732A54/TDDE31 Big Data Analytics Lecture 8: Machine Learning with MapReduce

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Literature

- Main sources
 - Dean, J. and Ghemawat, S. MapReduce: Simplified Data Processing on Large Clusters. *Communications of the ACM*, 51(1):107-113, 2008.
 - Chu, C.-T. et al. Map-Reduce for Machine Learning on Multicore. In Proceedings of the 19th International Conference on Neural Information Processing Systems, 281-288, 2006.
- Additional sources
 - Dean, J. and Ghemawat, S. MapReduce: Simplified Data Processing on Large Clusters. In Proceedings of the 6th Symposium on Operating Systems Design and Implementation, 2004.
 - Gillick, D., Faria, A. and DeNero, J. MapReduce: Distributed Computing for Machine Learning. Technical Report, Berkley, 2006.
 - Slides for 732A99/TDDE01 Machine Learning.

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- Moreover, it is a straightforward way to adapt some machine learning algorithms to cope with big data.
- Apache Mahout is a project to produce distributed implementations of machine learning algorithms. It builds on Hadoop's MapReduce. However, these implementations are now deprecated, in favor of Apache Spark.

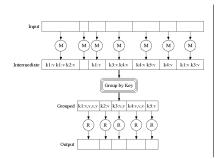
- The user only has to implement the following two functions:
 - Map function:
 - Input: A pair (in_key, in_value).
 - Output: A list list(out_key, intermediate_value).
 - Reduce function:
 - Input: A pair (out_key, list(intermediate_value)).
 - Output: A list list(out_value).

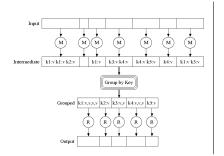
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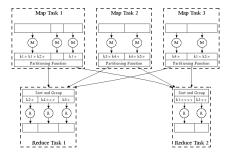
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- Example for counting word occurrences in a collection of documents:

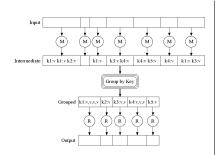
map(String key, String value):
 // key: document name
 // value: document contents
 for each word w in value:
 EmitIntermediate(w, "1");

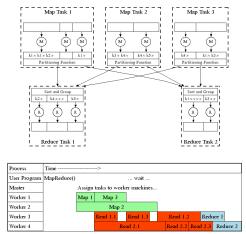
reduce(String key, Iterator values):
 // key: a word
 // values: a list of counts
 int result = 0;
 for each v in values:
 result += ParseInt(v);
 Emit(AsString(result));











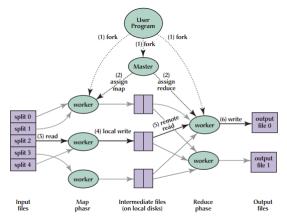


Fig. 1. Execution overview.

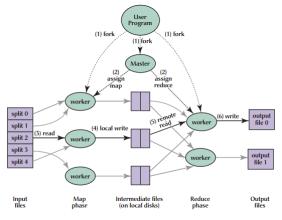


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1. Split the input file in *M* pieces and store them on the local disks of the nodes of the cluster. Start up many copies of the user's program on the nodes.

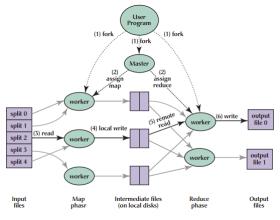
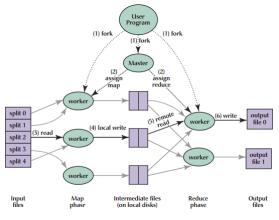


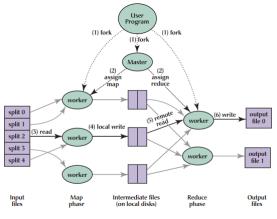
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- 1. Split the input file in *M* pieces and store them on the local disks of the nodes of the cluster. Start up many copies of the user's program on the nodes.
- 2. One copy (the master) assigns tasks to the rest of the copies (the workers). To reduce communication, it tries to assign map workers to nodes with input data.





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- 4. The buffered results are written to **local** disk. The disk is **partitioned** in *R* pieces, e.g. *hash(out_key) mod R*. The location of the partitions on disk are passed back to the master so that they can be forwarded to the reduce workers.

