Optimization in Data Analysis

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Outline

- Data Analysis and Machine Learning
 - The Setup
 - About Ten Applications / Examples
- Optimization in Data Analysis
 - Basic Formulations
 - Relevant Algorithms

Data Analysis

Also: Machine Learning, Statistical Inference, Data Mining.

- Extract meaning from data: Understand statistical properties, important features, fundamental structures in the data.
- Use this knowledge to make predictions about other, similar data.

Foundations in Statistics, then Computer Science (AI, Machine Learning, Databases, Parallel Systems), more recently Optimization.

For a recent Statistics perspective on past, present, and future, see David Donoho: *50 Years of Data Science*, September 2015.

Modeling and domain-specific knowledge is vital: "80% of data analysis is spent on the process of cleaning and preparing the data." [Dasu and Johnson, 2003].

(Most academic research deals with the other 20%.)

"Big Data"

New "Data Science Centers" at many institutions, new degree programs (e.g. Masters in Data Science), new funding initiatives.

- Huge amounts of data are being collected, routinely.
 - Consumer and citizen data: phone logs (call and text), email, surveillance cameras, web activity, online shopping activity,...
 - Scientific data (particle colliders, satellites, biological / genomic, astronomical,...)
- Affects everyone directly!
- Powerful computers make it possible to handle larger data sets and analyze more thoroughly.
- Methodological innovations in some areas. e.g. Deep Learning.
 - Speech recognition in smart phones
 - AlphaGo: Al playing Go

Typical Setup

After cleaning and formatting, obtain a data set of m objects:

- Vectors of features: a_j , $j = 1, 2, \ldots, m$.
- Outcome / observation / label y_j for each feature vector.

The outcomes y_j could be:

- a real number: regression
- a label indicating that a_j lies in one of M classes (for $M \ge 2$): classification
- multiple labels: classify *a_j* according to multiple criteria.
- no labels (y_j is null):
 - subspace identification: Locate low-dimensional subspaces that approximately contain the (high-dimensional) vectors a_j;
 - **clustering**: Partition the *a_j* into a few clusters.

(Structure may reveal which features in the a_j are important / distinctive, or enable predictions to be made about new vectors a.)

Fundamental Data Analysis Task

Seek a function ϕ that:

- approximately maps a_j to y_j for each $j: \phi(a_j) \approx y_j$ for j = 1, 2, ..., m, if labels y_j are available.
- if there are no labels y_j , or if some labels are missing, seek ϕ that does something useful with the data $\{a_j\}$, e.g. assigns each a_j to an appropriate cluster or subspace.
- satisfies some additional properties simplicity, structure that make it "plausible" for the application, robust to perturbations in the data, generalizable to other data samples.

Can usually define ϕ in terms of some parameter vector x — thus identification of ϕ becomes a data-fitting problem.

Objective function in this problem often built up of m terms that capture mismatch between predictions and observations for each (a_j, y_j) .

The process of finding ϕ is often called learning or training.

Why?

What's the use of ϕ ?

- - Feature selection: reveal the components of vectors a_j that are most important in determining the outputs y_j, and quantifies the importance of these features.
 - Uncovers some hidden structure, e.g.
 - finds some low-dimensional subspaces that contain the a_i;
 - * find clusters that contain the a_i ;
 - ★ find a decision tree that builds intuition about how y_j depend on a_j.

• Prediction: Given new data vectors a_k , predict outputs $y_k \leftarrow \phi(a_k)$.

Complications

- noise or errors in a_j and y_j . Would like ϕ (and x) to be robust to this. Thus, generalization / regularization.
- avoid overfitting: Observed data is viewed as an empirical, sampled representation of some underlying reality. Want to avoid overfitting to the particular sample. (Ideally, the training process should produce a similar result for other samples from the same data set.) Again, generalization / regularization.
- **missing data**: Vectors *a_j* may be missing elements (but may still contain useful information).
- **missing labels:** Some or all *y_j* may be missing or null semi-supervised or unsupervised learning.
- online learning: Data (a_j, y_j) is arriving in a stream rather than all known up-front.

Application I: Least Squares

$$\min_{x} f(x) := \frac{1}{2} \sum_{j=1}^{m} (a_{j}^{T} x - y_{j})^{2} = \frac{1}{2} \|Ax - y\|_{2}^{2}.$$

[Gauss, 1799], [Legendre, 1805]; see [Stigler, 1981].

Here the function mapping data to output is linear: $\phi(a_j) = a_j^T x$.

• ℓ_2 regularization reduces sensitivity of the solution x to noise in y.

$$\min_{x} \frac{1}{2} \|Ax - y\|_{2}^{2} + \lambda \|x\|_{2}^{2}.$$

• ℓ_1 regularization yields solutions x with few nonzeros:

$$\min_{x} \frac{1}{2} \|Ax - y\|_{2}^{2} + \lambda \|x\|_{1}.$$

Feature selection: Nonzero locations in x indicates important components of a_j .

• Nonconvex separable regularizers (SCAD, MCP) have nice statistical properties, but lead to nonconvex optimization formulations.

