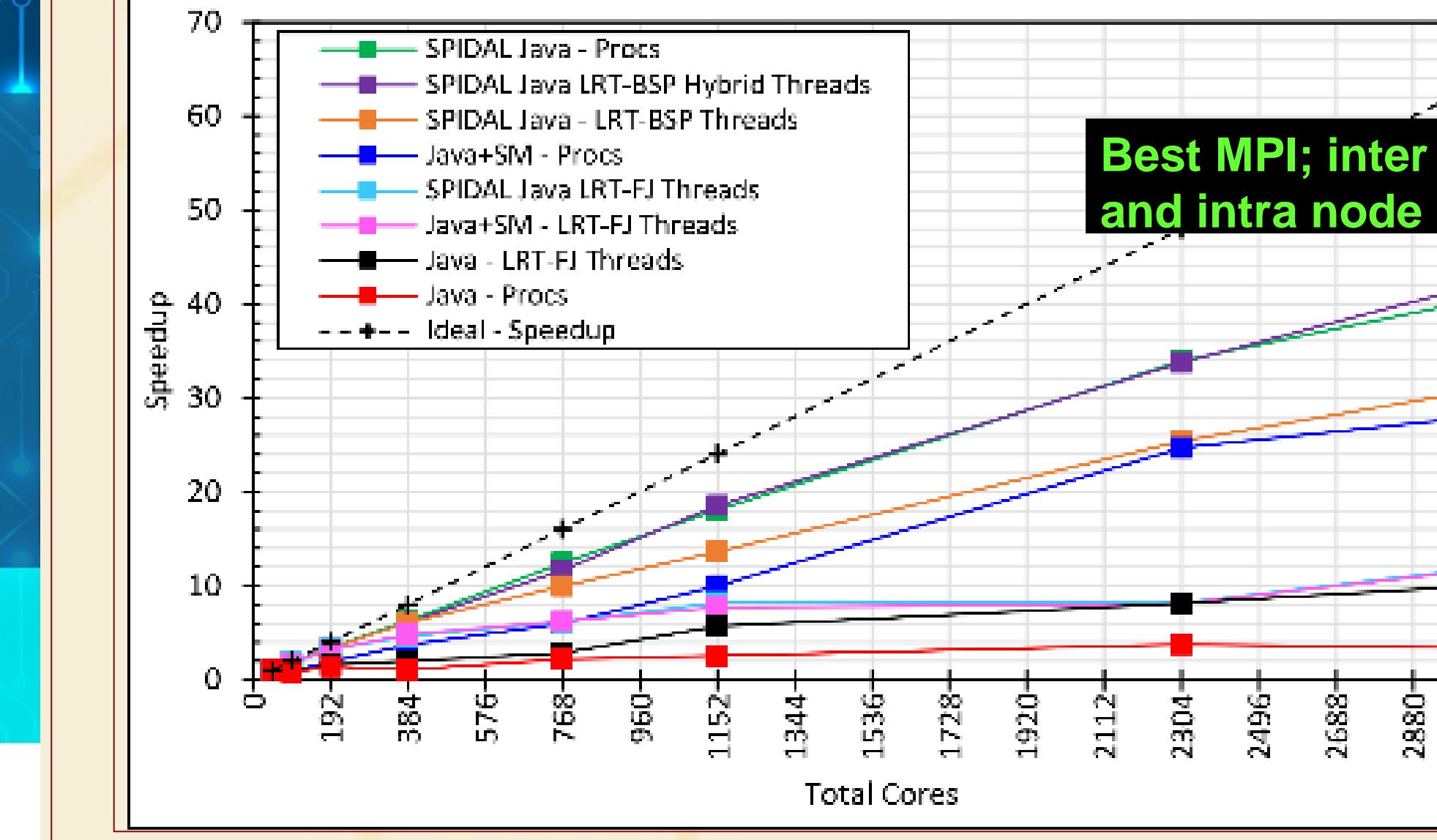
### 64 Features in 4 views for HPC-Big Data Converged Classification



