A majorized proximal point dual Newton algorithm for nonconvex statistical optimization problems

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The Sixth International Conference on Continuous Optimization Technical University (TU) of Berlin, August 3-8, 2019.

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Second Order Methods with First Order Costs

Newton's methods

Our approach — Newton's methods (second order methods) with low costs [can be very low]



Figure: Sir Isaac Newton (Niu Dun) (4 January 1643 - 31 March 1727)

Solving a sparse + low rank linear equation problem: a basic idea

• One core mathematical problem is to solve the following linear equation

$$Bx = b,$$

where $B \in \Re^{n \times n}$ and $b \in \Re^n$

Assume that B is non-singular. One can use the Gaussian elimination method [the ancient Chinese (Jiu Zhang Suan Shu) and Indians discovered this method thousand years ago] to get

$$x = B^{-1}b$$

with the cost of $O(n^3)$ flops – way too high for a big n.

● If B is "sparse", e.g., $B = I + uv^T$, where $u, v \in \Re^n$, one can reduce the cost from $O(n^3)$ to O(n) via the Sherman-Morrison-Woodbury formula

$$x = B^{-1}b = (I + uv^T)^{-1}b = b - v \underbrace{(1 + v^T u)^{-1}}_{\text{number}} (u^T b).$$

The Setting

Let $X \in \Re^{m \times n}$ be the input data and b be the response variables with a noise vector $\varepsilon = b - X\ddot{\beta}$. Let $\lambda > 0$. One of the most commonly used models to control the overfitting and/or variable selection is the Lasso model

$$\min_{\beta \in \Re^n} \left\{ \frac{1}{2} \| X\beta - b \|^2 + \lambda \| \beta \|_1 \right\}$$

which relies on knowing the standard deviation of the noise. Here in the convex case we are interested in the more general model

$$\min_{\beta \in \Re^n} \left\{ \underbrace{h(X\beta)}_{f(\beta)} + p(\beta) \right\}$$

where both $h(\cdot)$ and $p(\cdot)$ are proper and closed convex functions, which can be nonsmooth or non-Lipschitzian. Here, h is not assumed to be differentiable!!!

Simple convex examples

One interesting example is the square-root Lasso (srLasso) model (Alex Belloni et al. (2011))

$$\min_{\beta \in \Re^n} \left\{ \|X\beta - b\| + \lambda \|\beta\|_1 \right\}$$

which is equivalent to the robust least regression (Huan Xu et al. 2010)

$$\min_{\beta \in \Re^n} \left\{ \max_{\Delta X \in \mathcal{U}} \| b - (X + \Delta X) \beta \| \right\}$$

with the uncertainty set

$$\mathcal{U} := \{ (\Delta_1, \dots, \Delta_n) \mid \|\Delta_i\| \le \lambda, \ i = 1, \dots, n \}$$

Another commonly used example is the constrained Lasso model

 $\min_{\beta \in \Re^n} \left\{ \|\beta\|_1 \mid \|X\beta - b\| \le \tau \right\}$

Here $h(\cdot)$ is the indicator function

 $h(y) := \delta_{B^{\tau}}(y) \quad \forall y \in \Re^m$

over the ball $B^\tau:=\{y\in\Re^m\,|\,||y||\leq\tau\}$ centered at 0 with radius $\tau>0$

More Lasso-type problems

There are many more convex Lasso-type models:

(LASSO)
$$\min\left\{\frac{1}{2}\|X\beta - b\|^2 + \lambda\|\beta\|_1 \mid \beta \in \Re^n\right\}$$

where $\lambda > 0$.

(Fused LASSO) $\min\left\{\frac{1}{2}\|X\beta - b\|^2 + \lambda\|\beta\|_1 + \lambda_2\|B\beta\|_1\right\}$ $B = \begin{pmatrix} 1 & -1 \\ & 1 & -1 \\ & & \ddots & \ddots \\ & & & 1 & -1 \end{pmatrix}$

(Clustered LASSO)
$$\min\left\{\frac{1}{2}\|X\beta - b\|^2 + \lambda\|\beta\|_1 + \lambda_2 \sum_{i=1}^n \sum_{j=i+1}^n |\beta_i - \beta_j|\right\}$$

Note that the above problem is not numerically solvable if n is large as the objective function value computation itself would cost $O(n^2)$ flops.

Fortunately, [Lin-Liu-S.-Toh 2018] showed

$$\sum_{1 \le i < j \le n} |\beta_i - \beta_j| = \langle w, \beta^{\downarrow} \rangle,$$

where β^{\downarrow} is the vector whose components are those of β sorted in a non-increasing order, i.e. $\beta_1^{\downarrow} \geq \beta_2^{\downarrow} \geq \cdots \geq \beta_n^{\downarrow}$ [costs $O(n \log n)$] and the weight vector $w \in \Re^n$ is defined by

$$w_k = n - 2k + 1, \ k = 1, \cdots, n.$$

(SLOPE, Ordered Lasso)

$$\min\left\{\frac{1}{2}\|X\beta - b\|^2 + \sum_{i=1}^n \lambda_i |\beta|_i^{\downarrow}\right\}$$

with parameters $\lambda_1 \geq \lambda_2 \geq \cdots \geq \lambda_n$ and $\lambda_1 > 0$.

