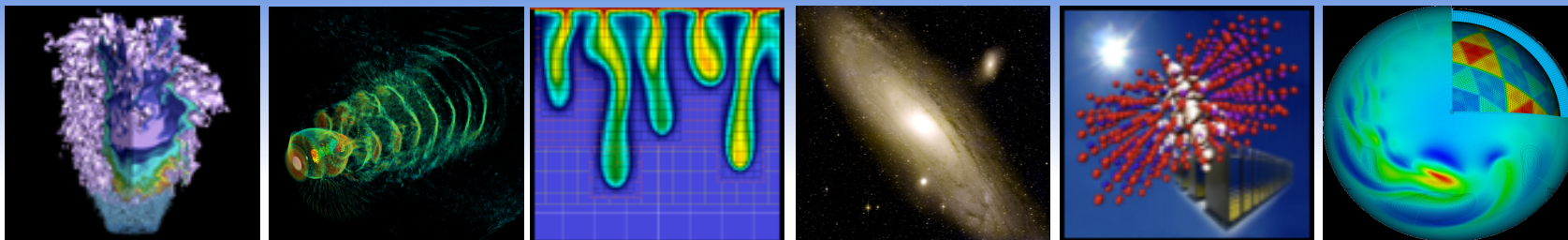


# Outline of Tutorial

- **Hadoop and Pig Overview**
- **Hands-on**



# Hadoop and Pig Overview

Lavanya Ramakrishnan  
Shane Canon

*Lawrence Berkeley National Lab*

**October 2011**

# Overview

- **Concepts & Background**
  - **MapReduce and Hadoop**
- **Hadoop Ecosystem**
  - **Tools on top of Hadoop**
- **Hadoop for Science**
  - **Examples, Challenges**
- **Programming in Hadoop**
  - **Building blocks, Streaming, C-HDFS API**

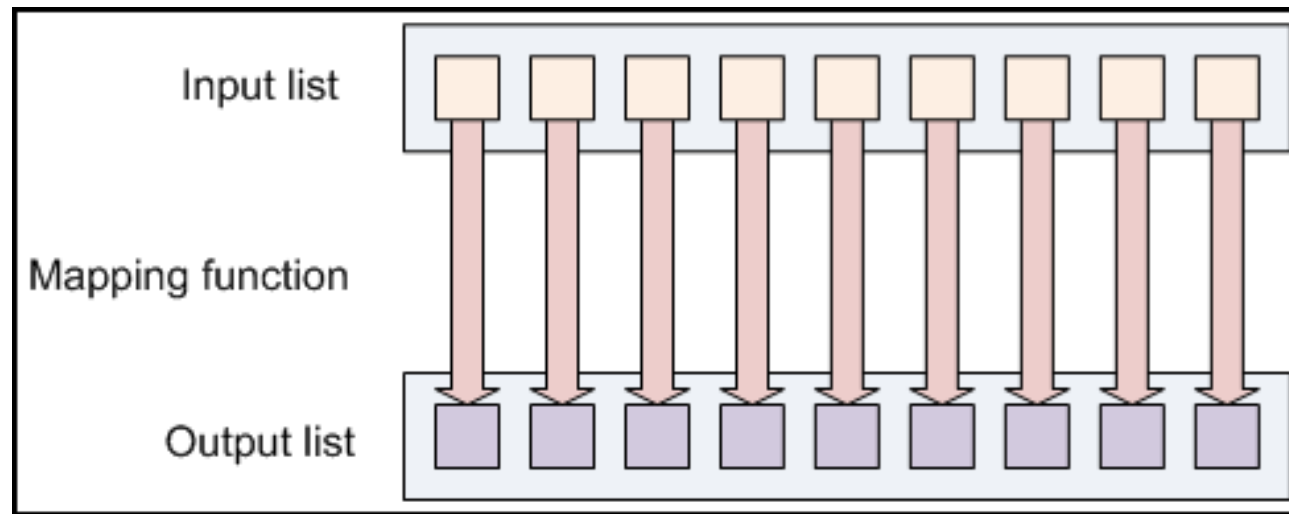
# Processing Big Data

- **Internet scale generates BigData**
  - **Terabytes of data/day**
  - **just reading 100 TB can be overwhelming**
    - **using clusters of standard commodity computers for linear scalability**
- **Timeline**
  - **Nutch open source search project (2002-2004)**
  - **MapReduce & DFS implementation and Hadoop splits out of Nutch (2004-2006)**