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Optimization in Data Analysis

Application II: Matrix Completion

Regression over a structured matrix: Observe a matrix X by probing it with linear operators $A_j(X)$, giving observations y_j , j = 1, 2, ..., m. Solve a regression problem:

$$\min_{X} \frac{1}{2m} \sum_{j=1}^{m} (\mathcal{A}_{j}(X) - y_{j})^{2} = \frac{1}{2m} \|\mathcal{A}(X) - y\|_{2}^{2}.$$

Each A_j may observe a single element of X, or a linear combination of elements. Can be represented as a matrix A_j , so that $A_j(X) = \langle A_j, X \rangle$. Seek the "simplest" X that satisfies the observations, e.g. low-rank, or low-rank plus sparse. A nuclear-norm (sum-of-singular-values) regularization term induces low rank on X:

$$\min_X rac{1}{2m} \|\mathcal{A}(X) - y\|_2^2 + \lambda \|X\|_*, \quad ext{for some } \lambda > 0.$$

[Recht et al., 2010]

Explicit Low-Rank Parametrization

Compact, nonconvex formulation is obtained by parametrizing X directly:

$$X = LR^T$$
, where $L \in \mathbb{R}^{m \times r}$, $R \in \mathbb{R}^{n \times r}$,

where r is known (or suspected) rank.

$$\min_{L,R} \frac{1}{2m} \sum_{j=1}^m (\mathcal{A}_j(LR^T) - y_j)^2.$$

For symmetric X, write $X = ZZ^T$, where $Z \in \mathbb{R}^{n \times r}$, so that

$$\min_{Z} \frac{1}{2m} \sum_{j=1}^{m} (\mathcal{A}_j(ZZ^{T}) - y_j)^2.$$

(No need for regularizer — rank is hard-wired into the formulation.)

Despite the nonconvexity, near-global minima can be found when A_j are incoherent. Use appropriate initialization [Candès et al., 2014], [Zheng and Lafferty, 2015] or the observation that all local minima are near-global [Bhojanapalli et al., 2016].

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Application III: Nonnegative Matrix Factorization

Given $m \times n$ matrix Y, seek factors $L(m \times r)$ and $R(n \times r)$ that are element-wise positive, such that $LR^T \approx Y$.

$$\min_{L,R} \frac{1}{2} \|LR^T - Y\|_F^2 \text{ subject to } L \ge 0, R \ge 0.$$

Applications in computer vision, document clustering, chemometrics, ...

Could combine with matrix completion, when not all elements of Y are known, if it makes sense on the application to have nonnegative factors.

If positivity constraint were not present, could solve this in closed form with an SVD, since Y is observed completely.

Application IV: Sparse Inverse Covariance

Let $Z \in \mathbb{R}^p$ be a (vector) random variable, whose entries have zero mean. Let z_1, z_2, \ldots, z_N be samples of Z. Sample covariance matrix (estimates covariance between components of Z):

$$\mathcal{S} := rac{1}{N-1}\sum_{\ell=1}^N z_\ell z_\ell^T.$$

Seek a sparse inverse covariance matrix: $X \approx S^{-1}$.

X reveals dependencies between components of Z: $X_{ij} = 0$ if the *i* and *j* components of Z are conditionally independent.

(Nonzeros in X indicate arcs in the dependency graph.)

Obtain X from the regularized formulation:

$$\min_X \left\langle S,X
ight
angle - \log \det(X) + \lambda \|X\|_1, \quad ext{where } \|X\|_1 = \sum_{i,j} |X_{ij}|.$$

[Friedman et al., 2008, Scheinberg and Ma, 2012].

Application V: Subspace Identification

Given vectors $a_j \in \mathbb{R}^n$ with missing entries, find a subspace of \mathbb{R}^n such that all "completed" vectors a_j lie approximately in this subspace.

If $\Omega_j \subset \{1, 2, ..., n\}$ is the set of observed elements in a_j , seek $X \in \mathbb{R}^{n \times d}$ such that

$$[a_j - Xs_j]_{\Omega_j} \approx 0,$$

for some $s_j \in \mathbb{R}^d$ and all j = 1, 2, ...[Balzano et al., 2010, Balzano and Wright, 2014].

Application: Structure from motion. Reconstruct opaque object from planar projections of surface reference points.



Application VI: Linear Support Vector Machines

Each item of data belongs to one of two classes: $y_j = +1$ and $y_j = -1$. Seek (x, β) such that

$$a_j^T x - \beta \ge 1$$
 when $y_j = +1;$
 $a_j^T x - \beta \le -1$ when $y_j = -1.$

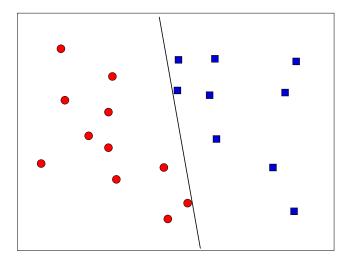
The mapping is $\phi(a_j) = \operatorname{sign}(a_j^T x - \beta)$.

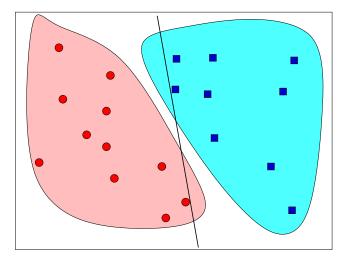
Design an objective so that the *j*th loss term is zero when $\phi(a_j) = y_j$, positive otherwise. A popular one is hinge loss:

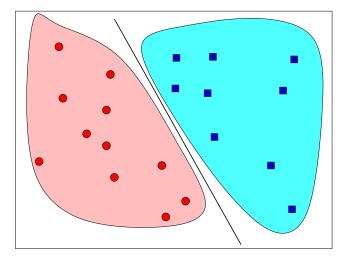
$$H(x) = \frac{1}{m} \sum_{j=1}^{m} \max(1 - y_j(a_j^T x - \beta), 0).$$

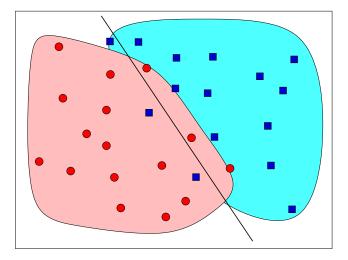
Add a regularization term $(\lambda/2)||x||_2^2$ for some $\lambda > 0$ to maximize the margin between the classes.