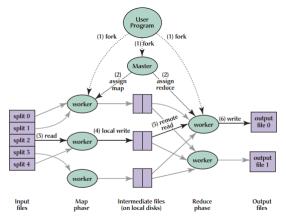


Fig. 1. Execution overview.

5. The reduce worker reads its partition remotely (a.k.a shuffle) and sorts it by key.

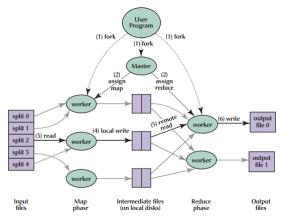
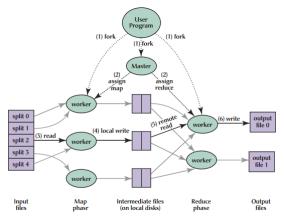


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- 5. The reduce worker reads its partition remotely (a.k.a shuffle) and sorts it by key.
- 6. The reduce worker processes each key using the user's reduce function. The result is written to the **global** file system.

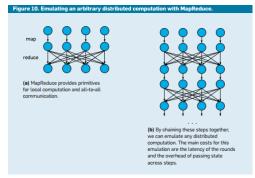




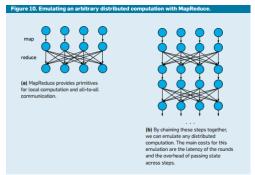
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- 7. The output of a MapReduce call may be the input to another. Note that we have performed M map tasks and R reduce tasks.

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However, the emulation may be inefficient since the message exchange relies on external storage, e.g. disk.

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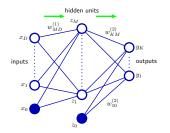
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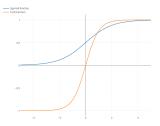
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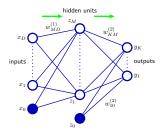
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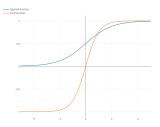
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 - Too large values may imply too many scheduling decisions, and too many output files.
 - For instance, M = 200000 and R = 5000 for 2000 available nodes.

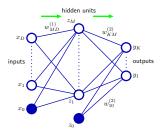


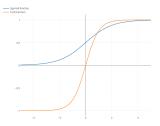




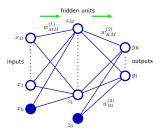


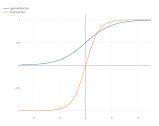
• Activations: $a_j = \sum_i w_{ji}^{(1)} x_i + w_{j0}^{(1)}$



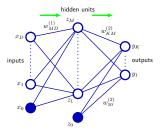


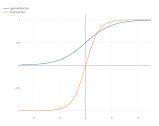
- Activations: a_j = ∑_i w_{ji}⁽¹⁾ x_i + w_{j0}⁽¹⁾
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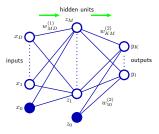


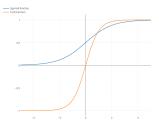
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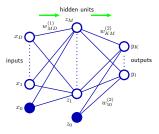


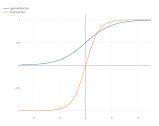
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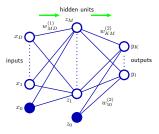


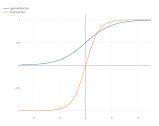
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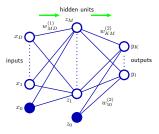
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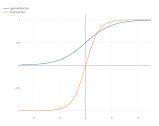




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$$y_k(\mathbf{x}) = \sigma\left(\sum_{j} w_{kj}^{(2)} h\left(\sum_{i} w_{ji}^{(1)} x_i + w_{j0}^{(1)}\right) + w_{k0}^{(2)}\right)$$

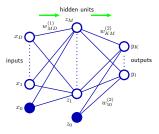


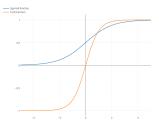


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- All the previous is, of course, generalizable to more layers.

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- Consider a training set {(x_n, t_n)} of size N. Consider minimizing the error function

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The weight space is highly multimodal and, thus, we have to resort to approximate iterative methods to minimize the previous expression.