We are interested in n (number of features) large and/or m (number of samples) large. Note that the regularization term in SLOPE (ordered Lasso) is not separable.

More on the loss functions

In the loss function part, f can be the logistic regression function, defined as below: given $b \in \Re^m$ and $X \in \Re^{m \times n}$,

$$f(\beta) = \sum_{i=1}^{m} \log(1 + \exp(-b_i (X\beta)_i))$$
(1)

• Define $h: \Re^m \to \Re$ as follows

$$h(z) = \sum_{i=1}^{m} \log(1 + \exp(-b_i z_i)) \quad \forall z \in \Re^m$$

The function f defined by (1) can be written as

 $f(\beta) = h(X\beta)$

In general, the loss function $f: \Re^{pK} \to \Re$ can take as the multinomial logistic regression function: given $A := (A_1, \ldots, A_N)^T \in \Re^{N \times p}$,

$$f(\beta) = -\sum_{i=1}^{N} \left(\sum_{k=1}^{K} y_{ik} A_i^T \beta_k - \log \sum_{k=1}^{K} \exp(A_i^T \beta_k) \right)$$
(2)

• Define $h: \Re^{NK} \to \Re$ as follows:

$$h(z) := -\sum_{i=1}^{N} \left(\sum_{k=1}^{K} y_{ik} Z_{ik} - \log \sum_{k=1}^{K} \exp(Z_{ik}) \right), \text{ with } Z := \operatorname{mat}(z)$$

Then the function f defined by (2) can be written as

 $f(\beta) := h(X\beta), \text{ with } X := I \otimes A \in \Re^{NK \times pK}$

A proper nonconvex regularization can achieve a sparse estimation with fewer measurements, faster convergence and more robust against noises.

In this talk, we aim to develop an efficient and robust algorithm for solving the following nonconvex problem (\mathbf{P}) :

$$\min_{\beta \in \Re^n} \left\{ g(\beta) := \underbrace{h(X\beta)}_{f(\beta)} + \underbrace{p(\beta) - q(\beta)}_{r(\beta)} \right\}$$
(3)

Here $p: \Re^n \to (-\infty, +\infty]$ is a proper closed convex function and $q: \Re^n \to \Re$ is a finite-valued (smooth, not essential) convex function. Moreover, we require the proximal functions of h and p to be (strongly) semismooth.

For $\lambda > 0$, the SCAD regularization is defined by $r(\beta) = p(\beta) - q(\beta)$ with

$$\begin{split} p(\beta) &= \lambda \|\beta\|_1 \\ q(\beta) &= \sum_{i=1}^n \left\{ \begin{array}{ll} 0, & \text{if } |\beta_i| < \lambda \\ \frac{(|\beta_i| - \lambda)^2}{2(a_s - 1)}, & \text{if } \lambda \leq |\beta_i| \leq a_s \lambda \\ \lambda |\beta_i| - \frac{a_s + 1}{2} \lambda^2, & \text{if } |\beta_i| > a_s \lambda \end{array} \right. \end{split}$$

Note that $q(\cdot)$ is continuously differentiable. In our numerical experiments, we take $a_s = 3.7$.

Nonconvex regularizers: MCP

For two positive parameters $a_m>2$ and $\lambda,$ the MCP regularization can be defined as $r(\beta)=p(\beta)-q(\beta)$ with

$$\begin{split} p(\beta) &= \lambda \|\beta\|_1 \\ q(\beta) &= \sum_{i=1}^n \begin{cases} \frac{\beta_i^2}{a_m}, & \text{if } |\beta_i| \le a_m \lambda, \\ 2\lambda |\beta_i| - a_m \lambda^2, & \text{if } |\beta_i| > a_m \lambda \end{cases} \end{split}$$

The function $q(\cdot)$ is continuously differentiable with its derivative given by

$$\frac{\partial q(\beta)}{\partial \beta_i} = \begin{cases} \frac{2\beta_i}{a_m}, & \text{if } |\beta_i| \le a_m \lambda, \\ 2\lambda \operatorname{sign}(\beta_i), & \text{if } |\beta_i| > a_m \lambda \end{cases}$$

In our numerical experiments, we take $a_m = 3.7$.

Nonconvex regularizers: Difference of Ky Fan norms

For any positive integer k, let $\|\cdot\|_{(k)}$ denote Ky Fan's k-norm function, i.e., for any $\beta \in \Re^n$, $\|\beta\|_{(k)}$ is the sum of the first k largest absolute values of β .

By noting that the cardinality constraint

$$\|\beta\|_0 \leq k$$

can be written equivalently as

$$\|\beta\|_{(k)} = \|\beta\|_1,$$

in the majorized penalty method (Gao & S., 2010) we define the regularization term $r(\beta)=p(\beta)-q(\beta)$ with

$$p(\beta) = ||\beta||_1 \quad \& \quad q(\beta) = ||\beta||_{(k)} \quad \forall \beta \in \Re^n$$

or

$$p(\beta) = \|\beta\|_{(k+1)} \quad \& \quad q(\beta) = \|\beta\|_{(k)} \quad \forall \beta \in \Re^n$$

Note that $q(\cdot)$ is continuously differentiable near any β with $\|\beta\|_0 = k$.

