Outline

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- Computations

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Sources: Recent Conference Papers / Submissions

- Alacaoglu, A., Cevher, V., and Wright, S. J., On the Complexity of a Practical Primal-Dual Coordinate Method, arXiv preprint arXiv:2201.07684 (2022).
- Song, C., Lin, C. Y., Wright, S. J., and Diakonikolas, J., Coordinate Linear Variance Reduction for Generalized Linear Programming, arXiv preprint arXiv:2111.01842 (2021).
- Song, C., Wright, S. J. and Diakonikolas, J., Variance reduction via primal-dual accelerated dual averaging for nonsmooth convex finite-sums, In International Conference on Machine Learning, pp. 9824-9834. PMLR (2021).

Convex-Concave Min-Max

 $\min_{x \in \mathbb{R}^d} \max_{y \in \mathbb{R}^n} L(x, y)$ (Min-Max)

where

$$\begin{split} L(x,y) &= \sum_{i=1}^n \left[\langle A_i x, y^{(i)} \rangle - h_i^*(y^{(i)}) \right] + g(x) \\ &= \langle A x, y \rangle - h^*(y) + g(x), \end{split}$$

- $h_i^* : \mathbb{R} \to \mathbb{R} \cup \{+\infty\}$ is convex conjugate of h_i defined by $h_i^*(t) := \sup_s(st h_i(s))$ (convex and extended-valued);
- $g : \mathbb{R}^d \to \mathbb{R} \cup \{+\infty\}$ (convex and extended-valued);

•
$$h^*(y) = \sum_{i=1}^n h_i^*(y^{(i)})$$
 (separable);

- $A_i \in \mathbb{R}^d$ is a row vector;
- A is the $n \times d$ matrix with rows A_i .

More Specs

We consider cases in which *A* is dense and *A* is sparse.

In the case of sparse A, we assume for analysis that g is separable, that is,

$$g(x) = \sum_{j=1}^d g_j(x^{(j)}).$$

All algorithms make use of the prox-operator denoted for diagonal weighting matrix $T \succ 0$ and function g by $\operatorname{prox}_{T,g}$ and defined

$$\begin{split} \mathsf{prox}_{\mathrm{T},g}(x) &:= \arg\min_{u} \ \frac{1}{2} \|u - x\|_{\mathrm{T}^{-1}}^{2} + g(u) \\ &= \arg\min_{u} \ \frac{1}{2} \sum_{i=1}^{d} \frac{(x^{(i)} - u^{(i)})^{2}}{\mathrm{T}_{ii}} + g(u). \end{split}$$

Assume that we can compute prox-operators for g and h_i^* "easily."

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Primal-Dual Coordinate Methods

Case I: Empirical Risk Minimization (ERM)

Ubiquitous in statistics and machine learning.

$$\min_{x \in \mathbb{R}^d} \sum_{i=1}^n h_i(A_i x) + g(x).$$
 (ERM)

• Least squares:
$$h_i(t) = \frac{1}{2}t^2$$
;

- ℓ_1 regression: $h_i(t) = |t|;$
- Hinge loss: $h_i(t) = \max(t, 0)$ (used in SVM, neural nets);
- Regularization: Tikhonov $g(x) = \lambda ||x||_2^2$, ℓ_1 : $g(x) = ||x||_1$;
- TV regularization: $h_i(t) = ||t||_2$, $A_i \in \mathbb{R}^{2 \times d}$;
- logistic regression, least absolute deviation, ...

Case II: Linearly Constrained Optimization

min
$$g(x)$$
 s.t. $A_i x \in C_i$, $i = 1, 2, ..., n$, (LinOpt)

where $C_i = \{b_i\}$ (equality constraints) or $C_i = \{t : t \ge b_i\}$ (inequality).

Recall that we need to compute prox-operators involving convex g.

- Trivial if g is linear.
- Can be arranged (possibly via reformulation) if g is convex quadratic.

For algorithm output x_{out} , we could bound the number of iterates to attain

•
$$\mathbb{E}|g(x_{ ext{out}}) - g(x_{\star})| \leq arepsilon$$
, and

• $\mathbb{E} \operatorname{dist}(Ax_{\operatorname{out}}, C) \leq \varepsilon$.