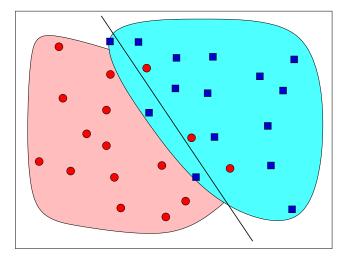
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Application VII: Nonlinear SVM

Data a_j , j = 1, 2, ..., m may not be *separable* neatly into two classes $y_j = +1$ and $y_j = -1$. Apply a nonlinear transformation $a_j \rightarrow \psi(a_j)$ ("lifting") to make separation more effective. Seek (x, β) such that

$$\psi(a_j)^T x - \beta \ge 1$$
 when $y_j = +1$;
 $\psi(a_j)^T x - \beta \le -1$ when $y_j = -1$.

Leads to the formulation:

$$\frac{1}{m}\sum_{j=1}^{m}\max(1-y_{j}(\psi(a_{j})^{T}x-\beta),0)+\frac{1}{2}\lambda\|x\|_{2}^{2}.$$

Can avoid defining ψ explicitly by using instead the dual of this QP.

Nonlinear SVM: Dual

Dual is a quadratic program in m variables, with simple constraints:

$$\min_{\alpha \in \mathbb{R}^m} \frac{1}{2} \alpha^T Q \alpha - e^T \alpha \text{ s.t. } 0 \le \alpha \le (1/\lambda)e, \ y^T \alpha = 0.$$

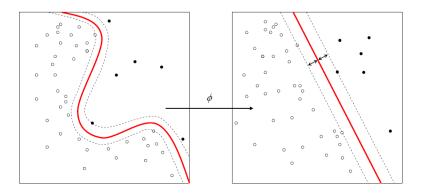
where $Q_{k\ell} = y_k y_\ell \psi(a_k)^T \psi(a_\ell)$, $y = (y_1, y_2, \dots, y_m)^T$, $e = (1, 1, \dots, 1)^T$. No need to choose $\psi(\cdot)$ explicitly. Instead choose a kernel K, such that

$$K(a_k, a_\ell) \sim \psi(a_k)^T \psi(a_\ell).$$

[Boser et al., 1992, Cortes and Vapnik, 1995]. "Kernel trick." Gaussian kernels are popular:

$$K(a_k, a_\ell) = \exp(-\|a_k - a_\ell\|^2/(2\sigma)), \quad \text{for some } \sigma > 0.$$

Nonlinear SVM



Application VIII: Logistic Regression

Binary logistic regression is similar to binary SVM, except that we seek a function p that gives odds of data vector a being in class 1 or class 2, rather than making a simple prediction.

Seek odds function p parametrized by $x \in \mathbb{R}^n$:

$$p(a; x) := (1 + e^{a^T x})^{-1}.$$

Choose x so that $p(a_j; x) \approx 1$ when $y_j = 1$ and $p(a_j; x) \approx 0$ when $y_j = 2$. Choose x to minimize a negative log likelihood function:

$$\mathcal{L}(x) = -\frac{1}{m} \left[\sum_{y_j=2} \log(1 - p(a_j; x)) + \sum_{y_j=1} \log p(a_j; x) \right]$$

mapping function: $\phi(a; x) = \text{index of max element of} \quad \begin{bmatrix} p(a; x) \\ 1 - p(a; x) \end{bmatrix}$.

Sparse solutions x are interesting because the indicate which components of a_j are critical to classification. Can solve: $\min_z \mathcal{L}(z) + \lambda ||z||_1$.

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Application IX: Multiclass Logistic Regression

Have M classes instead of just 2. M can be large e.g. identify phonemes in speech, identify line outages in a power grid.

Labels $y_{j\ell} = 1$ if data point j is in class ℓ ; $y_{j\ell} = 0$ otherwise; $\ell = 1, ..., M$. Find subvectors $x_{[\ell]}$, $\ell = 1, 2, ..., M$ such that if a_j is in class k we have

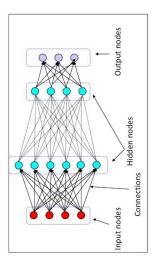
 $a_j^T x_{[k]} \gg a_j^T x_{[\ell]}$ for all $\ell \neq k$.

Find $x_{[\ell]}$, $\ell = 1, 2, \dots, M$ by minimizing a negative log-likelihood function:

$$f(x) = -\frac{1}{m} \sum_{j=1}^{m} \left[\sum_{\ell=1}^{M} y_{j\ell}(a_j^T x_{[\ell]}) - \log \left(\sum_{\ell=1}^{M} \exp(a_j^T x_{[\ell]}) \right) \right]$$

Can use group LASSO regularization terms to select important features from the vectors a_j , by imposing a common sparsity pattern on all $x_{[\ell]}$.

Application X: Deep Learning



Inputs are the vectors a_j , outputs are odds of a_j belonging to each class (as in multiclass logistic regression).

At each layer, inputs are converted to outputs by a linear transformation composed with an element-wise function:

$$a^{\ell+1} = \sigma(W^{\ell}a^{\ell} + g^{\ell}),$$

where a^{ℓ} is node values at layer ℓ , (W^{ℓ}, g^{ℓ}) are *parameters* in the network, σ is the element-wise function.

Deep Learning

The element-wise function σ makes transformations to scalar input:

- Logistic function: $t \rightarrow 1/(1 + e^{-t})$;
- Hinge: $t \rightarrow \max(t, 0)$;
- Bernoulli: random! $t \to 1$ with probability $1/(1 + e^{-t})$ and $t \to 0$ otherwise (inspired by neuron behavior).

The example depicted shows a completely connected network — but more typically networks are engineered to the application (speech processing, object recognition, \dots).

