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

## Defeng Sun

Department of Applied Mathematics
The Hong Kong
POLYTECHNIC UNIVERSITY
香港理工大學

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Based on joint works with Kim－Chuan Toh（NUS），Xudong Li（Fudan） Peipei Tang（ZUCC），Chengjing Wang（SWJTU），Ning Zhang（PolyU）

# Second Order Methods 

with

## First Order Costs

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)
(1) One core mathematical problem is to solve the following linear equation

$$
B x=b,
$$

where $B \in \Re^{n \times n}$ and $b \in \Re^{n}$
(2) 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 $O\left(n^{3}\right)$ flops - way too high for a big $n$.
(3) If $B$ is "sparse", e.g., $B=I+u v^{T}$, where $u, v \in \Re^{n}$, one can reduce the cost from $O\left(n^{3}\right)$ to $O(n)$ via the Sherman-Morrison-Woodbury formula

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

( ( Solving structured linear equations can be cheap $\Longrightarrow$ Second order methods with first order computational costs possible!!!

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

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

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!!!

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

$$
\mathcal{U}:=\left\{\left(\Delta_{1}, \ldots, \Delta_{n}\right) \mid\left\|\Delta_{i}\right\| \leq \lambda, i=1, \ldots, n\right\}
$$

Another commonly used example is the constrained Lasso model

$$
\min _{\beta \in \Re^{n}}\left\{\|\beta\|_{1} \mid\|X \beta-b\| \leq \tau\right\}
$$

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

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

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

There are many more convex Lasso-type models:

$$
\min \left\{\left.\frac{1}{2}\|X \beta-b\|^{2}+\lambda\|\beta\|_{1} \right\rvert\, \beta \in \Re^{n}\right\}
$$

where $\lambda>0$.
(Fused LASSO)

$$
\min \left\{\frac{1}{2}\|X \beta-b\|^{2}+\lambda\|\beta\|_{1}+\lambda_{2}\|B \beta\|_{1}\right\}
$$

$$
B=\left(\begin{array}{ccccc}
1 & -1 & & & \\
& 1 & -1 & & \\
& & \ddots & \ddots & \\
& & & 1 & -1
\end{array}\right)
$$

(Clustered LASSO)

$$
\min \left\{\frac{1}{2}\|X \beta-b\|^{2}+\lambda\|\beta\|_{1}+\lambda_{2} \sum_{i=1}^{n} \sum_{j=i+1}^{n}\left|\beta_{i}-\beta_{j}\right|\right\}
$$

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

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

$$
\sum_{1 \leq i<j \leq n}\left|\beta_{i}-\beta_{j}\right|=\left\langle w, \beta^{\downarrow}\right\rangle,
$$

where $\beta^{\downarrow}$ is the vector whose components are those of $\beta$ 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

$$
w_{k}=n-2 k+1, k=1, \cdots, n
$$

(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.

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

$$
\begin{equation*}
f(\beta)=\sum_{i=1}^{m} \log \left(1+\exp \left(-b_{i}(X \beta)_{i}\right)\right) \tag{1}
\end{equation*}
$$

- Define $h: \Re^{m} \rightarrow \Re$ as follows

$$
h(z)=\sum_{i=1}^{m} \log \left(1+\exp \left(-b_{i} z_{i}\right)\right) \quad \forall z \in \Re^{m}
$$

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

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

In general, the loss function $f: \Re^{p K} \rightarrow \Re$ can take as the multinomial logistic regression function: given $A:=\left(A_{1}, \ldots, A_{N}\right)^{T} \in \Re^{N \times p}$,

$$
\begin{equation*}
f(\beta)=-\sum_{i=1}^{N}\left(\sum_{k=1}^{K} y_{i k} A_{i}^{T} \beta_{k}-\log \sum_{k=1}^{K} \exp \left(A_{i}^{T} \beta_{k}\right)\right) \tag{2}
\end{equation*}
$$

- Define $h: \Re^{N K} \rightarrow \Re$ as follows:

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

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

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

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 (P):

$$
\begin{equation*}
\min _{\beta \in \Re^{n}}\{g(\beta):=\underbrace{h(X \beta)}_{f(\beta)}+\underbrace{p(\beta)-q(\beta)}_{r(\beta)}\} \tag{3}
\end{equation*}
$$