Case III: Generalized LP

min
$$c^T x + r(x)$$
 s.t. $Ax = b, x \in \mathcal{X}$, (GLP)

which can be written in min-max form as

$$\min_{x\in\mathcal{X}}\max_{y\in\mathbb{R}^n}L(x,y)=\langle Ax,y\rangle+c^Tx+r(x)-b^Ty.$$

 $\mathcal{X} \subset \mathbb{R}^d$ is closed and convex, r is convex. We assume that the following modified prox-operator is easy to compute:

$$\operatorname{prox}_{\mathcal{X},r}(\hat{x}) := \arg\min_{z \in \mathcal{X}} \frac{1}{2} \|z - \hat{x}\|_2^2 + r(z).$$

- Ordinary LP: $\mathcal{X} = \mathbb{R}^d_{\geq 0}$ and regularized LP: $\mathcal{X} = \mathbb{R}^d_{\geq 0}$, $r(x) = \lambda \|x\|_2^2$;
- Reinforcement Learning [De Farias and Van Roy, 2003]
- Optimal Transport [Villani, 2009]
- DRO (f-divergence, Wasserstein) (see below)
- relaxed Neural Net verification [Liu et al., 2020].

Case IIIa: Distributionally Robust Optimization (DRO)

Setup: sample vectors $\{a_1, a_2, \ldots, a_n\}$ in \mathbb{R}^d with labels $\{b_1, b_2, \ldots, b_n\}$, where $b_i \in \{1, -1\}$. Usual ERM problem is

$$\min_{w} \frac{1}{n} \sum_{i=1}^{n} h(b_i a_i^T w)$$

where $h : \mathbb{R} \to \mathbb{R} \cup \{+\infty\}$ is convex (e.g. hinge loss).

 Wasserstein metric defines a distance between distributions ℙ and ℚ over ℝ^d × {−1,1}, based on cost

$$\zeta((\mathsf{a},\mathsf{b}),(\mathsf{a}',\mathsf{b}')) = \|\mathsf{a}-\mathsf{a}'\|_1 + \kappa |\mathsf{b}-\mathsf{b}'|$$

for some $\kappa > 0$;

P_n = ¹/_n Σⁿ_{i=1} δ_(a_i,b_i) is the empirical distribution defined by the data;
 Seek sup of the objective over the ball of radius ρ around P_n (in space)

of distributions over (a, b) defined by the Wasserstein metric:

$$\min_{w \in \mathbb{R}^d} \sup_{\text{dist}(\mathbb{P},\mathbb{P}_n) \leq \rho} \mathbb{E}^{\mathbb{P}}[h(ba^T w)].$$

GLP formulation of the DRO problem above is

$$\begin{split} \min_{\substack{w,\lambda,u,v,s,t}} & \rho\lambda + \frac{1}{n}\sum_{i=1}^{n}s_i \\ \text{s.t.} & u_i = b_i a_i^T w, \quad i = 1, 2, \dots, n, \\ & v_i = -u_i, \quad i = 1, 2, \dots, n, \\ & t_i = 2\kappa\lambda + s_i, \quad i = 1, 2, \dots, n, \\ & h(u_i) \leq s_i, \quad i = 1, 2, \dots, n, \\ & h(v_i) \leq t_i, \quad i = 1, 2, \dots, n, \\ & \|w\|_{\infty} \leq \lambda/M. \end{split}$$

 \mathcal{X} is defined by the last 3 constraints. The corresponding prox operation is separable so can be implemented easily.

Algorithms: The Basics

Formulation (reminder):

 $\min_{x \in \mathbb{R}^d} \max_{y \in \mathbb{R}^n} L(x, y)$ (Min-Max)

$$\begin{split} L(x,y) &= \sum_{i=1}^n \left[\langle A_i x, y^{(i)} \rangle - h_i^*(y^{(i)}) \right] + g(x) \\ &= \langle A x, y \rangle - h^*(y) + g(x), \end{split}$$

Gradient Ascent-Descent (GDA):

$$\bar{x}_{k+1} = \operatorname{prox}_{\tau,g}(\bar{x}_k - \tau A^\top \bar{y}_k)$$

$$\bar{y}_{k+1} = \operatorname{prox}_{\sigma,h^*}(\bar{y}_k + \sigma A \bar{x}_{k+1}),$$
(GDA)

for positive step sizes τ and σ . ¹

 $^{{}^{1}\}mathrm{T} = \tau I$ in our earlier definition of prox.