Next, we shall consider the promised majorized proximal point dual Newton algorithm (mPPDNA) to solve the following problem [e.g., the srLasso problem for example]

$$\min_{\beta \in \Re^n} \left\{ h(X\beta) + p(\beta) - q(\beta) \right\}$$

- In Stage 1, replace q by its linear approximation at the origin [when q(0) = 0 and $0 \in \partial q(0)$, which hold true for many interesting cases of q, we just delete q from the original problem] and add "proper" proximal terms to obtain an initial point for the second stage.
- In Stage 2, a series of majorized proximal subproblems are solved to obtain an approximate solution point.

Given $\sigma > 0$, $\tau > 0$, $\tilde{\beta} \in \Re^n$, $\tilde{v} \in \Re^n$, $\tilde{b} \in \Re^m$, in our main algorithm (mPPDNA) we need to solve the following minimization subproblem

$$\min_{\beta \in \Re^n} \left\{ g(\beta; \sigma, \tau, \tilde{\beta}, \tilde{v}, \tilde{b}) := \underbrace{h(X\beta) + p(\beta)}_{\text{convex}} - \underbrace{(q(\tilde{\beta}) + \langle \tilde{v}, \beta - \tilde{\beta} \rangle)}_{\text{linear}} + \frac{\sigma}{2} \|\beta - \tilde{\beta}\|^2 + \frac{\tau}{2} \|X\beta - \tilde{b}\|^2 \right\}$$
(4)

Here, $\tilde{v} \in \partial q(\tilde{\beta})$. Obviously, $g(\cdot; \sigma, \tau, \tilde{\beta}, \tilde{v}, \tilde{b})$ is a strongly convex function albeit nonsmooth or non-Lipschitian.

- The big question is how one can solve (4) in a fast and robust way!!!
- For the convex case: $q \equiv 0$
- If h is strongly convex on $\mathcal E$, we can take $\tau=0$ though not necessary

The dual of the subproblem

The dual of (4), after converting it into the minimization form and ignoring the constant term, is

$$\min_{u \in \Re^m} \left\{ \phi(u; \sigma, \tau) := \frac{\tau}{2} \|\tilde{b} + \tau^{-1}u\|^2 - e_{\tau}h(\tilde{b} + \tau^{-1}u) + \frac{\sigma}{2} \|\tilde{\beta} + \sigma^{-1}(\tilde{v} - X^*u)\|^2 - e_{\sigma}p(\tilde{\beta} + \sigma^{-1}(\tilde{v} - X^*u)) \right\}.$$

Recall that for any $t>0, \, e_tf(\cdot)$ is the Moreau envelope of a closed proper convex function f, associated with t, given by

$$e_t f(x) := \min_{z \in \Re^n} \left\{ f(z) + \frac{t}{2} \|z - x\|^2 \right\}, \quad \forall \ x \in \Re^n.$$
(5)

Here $e_t f(\cdot)$ is continuously differentiable with

$$\nabla e_t f(x) = t[x - \mathsf{P}_t f(x)], \quad \forall x \in \Re^n,$$

where $P_t f(x)$ is the unique optimal solution to problem (5). $P_t f(\cdot)$, called the proximal mapping of f, is globally Lipschitz continuous with modulus 1.

A semismooth Newton method for the subproblem

We shall apply the superlinearly (quadratically) convergent sparse semismooth Newton method to find the solution \bar{u} of the nonsmooth equations

$$\nabla \phi(u;\sigma,\tau) = \mathsf{P}_{\tau} h(\tilde{b} + \tau^{-1}u) - X \mathsf{P}_{\sigma} p(\tilde{\beta} + \sigma^{-1}(\tilde{v} - X^*u)) = 0.$$

Then the unique optimal solution $\bar{\beta}$ to problem (4) is

$$\bar{\beta} = \mathsf{P}_{\sigma} p(\tilde{\beta} + \sigma^{-1}(\tilde{v} - X^* \bar{u})).$$

Proposition

Suppose that problem (4) is nondegenerate, which holds true if $f(\cdot) \equiv h(X \cdot)$ is continuously differentiable near $\bar{\beta}$ (this is the no-overfitting assumption for the squared root Lasso problem). Then all the elements in Clarke's generalized Jacobian $\partial^2 \phi(\bar{u})$ are self-adjoint and positive definite.

A globalized SSN algorithm for the subproblem (4)

Algorithm SSN (SSN(σ, τ)): Given $\mu \in (0, \frac{1}{2})$, $\overline{\eta} \in (0, 1)$, $\overline{\tau} \in (0, 1]$, ν_1 , $\nu_2 \in (0, 1)$, and $\delta \in (0, 1)$, choose $u^0 \in \Re^m$. Set j = 0 and iterate the following steps.

