

# Large Scale Learning

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Analyses we have done in class are for in-memory data:

• Datasets can be loaded onto memory of a single computing node.

Database systems can execute SQL queries, which can be used for efficient learning of some models (e.g. decision trees) over data on disk

• Operations are usually performed by a single computing node.

In the 90s database systems that operate over multiple computing nodes became available

Basis of the first generation of large data warehousing.

In the last decade, systems that manipulate data over multiple nodes have become standard.

#### **Basic observation**

for very large datasets, many of the operations for aggregation and summarization, which also form the basis of many learning methods, can be parallelized.

#### For example:

- partition observations and perform transformation on each partition as a parallel process
- partition variables and perform transformation on each variable as a parallel process
- for summarization (group\_by and summarize), partition
   observations based on group\_by expression, perform summarize
   on each partition.

Efficiency of implementation of this type of parallelism depends on underlying architecture:

Shared memory vs. Shared storage vs. Shared nothing

For massive datasets, *shared nothing* is usually preferred since fault tolerance is perhaps the most important consideration.

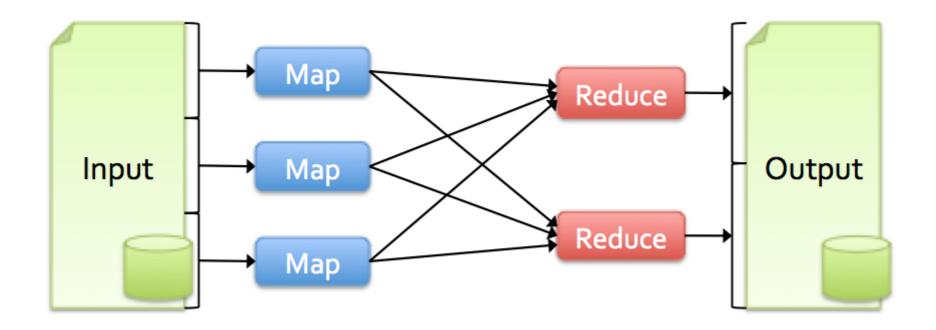
Map-Reduce is an implementation idea for a shared nothing architecture. It is based on:

- distributed storage
- data proximity (perform operaations on data that is physically close)
- fault tolerance.

Basic computation paradigm is based on two operations:

- reduce: perform operation on subset of observations in parallel
- map: decide which parallel process (node) should operate on each observation

The fundamental operations that we have learned very well in this class are nicely represented in this framework: group\_by clause corresponds to map, and summarize function corresponds to reduce.



Map-reduce is most efficient when computations are organized in an acyclic graph.

Data is moved from stable storage to computing process and the result moved to stable storage without much concern for operation ordering.

This architecture provides runtime benefits due to flexible resource allocation and strong failure recovery.

Existing implementations of Map-reduce systems do not support interactive use, or workflows that are hard to represent as acyclic graphs.

Recent system based on the general map-reduce framework

Designed for ultra-fast data analysis.

Provides efficient support for interactive analysis (the kind we do in Jupyter)

Designed to support iterative workflows needed by many Machine Learning algorithms.

The basic data abstraction in Spark is the resilient distributed dataset (RDD).

Applications keep working sets of data in memory and support iterative algorithms and interactive workflows.

#### RDDs are

- (1) inmutable and *partitioned* collections of objects,
- (2) created by parallel *transformations* on data in stable storage (e.g., map, filter, group\_by, join, ...)
- (3) cached for efficient reuse
- (4) operated upon by actions defeind on RDDs (count, reduce, collect, save, ...)

#### **Fault Tolerance**

RDDs maintain *lineage*, so partitions can be reconstructed upon failure.

The components of a SPARK workflow

**Transformations**: Define new RDDs

https://spark.apache.org/docs/latest/programming-guide.html#transformations

Actions: Return results to driver program

https://spark.apache.org/docs/latest/programming-guide.html#actions

Spark was designed first for Java with an interactive shell based on Scala. It has strong support in Python and increasing support in R SparkR.