Algorithms: The Basics

Primal-Dual Hybrid Gradient (PDHG) [Chambolle and Pock, 2011] uses extrapolation in the x step:

$$\bar{x}_{k+1} = \operatorname{prox}_{\tau,g}(\bar{x}_k - \tau A^\top (2\bar{y}_k - \bar{y}_{k-1}))$$

$$\bar{y}_{k+1} = \operatorname{prox}_{\sigma,h^*}(\bar{y}_k + \sigma A \bar{x}_{k+1}),$$
(PDHG)

Equivalent form of PDHG:

$$\bar{x}_{k+1} = \mathsf{prox}_{\tau, g}(\hat{x}_k - \tau A^\top \bar{y}_k) \tag{1a}$$

$$\bar{y}_{k+1} = \operatorname{prox}_{\sigma,h^*}(\bar{y}_k + \sigma A \bar{x}_{k+1})$$
(1b)

$$\hat{x}_{k+1} = \bar{x}_{k+1} - \tau A^{\top} (\bar{y}_{k+1} - \bar{y}_k).$$
 (1c)

[Chambolle and Pock, 2011] discusses connections to Douglas-Rachford, Extrapolated gradient, ADMM.

[Aragón-Artacho et al., 2020] show application of DR to finding intersection of sets, pictures show the benefits of extrapolation.

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Primal-Dual Coordinate Methods

Algorithms: Additional Features

Theoretical convergence / complexity properties of these algorithms can be improved (in some cases, including strong convexity / concavity and sparsity) by adding extra features.

- Coordinate descent: e.g. update random element(s) of y in (1b) instead of the whole vector.
- Variance Reduction: Adjust the update formula for x to account for noise arising from coordinate update of y.
- Dual Averaging: At step k, use a gradient term that is a weighted average over all previous iterations.
- Importance sampling: Apply different weights to different components of each update (e.g. weight matrix T in definition of prox).
- Iterate averaging: Output a weighted average of iterates, rather than the final iterate for *x*.

Some are used by PURE-CD and VRPDA².

Measuring Approximate Optimality

Algorithms can be compared using rate of reduction of a gap function that measures near-optimality. Define G as follows:

$$G(x',y',x,y) := L(x',y) - L(x,y')$$

= $[g(x') + \langle Ax',y \rangle - h^*(y)] - [g(x) + \langle Ax,y' \rangle - h^*(y')],$

and for compact set $\mathcal{Z} \subset \mathbb{R}^d \times \mathbb{R}^n$ define the gap function

$$\mathsf{Gap}(x',y') := \max_{(x,y)\in\mathcal{Z}} G(x',y',x,y).$$

Suppose output (x^{K}, y^{K}) is generated by an algorithm after K iterations. If the algorithm is *random*, can measure:

- "max of expectation" $\max_{(x,y)\in\mathcal{Z}} \mathbb{E}G(x^{K}, y^{K}, x, y);$
- "expected gap" (stronger)

$$\mathbb{E} \max_{(x,y)\in\mathcal{Z}} G(x^{K}, y^{K}, x, y) = \mathbb{E} \operatorname{Gap}(x^{K}, y^{K}).$$

Complexity Analysis

Find upper bounds on the number of flops needed to reduce (expected) gap measures below a given threshold $\varepsilon > 0$. Particularly interested in dependence on ε as well as

- Dimensions d (for primal x) and n (for dual y);
- size of A: e.g. ||A||, $\max_{i=1,2,...,n} ||A_i||$, or $\sum_{i=1}^n ||A_i||$;
- nnz(A) (for sparse A);
- Distance between (x_0, y_0) and the optimum (x^*, y^*) .

Some algorithms (e.g. stochastic PDHG [Chambolle et al., 2018]) have less impressive bounds yet perform well for some types of problems.

PURE-CD: Dense A [Alacaoglu et al., 2020]

Define coordinate selection probabilities $p^{(i)}$, i = 1, 2, ..., n, with $P = \text{diag}(p^{(1)}, ..., p^{(n)})$

- 1: Initialize $x_0 \in \operatorname{dom} g, y_0 \in \operatorname{dom} h^*$
- 2: for $k \ge 0$ do

3:
$$\bar{x}_{k+1} = \operatorname{prox}_{\mathrm{T}_k,g}(x_k - \mathrm{T}_k A^{\top} y_k)$$

- 4: Pick $i_k \in [n]$ with $Pr(i_k = i) = p^{(i)}$
- 5: $[y_{k+1} = \operatorname{prox}_{\sigma_k, h^*}(y_k + \sigma_k A \bar{x}_{k+1})]_{i_k}, [y_{k+1} = y_k]_{i_k}$
- 6: $x_{k+1} = \bar{x}_{k+1} T_k \Theta_k A^\top P^{-1}(y_{k+1} y_k)$

7: end for

Notation:

- [·] *J* means that the formula is executed on only the components indexed by the set *J*.
- $[\cdot]_{\backslash J}$ means that the formula is executed on all components *except* those indexed by the set *J*.