1. Choose $V^j \in \partial \mathsf{P}_{\tau} h(\tilde{b} + \tau^{-1} u^j)$ and $U^j \in \partial \mathsf{P}_{\sigma} p(\tilde{\beta} + \sigma^{-1}(\tilde{v} - X^* u^j))$. Let $H^j = \tau^{-1} V^j + \sigma^{-1} X U^j X^*$ and find the exact solution Δu^j or apply the PCG method to find an approximate solution Δu^j to

$$(H^j + \varepsilon_j I)\Delta u = -\nabla \phi(u^j; \sigma, \tau)$$

such that

$$\|H^{j}\Delta u^{j} + \nabla \phi(u^{j};\sigma,\tau)\| \le \eta_{j} := \min(\overline{\eta}, \|\nabla \phi(u^{j};\sigma,\tau)\|^{1+\overline{\tau}})$$

where $\varepsilon_j := \nu_1 \min \left\{ \nu_2, \| \nabla \phi(u^j; \sigma, \tau) \| \right\}$

2. Set $\alpha_j = \delta^{l_j}$, where l_j is the first nonnegative integer l for which

$$\phi(u^j + \delta^l \Delta u^j; \sigma, \tau) \leq \phi(u^j; \sigma, \tau) + \mu \delta^l \langle \nabla \phi(u^j; \sigma, \tau), (\Delta u^j) \rangle$$

3. Set
$$u^{j+1} = u^j + \alpha_j \Delta u^j$$

Theorem

Assume that $P_{\tau}h(\cdot)$ and $P_{\sigma}p(\cdot)$ are strongly semismooth. If problem (4) is nondegenerate, in particular if $f(\cdot) \equiv h(X \cdot)$ is continuously differentiable near $\bar{\beta}$, then $\{u^j\}$ converges to the unique optimal solution \bar{u} and

$$||u^{j+1} - \bar{u}|| = O(||u^j - \bar{u}||^{1+\bar{\tau}}).$$

Note that if $\bar{\tau} = 1$, we get the quadratic convergence.

mPPDNA

Algorithm. Let $\sigma^0, \sigma^1>0, \tau^0, \tau^1>0$ be given parameters 1. Compute

$$\beta^1 \approx \operatorname*{argmin}_{\beta \in \Re^n} \left\{ g(\beta; \sigma^0, \tau^0, 0, 0, b) \right\}$$

via solving its dual problem such that a prescribed stopping criterion is satisfied. Let k=1 and go to Step 2.1.

2.1 Choose
$$v^k \in \partial q(\beta^k)$$
 and compute

$$\beta^{k+1} = \operatorname*{argmin}_{\beta \in \Re^n} \left\{ g(\beta; \sigma^k, \tau^k, \beta^k, v^k, X\beta^k) + \langle \delta^k, \beta - \beta^k \rangle \right\}$$

via solving its dual problem such that the vector δ^k satisfies a prescribed accuracy condition.

2.2. If β^{k+1} satisfies a prescribed stopping condition, terminate; otherwise update $\sigma^{k+1} = \rho_k \sigma^k$, $\tau^{k+1} = \rho_k \tau^k$ with $\rho_k \in (0,1)$ and return to Step 2.1 with k = k + 1.

PPDNA: for the convex case of $q(\cdot) \equiv 0$

Algorithm. Let $\sigma^0, \sigma^1 > 0, \tau^0, \tau^1 > 0$ be given parameters. $\beta^0 \in \text{dom}(p)$.

1. Compute

$$\beta^1 \approx \operatorname*{argmin}_{\beta \in \Re^n} \left\{ f(\beta) + p(\beta) + \frac{\sigma^0}{2} \|\beta - \beta^0\|^2 + \frac{\tau^0}{2} \|X\beta - b\|^2 \right\}$$

via solving its dual problem such that a prescribed stopping criterion is satisfied. Let k = 1 and go to Step 2.1.

2.1 Compute

$$\beta^{k+1} \approx \underset{\beta \in \Re^n}{\operatorname{argmin}} \left\{ f(\beta) + p(\beta) + \frac{1}{2} \left\| \beta - \beta^k \right\|_{\sigma^k I + \tau^k X^* X}^2 \right\}$$

via solving its dual problem satisfying a prescribed accuracy condition. 2.2. If β^{k+1} satisfies a prescribed stopping condition, terminate; otherwise update $\sigma^{k+1} = \rho_k \sigma^k$, $\tau^{k+1} = \rho_k \tau^k$ with $\rho_k \in (0, 1)$ and return to Step 2.1 with k = k + 1. For simplicity, assume that we take for some constant c > 0 that

$$\tau_k \equiv c\sigma_k \quad \forall k.$$

Then the k-th subproblem of PPDNA can be written as

$$\beta^{k+1} \approx \operatorname*{argmin}_{\beta \in \Re^n} \left\{ g_k(\beta) := f(\beta) + p(\beta) + \frac{\sigma_k}{2} \big\| \beta - \beta^k \big\|_M^2 \right\},$$

where

$$M:=I+cX^*X\succ 0.$$

The stopping criterion for inner subproblems

(A)
$$g_k(\beta_{k+1}) - \inf g_k \le \sigma_k \varepsilon_k^2/2, \quad \sum \varepsilon_k < \infty.$$

Theorem (Global convergence)

Suppose that the solution set to (**P**) is nonempty. Then, $\{\beta^k\}$ is bounded and converges to an optimal solution β^* of (**P**).

Assumption (Error bound)

For a maximal monotone operator $\mathcal{T}(\cdot)$ with $\mathcal{T}^{-1}(0) \neq \emptyset$, $\exists \varepsilon > 0$ and a > 0 s.t.