- Spark programming guide:
   https://spark.apache.org/docs/latest/programming-guide.html
- More info on python API: https://spark.apache.org/docs/0.9.1/python-programming-guide.html

Other learning methods we have seen, like regression and SVMs (or even PCA), are **optimization problems** 

We can design gradient-descent based optimization algorithms that process data efficiently.

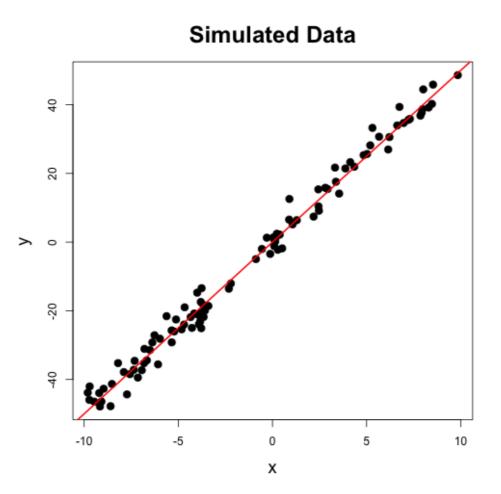
We will use linear regression as a case study of how this insight would work.

Let's use linear regression with one predictor, no intercept as a case study.

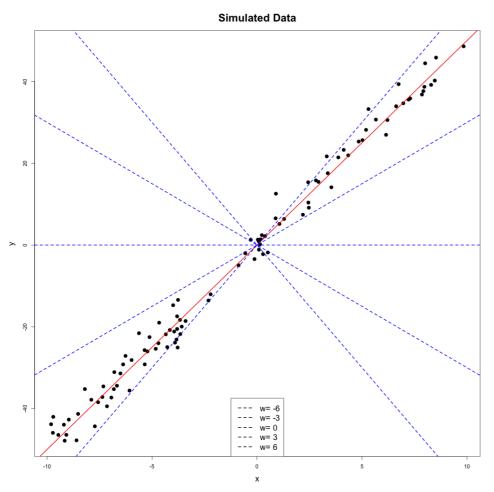
**Given**: Training set  $\{(x_1, y_1), \dots, (x_n, y_n)\}$ , with continuous response  $y_i$  and single predictor  $x_i$  for the i-th observation.

**Do**: Estimate parameter w in model y = wx to solve

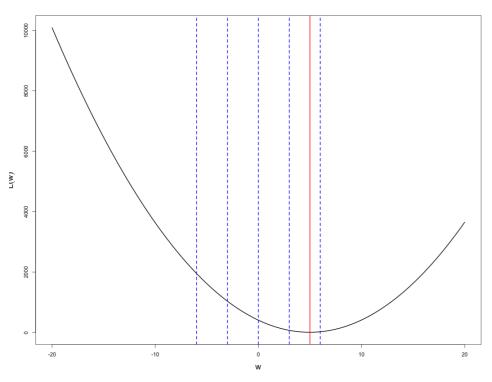
$$\min_w L(w) = rac{1}{2}\sum_{i=1}^n (y_i-wx_i)^2$$



Suppose we want to fit this model to the following (simulated) data:



Our goal is then to find the value of (w) that minimizes mean squared error. This corresponds to finding one of these many possible lines.

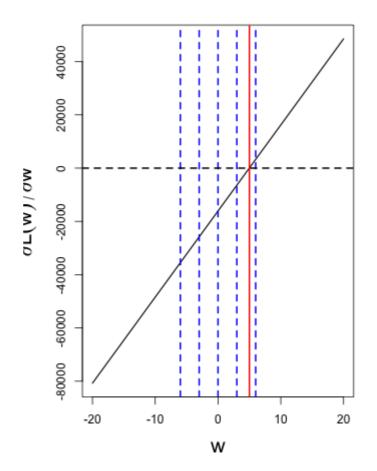


Each of which has a specific error for this dataset:

- 1) As we saw before in class, loss is minimized when the derivative of the loss function is 0
- 2) and, the derivative of the loss (with respect to w) at a given estimate w suggests new values of w with smaller loss!