# MapReduce

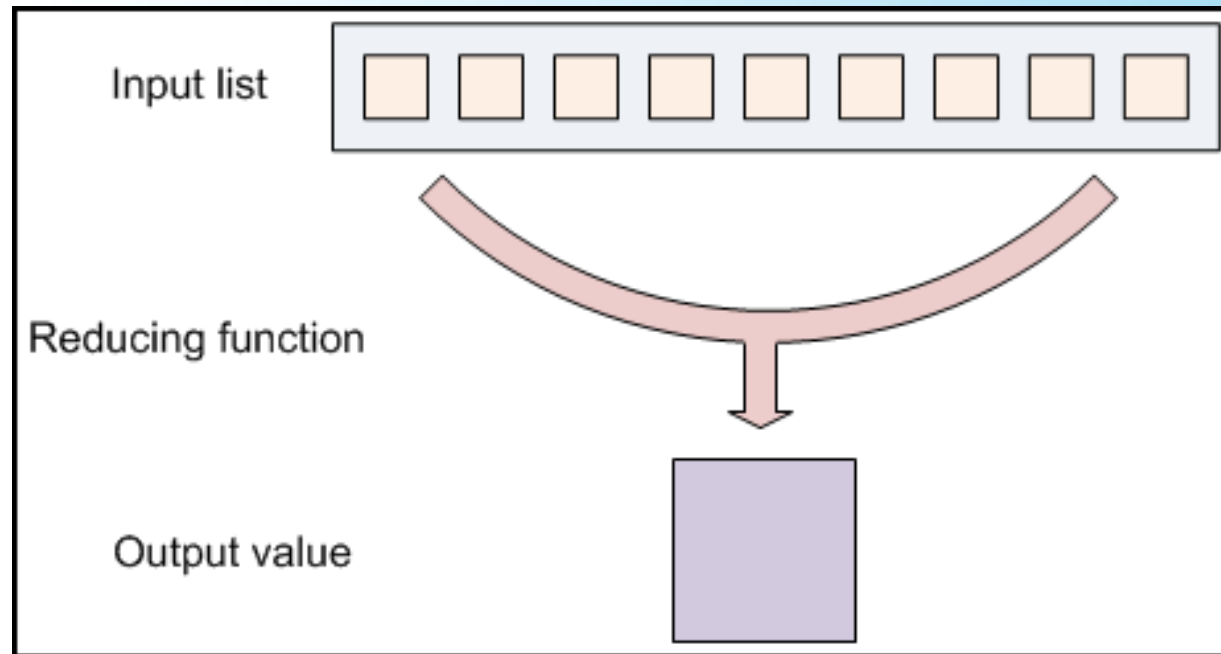
- **Computation performed on large volumes of data in parallel**
  - divide workload across large number of machines
  - need a good data management scheme to handle scalability and consistency
- **Functional programming concepts**
  - map
  - reduce

# Mapping



- Map input to an output using some function
- Example
  - string manipulation

# Reduces



- Aggregate values together to provide summary data
- Example
  - addition of the list of numbers

# Google File System

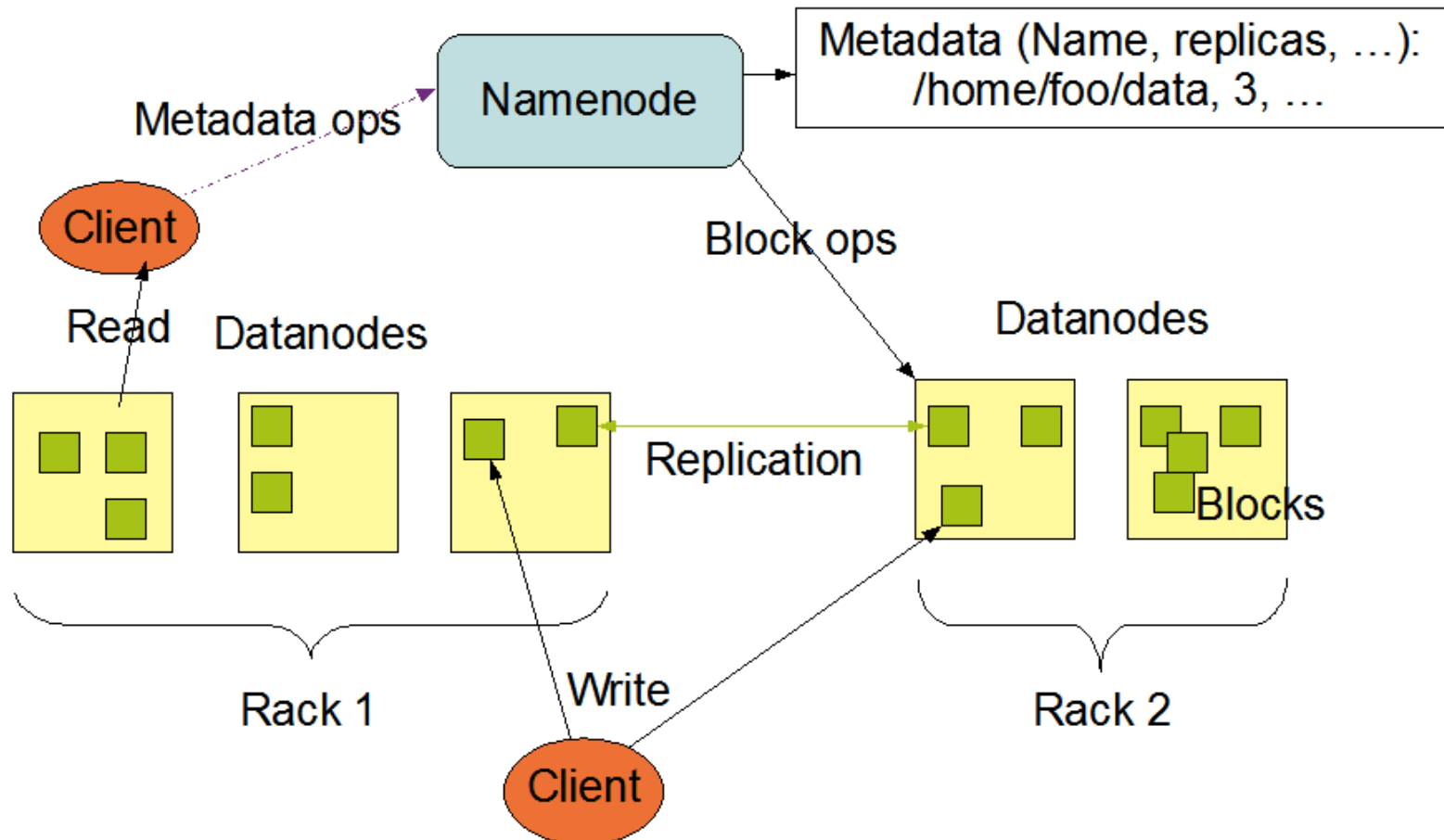
- **Distributed File System**
  - accounts for component failure
  - multi-GB files and billions of objects
- **Design**
  - single master with multiple chunkservers per master
  - file represented as fixed-sized chunks
  - 3-way mirrored across chunkservers



