Random Primal-Dual Method for Parallel MRI Reconstruction

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- Examples of Convex Minimization
- Stochastic Primal-Dual Hybrid Gradient (SPDHG)
- Parallel MRI Reconstruction



$$\hat{x} \in \operatorname{argmin}_{x \in X} \sum_{i=1}^{n} f_i(A_i x) + g(x)$$

• $f_i: Y_i \to \mathbf{R}_{\infty}, g: X \to \mathbf{R}_{\infty}$ convex, proper, lower-semicontinuous • $A_i: X \to Y_i$ linear

- f_i, g can be **non-smooth**
- \hat{x} can be **not unique**

Examples



$$\hat{x} \in \operatorname*{argmin}_{x \in X} \sum_{i=1}^{n} f_i(A_i x) + g(x)$$

Image Denoising:

$$\hat{x} \in \operatorname*{argmin}_{r} \| \nabla x \|_1 + \lambda \| u^0 - x \|_1$$



Noisy image u^0



Examples



$$\hat{x} \in \operatorname*{argmin}_{x \in X} \sum_{i=1}^{n} f_i(A_i x) + g(x)$$

Binary Classification:







$$\hat{x} \in \operatorname{argmin}_{x \in X} \sum_{i=1}^{n} f_i(A_i x) + g(x)$$

$$\hat{x} \in \operatorname{argmin}_{x} \sum_{i=1}^{n} \|(S \circ F \circ C_i)x - b_i\|^2 + \frac{\lambda}{2} \|x\|^2$$



C)

Data b_i in Fourier space



$$\begin{cases} A_i = S \circ F \circ C_i \\ f_i = \| \cdot - \mathbf{b}_i \|^2 \\ g = \frac{\lambda}{2} \| \cdot \|^2 \end{cases}$$

Reconstructed image \hat{x}

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$$\hat{x} \in \operatorname*{argmin}_{x \in X} \left\{ \Phi(x) := \sum_{i=1}^{n} f_i(A_i x) + g(x) \right\}$$

Gradient Descent

$$x^{k+1} = x^k - \alpha \nabla \Phi(x^k)$$

requires smooth functionals f_i, g .

• Proximal Point Algorithm

$$x^{k+1} = \operatorname{prox}_{\sigma\Phi}(x^k)$$

proximity operator $prox_{\sigma\Phi} = prox_{\sigma(f \circ A+g)}$ is generally not explicit:

$$\operatorname{prox}_{\sigma\Phi}(x) := \operatorname{argmin}_{u} \frac{\|u - x\|^2}{2\sigma} + \Phi(u)$$



$$\hat{x} \in \operatorname{argmin}_{x \in X} \left\{ \Phi(x) := \sum_{i=1}^{n} f_i(A_i x) + g(x) \right\}$$

 \Leftrightarrow

 $y = (y_1, ..., y_n)$:

$$(\hat{x}, \hat{y}) \in rgmin_x \max_{y} \sum_{i=1}^n \langle A_i x, y_i
angle - f_i^*(y_i) + g(x)$$

where f^* is the *fenchel dual*:

$$f^*(\mathbf{y}) := \sup_w \langle w, \mathbf{y} \rangle - f(w)$$



$$\hat{x} \in \operatorname{argmin}_{x \in X} \left\{ \Phi(x) := \sum_{i=1}^{n} f_i(A_i x) + g(x) \right\}$$
$$y = (y_1, \dots, y_n) : \qquad \Longleftrightarrow$$
$$(\hat{x}, \hat{y}) \in \operatorname{argmin}_x \max_{y} \sum_{i=1}^{n} \langle A_i x, y_i \rangle - f_i^*(y_i) + g(x)$$

A primal-dual algorithm with step-size parameters $\tau, \sigma > 0$ reads:

$$\begin{aligned} x^{k+1} &= \operatorname{prox}_{\tau g}(x^k - \tau A^* y^k) \\ y^{k+1}_i &= \operatorname{prox}_{\sigma_i f^*_i}(y^k_i + \sigma_i A_i x^k) \qquad i \in \{1, ..., n\} \end{aligned}$$

In general, $\operatorname{prox}_{\tau g}$ and $\operatorname{prox}_{\sigma_i f_i^*}$ are explicit, while $\operatorname{prox}_{(f \circ A + g)}$ is not



$$(\hat{x}, \hat{y}) \in rg \min_{x} \max_{y} \sum_{i=1}^{n} \langle A_i x, y_i
angle - f_i^*(y_i) + g(x)$$

PDHG:

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

 $y_i^{k+1} = \operatorname{prox}_{\sigma f_i^*}(y_i^k + \sigma_i A_i x^{k+1})$ $i \in \{1, ..., n\}$
 $\bar{y}^k = 2y^{k+1} - y^k$

Theorem (Chambolle & Pock 2011)

Let f^*,g be proper, convex and lower-semicontinuous, and let τ,σ satisfy

$$\|S^{1/2}A\tau^{1/2}\|^2 < 1$$

where $S = \text{diag}(\sigma_1, ..., \sigma_n)$. Then PDHG converges to a saddle point.



$$(\hat{x},\hat{y}) \in rgmin_x \max_{y} \sum_{i=1}^n \langle A_i x, y_i
angle - f_i^*(y_i) + g(x)$$

$$\begin{array}{ll} \mathsf{PDHG}: & x^{k+1} = \mathrm{prox}_{\tau g}(x^k - \tau A^* \bar{y}^k) \\ & y_i^{k+1} = \mathrm{prox}_{\sigma f_i^*}(y_i^k + \sigma_i A_i x^{k+1}) & i \in \{1, ..., n\} \\ & \bar{y}^k = 2y^{k+1} - y^k \end{array}$$

Idea: Update only one dual variable y_i at random.



$$(\hat{x},\hat{y}) \in rg \min_{x} \max_{y} \sum_{i=1}^{n} \langle A_i x, y_i
angle - f_i^*(y_i) + g(x)$$

SPDHG: (serial sampling)

select $j \in \{1, \dots, n\}$ with probabilities $p_i = \mathbb{P}(i=j) > 0$

$$\begin{aligned} x^{k+1} &= \operatorname{prox}_{\tau g}(x^k - \tau A^* \bar{y}^k) \\ y^{k+1}_i &= \begin{cases} \operatorname{prox}_{\sigma_i f^*_i}(y^k_i + \sigma_i A_i x^{k+1}) & \text{if } i = j \\ y^k_i & \text{else} \end{cases} \\ \bar{y}^k &= y^{k+1} + \theta p_j^{-1}(y^{k+1} - y^k) \end{aligned}$$

1 iteration of PDHG ~pprox~n iterations of SPDHG

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$$(\hat{x}, \hat{y}) \in rgmin_x \max_y \sum_{i=1}^n \langle A_i x, y_i \rangle - f_i^*(y_i) + g(x)$$

SPDHG (arbitrary sampling S):

select $\mathbb{S}^k \in \{1, \dots, n\}$ with probabilities $p_i = \mathbb{P}(i \in \mathbb{S}^k) > 0$

$$\begin{aligned} x^{k+1} &= \operatorname{prox}_{\tau g}(x^k - \tau \bar{z}^k) \\ y_i^{k+1} &= \begin{cases} \operatorname{prox}_{\sigma_i f_i^*}(y_i^k + \sigma_i A_i x^{k+1}) & \text{if } i \in \mathbb{S}^k \\ y_i^k & \text{else} \end{cases} \\ \delta_i^k &= A_i^*(y_i^{k+1} - y_i^k) \text{ for all } i \in \mathbb{S}^k \\ z^{k+1} &= z^k + \sum_{i \in \mathbb{S}^k} \delta_i^k \\ \bar{z}^{k+1} &= z^{k+1} + \theta \sum_{i \in \mathbb{S}^k} p_i^{-1} \delta_i^k \end{aligned}$$

only A_i and A_i^* such that $i \in \mathbb{S}^k$ are evaluated

Stochastic Primal-Dual Hybrid Gradient



Previous result:

Convergence in Bregman distance

Theorem (Chambolle et al. 2018):

Let f_i^*, g be convex, proper and lower-semicontinuous, and

$$\tau \sigma_i \|A_i\|^2 < p_i, \quad i \in \{1, ..., n\}$$

then the **Bregman distance** of the iterates (x^k, y^k) to a solution (\hat{x}, \hat{y}) converges almost surely to zero, i.e.

$$\left[\langle A^* \hat{y}, x^k \rangle - f^*(\hat{y}) + g(x^k)\right] - \left[\langle A \hat{x}, y^k \rangle - f^*(y^k) + g(\hat{x})\right] \stackrel{a.s.}{\to} 0$$

Convergence in Bregman distance does not imply convergence in the norm



New result:

Convergence for serial sampling and finite dimension

Theorem (Gutiérrez, Delplancke, Ehrhardt 2021)

Let f_i^\ast,g be convex, proper and lower-semicontinuous in finite dimensional separable Hilbert spaces, and

$$\tau \sigma_i \|A_i\|^2 < p_i, \quad i \in \{1, ..., n\}$$

then SPDHG with serial sampling converges a.s. to a saddle point.