Here $p: \Re^{n} \rightarrow(-\infty,+\infty]$ is a proper closed convex function and $q: \Re^{n} \rightarrow \Re$ 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

$$
\begin{aligned}
& p(\beta)=\lambda\|\beta\|_{1} \\
& q(\beta)=\sum_{i=1}^{n}\left\{\begin{array}{lll}
0, & \text { if } & \left|\beta_{i}\right|<\lambda \\
\frac{\left(\left|\beta_{i}\right|-\lambda\right)^{2}}{2\left(a_{s}-1\right)}, & \text { if } & \lambda \leq\left|\beta_{i}\right| \leq a_{s} \lambda \\
\lambda\left|\beta_{i}\right|-\frac{a_{s}+1}{2} \lambda^{2}, & \text { if } & \left|\beta_{i}\right|>a_{s} \lambda
\end{array}\right.
\end{aligned}
$$

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

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

$$
\begin{aligned}
& p(\beta)=\lambda\|\beta\|_{1} \\
& q(\beta)=\sum_{i=1}^{n} \begin{cases}\frac{\beta_{i}^{2}}{a_{m}}, & \text { if }\left|\beta_{i}\right| \leq a_{m} \lambda, \\
2 \lambda\left|\beta_{i}\right|-a_{m} \lambda^{2}, & \text { if }\left|\beta_{i}\right|>a_{m} \lambda\end{cases}
\end{aligned}
$$

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

$$
\frac{\partial q(\beta)}{\partial \beta_{i}}= \begin{cases}\frac{2 \beta_{i}}{a_{m}}, & \text { if }\left|\beta_{i}\right| \leq a_{m} \lambda, \\ 2 \lambda \operatorname{sign}\left(\beta_{i}\right), & \text { if }\left|\beta_{i}\right|>a_{m} \lambda\end{cases}
$$

In our numerical experiments, we take $a_{m}=3.7$.

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 $\beta$.
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 $\beta$ 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]

$$
\min _{\beta \in \Re^{n}}\{h(X \beta)+p(\beta)-q(\beta)\}
$$

- 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

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

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

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

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

$$
\begin{equation*}
e_{t} f(x):=\min _{z \in \Re^{n}}\left\{f(z)+\frac{t}{2}\|z-x\|^{2}\right\}, \quad \forall x \in \Re^{n} \tag{5}
\end{equation*}
$$

Here $e_{t} f(\cdot)$ is continuously differentiable with

$$
\nabla e_{t} f(x)=t\left[x-\mathrm{P}_{t} f(x)\right], \quad \forall x \in \Re^{n},
$$

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

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

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

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

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

## 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.

Algorithm SSN (SSN $(\sigma, \tau)$ ): Given $\mu \in\left(0, \frac{1}{2}\right), \bar{\eta} \in(0,1), \bar{\tau} \in(0,1], \nu_{1}$, $\nu_{2} \in(0,1)$, and $\delta \in(0,1)$, choose $u^{0} \in \Re^{m}$. Set $j=0$ and iterate the following steps.

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

$$
\left(H^{j}+\varepsilon_{j} I\right) \Delta u=-\nabla \phi\left(u^{j} ; \sigma, \tau\right)
$$

such that

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

where $\varepsilon_{j}:=\nu_{1} \min \left\{\nu_{2},\left\|\nabla \phi\left(u^{j} ; \sigma, \tau\right)\right\|\right\}$
2. Set $\alpha_{j}=\delta^{l_{j}}$, where $l_{j}$ is the first nonnegative integer $l$ for which

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

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 $\left\{u^{j}\right\}$ converges to the unique optimal solution $\bar{u}$ and

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

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

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

1. Compute

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

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 $v^{k} \in \partial q\left(\beta^{k}\right)$ and compute

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

via solving its dual problem such that the vector $\delta^{k}$ satisfies a prescribed accuracy condition.
2.2. If $\beta^{k+1}$ 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 $k=k+1$.

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

1. Compute

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

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 $\beta^{k+1}$ 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

$$
\beta^{k+1} \approx \underset{\beta \in \Re^{n}}{\operatorname{argmin}}\left\{g_{k}(\beta):=f(\beta)+p(\beta)+\frac{\sigma_{k}}{2}\left\|\beta-\beta^{k}\right\|_{M}^{2}\right\},
$$

where

$$
M:=I+c X^{*} X \succ 0 .
$$

The stopping criterion for inner subproblems

$$
\text { (A) } \quad g_{k}\left(\beta_{k+1}\right)-\inf g_{k} \leq \sigma_{k} \varepsilon_{k}^{2} / 2, \quad \sum \varepsilon_{k}<\infty .
$$