# Hadoop

- **Open source reliable, scalable distributed computing platform**
  - implementation of MapReduce
  - Hadoop Distributed File System (HDFS)
  - runs on commodity hardware
- **Fault Tolerance**
  - restarting tasks
  - data replication
- **Speculative execution**
  - handles stragglers

# HDFS Architecture



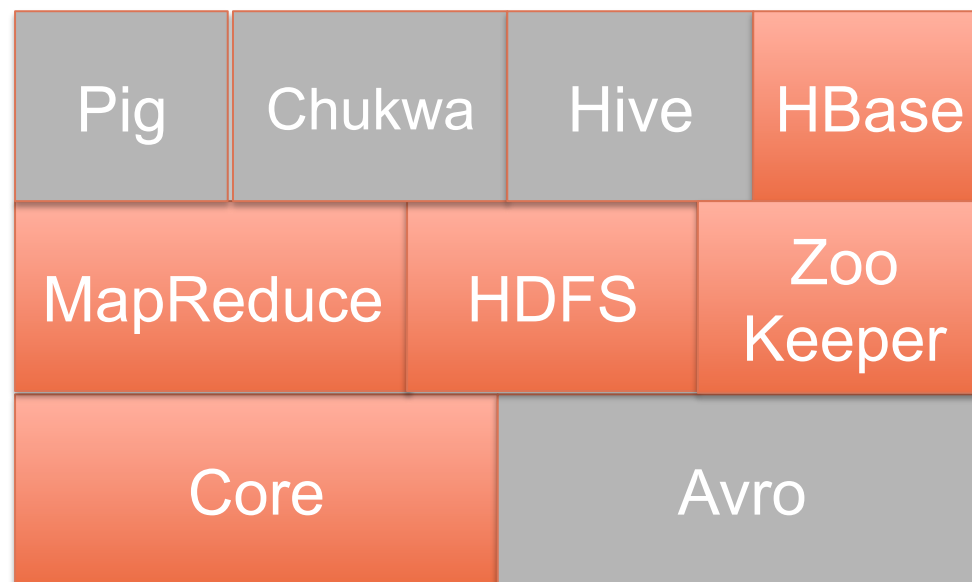
# HDFS and other Parallel Filesystems

	HDFS	GPFS and Lustre
Typical Replication	3	1
Storage Location	Compute Node	Servers
Access Model	Custom (except with Fuse)	POSIX
Stripe Size	64 MB	1 MB
Concurrent Writes	No	Yes
Scales with	# of Compute Nodes	# of Servers
Scale of Largest Systems	O(10k) Nodes	O(100) Servers
User/Kernel Space	User	Kernel

# Who is using Hadoop?

- **A9.com**
- **Amazon**
- **Adobe**
- **AOL**
- **Baidu**
- **Cooliris**
- **Facebook**
- **NSF-Google  
university initiative**
- **IBM**
- **LinkedIn**
- **Ning**
- **PARC**
- **Rackspace**
- **StumbleUpon**
- **Twitter**
- **Yahoo!**

# Hadoop Stack



Source: Hadoop: The Definitive Guide

Constantly evolving!

# Google Vs Hadoop

Google	Hadoop
MapReduce	Hadoop MapReduce
GFS	HDFS
Sawzall	Pig, Hive
BigTable	Hbase
Chubby	Zookeeper
Pregel	Hama, Giraph

# Pig

- **Platform for analyzing large data sets**
- **Data-flow oriented language “Pig Latin”**
  - data transformation functions
  - datatypes include sets, associative arrays, tuples
  - high-level language for marshalling data
- **Developed at Yahoo!**

# Hive

- **SQL-based data warehousing application**
  - features similar to Pig
  - more strictly SQL-type
- **Supports SELECT, JOIN, GROUP BY, etc**
- **Analyzing very large data sets**
  - log processing, text mining, document indexing
- **Developed at Facebook**



# HBase

- **Persistent, distributed, sorted, multidimensional, sparse map**
  - based on Google BigTable
  - provides interactive access to information
- **Holds extremely large datasets (multi-TB)**
- **High-speed lookup of individual (row, column)**

# ZooKeeper

- **Distributed consensus engine**
  - runs on a set of servers and maintains state consistency
- **Concurrent access semantics**
  - leader election
  - service discovery
  - distributed locking/mutual exclusion
  - message board/mailboxes
  - producer/consumer queues, priority queues and multi-phase commit



## Other Related Projects [1/2]

- **Chukwa – Hadoop log aggregation**
- **Scribe – more general log aggregation**
- **Mahout – machine learning library**
- **Cassandra – column store database on a P2P backend**
- **Dumbo – Python library for streaming**
- **Spark – in memory cluster for interactive and iterative**
- **Hadoop on Amazon – Elastic MapReduce**

## Other Related Projects [2/2]

- **Sqoop – import SQL-based data to Hadoop**
- **Jaql – JSON (JavaScript Object Notation) based semi-structured query processing**
- **Oozie – Hadoop workflows**
- **Giraph – Large scale graph processing on Hadoop**
- **Hcatlog – relational view of HDFS**
- **Fuse-DS – POSIX interface to HDFS**

# Hadoop for Science



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# Magellan and Hadoop

- **DOE funded project to determine appropriate role of cloud computing for DOE/SC midrange workloads**
- **Co-located at Argonne Leadership Computing Facility (ALCF) and National Energy Research Scientific Center (NERSC)**
- **Hadoop/Magellan research questions**
  - Are the new cloud programming models useful for scientific computing?