- local aggregation of inputs: pooling;
- restricted connectivity + constraints on weights (elements of W^ℓ matrices): convolutions.

Training Deep Learning Networks

The network contains many parameters — (W^{ℓ}, g^{ℓ}) , $\ell = 1, 2, ..., L$ in the notation above — that must be selected by training on the data (a_j, y_j) , j = 1, 2, ..., m.

Objective has the form:

$$\sum_{j=1}^m h(x; a_j, y_j)$$

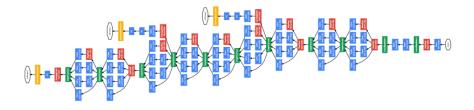
where $x = (W^1, g^1, W^2, g^2, ...)$ are the parameters in the model and h measures the mismatch between observed output y_j and the outputs produced by the model (as in multiclass logistic regression).

Nonlinear, Nonconvex. It's also random in some cases (but then we can work with expectation).

Composition of many simple functions.

GoogLeNet

Visual object recognition: Google's State of the Art network from 2014 [Szegedy et al., 2015].



A Basic Paradigm

Many optimization formulations in data analysis have this form:

$$\min_{x} \frac{1}{m} \sum_{j=1}^{m} h_j(x) + \lambda \Omega(x),$$

where

- h_j depends on parameters x of the mapping ϕ , and data items (a_j, y_j) ;
- Ω is the regularization term, often nonsmooth, convex, and separable in the components of x (but not always!).
- $\lambda \ge 0$ is the regularization parameter.
- (Ω could also be an indicator for simple set e.g. $x \ge 0$.)

Alternative formulation:

$$\min_{x} \frac{1}{m} \sum_{j=1}^{m} h_j(x)$$
 s.t. $\Omega(x) \leq \tau$.

Structure in $h_j(x)$ and Ω strongly influences the choice of algorithms.

Optimization in Data Analysis: Some Background

Optimization formulations of data analysis / machine learning problems were popular, and becoming more so... but not universal:

- Heuristics are popular: k-means clustering, random forests.
- Linear algebra approaches suffice in some applications.
- Greedy feature selection.

Algorithmic developments have sometimes happened independently in the machine learning and optimization communities, e.g. stochastic gradient from 1980s-2009.

Occasional contacts in the past:

- Mangasarian solving SVM formulations [Mangasarian, 1965, Wohlberg and Mangasarian, 1990, Bennett and Mangasarian, 1992]
- Backpropagation in neural networks equivalent to gradient descent [Mangasarian and Solodov, 1994]
- Basis pursuit [Chen et al., 1998].

...but now, crossover is much more frequent and systematic.

Optimization in Learning: Context

Naive use of "standard" optimization algorithms is usually not appropriate.

Large data sets make some standard operations expensive: Evaluation of gradients, Hessians, functions, Newton steps, line searches.

The optimization formulation is viewed as a sampled, empirical approximation of some underlying "infinite" reality.

- Don't need or want an exact minimizer of the stated objective this would be overfitting the empirical data. An approximate solution suffices.
- Choose the regularization parameter λ so that ϕ gives best performance on some "holdout" set of data: validation, tuning.

Low-Dimensional Structure

Generalizability often takes the form of low-dimensional structure, when the function is parametrized appropriately.

- Variable vector x may be sparse (feature selection, compressed sensing).
- Variable matrix X may be low-rank.
- Data objects *a_j* lie approximately in a low-dimensional subspace.

Example: Identify genetic risk factors for a disease by identifying the crucial locations on the genome (alleles).

- Convex formulations with regularization are often tractable and efficient in practice.
- Discrete or nonlinear formulations are more natural, but harder to solve and analyze. Recent advances make discrete formulations more appealing (Bertsimas et al.)

ALGORITHMS

Seek optimization algorithms — mostly elementary — that can exploit the structure of the formulations above.

- Full-gradient algorithms.
 - with projection and shrinking to handle Ω
- Accelerated gradient
- Stochastic gradient
 - and hybrids with full-gradient
- Coordinate descent
- Conditional gradient
- Augmented Lagrangian / ADMM

Everything Old is New Again

"Old" approaches from the optimization literature have become popular:

- Nesterov acceleration of gradient methods [Nesterov, 1983].
- Alternating direction method of multipliers (ADMM) [Eckstein and Bertsekas, 1992].
- Parallel coordinate descent and incremental gradient algorithms [Bertsekas and Tsitsiklis, 1989]
- Stochastic gradient [Robbins and Monro, 1951]
- Frank-Wolfe / conditional gradient [Frank and Wolfe, 1956, Dunn, 1979].

Many extensions have been made to these methods and their convergence analysis. Many variants and adaptations proposed.

Gradient Methods

Basic formulation, without regularization:

$$\min_{x} H(x) := \frac{1}{m} \sum_{j=1}^{m} h_j(x).$$

Steepest descent takes steps in the negative gradient direction:

$$x^{k+1} \leftarrow x^k - \alpha_k \nabla H(x^k).$$

Classical analysis applies for H smooth, convex, and

Lipschitz:
$$\|\nabla H(x) - \nabla H(z)\| \le L \|x - z\|$$
, for some $L > 0$,
Lojasiewicz: $\|\nabla H(x)\|^2 \ge 2\mu [H(x) - H^*]$, for some $\mu \ge 0$.