 $\forall \eta \in \mathcal{B}(0,\varepsilon) \quad \text{and} \quad \forall \xi \in \mathcal{T}^{-1}(\eta), \quad \mathsf{dist}_M(\xi,\mathcal{T}^{-1}(0)) \leq a \|\eta\|_M,$

where $\mathcal{B}(0,\varepsilon) = \{y \in \mathcal{Y} \mid ||y|| \le \varepsilon\}$. The constant a is called the error bound modulus associated with \mathcal{T} .

- **(**) In many cases, \mathcal{T} is a polyhedral multifunction [Robinson, 1981].
- ② *T_g* (∂g) of LASSO, fused LASSO and elastic net regularized LS problems (piecewise linear-quadratic programming problems [J. Sun, PhD thesis, 1986] +1 ⇒ error bound).
- T_g of l₁ or elastic net regularized logistic regression [Luo and Tseng, 1992; Tseng and Yun, 2009].

PPDNA: Fast linear local convergence

Stopping criterion for the local convergence analysis

(B)
$$g_k(\beta^{k+1}) - \inf g_k$$

 $\leq \min\{1, (\sigma_k \delta_k^2/2)\} \|\beta^{k+1} - \beta^k\|_M^2, \quad \sum \delta_k < \infty$

Theorem

Assume that the solution set Ω to (**P**) is nonempty. Assume that error bound condition holds for \mathcal{T}_g with modulus l_g . Then, $\{\beta^k\}$ is convergent and, for all k sufficiently large,

$$\mathsf{dist}_M(eta^{k+1},\Omega) \ \le \ heta_k \mathsf{dist}_M(eta^k,\Omega),$$

where $\theta_k \approx \left(l_g(l_g^2 + \sigma_k^{-2})^{-1/2} + 2\delta_k\right) \rightarrow \theta_\infty = (l_g\sigma_\infty)/\sqrt{1 + (l_g\sigma_\infty)^2} < 1$ as $k \rightarrow \infty$.

Note that $\theta_{\infty} << 1$ when $l_g \sigma_{\infty}$ is close to zero. Thus, PPDNA can be treated as an approximate Newton's method!!! (arbitrary linear convergence rate, a name coined by M.J.D. Powell in 1969).

So far we have

- Outer iterations (PPA): asymptotically superlinear (arbitrary rate of linear convergence)
- Inner iterations (nonsmooth Newton): superlinear + cheap

Essentially, we have a " $fast^{2}$ " algorithm.

The mPPDNA for the nonconvex case: KL property

- Let $\eta > 0$ and Φ_{η} be a set of all concave functions $\phi : [0, \eta) \to \Re_+$ such that $\phi(0) = 0$, ϕ is continuous at 0 and continuously differentiable on $(0, \eta)$ and $\phi'(x) > 0$, for $\forall x \in (0, \eta)$.
- The function f is said to have the Kurdyka-Łojasiewicz (KL) property at \bar{x} if there exists $\eta > 0$, a neighbourhood \mathcal{U} of \bar{x} and a concave function $\phi \in \Phi_{\eta}$ such that

 $\phi'(f(x) - f(\bar{x})) \mathsf{dist}(0, \partial f) \geq 1, \ \forall x \in \mathcal{U} \ \text{and} \ f(\bar{x}) < f(x) < f(\bar{x}) + \eta,$

where dist $(x, C) := \min_{y \in C} ||y - x||$ is the distance from a point x to a nonempty closed set C.

A function is said to have the KL property at \bar{x} with an exponent α if the function ϕ in the definition of the KL property takes the form as $\phi(x) = \gamma x^{1-\alpha}$ with $\gamma > 0$ and $\alpha \in [0, 1)$.

Theorem

Suppose that the function $g(\cdot)$ is bounded below and that q is continuously differentiable near \mathcal{B}^{∞} , the set of all cluster points of the sequence $\{\beta^k\}$ generated by mPPDNA. Then every cluster point in \mathcal{B}^{∞} , if exists, is a d-stationary point of (3).

Theorem

Suppose that the function $g(\cdot)$ is bounded below and that q is continuously differentiable near \mathcal{B}^{∞} , the set of all cluster points of the sequence $\{\beta^k\}$ generated by mPPDNA. If either one of the following two conditions holds,

- (a) \mathcal{B}^{∞} contains an isolated element;
- (b) The sequence $\{\beta^k\}$ is bounded; for all $\beta \in \mathcal{B}^{\infty}$, $\nabla q(\cdot)$ is locally Lipschitz continuous near β ; and the function g has the KL property at all $\beta \in \mathcal{B}^{\infty}$;

then the whole sequence $\{\beta^k\}$ converges to a unique element of \mathcal{B}^{∞} .

Theorem

Moreover, if the condition (b) is satisfied and $\{\beta^k\}$ converges to $\beta^{\infty} \in \mathcal{B}^{\infty}$, the function g has the KL property at β^{∞} with an exponent $\alpha \in [0, 1)$, then we have

- (i) if $\alpha = 0$, then the sequence $\{\beta^k\}$ converges in a finite number of steps;
- (ii) if $\alpha \in (0, \frac{1}{2}]$, then the sequence $\{\beta^k\}$ converges R-linearly, that is, for all $k \ge 1$ there exist $\nu > 0$ and $\eta \in [0, 1)$ such that $\|\beta^k \beta^\infty\| \le \nu \eta^k$;
- (iii) if $\alpha \in (\frac{1}{2}, 1)$, then the sequence $\{\beta^k\}$ converges R-sublinearly, that is, for all $k \ge 1$ there exists $\nu > 0$ such that $\|\beta^k \beta^{\infty}\| \le \nu k^{-\frac{1-\alpha}{2\alpha-1}}$.