# Data Intensive Science

- **Evaluating hardware and software choices for supporting next generation data problems**
- **Evaluation of Hadoop**
  - **using mix of synthetic benchmarks and scientific applications**
  - **understanding application characteristics that can leverage the model**
    - **data operations: filter, merge, reorganization**
    - **compute-data ratio**



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(collaboration w/ Shane Canon, Nick Wright, Zacharia Fadika)



# MapReduce and HPC

- **Applications that can benefit from MapReduce/Hadoop**
  - Large amounts of data processing
  - Science that is scaling up from the desktop
  - Query-type workloads
- **Data from Exascale needs new technologies**
  - Hadoop On Demand lets one run Hadoop through a batch queue



# Hadoop for Science

- **Advantages of Hadoop**
  - transparent data replication, data locality aware scheduling
  - fault tolerance capabilities
- **Hadoop Streaming**
  - allows users to plug any binary as maps and reduces
  - input comes on standard input

# BioPig

- **Analytics toolkit for Next-Generation Sequence Data**
- **User defined functions (UDF) for common bioinformatics programs**
  - **BLAST, Velvet**
  - **readers and writers for FASTA and FASTQ**
  - **pack/unpack for space conservation with DNA sequences**

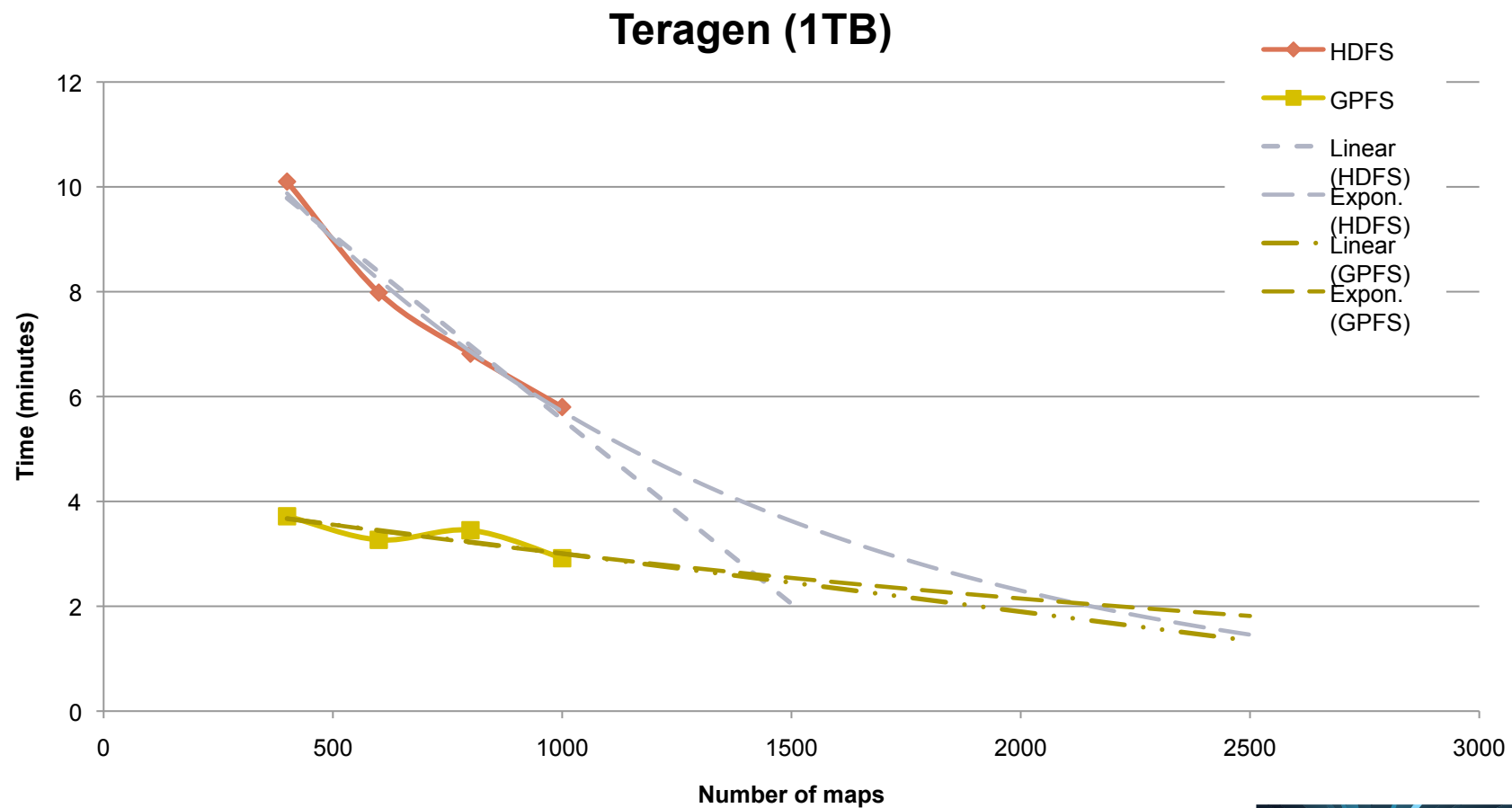
# Application Examples

- **Bioinformatics applications (BLAST)**
  - parallel search of input sequences
  - Managing input data format
- **Tropical storm detection**
  - binary file formats can't be handled in streaming
- **Atmospheric River Detection**
  - maps are differentiated on file and parameter

# “Bring your application” Hadoop workshop

- **When: TBD**
- **Send us email if you are interested**
  - [LRamakrishnan@lbl.gov](mailto:LRamakrishnan@lbl.gov)
  - [Scanon@lbl.gov](mailto:Scanon@lbl.gov)
- **Include a brief description of your application.**