 $\mu = 0$: sublinear convergence for $\alpha_k \equiv 1/L$: $H(x^k) - H^* = O(1/k)$. $\mu > 0$: linear convergence with $\alpha_k \equiv 1/L$:

$$H(x^{k}) - H^{*} \leq \left(1 - \frac{\mu}{L}\right)^{k} [H(x^{0}) - H^{*}].$$

Shrinking, Projecting for $\boldsymbol{\Omega}$

In the regularized form:

$$\min_{x} H(x) + \lambda \Omega(x),$$

the gradient step in H can be combined with a shrink operation in which the effect of Ω is accounted for exactly:

$$x^{k+1} = rg\min_{z} \nabla H(x^k)^T (z - x^k) + rac{1}{2lpha_k} \|z - x_k\|^2 + \lambda \Omega(z).$$

When Ω is the indicator function for a convex set, x^{k+1} is the projection of $x^k - \alpha_k \nabla H(x^k)$ onto this set: gradient projection.

For many Ω of interest, this problem can be solved quickly (e.g. O(n)). Algorithms and convergence theory for steepest descent on smooth H usually extend to this setting (for convex Ω).

Accelerated Gradient and Momentum

Accelerated gradient methods [Nesterov, 1983] highly influential.

Fundamental idea: **Momentum!** Search direction at iteration k depends on the latest gradient $\nabla H(x^k)$ and also the search direction at iteration k - 1, which encodes gradient information from earlier iterations.

Heavy-ball & conjugate gradient (incl. nonlinear CG) also use momentum. Heavy-Ball for $\min_x H(x)$:

$$x^{k+1} = x^k - \alpha \nabla H(x^k) + \beta (x^k - x^{k-1}).$$

Nesterov's optimal method:

$$x^{k+1} = x^k - \alpha_k \nabla H(x^k + \beta_k(x^k - x^{k-1})) + \beta_k(x^k - x^{k-1}).$$

Typically $\alpha_k \approx 1/L$ and $\beta_k \approx 1$.

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Typically $\alpha_k \approx 1/L$ and $\beta_k \approx 1$.

Accelerated Gradient Convergence

Typical convergence:

Weakly convex
$$\mu = 0$$
: $H(x^k) - H^* = O(1/k^2)$;
Strongly convex $\mu > 0$: $H(x^k) - H^* \le M \left(1 - \sqrt{\frac{\mu}{L}}\right)^k [H(x^0) - H^*]$.

- Approach can be extended to regularized functions $H(x) + \lambda \Omega(x)$ [Beck and Teboulle, 2009].
- Partial-gradient approaches (stochastic gradient, coordinate descent) can be accelerated in similar ways.

Are Gradient Methods Useful?

They're useful for some problems in which X is a matrix, in which full gradients are practical to compute.

- Matrix completion, including explicitly parametrized problems with $X = LR^{T}$ or $X = ZZ^{T}$; and Nonnegative matrix factorization.
- Subspace identification;
- Sparse covariance estimation;

They are less appealing when m is large. To calculate

$$abla H(x) = rac{1}{m} \sum_{j=1}^m
abla h_j(x),$$

generally need to make a full pass through the data.

Often not practical for massive data sets. But can be hybridized with stochastic gradient methods (see below).

Stochastic Gradient (SG)

For $H(x) = (1/m) \sum_{j=1}^{m} h_j(x)$, iteration k has the form:

• Choose $j_k \in \{1, 2, \dots, m\}$ uniformly at random;

• Set
$$x^{k+1} \leftarrow x^k - \alpha_k \nabla h_{j_k}(x^k)$$
.

 $\nabla h_{j_k}(x^k)$ is a proxy for $\nabla H(x^k)$ but it depends on just one data item a_{j_k} and is much cheaper to evaluate.

Unbiased — $\mathbb{E}_j \nabla h_j(x) = \nabla H(x)$ — but the variance may be very large.

• Average the iterates for more robust convergence:

$$\bar{x}^k = \frac{\sum_{\ell=1}^k \gamma_\ell x^\ell}{\sum_{\ell=0}^k \gamma_\ell}$$
, where γ_ℓ are positive weights.

- Minibatch: Use a set J_k ⊂ {1, 2, ..., m} rather than a single item. (Smaller variance in the gradient estimate.)
- See [Nemirovski et al., 2009] and many other works.

Stochastic Gradient (SG)

Convergence results for h_j convex require bounds on the variance of the gradient estimate:

$$\frac{1}{m}\sum_{j=1}^m \|\nabla h_j(x)\|_2^2 \leq B^2 + L_g \|x - x^*\|^2.$$

Analyze expected convergence, e.g. $\mathbb{E}(f(x^k) - f^*)$ or $\mathbb{E}(f(\bar{x}^k) - f^*)$, where the expectation is over the sequence of indices j_0, j_1, j_2, \ldots .

Sample results:

- *H* strongly convex, $\alpha_k \sim 1/k$: $\mathbb{E}(H(x^k) H^*) = O(1/k)$;
- *H* weakly convex, $\alpha_k \sim 1/\sqrt{k}$: $\mathbb{E}(H(\bar{x}^k) H^*) = O(1/\sqrt{k})$.
- B = 0, H strongly convex, $\alpha_k = \text{const: } \mathbb{E}(||x^k x^*||_2^2) = O(\rho^k)$ for some $\rho \in (0, 1)$.

Generalizes beyond finite sums, to $H(x) = \mathbb{E}_{\xi} f(x; \xi)$, where ξ is random.

Stochastic Gradient Applications

SG fits well the summation form of H (with large m), so has widespread applications:

- SVM (primal formulation).
- Logistic regression: binary and multiclass.
- Deep Learning. The Killer App! (Nonconvex) [LeCun et al., 1998]
- Subspace Identification (GROUSE): Project stochastic gradient searches onto subspace [Balzano and Wright, 2014].

Incremental Gradient (Finite Sum, Cyclic Order)

$$H(x)=\frac{1}{m}\sum_{j=1}^m h_j(x).$$

Steps: $x^{k+1} \leftarrow x^k - \alpha_k \nabla h_{j_k}(x^k)$, where $j_k = 1, 2, \dots, m, 1, 2, \dots$.