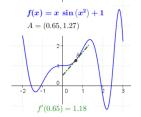
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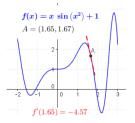
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- Batch gradient descent

$$\boldsymbol{w}^{t+1} = \boldsymbol{w}^t - \eta \nabla E(\boldsymbol{w}^t) = \boldsymbol{w}^t - \eta \sum_n \nabla E_n(\boldsymbol{w}^t)$$

where $\eta > 0$ is the learning rate, and $\nabla E_n(\boldsymbol{w}^t)$ can be computed efficiently thanks to the backpropagation algorithm.





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- What is the key and what is the value ? What needs to be broadcasted ?

• Consider binary classification with input space \mathbb{R}^D . Consider a training set $\{(\mathbf{x}_n, t_n)\}$ where $t_n \in \{-1, +1\}$. Consider using the linear model

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The motivation is that the larger the margin, the smaller the generalization error.

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where C is a user-defined parameter, and $n \in E$ if and only if $t_n y(\mathbf{x}_n) < 1$.

Note that the previous expression is a quadratic function and, thus, it is concave (up) and, thus, "easy" to minimize. For instance, we can use again batch gradient descent.

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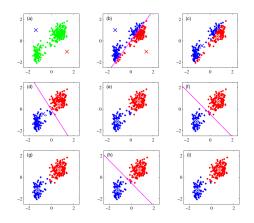
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- Consider data clustering (a.k.a. unsupervised learning) via the K-means algorithm.
 - 1 Assign each point to a cluster (a.k.a. subpopulation) at random
 - 2 Compute the cluster centroids as the averages of the points assigned to each cluster
 - 3 Repeat until the centroids do not change
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The K-means algorithm partitions the data, i.e. it hard-assigns instances to subpopulations. Model-based clustering on the other hand aims to soft-assign instances to the subpopulations by applying Bayes theorem as follows:

$$p(k|\mathbf{x},\boldsymbol{\theta},\boldsymbol{\pi}) = \frac{\pi_k p(\mathbf{x}|\boldsymbol{\theta}_k)}{\sum_k \pi_k p(\mathbf{x}|\boldsymbol{\theta}_k)}$$

where $p(\mathbf{x}|\boldsymbol{\theta}_k)$ are called mixture components, and $\pi_k = p(k)$ are called mixing coefficients. A component models the data distribution for a chosen subpopulation, and a coefficient represents the probability of a subpopulation being chosen.

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More specifically, for components modeled as multivariate Gaussian distributions, we have that:

$$\boldsymbol{p}(\boldsymbol{x}|\boldsymbol{\theta}_{k}) = \mathcal{N}(\boldsymbol{x}|\boldsymbol{\mu}_{k},\boldsymbol{\Sigma}_{k}) = \frac{1}{2\pi^{D/2}} \frac{1}{|\boldsymbol{\Sigma}_{k}|^{1/2}} e^{-\frac{1}{2}(\boldsymbol{x}-\boldsymbol{\mu}_{k})^{T}\boldsymbol{\Sigma}_{k}^{-1}(\boldsymbol{x}-\boldsymbol{\mu}_{k})}.$$

To solve model-based clustering, we have to estimate the model parameters (θ, π) from data. To this end, we use the EM algorithm.

Given a sample {x_n} of size N from a mixture of multivariate Gaussian distributions, the expected log likelihood function is maximized when

$$\pi_{k}^{ML} = \frac{\sum_{n} p(z_{nk} | \mathbf{x}_{n}, \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma})}{N}$$
$$\mu_{k}^{ML} = \frac{\sum_{n} \mathbf{x}_{n} p(z_{nk} | \mathbf{x}_{n}, \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma})}{\sum_{n} p(z_{nk} | \mathbf{x}_{n}, \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma})}$$
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where z_n is a *K*-dimensional binary vector indicating component memberships (one-hot encoding):

$$p(z_{nk}|\mathbf{x}_n, \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \frac{p(\mathbf{x}_n | z_{nk}, \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) p(z_{nk} | \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma})}{\sum_k p(\mathbf{x}_n | z_{nk}, \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) p(z_{nk} | \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma})} = \frac{\pi_k p(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_k \pi_k p(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}$$