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

## 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.

$$
\forall \eta \in \mathcal{B}(0, \varepsilon) \quad \text { and } \quad \forall \xi \in \mathcal{T}^{-1}(\eta), \quad \operatorname{dist}_{M}\left(\xi, \mathcal{T}^{-1}(0)\right) \leq a\|\eta\|_{M},
$$

where $\mathcal{B}(0, \varepsilon)=\{y \in \mathcal{Y} \mid\|y\| \leq \varepsilon\}$. The constant $a$ is called the error bound modulus associated with $\mathcal{T}$.
(1) In many cases, $\mathcal{T}$ is a polyhedral multifunction [Robinson, 1981].
(2) $\mathcal{T}_{g}(\partial g)$ of LASSO, fused LASSO and elastic net regularized LS problems (piecewise linear-quadratic programming problems [J. Sun, PhD thesis, 1986] $+1 \Rightarrow$ error bound).
(3) $\mathcal{T}_{g}$ of $\ell_{1}$ or elastic net regularized logistic regression [Luo and Tseng, 1992; Tseng and Yun, 2009].

Stopping criterion for the local convergence analysis

$$
\text { (B) } \begin{aligned}
& g_{k}\left(\beta^{k+1}\right)-\inf g_{k} \\
\leq & \min \left\{1,\left(\sigma_{k} \delta_{k}^{2} / 2\right)\right\}\left\|\beta^{k+1}-\beta^{k}\right\|_{M}^{2}, \quad \sum \delta_{k}<\infty .
\end{aligned}
$$

## Theorem

Assume that the solution set $\Omega$ to $(\mathbf{P})$ is nonempty. Assume that error bound condition holds for $\mathcal{T}_{g}$ with modulus $l_{g}$. Then, $\left\{\beta^{k}\right\}$ is convergent and, for all $k$ sufficiently large,

$$
\operatorname{dist}_{M}\left(\beta^{k+1}, \Omega\right) \leq \theta_{k} \operatorname{dist}_{M}\left(\beta^{k}, \Omega\right)
$$

where $\theta_{k} \approx\left(l_{g}\left(l_{g}^{2}+\sigma_{k}^{-2}\right)^{-1 / 2}+2 \delta_{k}\right) \rightarrow \theta_{\infty}=\left(l_{g} \sigma_{\infty}\right) / \sqrt{1+\left(l_{g} \sigma_{\infty}\right)^{2}}<1$ as $k \rightarrow \infty$.

Note that $\theta_{\infty} \ll 1$ 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
(1) Outer iterations (PPA): asymptotically superlinear (arbitrary rate of linear convergence)
(2) Inner iterations (nonsmooth Newton): superlinear + cheap

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

- Let $\eta>0$ and $\Phi_{\eta}$ be a set of all concave functions $\phi:[0, \eta) \rightarrow \Re_{+}$ such that $\phi(0)=0, \phi$ is continuous at 0 and continuously differentiable on $(0, \eta)$ and $\phi^{\prime}(x)>0$, 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
$\phi^{\prime}(f(x)-f(\bar{x})) \operatorname{dist}(0, \partial f) \geq 1, \forall x \in \mathcal{U}$ and $f(\bar{x})<f(x)<f(\bar{x})+\eta$,
where $\operatorname{dist}(x, C):=\min _{y \in C}\|y-x\|$ 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 $\alpha$ if the function $\phi$ 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 $\mathcal{B}^{\infty}$, the set of all cluster points of the sequence $\left\{\beta^{k}\right\}$ generated by mPPDNA. Then every cluster point in $\mathcal{B}^{\infty}$, 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 $\mathcal{B}^{\infty}$, the set of all cluster points of the sequence $\left\{\beta^{k}\right\}$ generated by mPPDNA. If either one of the following two conditions holds,
(a) $\mathcal{B}^{\infty}$ contains an isolated element;
(b) The sequence $\left\{\beta^{k}\right\}$ is bounded; for all $\beta \in \mathcal{B}^{\infty}, \nabla q(\cdot)$ is locally Lipschitz continuous near $\beta$; and the function $g$ has the KL property at all $\beta \in \mathcal{B}^{\infty}$;
then the whole sequence $\left\{\beta^{k}\right\}$ converges to a unique element of $\mathcal{B}^{\infty}$.

## Theorem

Moreover, if the condition (b) is satisfied and $\left\{\beta^{k}\right\}$ converges to $\beta^{\infty} \in \mathcal{B}^{\infty}$, the function $g$ has the $K L$ property at $\beta^{\infty}$ with an exponent $\alpha \in[0,1)$, then we have
(i) if $\alpha=0$, then the sequence $\left\{\beta^{k}\right\}$ converges in a finite number of steps;
(ii) if $\alpha \in\left(0, \frac{1}{2}\right]$, then the sequence $\left\{\beta^{k}\right\}$ converges $R$-linearly, that is, for all $k \geq 1$ there exist $\nu>0$ and $\eta \in[0,1)$ such that $\left\|\beta^{k}-\beta^{\infty}\right\| \leq \nu \eta^{k} ;$
(iii) if $\alpha \in\left(\frac{1}{2}, 1\right)$, then the sequence $\left\{\beta^{k}\right\}$ converges $R$-sublinearly, that is, for all $k \geq 1$ there exists $\nu>0$ such that $\left\|\beta^{k}-\beta^{\infty}\right\| \leq \nu k^{-\frac{1-\alpha}{2 \alpha-1}}$.