PURE-CD: Dense A: Notes

Cost per iteration: O(d)

- $\operatorname{prox}_{T_k,g}$,
- calculate $A_{i_k} \bar{x}_{k+1}$,
- update $A^{\top}y_{k+1}$.

After K iterations, output x^{K} = average of $\bar{x}_{1}, \bar{x}_{2}, \ldots, \bar{x}_{K}$. (If needed, also output a weighted average of the y_{k} .)

Tracks PDHG with key differences:

- diagonal scalings: T_k , P, Θ_k ;
- Update just one (random) component of y in lines 4 and 5.

PURE-CD Dense: Complexity Result for LinOpt

min
$$g(x)$$
 s.t. $A_i x \in C_i, i = 1, 2, \ldots, n$.

Assume that primal-dual solution (x_{\star}, y_{\star}) exists. For scaling matrices:

$$T_{k} = T = \tau I, \quad \tau = \frac{1}{\sum_{i=1}^{n} \|A_{i}\|},$$

$$\Sigma_{k} = \Sigma = \operatorname{diag}(\sigma^{(1)}, \dots, \sigma^{(n)}), \quad \sigma^{(i)} = \frac{\gamma}{\|A_{i}\|}, \quad \gamma \in (0, 1),$$

$$p^{(i)} = \frac{\|A_{i}\|}{\sum_{i=1}^{n} \|A_{i}\|}, \quad \Theta_{k} = \theta I = I.$$

Define $x^{\mathcal{K}} = \frac{1}{\mathcal{K}} \sum_{k=1}^{\mathcal{K}} \bar{x}_k$. Then for D_1 , D_2 dep. on (x_0, y_0) and (x_\star, y_\star) :

$$\mathbb{E}|g(x^{\mathcal{K}}) - g(x_{\star})| \leq rac{8D_2}{\gamma(1-\gamma)}rac{\sum_{i=1}^n \|\mathcal{A}_i\|}{\mathcal{K}}, \ \mathbb{E}\left[\operatorname{dist}(\mathcal{A}x^{\mathcal{K}}, \mathcal{C})
ight] \leq rac{8D_1}{\gamma(1-\gamma)}rac{\sum_{i=1}^n \|\mathcal{A}_i\|}{\mathcal{K}},$$

with cost O(d) per iteration.

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PURE-CD Dense: Complexity Result for Min-Max

Here need more complicated settings of T_k (used in the algorithm) and Σ_k (used only in analysis). Set $\Theta_k \equiv I$ and $\sigma^{(i)} = \gamma/||A_i||$, some $\gamma \in (0, 1)$. Weighted averages of \bar{x}_k and y_k for k = 1, 2, ..., K yield outputs (x^K, y^K) . Use a special choice of starting points \bar{x}_1 , y_1 due to [Song et al., 2021b] Obtain a complicated bound for \mathbb{E} Gap (x^K, y^K) as a function of K – ultimately, arithmetic convergence at O(1/K) rate, after a burn-in period of slow linear convergence.

Can guarantee $\mathbb{E}\operatorname{Gap}(x^{K}, y^{K}) \leq \varepsilon$ with expected complexity

$$K = \tilde{O}\left(nd + D_{\mathcal{Z}}\frac{nd\max_{i} \|A_{i}\|}{\varepsilon}\right) \text{ iterations,}$$

where $D_{\mathcal{Z}}$ depends on set \mathcal{Z} and (x_0, y_0) and \tilde{O} indicates "ignoring log terms."