# HDFS vs GPFS (Time)



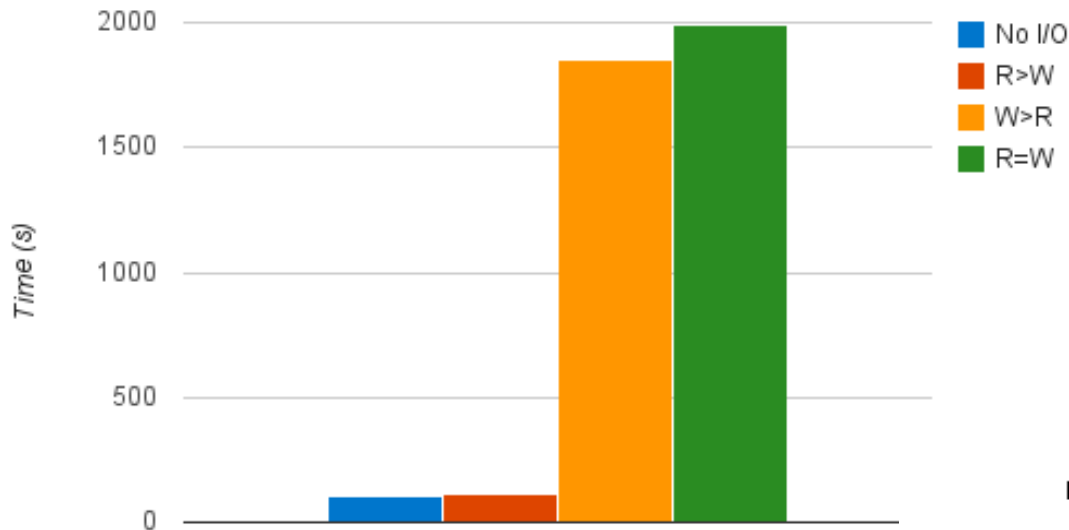
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# Application Characteristic Affect Choices

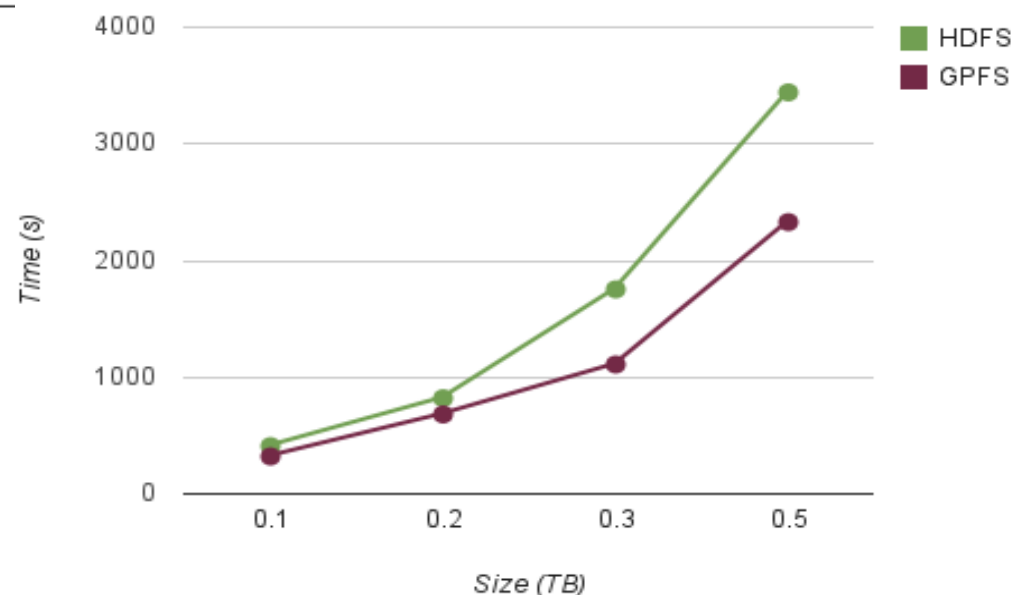
Square Number Check: 0.1TB (10,000,000,000 Numbers)



- Identical data loads and processing load
- Amount of writing in application affects performance