Numerical experiments

- Our numerical experiments are implemented on a PC (Intel Core 2 Duo 2.6 GHz with 4 GB RAM).
- The parameter λ is defined by $\lambda = \lambda_c \Lambda$, $\Lambda = 1.1 \Phi^{-1}(1 0.05/(2n))$ with Φ the cumulative normal distribution function.
- The number of nonzero elements of a vector is defined by the minimal \boldsymbol{k} such that

$$\sum_{i=1}^k |\tilde{\beta}_i| \ge 0.9999 \|\beta\|_1$$

where $\tilde{\beta}$ is obtained by sorting β such that $|\tilde{\beta}_1| \ge |\tilde{\beta}_2| \ge \ldots \ge |\tilde{\beta}_n|$

The step 1 of the mPPNDA algorithm will be terminated if the relative $\mathsf{KKT}\xspace$ residual 1 satisfies

$$\eta_{kkt} := \frac{\left\| \beta - \mathsf{P}_{\lambda} p\left(\beta - \frac{X^*(X\beta - b)}{\|X\beta - b\|}\right) \right\|}{1 + \|\beta\| + \frac{\|X^*(X\beta - b)\|}{\|X\beta - b\|}} < 10^{-6},$$
(6)

or the number of iterations reaches the maximum 200 while the ADMMs will be terminated if (6) is satisfied or the number of iterations reaches the maximum 5000.

¹Whenever possible, try to avoid using the "fast convergence criteria" such as the relative distance of two consecutive iterates. Instead, try to design fast convergent algorithms independent of the "fast convergence criteria", which may only indicate that the employed algorithm is slow for an earlier termination.

Table: The performances of the Flare package and pADMM on synthetic datasets for the srLasso problem.

probname	λ_c	pobj	time
m; n		Flare pADMM	Flare pADMM
exmp1	1.0	3.8876+3 3.5799+3	11:26 12
8000;800	0.5	3.0501+3 1.9174+3	21:09 13
	0.1	1.0487+3 5.8738+2	28:42 16
exmp2	1.0	2.2422+3 2.2419+3	14:09 19
8000;800	0.5	1.8050+3 1.2811+3	27:18 11
	0.1	5.6150+2 4.6013+2	27:37 09
exmp3	1.0	2.4758+3 2.4569+3	10:05 07
8000;400	0.5	1.9819+3 1.9421+3	7:26 07
	0.1	1.4888+3 1.4438+3	7:14 05
exmp4	1.0	1.1210+4 1.1205+4	29:11 20:16
8000;4000	0.5	$1.0165 + 4 \mid 1.0165 + 4$	1:43:48 21:48
	0.1	7.6846+3 3.4069+3	3:11:27 5:12

Table: The performances of the Flare package and pADMM on UCI datasets for the srLasso problem.

probname	λ_c	pobj	time	
m; n		Flare pADMM	Flare pADMM	
abalone.scale.expanded7	1.0	- 2.3852+2	- 25:57	
4177;6435	0.5	- 2.0312+2	- 25:32	
	0.1	- 1.5586+2	- 26:29	
mpg.scale.expanded7	1.0	2.3550+2 2.3544+2	1:00 04	
392;3432	0.5	$1.5856 + 2 \mid 1.5831 + 2$	57 03	
	0.1	7.8656+1 7.8616+1	1:06 03	
space.ga.scale.expanded9	1.0	$1.3113 + 1 \mid 1.3113 + 1$	12:59 5:19	
3107;5005	0.5	2.2419+1 2.1607+1	9:01 2:00	
	0.1	$1.2950 + 1 \mid 1.1999 + 1$	6:13 3:00	

The "-" in the Table means that the Flare package fails to solve the problem due to being out of memory.

Table: The performances of ADMMs and PPDNA on synthetic datasets for the srLasso problem. In the table, "a"=PPDNA, "b"=pADMM, "c"=dADMM.

probname m; n	λ_c	nnz	η_{kkt} $a \mid b \mid c$	η_G $a \mid b \mid c$	pobj a b c	time a b c	testerror
exmp1 8000;800	0.127	499	9.4-7 9.9-7 9.9-7	1.9-8 3.1-7 1.7-8	6.6996+2 6.6996+2 6.6996+2	30 3:49 2:44	9.4085+0
exmp2 8000;800	0.081	627	9.8-7 9.9-7 9.9-7	2.6-9 1.8-7 7.2-9	4.1824+2 4.1824+2 4.1824+2	32 1:49 2:26	9.4807+0
exmp3 8000;400	0.124	298	7.4-7 9.1-7 9.9-7	4.8-9 1.1-7 3.7-9	1.4476+3 1.4476+3 1.4476+3	08 17 1:05	2.3420+2
exmp4 8000;4000	0.117	2845	7.7-7 9.9-7 9.5-7	8.6-10 5.2-7 5.3-7	3.6799+3 3.6799+3 3.6799+3	4:41 5:41 16:44	3.6759+2