PURE-CD: Sparse A [Alacaoglu et al., 2020]

Recall notation
$$J(i) := \{j \in [d] : A_{i,j} \neq 0\}$$

1: Initialize
$$x_0 \in \text{dom } g, y_0 \in \text{dom } h^*$$
;
2: for $k \ge 0$ do
3: Pick $i_k \in [n]$ with $\Pr(i_k = i) = \frac{1}{n}$
4: $\left[\bar{x}_{k+1} = \operatorname{prox}_{\tau_k,g_j} (x_k - \tau_k (A^\top y_k))\right]_{J(i_k)}$
5: $\left[y_{k+1} = \operatorname{prox}_{\sigma_k,h^*} (y_k + \sigma_k A \bar{x}_{k+1})\right]_{i_k}$
6: $\left[y_{k+1} = y_k\right]_{\setminus i_k}$
7: $\left[x_{k+1} = \bar{x}_{k+1} - \tau_k \theta_k A_{i_k}^T (y_{k+1}^{(i_k)} - y_k^{(i_k)})\right]_{J(i_k)}$
8: $\left[x_{k+1} = x_k\right]_{\setminus J(i_k)}$
9: end for

PURE-CD: Sparse A: Comments

- Assumes that g is separable: $g(x) = \sum_{j=1}^{d} g_j(x^{(j)})$.
- Similar but not identical to the dense version of PURE-CD.
 - Calculate only the $J(i_k)$ components of the intermediate vector \bar{x}_{k+1} , which are needed to update the i_k component of y.
 - Since x_{k+1} depends componentwise on x̄_{k+1}, we calculate only the J(i_k) components of x̄_{k+1}.
 - ▶ But in the dense version, all components of x
 {k+1} are required to obtain x{k+1}.
 - Thus PURE-CD: Sparse needs its own convergence theory.
- Cost for iteration k is $O(|J(i_k)|)$ not O(d).

• (Expected cost is O(nnz(A)/n).)

PURE-CD Sparse: Complexity Results for Min-Max

Focus on results where strong convexity is present in g and/or h^* (both separable functions).

- Each g_j has modulus of convexity $\mu_g \ge 0$;
- Each h_i^* has modulus of convexity $\mu_h \ge 0$,

Results are for last iterates x_K and/or y_K , not averaged iterates.

When $\mu_g > 0$ and $\mu_h > 0$, we have $\mathbb{E}\left[\|x_K - x_\star\|^2 + \|y_K - y_\star\|^2\right] \le \varepsilon$ with expected complexity

$$ilde{O}\left(\left(\mathrm{nnz}(\mathsf{A})\left(1+rac{\max_{i}\|\mathsf{A}_{i}\|}{\sqrt{\mu_{h}\mu_{g}}}
ight)
ight)\logarepsilon^{-1}
ight).$$

Choices of Θ_k , $\sigma_k^{(i)}$, T_k do not depend on k, but require knowledge of μ_g and μ_h

PURE-CD Sparse: Complexity Results for Min-Max

When $\mu_g > 0$ but possibly $\mu_h = 0$ (strong convexity in g only) can make a (complicated) choice of parameters to ensure that $\mathbb{E}\left[\|x_{\mathcal{K}} - x_{\star}\|^2\right] \leq \varepsilon$ with expected complexity

$$O\left(\mathrm{nnz}(A) \left(1 + \sqrt{rac{D_{\star}}{arepsilon}} \max\left(1, rac{\max_i \|A_i\|}{\mu_g}
ight)
ight)
ight),$$

When $\mu_h > 0$ but possibly $\mu_g = 0$ (strong convexity in *h* only) a different (still complicated) choice of parameters $\sigma_k^{(j)}$, $\tau_k^{(j)}$, Θ_k ensures that $\mathbb{E}\left[\|y_K - y_\star\|^2\right] \le \varepsilon$ with expected complexity

$$O\left(\operatorname{nnz}(A)\left(1+\sqrt{\frac{D_{\star}}{\varepsilon}}\max\left(1,\frac{\max_{i}\|A_{i}\|}{\mu_{h}}\right)\right)\right),$$

Complexity Comparisons

The PURE-CD complexity bounds are compared with various other algorithms for Min-Max, or special cases of it:

- PDHG [Chambolle and Pock, 2011]
- SPDHG [Chambolle et al., 2018]
- VRPDA [Song et al., 2021b]
- CLVR [Song et al., 2021a]
- SPDAD [Tan et al., 2020]
- VRVI [Carmon et al., 2019, Alacaoglu and Malitsky, 2021]
- Katyusha [Allen-Zhu, 2017]
- SPDC [Zhang and Lin, 2015]

In each case, PURE-CD matches or improves the complexities of these alternatives, in terms of their dependence on n, d, measures of A, ε .