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then SPDHG with serial sampling converges a.s. to a saddle point.

Theorem (Alacaoglu et al. 2020)



Even newer result:

Convergence for any sampling and infinite dimensions

Theorem

Let f_i^*, g be convex, proper and lower-semicontinuous in separable Hilbert spaces of **arbitrary dimension** and let \mathbb{S} denote **any random sampling** of $\{1, ..., n\}$. Let $\{v_1, ..., v_n\} > 0$ be such that

$$\mathbb{E}_{\mathbb{S}} \Big\| \sum_{i \in \mathbb{S}} \tau^{1/2} \sigma_i^{1/2} A_i^* z_i \Big\|^2 \le \sum_{i=1}^n p_i v_i \|z_i\|^2 \quad \text{and} \quad v_i < p_i \quad i \in \{1, ..., n\}$$

Then SPDHG converges a.s. to a saddle point.

Furthermore, for different samplings we propose step size parameters for optimal convergence speed.





MRI Scanner

Original:

 x^0

Encoded Data:

$$\mathbf{b}_i = (S \circ F \circ C_i) \mathbf{x}^0 + \eta_i$$



Sensitivity of each coil

Reconstructed Solution:

$$\hat{x} \in \operatorname{argmin}_{x} \sum_{i=1}^{n} \|(S \circ F \circ C_i)x - b_i\|^2 + \frac{\lambda}{2} \|x\|^2$$





 $\lambda = 1$

 $\lambda = 1$



 $\lambda = 10^{-2}$



 $\lambda = 10^{-4}$

Dataset 2:

Dataset 1:



$$\lambda = 10^{-2}$$



 $\lambda = 10^{-4}$











PDHG



solution

1 iteration of PDHG $\approx n$ iterations of SPDHG ≈ 1 epoch





1 iteration of PDHG $\approx n$ iterations of SPDHG ≈ 1 epoch



full sampling: 1 block of size $n \implies 1$ epoch ≈ 1 iteration

$$\Big\{\,(1,...,n)\,\Big\}$$

serial sampling: n blocks of size $1 \implies 1$ epoch $\approx n$ iterations

$$\left\{\,(1),(2),...,(n)\,\right\}$$

b-serial sampling: n/b blocks of size $b \implies 1$ epoch $\approx n/b$ iterations

$$\left\{\,(1,...,b),(b+1,...,2b),\,\ldots,(n-b+1,...,n)\,\right\}$$



Reconstruction error for $n = 12, \lambda = 10^{-2}$



The converge speed depends on the batch size

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Theorem (Chambolle et al. 2018)

Let f_i^{\ast},g be $\ensuremath{\mathsf{strongly}}$ convex, proper and lower-semicontinuous, and

 $\tau \sigma_i \|A_i\|^2 < p_i, \quad i \in \{1, ..., n\}$

then $(x^k, y^k) \stackrel{a.s.}{\to} (\hat{x}, \hat{y})$ with linear rate $\mathcal{O}(\theta^k)$.

Furthermore, for each sampling (full, serial, *b*-serial, etc...) we can compute the **optimal** convergence rate $\theta \in (0, 1)$.





The converge rate ϑ depends on the batch size



Given a fixed b, how many partitions of $\{1, ..., n\}$ exist of subsets of size b?



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A lot.



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Example: n = 12, b = 6, there are 462 different partitions of $\{1, ..., n\}$ into two subsets of size 6.

$$\left\{ (1,2,3,4,5,6), (7,8,9,10,11,12) \right\}$$
$$\left\{ (1,2,3,4,5,7), (6,8,9,10,11,12) \right\}$$
$$\vdots$$
$$\left\{ (1,8,9,10,11,12), (2,3,4,5,6,7) \right\}$$



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$$\vdots$$
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Example: For n = 12, b = 2, the number of partitions is 10,395.



Example: n = 12, b = 6

For each possible sampling of size b = 6, we compute ϑ :



The convergence rate ϑ depends on which partition we sample from

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$$n = 12, b = 6$$



The location of the coils gives us a clue on how to find the best partition



SPDHG

- Wide variety of applications
- Non-smooth functionals, large number of data
- Faster than PDHG

Our Contributions

- We prove convergence for any sampling
- We propose optimal step size parameters for different samplings
- For parallel MRI we show how convergence speed is affected by
 - Batch size b
 - The partition of $\{1,...,n\}$
 - The physical location of the receiver coils