### HPC-ABDS Apache Big Data Stack



### DA-MDS speedup for 200K with different optimization techniques

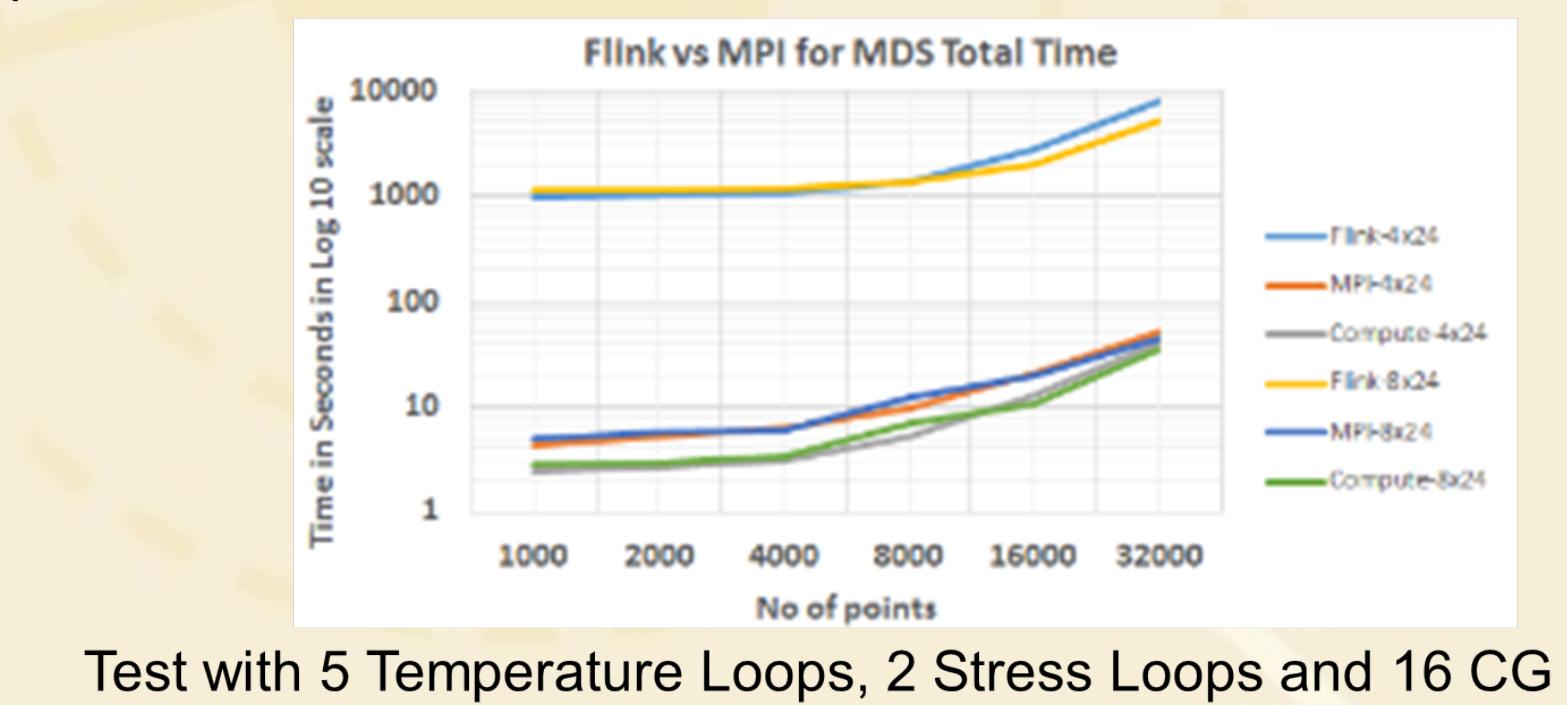


### Multidimensional Scaling with Flink

- Projects NxN distance matrix to NxD matrix where N is the number of points and D is the target dimension.
- Row wise partitioning of NxN matrix for parallelization
- Our algorithm uses deterministic annealing and has three nested loops called Temperature, Stress and CG (Conjugate Gradient) in that order.

#### Flink vs MPI DA-MDS Performance

Total time of MPI Java and Flink MDS implementations for 96 and 192 parallel tasks.