**Wikipedia data set**  
**On ~ 75 nodes,**  
**GPFS performs**  
**better with large**  
**nodes**

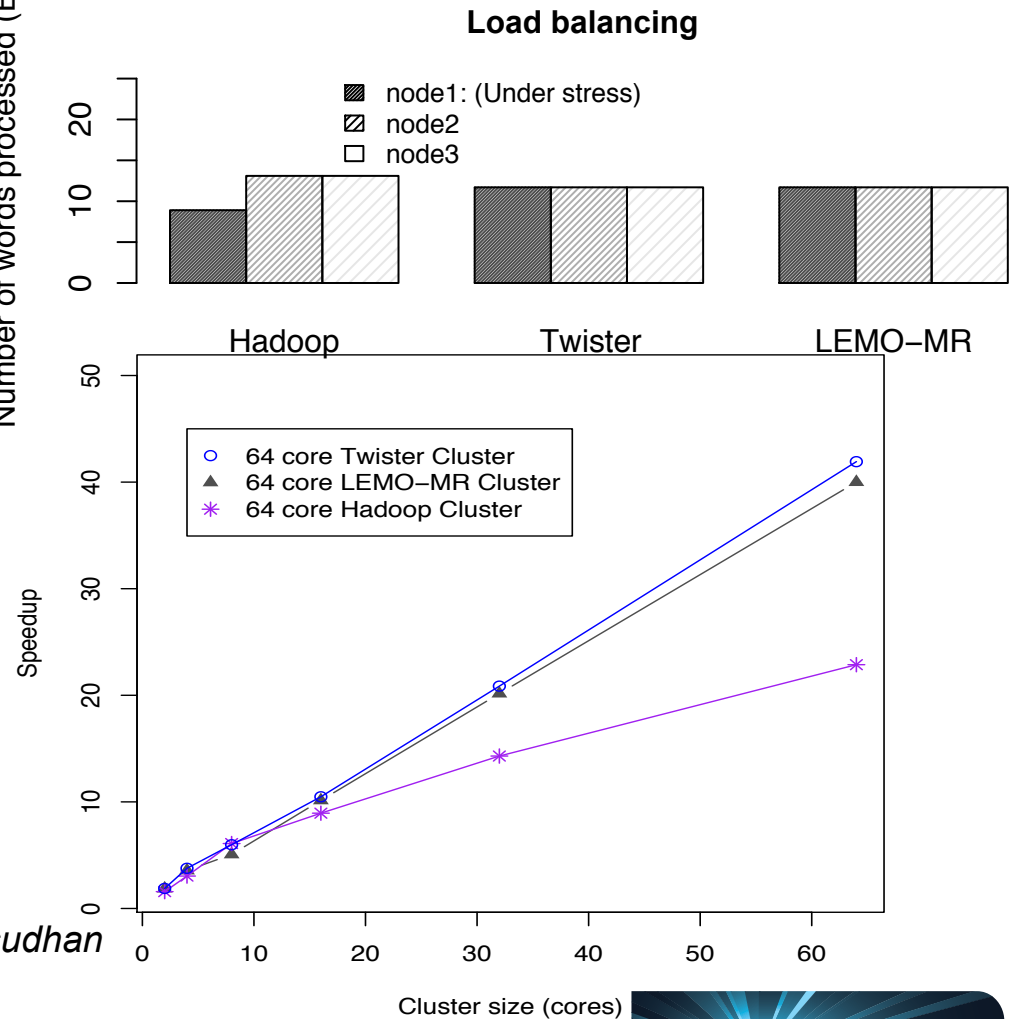
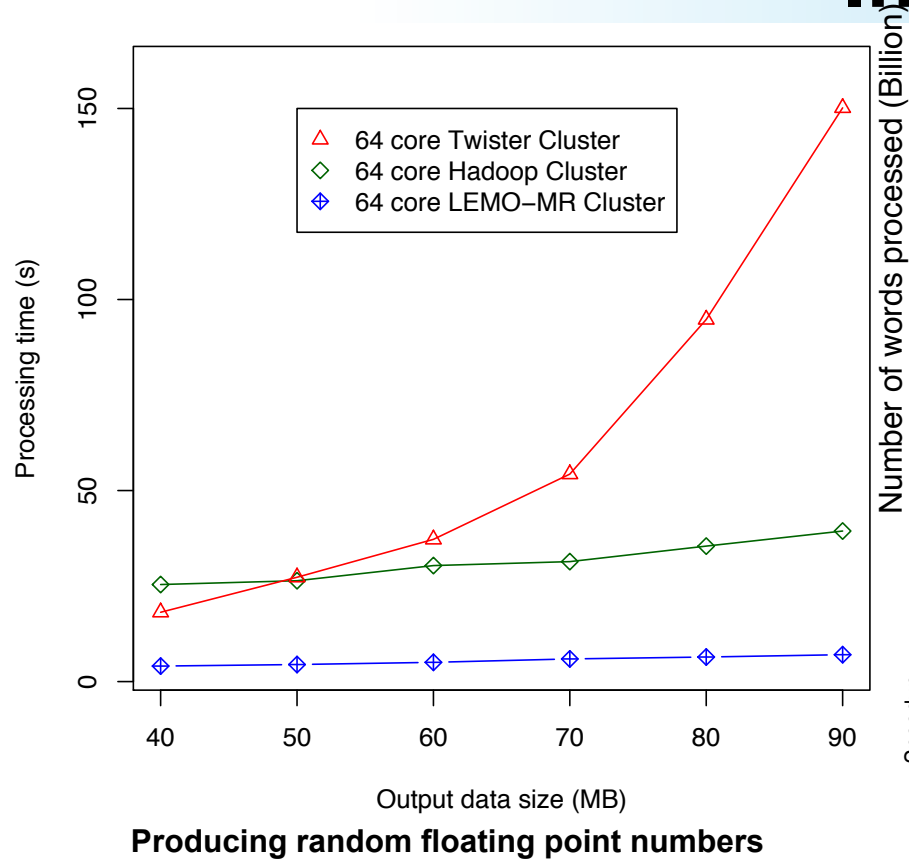
Hadoop (R=W): HDFS vs GPFS



# Hadoop: Challenges

- **Deployment**
  - all jobs run as user “hadoop” affecting file permissions
  - less control on how many nodes are used - affects allocation policies
- **Programming: No turn-key solution**
  - using existing code bases, managing input formats and data
- **Additional benchmarking, tuning needed, Plug-ins for Science**

# Comparison of MapReduce Implementations



Collaboration w/ Zacharia Fadika, Elif Dede, Madhusudhan Govindaraju, SUNY Binghamton



Processing 5 million 33 x 33 matrices  
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# Programming Hadoop



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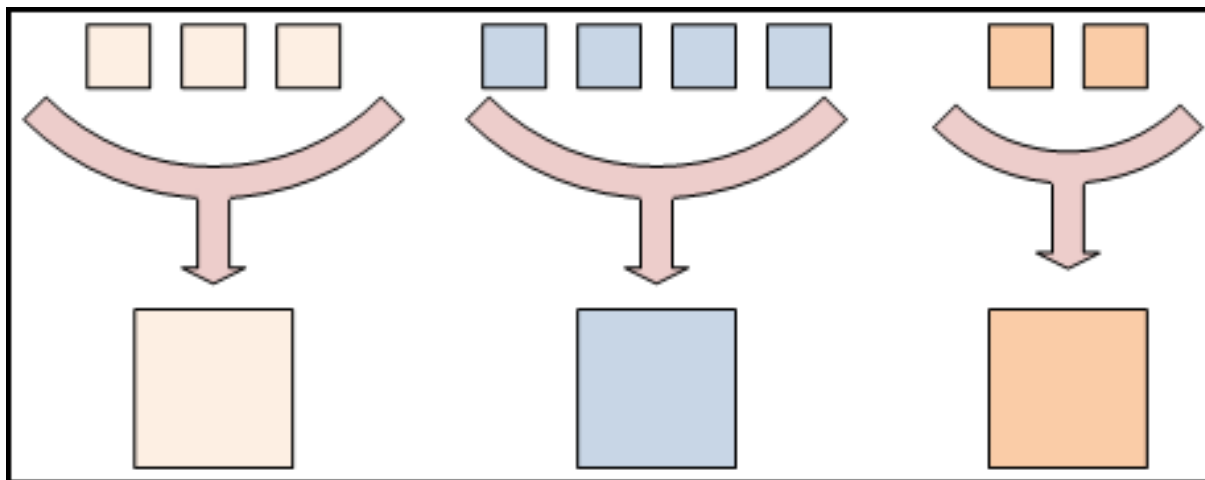
# Programming with Hadoop