# Let's take a look at the derivative:

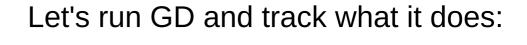
$$egin{aligned} rac{\partial}{\partial w}L(w) = \ rac{\partial}{\partial w}rac{1}{2}\sum_{i=1}^n(y_i-wx_i)^2 \ = \sum_{i=1}^n(y_i-wx_i)(-x_i) \end{aligned}$$

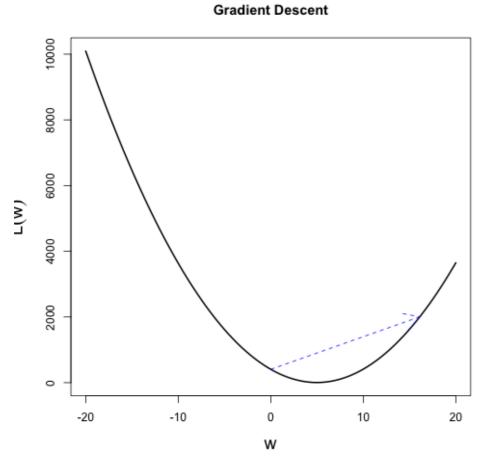


This is what motivates the Gradient Descent algorithm

- 1. Initialize w = normal(0, 1)
- 2. Repeat until convergence
  - $\circ$  Set  $w=w+\eta\sum_{i=1}^n(y_i-f(x_i))x_i$

The basic idea is to move the current estimate of w in the direction that minimizes loss the *fastest*.





"Batch" gradient descent: take a step (update w) by calculating derivative with respect to *all* n observations in our dataset.

$$w=w+\eta \sum_{i=1}^n (y_i-f(x_i,w))x_i$$

where  $f(x_i) = wx_i$ .

For multiple predictors (e.g., adding an intercept), this generalizes to the *gradient* 

$$\mathbf{w} = \mathbf{w} + \eta \sum_{i=1}^n (y_i - f(\mathbf{x}_i, \mathbf{w})) \mathbf{x}_i$$

where  $f(\mathbf{x}_i, \mathbf{w}) = w_0 + w_1 x_{i1} + \cdots + w_p x_{ip}$ 

Gradiest descent falls within a family of optimization methods called *first-order methods* (first-order means they use derivatives only). These methods have properties amenable to use with very large datasets:

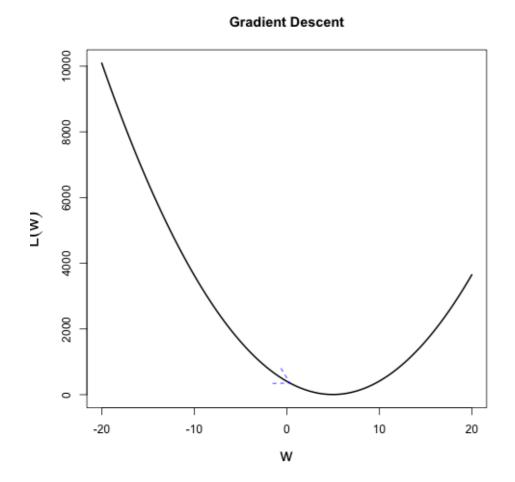
- 1. Inexpensive updates
- 2. "Stochastic" version can converge with few sweeps of the data
- 3. "Stochastic" version easily extended to streams
- 4. Easily parallelizable

Drawback: Can take many steps before converging

**Key Idea**: Update parameters using update equation *one observation at a time*:

- 1. Initialize  $\beta = \text{normal}(0, \sqrt{p})$ , i = 1
- 2. Repeat until convergence
  - $\circ$  For i=1 to n
  - $\circ$  Set  $w = w + \eta(y_i f(\mathbf{x}_i, \mathbf{w}))\mathbf{x}_i$

Let's run this and see what it does:



Why does SGD make sense?

For many problems we are minimizing a cost function of the type

$$rg \min_f rac{1}{n} \sum_i L(y_i, f_i) + \lambda R(f)$$

Which in general has gradient

$$rac{1}{n}\sum_{i}
abla_{f}L(y_{i},f_{i})+\lambda
abla_{f}R(f)$$

$$rac{1}{n}\sum_{i}
abla_{f}L(y_{i},f_{i})+\lambda
abla_{f}R(f)$$

The first term looks like an empirical estimate (average) of the gradient at  $f_i$ 

SGD then uses updates provided by a different *estimate* of the gradient based on a single point.