Typical requirements on step lengths α_k , for convergence:

$$\sum_{k=1}^{\infty} \alpha_k = \infty, \quad \sum_{k=1}^{\infty} \alpha_k^2 < \infty.$$

(The last condition required only for convergence of iterates $\{x^k\}$ as well as function values $\{f(x^k)\}$.)

Sublinear convergence rates proved in [Nedić and Bertsekas, 2000] for steplength $\alpha_k = C/(k+1)$.

Random Reshuffling [Gürbüzbalaban et al., 2015] (reordering order of functions between cycles) still yields sublinear convergence, but with constants reduced (sometimes dramatically) below worst-case values.

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Hybrids of Full Gradient and Stochastic Gradient

Stabilize SG by hybridizing with steepest descent (full-gradient). Typically get *linear convergence* for strongly convex functions, sublinear for weakly convex.

SAG: [LeRoux et al., 2012] Maintain approximations g_j to ∇h_j , use search direction $-(1/m)\sum_{j=1}^m g_j$. At iteration k, choose j_k at random, and update $g_{j_k} = \nabla h_{j_k}(x^k)$.

SAGA: [Defazio et al., 2014] Similar to SAG, but use search direction

$$-
abla h_{j_k}(x^k)+g_{j_k}-rac{1}{m}\sum_{j=1}^m g_j.$$

SVRG: [Johnson and Zhang, 2013] Similar again, but periodically do a full gradient evaluation to refresh all g_j .

Too much storage, **BUT** when h_j have the "ERM" form $h_j(a_j^T x)$ (linear least squares, linear SVM), all gradients can be stored in a scalar:

$$\nabla_{\mathsf{x}} h_j(a_j^\mathsf{T} \mathsf{x}) = a_j h_j'(a_j^\mathsf{T} \mathsf{x}).$$

Coordinate Descent (CD) Framework

... for smooth unconstrained minimization: $\min_x H(x)$:

Set Choose
$$x^1 \in \mathbb{R}^n$$
;
for $\ell = 0, 1, 2, ..., n$ do
for $j = 1, 2, ..., n$ do
Define $k = \ell n + j$
Choose index $i = i(\ell, j) \in \{1, 2, ..., n\}$;
Choose $\alpha_k > 0$;
 $x^{k+1} \leftarrow x^k - \alpha_k \nabla_i H(x^k) e_i$;
end for
end for

where

- $e_i = (0, \ldots, 0, 1, 0, \ldots, 0)^T$: the *i*th coordinate vector;
- $\nabla_i H(x) = i$ th component of the gradient $\nabla H(x)$;
- $\alpha_k > 0$ is the step length.

Economics of Partial Gradients (Least Squares, SVM)

$$H(x) = \frac{1}{m} \sum_{j=1}^{m} h_j(a_j^T x) + \lambda \sum_{i=1}^{n} \Omega_i(x_i)$$

and define A as the $m \times n$ matrix with *j*th row a_i^T .

Maintain g = Ax and $\nabla h_j(g_j)$ for j = 1, 2, ..., m. A CD step on coordinate i_k proceeds as follows:

• Compute the *i_k* element of the gradient of the summation term:

$$\nabla_{i_k} H(x) = \frac{1}{m} \sum_{j: A_{j, i_k} \neq 0} A_{j, i_k} \nabla h_j(g_j);$$

- Use this information, along with $\lambda \Omega_{i_k}$, to update: $x_{i_k} \leftarrow x_{i_k} + d_{i_k}$;
- Update $g_j \leftarrow g_j + A_{j,i_k} d_{i_k}$ and $\nabla h_j(g_j)$ (only for j s.t. $A_{j,i_k} \neq 0$);

Cost: $O(\text{nonzeros in } A_{i_k})$, vs O(nonzeros in A) for a full-gradient method. Cheaper (by factor $\approx 1/n$) for one element than the full gradient.

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CD Variants

- CCD (Cyclic CD): $i(\ell, j) = j$.
- RCD (Randomized CD a.k.a. Stochastic CD): *i*(ℓ, *j*) is chosen uniformly at random from {1, 2, ..., n}.
- RCCD (Randomized Cyclic CD):
 - At the start of epoch ℓ , we choose a random permutation of $\{1, 2, \ldots, n\}$, denoted by π_{ℓ} .
 - Index $i(\ell, j)$ is chosen to be the *j*th entry in π_{ℓ} .

Important quantities in analysis:

• L_{\max} : componentwise Lipschitz constant for ∇H :

$$|
abla_i H(x+te_i) -
abla_i H(x)| \leq L_i |t|, \quad L_{\max} = \max_{i=1,2,\dots,n} L_i.$$

- L: usual Lipschitz constant: $|\nabla H(x+d) \nabla H(x)| \le L \|d\|$.
- Lojasiewicz constant μ : $\|\nabla H(x)\|^2 \ge 2\mu[H(x) H^*]$

Randomized CD Convergence

Of the three variants, convergence of the randomized form has by far the most elementary analysis [Nesterov, 2012].

Get convergence rates for quantity $\phi_k := \mathbb{E}(H(x^k) - x^*)$.

$$\mu > 0: \quad \phi_{k+1} \le \left(1 - \frac{\mu}{nL_{\max}}\right)\phi_k, \quad k = 1, 2, \dots,$$
$$\mu = 0: \quad \phi_k \le \frac{2nL_{\max}R_0^2}{k}, \qquad k = 1, 2, \dots,$$

where R_0 bounds distance from x^0 to solution set.

If the economics of evaluating gradient components are right, this can be a factor L/L_{max} faster than full-gradient steepest descent!