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

$$
\sum_{i=1}^{k}\left|\tilde{\beta}_{i}\right| \geq 0.9999\|\beta\|_{1}
$$

where $\tilde{\beta}$ is obtained by sorting $\beta$ such that $\left|\tilde{\beta}_{1}\right| \geq\left|\tilde{\beta}_{2}\right| \geq \ldots \geq\left|\tilde{\beta}_{n}\right|$

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

$$
\begin{equation*}
\eta_{k k t}:=\frac{\left\|\beta-\mathrm{P}_{\lambda} p\left(\beta-\frac{X^{*}(X \beta-b)}{\|X \beta-b\|}\right)\right\|}{1+\|\beta\|+\frac{\left\|X^{*}(X \beta-b)\right\|}{\|X \beta-b\|}}<10^{-6}, \tag{6}
\end{equation*}
$$

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.

[^0]Table: The performances of the Flare package and pADMM on synthetic datasets for the srLasso problem.

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

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

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

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 <br> m; n | $\lambda_{c}$ | nnz | $\eta_{\text {kkt }}$ | $\eta_{G}$ | $a\|b\| c$ | $a\|b\| c$ | pobj |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| exmp1 <br> $8000 ; 800$ | 0.127 |  | $9.4-7\|9.9-7\| 9.9-7$ | $1.9-8\|3.1-7\| 1.7-8$ | $6.6996+2\|6.6996+2\| 6.6996+2$ | $30\|3: 49\| 2: 44$ | $9.4085+0$ |
| exmp2 <br> $800 ; 800$ | 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 <br> $800 ; 400$ | 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 <br> $8000 ; 4000$ | 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 <br> m; n | $\lambda_{c}$ | nnz | $\eta_{\text {kkt }}$ | $\eta_{G}$ | $a\|b\| c$ | $a\|b\| c$ | pobj |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| exmp1 <br> $8000 ; 800$ | 0.127 |  | $9.4-7\|9.9-7\| 9.9-7$ | $1.9-8\|3.1-7\| 1.7-8$ | $6.6996+2\|6.6996+2\| 6.6996+2$ | $30\|3: 49\| 2: 44$ | $9.4085+0$ |
| exmp2 <br> $800 ; 800$ | 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 <br> $800 ; 400$ | 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 <br> $8000 ; 4000$ | 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 <br> $\mathrm{m} ; \mathrm{n}$ | $\lambda_{c}$ | nnz | $\eta_{\text {kkt }}$ | $\eta_{G}$ | pobj | time |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| E2006.test <br> $3308 ; 150358$ |  |  | $a\|b\| c$ | $a\|b\| c$ | $a\|b\| c$ | $a\|b\| c$ |
| log1p.E2006.test <br> $3308 ; 1771946$ | 0.260 | 201 | $7.9-7\|1.2-4\| 1.2-3$ | $2.8-6\|3.0-3\| 3.9-5$ | $2.1642+1\|2.1713+1\| 2.1642+1$ | $1: 49\|2: 17: 37\| 1: 22: 12$ |
| pyrim.scale.expanded5 <br> $74 ; 201376$ | 0.109 | 70 | $5.9-7\|2.0-5\| 3.7-3$ | $6.9-7\|4.7-3\| 3.8-4$ | $6.8301-1\|6.9094-1\| 6.8308-1$ | $18\|20: 31\| 12: 10$ |
| abalone.scale.expanded7 <br> 4177;6435 | 0.004 | 82 | $9.6-7\|9.9-7\| 8.8-7$ | $1.0-9\|3.0-7\| 6.4-9$ | $1.3495+2\|1.3495+2\| 1.3495+2$ | $07\|1: 32\| 7: 57$ |
| bodyfat.scale.expanded7 <br> $252 ; 116280$ | 0.012 | 51 | $7.4-7\|1.1-6\| 9.9-7$ | $2.2-8\|6.3-5\| 3.5-9$ | $8.5770-2\|8.5834-2\| 8.5767-2$ | $10\|27: 59\| 8: 38$ |