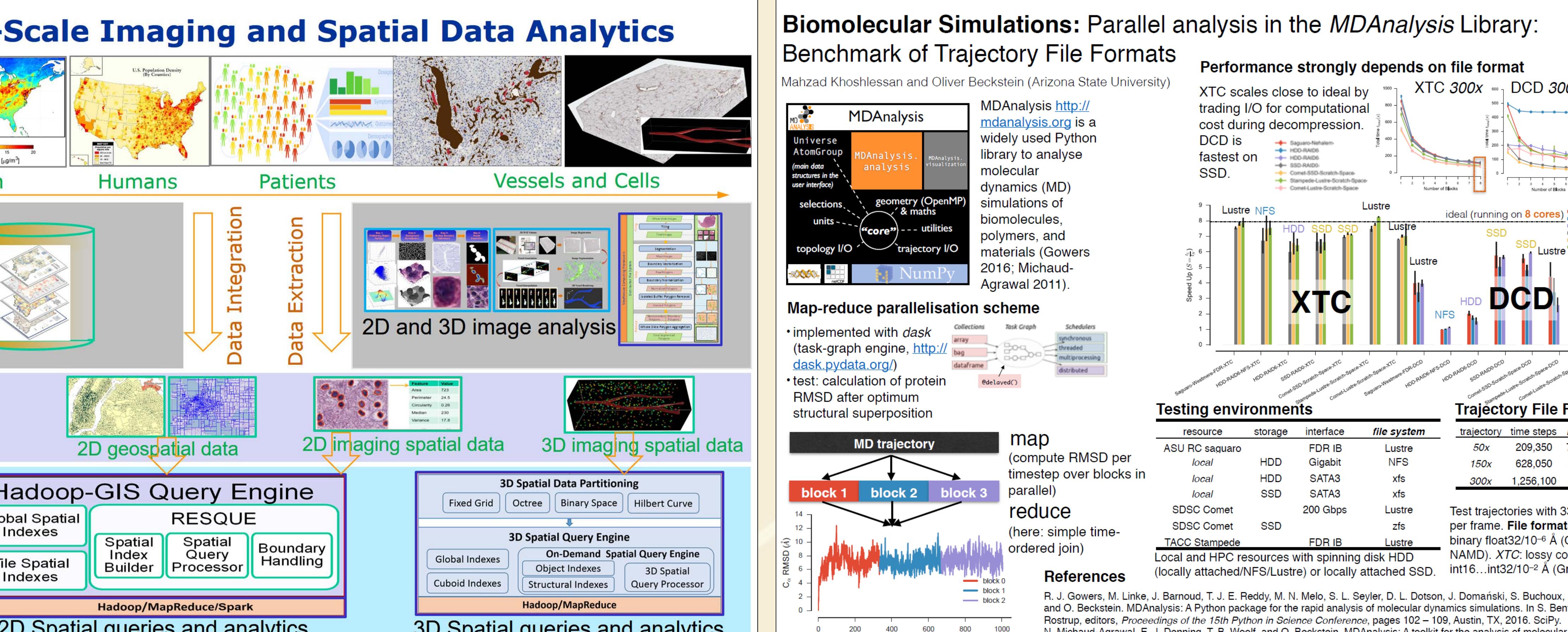


Test with 5 Temperature Loops, 2 Stress Loops and 16 CG Loops.

### MIDAS and SPIDAL Java High Performance Middleware and Language

[http://dsc.soic.indiana.edu/publications/SPIDAL-DIBBSReport\\_July2016.pdf](http://dsc.soic.indiana.edu/publications/SPIDAL-DIBBSReport_July2016.pdf)  
<http://spidal.org>

### Applications in action with MIDAS/SPIDAL



## Applications in action

### MIDAS and Biomolecular Simulations

- Previous Work (Middleware): We have created a hybrid execution framework by coupling the Pilot-abstraction with Apache Hadoop and Spark. This enhances the ability of applications to utilize HPC resources in conjunction with Hadoop frameworks concurrently or sequentially as needed.