This ratio is in range [1, n]. Maximized by $H(x) = (\mathbf{11}^T)x$.

Functions like this are good cases for RCD and RCCD, which are much faster than CCD or steepest descent.

Cyclic and Randomized Cyclic CD Convergence

Analysis of [Beck and Tetruashvili, 2013] treats CCD as an approximate form of Steepest Descent, bounding improvement in f over one cycle in terms of the gradient at the start of the cycle.

Get linear and sublinear rates that are slower than both RCD and Steepest Descent. This analysis is fairly tight — recent analysis of [Sun and Ye, 2016] confirms slow rates.

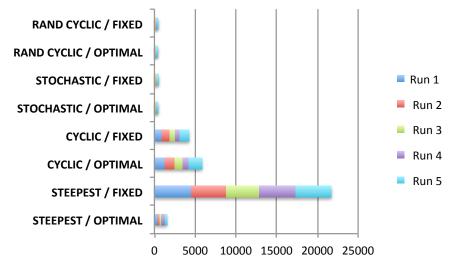
Same analysis applies to Randomized Cyclic (RCCD), but practical results for RCCD are much better, and usually at least as good as RCD. Results on quadratic function

$$H(x) = (1/2)x^T A x$$

demonstrate this behavior (see [Wright, 2015] and my talks in 2015).

We can explain good behavior of RCCD now! [Lee and Wright, 2016]

Computations: Large $\mathbf{11}^T$ component in A ($\zeta = .3$)



cond(A) \approx 3000, $L_{max} = 1$, $L \approx$ 50. Steepest Descent and CCD are poor.

CD Extensions

- Block CD: Replace single component *i* by block $I \subset \{1, 2, ..., n\}$.
- Dual nonlinear SVM [Platt, 1999]. Choose *two* components of α per iteration, to stay feasible w.r.t. constraint y^Tα = 0.
- Can be accelerated (efficiently) using "Nesterov" techniques: [Nesterov, 2012, Lee and Sidford, 2013].
- Adaptable to the separable regularized case $H(x) + \lambda \Omega(x)$.
- Parallel asynchronous variants, suitable for implementation on shared-memory multicore computers, have been proposed and analyzed. [Bertsekas and Tsitsiklis, 1989, Liu and Wright, 2015, Liu et al., 2015]

Conditional Gradient / "Frank-Wolfe"

 $\min_{x\in\Omega} f(x),$

where f is a convex function and Ω is a closed, bounded, convex set. Start at $x_0 \in \Omega$. At iteration k:

$$v_k := \arg\min_{v \in \Omega} v^T \nabla f(x_k);$$

$$x_{k+1} := x_k + \alpha_k (v_k - x_k), \quad \alpha_k = \frac{2}{k+2}.$$

- Potentially useful when it is easy to minimize a linear function over the *original* constraint set Ω;
- Admits an elementary convergence theory: 1/k sublinear rate.
- Same convergence rate holds if we use a line search for α_k .

Revived by [Jaggi, 2013].

Smooth Nonlinear Optimization: Avoiding Saddle Points

Much recent interest in methods for smooth nonlinear nonconvex problems $\min_x f(x)$ with guaranteed convergence to points satisfying approximate second-order optimality conditions:

$$abla f(x) = O(\epsilon^2), \quad
abla^2 f(x) \ge -\epsilon I, \quad \text{for small } \epsilon > 0.$$

Useful for matrix optimization problems (see earlier).

Also potentially useful for deep learning. (It's believed that saddle points are much more plentiful than local minima, so methods that can escape saddle points are interesting.)

A fundamental approach: cubic regularization: Given a Lipschitz constant L on the Hessian of f, form the following upper bounding cubic approx:

$$T_L(z;x) := f(x) + \nabla f(x)^T (z-x) + \frac{1}{2} (z-x)^T \nabla^2 f(x) (z-x) + \frac{L}{6} ||z-x||^3.$$

Basic algorithm:

$$x^{k+1} := \arg\min_{z} T_L(z; x^k).$$

Convergence to Second-Order Necessary Points

[Nesterov and Polyak, 2006]; see also [Griewank, 1981], [Cartis et al., 2011a, Cartis et al., 2011b].

For the cubic-regularized Newton method, we have

$$\|
abla f(x^{k+1})\| = O(\|x^{k+1} - x^k\|^2), \quad
abla^2 f(x^{k+1}) \ge -\|x^{k+1} - x^k\|I.$$

[Nesterov and Polyak, 2006, Theorem 1]: Given lower bound f^* on f and small $\epsilon > 0$, have

$$egin{aligned} \|
abla f(x^k)\| &\leq \epsilon & ext{within } k = O(\epsilon^{-3/2}) ext{ iterations;} \
abla^2 f(x^k) &\geq -\epsilon I & ext{within } k = O(\epsilon^{-3}) ext{ iterations,} \end{aligned}$$

where the constants in $O(\cdot)$ depend on $[f(x^0) - f^*]$ and L.

Variants and Extensions

- Trust-region methods (second-order model with constraint $||z x^k||_2^2 \le \Delta$) have similar complexity properties, when modified.
- Standard first-order method that evaluates Hessian only when ∇f(x^k) is small has slightly worse complexity.
- Extensions using third-derivative tensor can find third-order minima. Finding a fourth-order minimum is NP-hard. [Ananakumar and Ge, 2016]

But are saddle points really an issue in practice?

First-order methods are unlikely to converge to such points. An elementary application of the stable manifold theorem [Lee et al., 2016] shows that convergence of standard gradient descent converges to a strict saddle point only for x^0 in a measure-zero set. (No complexity guarantees.)