A typical improvement is $||A|| \to \max_i ||A_i|| - a$ factor of up to \sqrt{n} .

The proofs of these complexity results are extremely technical, involving mostly elementary manipulation of inequalities.

Telescoping sums over iterations k = 1, 2, ..., K is used often, and convexity is essential.

But considerable expertise is needed to choose the algorithmic parameters T_k , $\sigma_k^{(i)}$, Θ_k to achieve the desired cancellations.

VRPDA²: Another algorithm for Min-Max

[Song et al., 2021b]
1: Input:
$$(x_0, y_0) \in \mathcal{X} \times \mathcal{Y}, (u, v) \in \mathcal{X} \times \mathcal{Y}.$$

2: $\phi_0(\cdot) = \frac{1}{2} \|\cdot -x_0\|^2, \psi_0(\cdot) = \frac{1}{2} \|\cdot -y_0\|^2$. Initialize $y_1, x_1, \tilde{\psi}_1, \tilde{\phi}_1.$
3: $a_0 = B_0 = 0, \tilde{a}_1 = [2n \max_i ||A_i||]^{-1}$
4: $\psi_1 := n\tilde{\psi}_1, \phi_1 := n\tilde{\phi}_1, a_1 = B_1 = n\tilde{a}_1, z_1 = A^T y_1,$
5: for $k = 2, 3, ..., K$ do
6: $a_k = \min\left\{\tilde{a}_1\left(1 + \frac{1}{n-1}\right)^{k-1}, \frac{\sqrt{n(n+\sigma B_{k-1})}}{2R'}\right\}, B_k = B_{k-1} + a_k.$
7: Pick i_k uniformly at random in $[n]$.
8: $\bar{x}_{k-1} = x_{k-1} + \frac{a_{k-1}}{a_k}(x_{k-1} - x_{k-2}).$
9: $y_k = \arg\min_{y \in \mathbb{R}^n} \psi_k(y) := \psi_{k-1}(y) + na_k(\langle -A_{i_k}\bar{x}_{k-1}, y_{i_k} - v_{i_k} \rangle + h^*_{i_k}(y_{i_k})).$
10: $x_k = \arg\min_{x \in \mathbb{R}^d} \phi_k(x) := \phi_{k-1}(x) + na_k(x - u, z_{k-1} + (y_{k,i_k} - y_{k-1,i_k})A^T_{i_k} \rangle + g(x)).$
11: $z_k = z_{k-1} + (y_{k,i_k} - y_{k-1,i_k})A^T_{i_k}.$
12: end for
13: return weighted averages y^K and x^K .

VRPDA²: Notes

Also motivated by PDHG, similarities with PURE-CD.

- Special initialization for x_1 and y_1 . Costs one mult. by A and A^T .
- Extrapolation step in x (Line 8) with variable coefficient a_{k-1}/a_k .
- Update a single coordinate i_k of y (Line 9).
- Based on averaged gradients rather than steepest descent (as in PURE-CD
- Can't adapt to sparsity in A.
- Update steps (Lines 9-10) are prox-operations on $h_{i\nu}^*$ and g.
- Outputs weighted-average of x_k and y_k , k = 1, 2, ..., K as the result.

The algorithm and its analysis make heavy use of estimate sequences, in the style of [Nesterov, 2004].

$$\begin{split} \phi_k(x) &:= \phi_{k-1}(x) + a_k (n \langle x - u, z_{k-1} + (y_{k,i_k} - y_{k-1,i_k}) A_{i_k}^T \rangle + g(x)) \\ \psi_k(y) &:= \psi_{k-1}(y) + n a_k (\langle -A_{i_k} \bar{x}_{k-1}, y_{i_k} - v_{i_k} \rangle + h_{i_k}^*(y_{i_k})) \end{split}$$

VRPDA²: Convergence

For general case, attain $\mathbb{E}G(x^{K}, y^{K}, x^{\star}, y^{\star}) \leq \varepsilon$ (for weighted average iterates) in complexity

$$\tilde{O}\left(rac{nd\max_{i}\|A_{i}\|}{arepsilon}
ight).$$

When g is strongly convex with modulus μ_g , get $\mathbb{E} ||x^* - x_K||^2 \le \varepsilon$ (for last iterate x_K) in complexity

$$\tilde{O}\left(\frac{nd\max_i\|A_i\|}{\mu_g\sqrt{\varepsilon}}\right)$$

In PURE-CD we have the same bound ² but with *nd* replaced by nnz(A).