Table: The performances of ADMMs and PPDNA on synthetic datasets for the srLasso problem. In the table, "a"=PPDNA, "b"=pADMM, "c"=dADMM.

probname m; n	λ_c	nnz	η_{kkt} $a \mid b \mid c$	η_G $a \mid b \mid c$	pobj a b c	time a b c	testerror
exmp1 8000;800	0.127	499	9.4-7 9.9-7 9.9-7	1.9-8 3.1-7 1.7-8	6.6996+2 6.6996+2 6.6996+2	30 3:49 2:44	9.4085+0
exmp2 8000;800	0.081	627	9.8-7 9.9-7 9.9-7	2.6-9 1.8-7 7.2-9	4.1824+2 4.1824+2 4.1824+2	32 1:49 2:26	9.4807+0
exmp3 8000;400	0.124	298	7.4-7 9.1-7 9.9-7	4.8-9 1.1-7 3.7-9	1.4476+3 1.4476+3 1.4476+3	08 17 1:05	2.3420+2
exmp4 8000;4000	0.117	2845	7.7-7 9.9-7 9.5-7	8.6-10 5.2-7 5.3-7	3.6799+3 3.6799+3 3.6799+3	4:41 5:41 16:44	3.6759+2

Table: The performances of ADMMs and PPDNA on UCI datasets for the srLasso problem. In the table, "a"=PPDNA, "b"=pADMM, "c"=dADMM.

probname			$\eta_{\rm kkt}$	η_G	pobj	time	
m; n	A _c	11112	$a \mid b \mid c$	$a \mid b \mid c$	$a \mid b \mid c$	$a \mid b \mid c$	
E2006.test	0.019	1	57.7 32.7 97.7	25-7 38-8 81-9	2 1008+1 2 1008+1 2 1008+1	05 07 05	
3308;150358	0.010	-	0.1 1 0.2 1 0.1 1	2.0 1 0.0 0 0.2 0	2.1330 1 2.1330 1 2.1330 1	05 01 05	
log1p.E2006.test	0 260	201	7 9-7 1 2-4 1 2-3	28-6 30-3 39-5	2 1642+1 2 1713+1 2 1642+1	1.49 2.17.37 1.22.12	
3308;1771946	0.200		1.0 1 1.2 1 1.2 0	2.0 0 0.0 0 0.0 0			
pyrim.scale.expanded5	0 109	70	5 9 7 2 0 5 3 7 3	69.7 47.3 38.4	6 8301-1 6 9094-1 6 8308-1	18 20:31 12:10	
74;201376	0.105		0.01 2.00 0.10	0.01 1.10 0.01	0.00011 0.00011 0.00001	10 10:01 11:10	
abalone.scale.expanded7	0 004	82	96-7 99-7 88-7	1 0-9 3 0-7 6 4-9	1 3495+2 1 3495+2 1 3495+2	07 1.32 7.57	
4177;6435		02	5.01 5.51 6.61	2.0 5 0.0 1 0.1 5	1010012 1010012 11010012	01 2.02 1.01	
bodyfat.scale.expanded7	0.012	51	74-7 11-6 99-7	2 2-8 6 3-5 3 5-9	8 5770-2 8 5834-2 8 5767-2	10 27.59 8.38	
252;116280							

In our mPPDNA, the step 1 is used to generate an initial point for the step 2 and is stopped if $\eta_{kkt} < 10^{-4}$. The algorithms will be terminated if the relative KKT residual satisfies

$$\tilde{\eta}_{kkt} := \frac{\left\| \beta - \mathsf{P}_1(p-q) \left(\beta - \frac{X^*(X\beta - b)}{\|X\beta - b\|} \right) \right\|}{1 + \|\beta\| + \frac{\|X^*(X\beta - b)\|}{\|X\beta - b\|}} < 10^{-6}$$

Besides, the algorithms will also be stopped when they reach the pre-set maximum number of iterations (200 for the second step of mPPDNA and 5000 for ADMM).

For $\lambda > 0$, the SCAD regularization is defined by $r(\beta) = p(\beta) - q(\beta)$ with

$$\begin{split} p(\beta) &= \lambda \|\beta\|_1 \\ q(\beta) &= \sum_{i=1}^n \left\{ \begin{array}{ll} 0, & \text{if} \quad |\beta_i| < \lambda \\ \frac{(|\beta_i| - \lambda)^2}{2(a_s - 1)}, & \text{if} \quad \lambda \leq |\beta_i| \leq a_s \lambda \\ \lambda |\beta_i| - \frac{a_s + 1}{2} \lambda^2, & \text{if} \quad |x_i| > a_s \lambda \end{array} \right. \end{split}$$

In our numerical experiments, we take $a_s = 3.7$.