Augmented Lagrangian

Consider the linearly constrained problem,

min f(x) s.t. Ax = b,

where $f : \mathbb{R}^n \to \mathbb{R}$ is convex.

Define the Lagrangian function:

$$\mathcal{L}(x,\lambda) := f(x) + \lambda^{T}(Ax - b).$$

 x^* is a solution if and only if there exists a vector of Lagrange multipliers $\lambda^* \in \mathbb{R}^m$ such that

$$-A^T\lambda^* \in \partial f(x^*), \quad Ax^* = b,$$

or equivalently:

$$0\in \partial_x\mathcal{L}(x^*,\lambda^*), \quad
abla_\lambda\mathcal{L}(x^*,\lambda^*)=0.$$

Augmented Lagrangian

The augmented Lagrangian is (with $\rho > 0$)

$$\mathcal{L}(x,\lambda;\rho) := \underbrace{f(x) + \lambda^{T}(Ax - b)}_{\text{Lagrangian}} + \underbrace{\frac{\rho}{2} \|Ax - b\|_{2}^{2}}_{\text{"augmentation"}}$$

Basic Augmented Lagrangian (a.k.a. method of multipliers) is

$$\begin{aligned} x_k &= \arg\min_{x} \mathcal{L}(x, \lambda_{k-1}; \rho); \\ \lambda_k &= \lambda_{k-1} + \rho(Ax_k - b); \end{aligned}$$

[Hestenes, 1969, Powell, 1969]

Some constraints on x (such as $x \in \Omega$) can be handled explicitly:

$$x_{k} = \arg\min_{\mathbf{x}\in\Omega} \mathcal{L}(\mathbf{x}, \lambda_{k-1}; \rho);$$
$$\lambda_{k} = \lambda_{k-1} + \rho(Ax_{k} - b);$$

Alternating Direction Method of Multipliers (ADMM)

$$\min_{(x\in\Omega_x,z\in\Omega_z)} f(x) + h(z) \quad \text{s.t.} \quad Ax + Bz = c,$$

for which the Augmented Lagrangian is

$$\mathcal{L}(x,z,\lambda;\rho) := f(x) + h(z) + \lambda^{T}(Ax + Bz - c) + \frac{\rho}{2} \|Ax - Bz - c\|_{2}^{2}.$$

Standard AL would minimize $\mathcal{L}(x, z, \lambda; \rho)$ w.r.t. (x, z) jointly. However, since coupled in the quadratic term, separability is lost.

In ADMM, minimize over x and z separately and sequentially:

$$\begin{aligned} x_k &= \arg\min_{x\in\Omega_x} \mathcal{L}(x, z_{k-1}, \lambda_{k-1}; \rho); \\ z_k &= \arg\min_{z\in\Omega_z} \mathcal{L}(x_k, z, \lambda_{k-1}; \rho); \\ \lambda_k &= \lambda_{k-1} + \rho(Ax_k + Bz_k - c). \end{aligned}$$

Extremely useful framework for many data analysis / learning settings. Major references: [Eckstein and Bertsekas, 1992, Boyd et al., 2011]

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Optimization in Data Analysis

ADMM for Consensus Optimization

Given the unconstrained (but separable) problem

$$\min \sum_{i=1}^m f_i(x),$$

form m copies of the x, with the original x as a "master" variable:

$$\min_{x,x^1,x^2,...,x^m} \sum_{i=1}^m f_i(x^i) \text{ subject to } x^i - x = 0, \ i = 1, 2, ..., m.$$

Apply ADMM, with $z = (x^1, x^2, \dots, x^m)$. Get

$$\mathcal{L}(x, x^1, x^2, \dots, x^m, \lambda^1, \dots, \lambda^m; \rho) = \sum_{i=1}^m f_i(x^i) + (\lambda^i)^T (x^i - x) + \frac{\rho}{2} \|x^i - x\|_2^2.$$

The minimization w.r.t. $z = (x^1, x^2, \dots, x^m)$ is separable!

$$x_{k}^{i} = \arg\min_{x^{i}} f_{i}(x^{i}) + (\lambda_{k-1}^{i})^{T}(x^{i} - x_{k-1}) + \frac{\rho_{k}}{2} \|x^{i} - x_{k-1}\|_{2}^{2}, \ i = 1, 2, \dots, m.$$

Can be implemented in parallel.

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Consensus, continued

The minimization w.r.t. x can be done explicitly — averaging:

$$x_k = \frac{1}{m} \sum_{i=1}^m \left(x_k^i + \frac{1}{\rho_k} \lambda_{k-1}^i \right).$$

Update to λ^i can also be done in parallel, once the new x_k is known (and communicated):

$$\lambda_k^i = \lambda_{k-1}^i + \rho_k (x_k^i - x_k), \quad i = 1, 2, \dots, m.$$

If the initial λ_0^i have $\sum_{i=1}^m \lambda_0^i = 0$, we have $\sum_{i=1}^m \lambda_k^i = 0$ at all iterations k. Can simplify the update for x_k :

$$x_k = \frac{1}{m} \sum_{i=1}^m x_k^i.$$

Not Discussed!

Many interesting topics not mentioned, including

- quasi-Newton and approximate Newton methods.
- Linear equations Ax = b: Kaczmarz algorithms.
- Image processing: denoising and deblurring using regularization.
- Graphs: detect structure and cliques, consensus optimization,
- Integer and combinatorial formulations.
- Parallel variants: synchronous and asynchronous.
- Online learning.

Conclusions: Optimization in Data Analysis

- HUGE interest across multiple communities.
- Ongoing challenges because of increasing scale and complexity of data analysis problems, and the computational platforms on which they are being solved.
- Optimization techniques are meshed with the applications.
- The optimization / data analysis / machine learning research communities are becoming "meshed" too!

FIN

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