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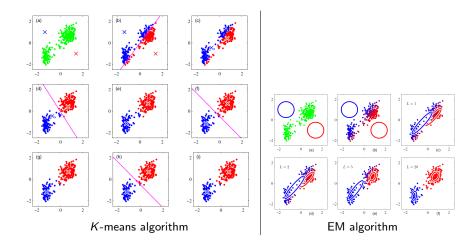
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EM algorithm

Set
$$\pi$$
, μ and Σ to some initial values
Repeat until π , μ and Σ do not change
Compute $p(z_{nk}|\mathbf{x}_n, \pi, \mu, \Sigma)$ for all n /* E step */
Set π_k to π_k^{ML} , μ_k to μ_k^{ML} , and Σ_k to Σ_k^{ML} for all k /* M step */



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$$p(z_{nk}|\mathbf{x}_n, \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma})$$
(3)

and

$$(\boldsymbol{x}_n - \boldsymbol{\mu}_k^{ML})(\boldsymbol{x}_n - \boldsymbol{\mu}_k^{ML})^T p(\boldsymbol{z}_{nk} | \boldsymbol{x}_n, \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}).$$
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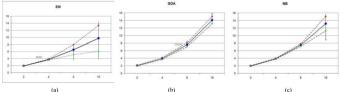
Machine Learning with MapReduce

Data Sets	samples (m)	features (n)
Adult	30162	14
Helicopter Control	44170	21
Corel Image Features	68040	32
IPUMS Census	88443	61
Synthetic Time Series	100001	10
Census Income	199523	40
ACIP Sensor	229564	8
KDD Cup 99	494021	41
Forest Cover Type	581012	55
1990 US Census	2458285	68

	lwlr	gda	nb	logistic	pca	ica	svm	nn	kmeans	em
Adult	1.922	1.801	1.844	1.962	1.809	1.857	1.643	1.825	1.947	1.854
Helicopter	1.93	2.155	1.924	1.92	1.791	1.856	1.744	1.847	1.857	1.86
Corel Image	1.96	1.876	2.002	1.929	1.97	1.936	1.754	2.018	1.921	1.832
IPUMS	1.963	2.23	1.965	1.938	1.965	2.025	1.799	1.974	1.957	1.984
Synthetic	1.909	1.964	1.972	1.92	1.842	1.907	1.76	1.902	1.888	1.804
Census Income	1.975	2.179	1.967	1.941	2.019	1.941	1.88	1.896	1.961	1.99
Sensor	1.927	1.853	2.01	1.913	1.955	1.893	1.803	1.914	1.953	1.949
KDD	1.969	2.216	1.848	1.927	2.012	1.998	1.946	1.899	1.973	1.979
Cover Type	1.961	2.232	1.951	1.935	2.007	2.029	1.906	1.887	1.963	1.991
Census	2.327	2.292	2.008	1.906	1.997	2.001	1.959	1.883	1.946	1.977
avg.	1.985	2.080	1.950	1.930	1.937	1.944	1.819	1.905	1.937	1.922

Table 3: Speedups achieved on a dual core processor, without load time. Numbers reported are dualcore time / single-core time. Super linear speedup sometimes occurs due to a reduction in processor idle time with multiple threads.

Machine Learning with MapReduce

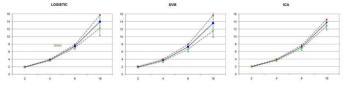








(f)



(e)



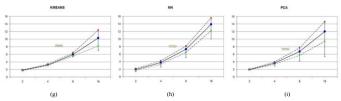


Figure 2: (a)-(i) show the speedup from 1 to 16 processors of all the algorithms over all the data sets. The Bold line is the average, error bars are the max and min speedups and the dashed lines are the variance.

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- Many machine learning algorithms (e.g. SVMs, NNs, MMs, K-means and EM algorithms) can easily be reformulated in terms of such functions.
- This does not apply for algorithms based on stochastic gradient descent.
- Moreover, MapReduce is inefficient for iterative tasks on the same dataset: Each iteration is a MapReduce call that loads the data anew from disk.
- Such iterative tasks are common in many machine learning algorithms, e.g. gradient descent, K-means and EM algorithms.
- Solution: Spark framework, in the next lecture.