- Current Work: (Applications) [i] We run Molecular Dynamics trajectory analysis using the hybrid execution framework, on a set of heterogeneous resources using task level parallel implementation and MapReduce programming through Spark. [ii] We are investigating the characteristics of these type of applications and the performance trade-off. Initial results show that these types of applications can greatly benefit from such an approach.

- Future Work (Streaming): Large volumes of streaming data that are real time or close to real time need to be processed or saved for later analysis and thus need high performance processing capabilities. We are integrating Kafka to handle messages, Spark for processing the data to RADICAL-Pilot to handle the deployment and the job submissions on HPC systems (Pilot-Streaming) and will employ these for HPC simulations and analysis.

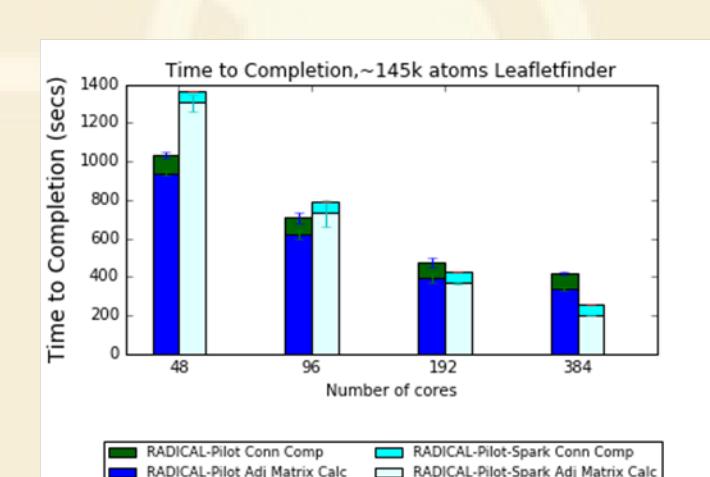
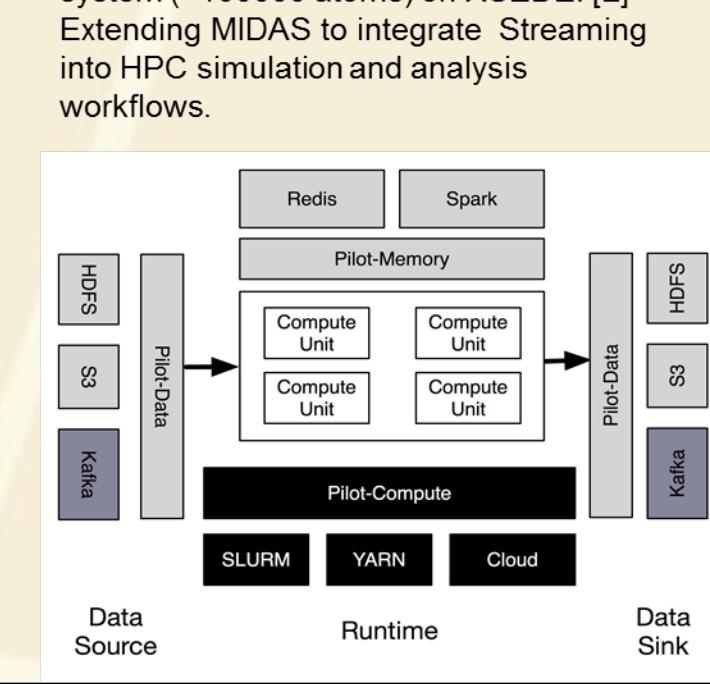
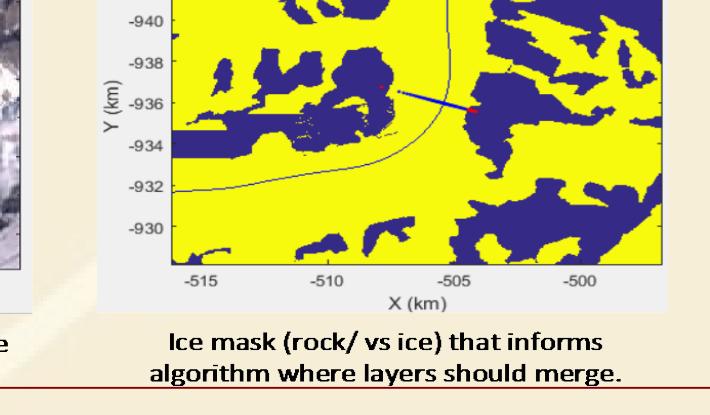
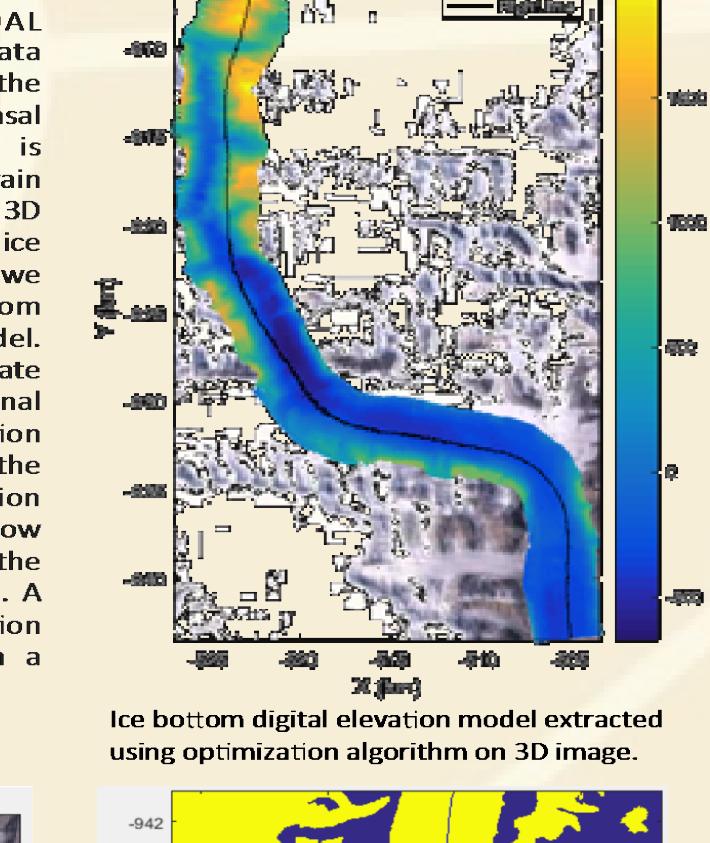