- **Map and reduce as Java programs using Hadoop API**
- **Pipes and Streaming can help with existing applications in other languages**
- **C- HDFS API**
- **Higher-level languages such as Pig might help with some applications**

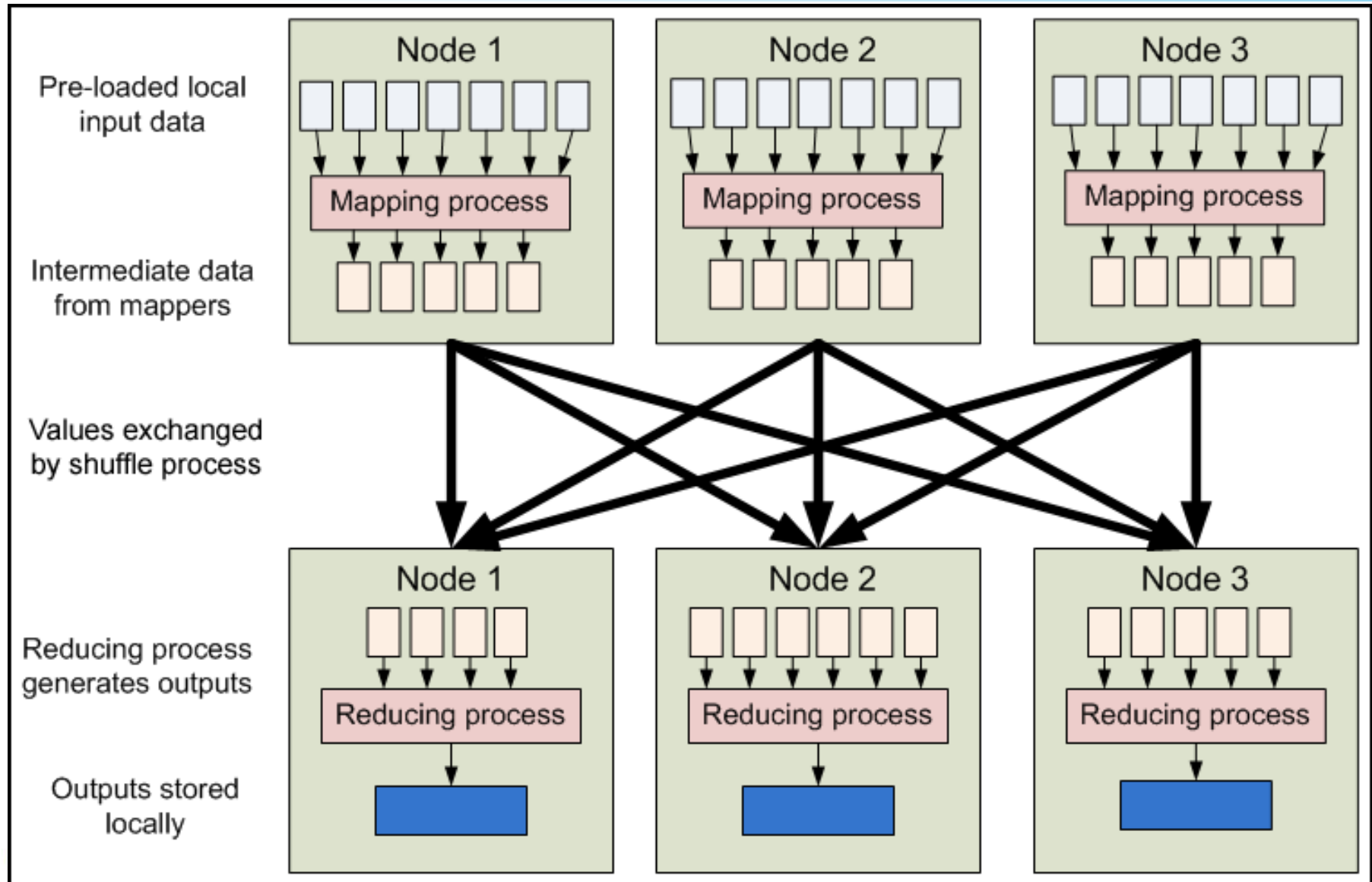
# Keys and Values

- **Maps and reduces produce key-value pairs**
  - arbitrary number of values can be output
  - may map one input to 0,1, ....100 outputs
  - reducer may emit one or more outputs
- **Example: Temperature recordings**
  - 94089 8:00 am, 59
  - 27704 6:30 am, 70
  - 94089 12:45 pm, 80
  - 47401 1 pm, 90

# Keys divide the reduce space



# Data Flow



# Mechanics[1/2]

- **Input files**
  - large 10s of GB or more, typically in HDFS
  - line-based, binary, multi-line, etc.
- **InputFormat**
  - function defines how input files are split up and read
  - **TextInputFormat** (default), **KeyValueInputFormat**, **SequenceFileInputFormat**
- **InputSplits**
  - unit of work that comprises a single map task
  - **FileInputFormat** divides it into 64MB chunks