Numerical results for the SCAD regularization

Table: The performances of ADMM and mPPDNA on synthetic datasets for the SCAD regularization. In the table, "a"=mPPDNA, "b"=ADMM.

probname m; n	λ_c	nnz	η_{kkt} $a \mid b$	pobj a b	time a b	testerror
exmp1 8000;800	0.145	460	3.9-7 5.9-1	5.9368+2 5.9392+2	20 3:39	9.2406+0
exmp2 8000;800	0.087	616	5.9-7 1.0-1	4.0760+2 4.0777+2	28 3:33	9.3745+0
exmp3 8000;400	0.230	293	7.8-7 2.7-1	1.5486+3 1.5529+3	10 2:02	2.3629+2
exmp4 8000;4000	0.153	2554	5.4-7 6.8-1	3.1837+3 3.1940+3	4:21 16:41	3.4480+2

Numerical results for the SCAD regularization

Table: The performances of ADMM and mPPDNA on UCI datasets for the SCAD regularization. In the table, "a"=mPPDNA, "b"=ADMM.

probname)	nnz	η_{kkt}	pobj	time	
m; n	Λ_c	11112	$a \mid b$	$a \mid b$	$a \mid b$	
E2006.test	0.071	1	2 2-8 9 0-7	$22165\pm1\pm22165\pm1$	08 12.51	
3308;150358	0.071	-	2.2 0 5.0 1	2.2103 1 2.2103 1	00 12.51	
log1p.E2006.test	0.257	207	2 1-7 5 9-3	2 1613+1 2 1366+2	3.50 2.36.14	
3308;1771946	0.201	207	2.1 1 5.5 5	2.1013 1 2.1300 2	0.00 2.00.11	
pyrim.scale.expanded5	0 109	70	1 4-7 4 3-3	6 8301-1 7 2608-1	13 21.26	
74;201376	0.105		1.17 1.5 5	0.0301 1 1.2000 1	10 21.20	
abalone.scale.expanded7	0.011	49	99-7 69-1	1 3292+2 1 3864+2	12 21.41	
4177;6435	0.011	15	5.57 0.51	1.5252 2 1.5001 2	12 21.11	
bodyfat.scale.expanded7	0 201	2	3 9-8 7 6-2	9 4125-1 9 5136-1	06 25.51	
252;116280	0.201	-	0.5 0 1.0 2	5.1120 1 5.0100 1	00 20.01	

Nonconvex regularizers: MCP

For two positive parameters $a_m>2$ and $\lambda,$ the MCP regularization can be defined as $r(\beta)=p(\beta)-q(\beta)$ with

$$\begin{split} p(\beta) &= \lambda \|\beta\|_1 \\ q(\beta) &= \sum_{i=1}^n \begin{cases} \frac{\beta_i^2}{a_m}, & \text{if } |\beta_i| \le a_m \lambda, \\ 2\lambda |\beta_i| - a_m \lambda^2, & \text{if } |\beta_i| > a_m \lambda \end{cases} \end{split}$$

The function $q(\cdot)$ is continuously differentiable with its derivative given by

$$\frac{\partial q(\beta)}{\partial \beta_i} = \begin{cases} \frac{2\beta_i}{a_m}, & \text{if } |\beta_i| \le a_m \lambda, \\ 2\lambda \text{sign}(\beta_i), & \text{if } |\beta_i| > a_m \lambda \end{cases}$$

In our numerical experiments, we take $a_m = 3.7$.

Numerical results for the MCP regularization

Table: The performances of ADMM and mPPDNA on synthetic datasets for the MCP regularization. In the table, "a" = mPPDNA, "b" = ADMM.

probname m; n	λ_c	nnz	$\frac{\eta_{kkt}}{a \mid b}$	pobj a b	time a b	testerror
exmp1 8000;800	0.209	380	5.2-8 1.9-2	5.5695+2 5.6091+2	29 3:33	9.3483+0
exmp2 8000;800	0.151	535	2.7-7 1.3-1	4.5225+2 4.5414+2	38 3:30	9.4916+0
exmp3 8000;400	0.081	267	9.3-7 1.5-1	1.3590+3 1.3617+3	1:11 2:01	2.3613+2
exmp4 8000;4000	0.293	1821	9.4-7 6.9-2	4.1362+3 4.2741+3	5:33 16:37	3.8471+2

Numerical results for the MCP regularization

Table: The performances of ADMM and mPPDNA on UCI datasets for the MCP regularization. In the table, "a" = mPPDNA, "b" = ADMM.

probname	1	nnz	η_{kkt}	pobj	time	
m; n	Λ_c	11112	$a \mid b$	$a \mid b$	$a \mid b$	
E2006.test	0.090	1	2.4-8 9.3-7	2.2077+1 2.2077+1	07 07	
3308;150358						
log1p.E2006.test	0.261	187	82-7 22-3	2 1455+1 3 6500+1	4.09 2.20.46	
3308;1771946	0.201	101	0.2 1 1 2.2 0	2.1.100 + 1 0.0000 + 1	1.000 2.201.10	
pyrim.scale.expanded5	0.221	43	99-7 70-3	1 1428+0 4 6112+0	18 19.36	
74;201376	0.221		5.5 1 1.0 0	1.1.1.20 + 0 + 1.0112 + 0	10 10.00	
abalone.scale.expanded7	0.012	55	7 1-7 1 6-5	1 3271+2 1 2693+2	09 21.32	
4177;6435	0.012		1.1 1 1 1.0 0	1.021112 1.2000 2	00 21.02	
bodyfat.scale.expanded7	0 183	2	28-7 53-6	5 7347-1 5 8278-1	06 25.11	
252;116280	0.100	-	2.0 1 0.0 0	0.1011 1 0.0210 1	00 20.11	

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