Figure caption: [U] Data from MD Trajectory analysis using MIDAS on SPARK for long trajectory for large system (>100000 atoms) on XSEDE. [L] Extending MIDAS to integrate Streaming into HPC simulation and analysis workloads.



### Polar Remote Sensing Algorithms

Many problems can be posed as mathematical optimization tasks, where the goal is to find the values for a set of variables that minimize a given objective or energy function. These often arise in the context of fitting a model to data. For instance, nearly all machine learning algorithms minimize an objective function that measures the error of the model on a set of labeled training data. Many optimization algorithms like K-means and LDA are similarly fitting model parameters to data to minimize some residual error. Many of these problems are non-convex. Many of them are simply finding a "simple" model to explain complex data, e.g., a model that compactly describes the data or an estimation of terms of ice layer structure, or a segmentation that compactly describes a high-resolution CT scan. This section describes our approach to many classes of optimization.

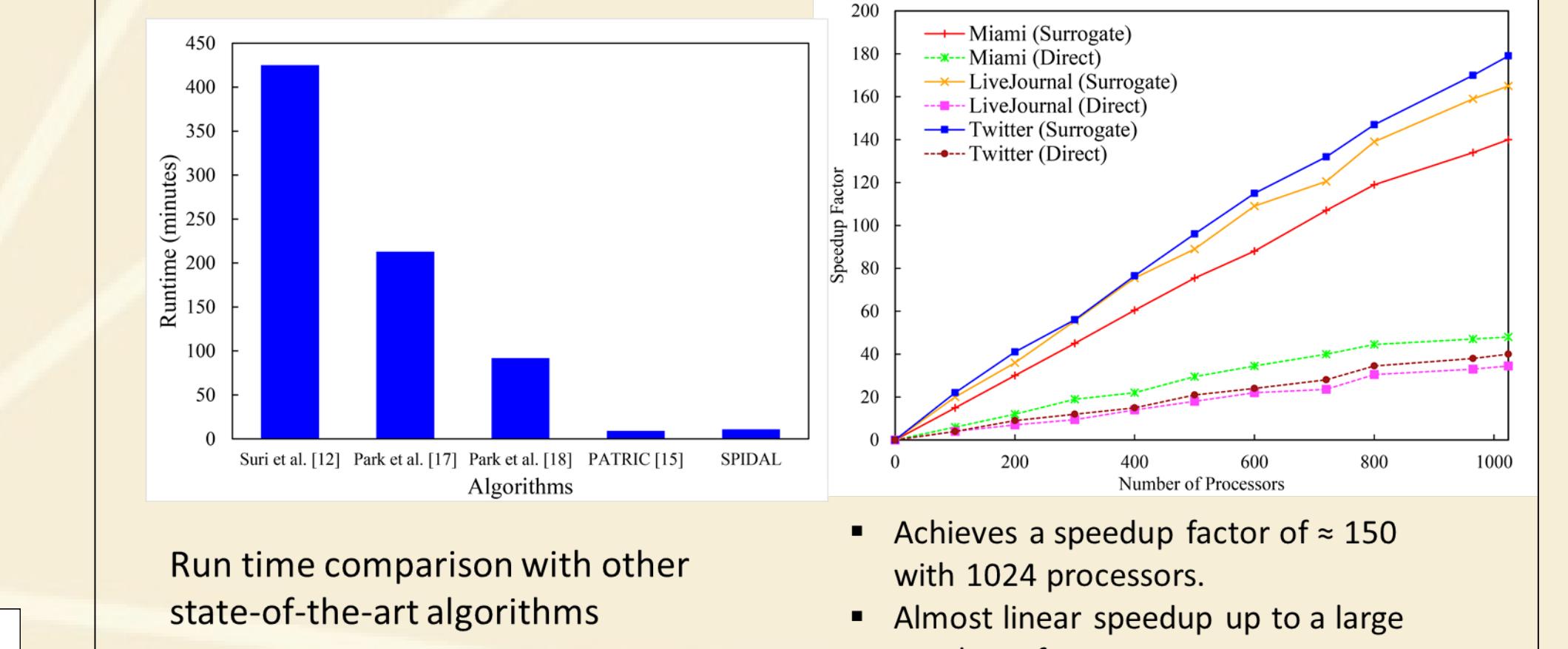
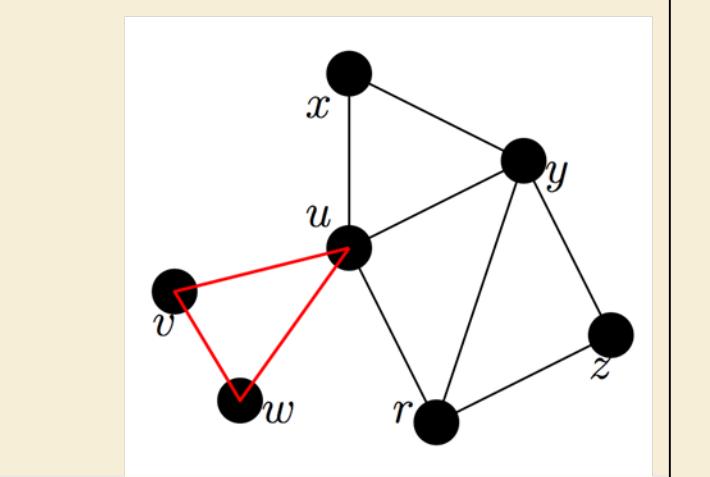


Map view showing projection of 3D image slice on a geographic map.

Ice mask (red) w/ ice) that informs algorithm where layers should merge.

### Counting Triangles in Massive Networks

Many applications in data mining, network analysis, social science, and database systems



### WebPlotViz – Browser Visualization of High Dimensional Data

WebPlotViz is a 2D/3D data point browser that can visualize very large volumes of 2D or 3D data, as points in a virtual space and enable users to explore the virtual space interactively. WebPlotViz also includes support for Time Series Data plots