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²when max_{*i*} $||A_i||/\mu_g > 1$

CLVR: Specialized to GLP [Song et al., 2021a]

min
$$c^T x + r(x)$$
 s.t. $Ax = b, x \in \mathcal{X}$. (GLP)

K.

Partition A into m row blocks – index partition $\{S^1, S^2, \ldots, S^m\}$.

1: Input:
$$x_0 \in \mathcal{X}, y_0 \in \mathbb{R}^n, z_0 = A^T y_0, \gamma > 0, \hat{L} > 0, \sigma \ge 0,$$

2: $a_1 = B_1 = \frac{1}{2\hat{L}m}, q_0 = a_1(z_0 + c).$
3: for $k = 1, 2, ..., K$ do
4: $x_k = \operatorname{prox}_{\frac{1}{\gamma}B_k r}(x_0 - \frac{1}{\gamma}q_{k-1}).$
5: Pick j_k uniformly at random in $\{1, 2, ..., m\}.$
6: $[y_k = y_{k-1}]_{\setminus S^{j_k}}; [y_k = y_{k-1} + \gamma m a_k (Ax_k - b)]_{S^{j_k}};$
7: $a_{k+1} = \frac{\sqrt{1 + \sigma B_k / \gamma}}{2\hat{L}m}, B_{k+1} = B_k + a_{k+1}.$
8: $z_k = z_{k-1} + A_{S^{j_k}}^T (y_k^{S^{j_k}} - y_{k-1}^{S^{j_k}}).$
9: $q_k = q_{k-1} + a_{k+1}(z_k + c) + m a_k(z_k - z_{k-1}).$
10: end for

11: **return** weighted averages x^{K} and y^{K} .

CLVR: Notes and Complexity

Again related to PDHD but with variations.

- Averaged gradients in x, block coordinate descent in y.
- Recall that specialized prox-operator involves constraint set \mathcal{X} .
- Can be implemented in a way that exploits sparsity in A
 -but this involves intermediate vectors and is more complicated than in Sparse PURE-CD.

Expected complexity for $\mathbb{E}G(x^{K}, y^{K}, x^{\star}, y^{\star}) < \varepsilon$ in Sparse CLVR is

$$O\left(\frac{\operatorname{nnz}(A) \max_{i=1,2,\dots,m} \|A_{S^i}\|}{\varepsilon}\right)$$

Computational Results: DRO

- Wasserstein DRO described above, with ℓ_1 norm and hinge loss.
- Several standard ML datasets (LIBSVM).
- Implemented in Julia. Use SparseArrays to support sparse vectors and matrices.
- CLVR uses blocks to improve utilization of multiple cores.







Comparing with General LP solvers (times)

Time (seconds)	Reformulated a9a	Reformulated gisette	Reformulated rcv1
	d = 130738, n = 97929	d = 44002, n = 28000	d=269914, n=155198
PDHG	2422	2772	935
SPDHG	$>4 imes10^4$	1820	$3.7 imes10^4$
JuMP+GLPK	899	$>4 imes10^4$	$>4 imes10^4$
$\overline{\rm JuMP+Gurobi(simplex)}$	893	2482	7008
JuMP+Gurobi(barrier)	26	1039.7	1039.5
CLVR	962	697	582

Adapting to sparsity with PURE-CD

Investigating adaptation to sparsity in Lasso problem, compared to other stochastic coordinate methods

- SPDHG [Chambolle et al., 2018]³: good for dense, but O(d) per iteration cost
- VC-CD [Fercoq and Bianchi, 2019]⁴: good for sparse, but *n* times worse step size for dense
- PURE-CD: good for dense & sparse



rcv1: 0.16% density

w8a: 3.9% density

covtype: 22.1% density

³Stochastic PDHG: Chambolle et al., SIOPT, 2018

⁴Vu-Condat with coordinate descent: Fercoq, Bianchi, SIOPT, 2019

Wright (UW-Madison

Primal-Dual Coordinate Method

Concluding Thoughts and Questions

Simplicity is a great virtue but it requires hard work to achieve it and education to appreciate it. And to make matters worse: complexity sells better.

- Edsger Wybe Dijkstra

Algorithms that are simple yet with optimal complexity properties have taken some time to arrive. The analysis is still highly technical.

Can we simplify the analysis? Or define slightly different (but still useful) measures of algorithm performance that admit simpler analysis?

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