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

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

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

$$
\begin{aligned}
& p(\beta)=\lambda\|\beta\|_{1} \\
& q(\beta)=\sum_{i=1}^{n}\left\{\begin{array}{lll}
0, & \text { if }\left|\beta_{i}\right|<\lambda \\
\frac{\left(\left|\beta_{i}\right|-\lambda\right)^{2}}{2\left(a_{s}-1\right)}, & \text { if } & \lambda \leq\left|\beta_{i}\right| \leq a_{s} \lambda \\
\lambda\left|\beta_{i}\right|-\frac{a_{s}+1}{2} \lambda^{2}, & \text { if } & \left|x_{i}\right|>a_{s} \lambda
\end{array}\right.
\end{aligned}
$$

In our numerical experiments, we take $a_{s}=3.7$.

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

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

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

| probname <br> $\mathrm{m} ; \mathrm{n}$ | $\lambda_{c}$ | nnz | $\eta_{k k t}$ | pobj | time |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  | $a \mid b$ | $a \mid b$ |  |
| E2006.test <br> $3308 ; 150358$ | 0.071 | 1 | $2.2-8 \mid 9.0-7$ | $2.2165+1 \mid 2.2165+1$ | $08 \mid 12: 51$ |
| log1p.E2006.test <br> $3308 ; 1771946$ | 0.257 | 207 | $2.1-7 \mid 5.9-3$ | $2.1613+1 \mid 2.1366+2$ | $3: 50 \mid 2: 36: 14$ |
| pyrim.scale.expanded5 <br> $74 ; 201376$ | 0.109 | 70 | $1.4-7 \mid 4.3-3$ | $6.8301-1 \mid 7.2608-1$ | $13 \mid 21: 26$ |
| abalone.scale.expanded7 <br> $4177 ; 6435$ | 0.011 | 49 | $9.9-7 \mid 6.9-1$ | $1.3292+2 \mid 1.3864+2$ | $12 \mid 21: 41$ |
| bodyfat.scale.expanded7 <br> $252 ; 116280$ | 0.201 | 2 | $3.9-8 \mid 7.6-2$ | $9.4125-1 \mid 9.5136-1$ | $06 \mid 25: 51$ |

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

$$
\begin{aligned}
& p(\beta)=\lambda\|\beta\|_{1} \\
& q(\beta)=\sum_{i=1}^{n} \begin{cases}\frac{\beta_{i}^{2}}{a_{m}}, & \text { if }\left|\beta_{i}\right| \leq a_{m} \lambda, \\
2 \lambda\left|\beta_{i}\right|-a_{m} \lambda^{2}, & \text { if }\left|\beta_{i}\right|>a_{m} \lambda\end{cases}
\end{aligned}
$$

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

$$
\frac{\partial q(\beta)}{\partial \beta_{i}}= \begin{cases}\frac{2 \beta_{i}}{a_{m}}, & \text { if }\left|\beta_{i}\right| \leq a_{m} \lambda, \\ 2 \lambda \operatorname{sign}\left(\beta_{i}\right), & \text { if }\left|\beta_{i}\right|>a_{m} \lambda\end{cases}
$$

In our numerical experiments, we take $a_{m}=3.7$.

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

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

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

| $\begin{gathered} \text { probname } \\ \mathrm{m} ; \mathrm{n} \end{gathered}$ | $\lambda_{c}$ | nnz | $\eta_{k k t}$ | pobj | time |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  | $a \mid b$ | $a \mid b$ | $a \mid b$ |
| $\begin{gathered} \text { E2006.test } \\ 3308 ; 150358 \end{gathered}$ | 0.090 | 1 | 2.4-8 \| 9.3-7 | $2.2077+1$ \| $2.2077+1$ | 07 \| 07 |
| $\begin{gathered} \hline \log 1 \text { p.E2006.test } \\ 3308 ; 1771946 \end{gathered}$ | 0.261 | 187 | 8.2-7 \| 2.2-3 | $2.1455+1$ \| $3.6500+1$ | 4:09 \| 2:20:46 |
| pyrim.scale.expanded5 74;201376 | 0.221 | 43 | 9.9-7 \| 7.0-3 | $1.1428+0$ \| $4.6112+0$ | 18\| 19:36 |
| abalone.scale.expanded7 4177;6435 | 0.012 | 55 | 7.1-7 \| 1.6-5 | $1.3271+2$ \| $1.2693+2$ | 09 \| 21:32 |
| bodyfat.scale.expanded7 $252 ; 116280$ | 0.183 | 2 | 2.8-7 \| 5.3-6 | 5.7347-1 \| 5.8278-1 | 06\| 25:11 |

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[^0]:    ${ }^{1}$ Whenever 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.