- Cheaper
- Potentially unstable

#### In practice

- Mini-batches: use ~100 or so examples at a time to estimate gradients
- Shuffle data order every pass (epoch)

SGD easily adapts to *data streams* where we receive observations one at a time and *assume* they are not stored.

This setting falls in the general category of *online* learning.

Online learning is extremely useful in settings with massive datasets

Parallelizing gradient descent

Gradient descent algorithms are easily parallelizable:

- Split observations across computing units
- For each step, compute partial sum for each partition (map), compute final update (reduce)

$$\mathbf{w} = \mathbf{w} + \eta * \sum_{ ext{partition } q} \sum_{i \in q} (y_i - f(\mathbf{x_i}, \mathbf{w})) \mathbf{x}_i$$

This observation has resulted in their implementation if systems for large-scale learning:

#### 1. Vowpal Wabbit

 Implements general framework of (sparse) stochastic gradient descent for many optimization problems

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#### 2. Spark MLlib

 Implements many learning algorithms using Spark framework we saw previously

For many ML algorithms prediction time-efficiency is determined by the number of predictors used in the model.

Reducing the number of predictors can yield huge gains in efficiency in deployment.

The amount of memory used to make predictions is also typically governed by the number of features. (Note: this is not true of kernel methods like support vector machines, in which the dominant cost is the number of support vectors.)

The idea behind sparse models, and in particular, sparse regularizers.

A disadvantage of optimizing problems of the form

$$\sum_i L(y_i,w) + \lambda {|w|}^2$$

That they tend to never produce weights that are exactly zero.

Instead, minimize problems of the form

$$\sum_i L(y_i,w) + \lambda |w|_1$$

where  $\|w\|_1 = \sum_j |w_j|$ 

This is a convex optimization problem.

Can use standard subgradient methods.

See CIML for further details.

For data sets with a large number of features/attributes an idea based on hashing can also help.

Suppose we are building a model over P features (P very large). We use hashing to reduce the number of features to smaller number P.

For each observation  $x \in \mathbb{R}^P$  we transform it into observation  $\tilde{x} \in \mathbb{R}^P$ .

We will use hash function  $h: P \rightarrow p$ 

Initialize  $\tilde{x} = \langle 0, 0, \dots, 0 \rangle$ 

For each  $j \in 1, ..., P$ 

- Hash j to position k = h(j)
- Update  $\tilde{x}_k \leftarrow \tilde{x}_k + x_j$

Return  $\tilde{x}$ 

We can think of this as a *feature mapping* 

$$\phi(x)_k = \sum_{j|j \in h^{-1}(k)} x_j$$

To see how what this does, we can see what the inner product between observations in the smaller feature space is:

$$egin{aligned} \phi(x)'\phi(z) &= \sum_k \left[\sum_{j|j\in h^{-1}(k)} x_j
ight] \left[\sum_{j'|j'\in h^{-1}(k)} z_{j'}
ight] \ &= \sum_k \sum_{j,j'|j,j'\in h^{-1}(k)} x_j z_{j'} \ &= \sum_j \sum_{j'|j'\in h^{-1}(h(j))} x_j z_{j'} \ &= \sum_j x_j z_j + \sum_{j'
eq j|j'\in h^{-1}(h(j))} x_j z_{j'} = x'z + \cdots \end{aligned}$$

So, we get the inner product in the original large dimensions plus an extra quadratic term

$$\phi(x)'\phi(z)=x'z+\sum_{j'
eq j|j'\in h^{-1}(h(j))}x_jz_{j'}$$

- We might get lucky and get a useful interaction between two features
- Nonetheless, the size of this sum is very small due to property of hash functions: *expected value* of product is  $\approx 0$

- Database operations for out-of-memory datasets
- Parallelization on shared-nothing architectures (map reduce)
- Spark as MR framework also supporting iterative procedures (optimization)
- Stochastic gradient descent (many low cost steps, easy to parallelize)
- Sparse models (build models with few features)
- Feature Hashing (build models over smaller number of features, retain inner product)