## Mechanics [2/2]

- **RecordReader**
  - loads data and converts to key value pair
- **Sort & Partiton & Shuffle**
  - intermediate data from map to reducer
- **Combiner**
  - reduce data on a single machine
- **Mapper & Reducer**
- **OutputFormat, RecordWriter**

# Word Count Mapper

```
public static class TokenizerMapper
    extends Mapper<Object, Text, Text, IntWritable>{
    private final static IntWritable one = new IntWritable(1);
    private Text word = new Text();

    public void map(Object key, Text value, Context context
        ) throws IOException, InterruptedException {
        StringTokenizer itr = new StringTokenizer(value.toString());
        while (itr.hasMoreTokens()) {
            word.set(itr.nextToken());
            context.write(word, one);
        }
    }
}
```



# Word Count Reducer

```
public static class IntSumReducer
    extends Reducer<Text,IntWritable,Text,IntWritable> {
    private IntWritable result = new IntWritable();

    public void reduce(Text key, Iterable<IntWritable> values,
Context context) throws IOException, InterruptedException {
        int sum = 0;
        for (IntWritable val : values) {
            sum += val.get();
        }
        result.set(sum);
        context.write(key, result);
    }
}
```

# Word Count Example

```
public static void main(String[] args) throws Exception {  
    Configuration conf = new Configuration();  
    String[] otherArgs = new GenericOptionsParser(conf, args).getRemainingArgs();  
    ....  
    Job job = new Job(conf, "word count");  
    job.setJarByClass(WordCount.class);  
    job.setMapperClass(TokenMapper.class);  
    job.setCombinerClass(IntSumReducer.class);  
    job.setReducerClass(IntSumReducer.class);  
    job.setOutputKeyClass(Text.class);  
    job.setOutputValueClass(IntWritable.class);  
    FileInputFormat.addInputPath(job, new Path(otherArgs[0]));  
    FileOutputFormat.setOutputPath(job, new Path(otherArgs[1]));  
    System.exit(job.waitForCompletion(true) ? 0 : 1);  
}
```

# Pipes

- **Allows C++ code to be used for Mapper and Reducer**
- **Both key and value inputs to pipes programs are provided as `std::string`**
- **\$ hadoop pipes**

# C-HDFS API

- **Limited C API to read and write from HDFS**

```
#include "hdfs.h"

int main(int argc, char **argv)
{
    hdfsFS fs = hdfsConnect("default", 0);
    hdfsFile writeFile = hdfsOpenFile(fs, writePath,
O_WRONLY|O_CREAT, 0, 0, 0);
    tSize num_written_bytes = hdfsWrite(fs, writeFile,
(void*)buffer, strlen(buffer)+1);
    hdfsCloseFile(fs, writeFile);
}
```

# Hadoop Streaming

- **Generic API that allows programs in any language to be used as Hadoop Mapper and Reducer implementations**
- **Inputs written to stdin as strings with tab character separating**
- **Output to stdout as key \t value \n**
- **\$ hadoop jar contrib/streaming/hadoop-[version]-streaming.jar**

# Debugging

- **Test core functionality separate**
- **Use Job Tracker**
- **Run “local” in Hadoop**
- **Run job on a small data set on a single node**
- **Hadoop can save files from failed tasks**

## Pig – Basic Operations

- **LOAD** – loads data into a relational form
- **FOREACH..GENERATE** – Adds or removes fields (columns)
- **GROUP** – Group data on a field
- **JOIN** – Join two relations
- **DUMP/STORE** – Dump query to terminal or file

There are others, but these will be used for the exercises today



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# Pig Example

**Find the number of gene hits for each model in an  
hmmsearch (>100GB of output, 3 Billion Lines)**

```
bash# cat * |cut -f 2|sort|uniq -c
```

```
> hits = LOAD '/data/bio/*' USING PigStorage() AS  
  (id:chararray,model:chararray, value:float);  
> amodels = FOREACH hits GENERATE model;  
> models = GROUP amodels BY model;  
> counts = FOREACH models GENERATE group,COUNT  
  (amodels) as count;  
> STORE counts INTO 'tcounts' USING PigStorage();
```



# Pig - LOAD

## Example:

```
hits = LOAD 'load4/*' USING PigStorage() AS  
      (id:chararray, model:chararray,value:float);
```

- **Pig has several built-in data types (chararray, float, integer)**
- **PigStorage can parse standard line oriented text files.**
- **Pig can be extended with custom load types written in Java.**
- **Pig doesn't read any data until triggered by a DUMP or STORE**

# Pig – FOREACH..GENERATE, GROUP

Example:

```
amodel = FOREACH model GENERATE hits;  
models = GROUP amodels BY model;  
counts = FOREACH models GENERATE group,COUNT  
          (amodels) as count;
```

- Use FOREACH..GENERATE to pick of specific fields or generate new fields. Also referred to as a projection
- GROUP will create a new record with the group name and a “bag” of the tuples in each group
- You can reference a specific field in a bag with <bag>.field (i.e. amodels.model)
- You can use aggregate functions like COUNT, MAX, etc on a bag

## Pig – Important Points

- **Nothing really happens until a DUMP or STORE is performed.**
- **Use FILTER and FOREACH early to remove unneeded columns or rows to reduce temporary output**
- **Use PARALLEL keyword on GROUP operations to run more reduce tasks**

## Questions?

- **Shane Canon**
  - [Scanon@lbl.gov](mailto:Scanon@lbl.gov)
- **Lavanya Ramakrishnan**
  - [LRamakrishnan@lbl.gov](mailto:LRamakrishnan@lbl.gov)