

# A WEIGHTED DIFFERENCE OF ANISOTROPIC AND ISOTROPIC TOTAL VARIATION MODEL FOR IMAGE PROCESSING

YIFEI LOU\*, TIEYONG ZENG†, STANLEY OSHER‡, AND JACK XIN§

**Abstract.** We propose a weighted difference of anisotropic and isotropic total variation (TV) as a regularization for image processing tasks, based on the well-known TV model and natural image statistics. Due to the difference form of our model, it is natural to compute via a difference of convex algorithm (DCA). We draw its connection to the Bregman iteration for convex problems, and prove that the iteration generated from our algorithm converges to a stationary point with the objective function values decreasing monotonically. A stopping strategy based on the stable oscillatory pattern of the iteration error from the ground truth is introduced. In numerical experiments on image denoising, image deblurring, and magnetic resonance imaging (MRI) reconstruction, our method improves on the classical TV model consistently, and is on par with representative start-of-the-art methods.

**Key words.** Anisotropic TV, Isotropic TV, Weighted Difference, Difference of Convex Algorithm, Convergence to Stationary Points, Stable Oscillatory Errors, Bregman and Split Bregman Iterations.

**AMS subject classifications.** 90C90, 65K10, 49N45, 49M20

**1. Introduction.** Many image processing tasks can be formulated as an inverse problem, in which the data  $f$  is assumed to be obtained approximately by applying a linear operator  $A$  on an image  $u$  with additive noise. For example,  $A$  is the identity matrix for image denoising, a convolution matrix for deblurring, and subsampling of Fourier transform for a magnetic resonance image (MRI) reconstruction problem. In most scenarios, solving  $u$  from  $Au = f$  is ill-posed in the sense that directly inverting  $A$  would result in bad and possibly multiple solutions. It is necessary and even desirable to constrain the solutions through regularization, with the help of prior knowledge of images that one wants to reconstruct. A general model for such inverse problem is

$$\hat{u} := \operatorname{argmin}_u J(u) + \frac{\mu}{2} \|Au - f\|_2^2, \quad (1.1)$$

where  $J(u)$  is the regularization term,  $\mu$  is a positive parameter to balance  $J(u)$  and the data fidelity term  $\|Au - f\|_2^2$ , and  $\hat{u}$  is an optimal solution of the model or a reconstructed result. A classical regularization is the total variation (TV) proposed by Rudin-Osher-Fatemi [33]. It is widely used in image processing applications, such as deconvolution [7, 16, 25], inpainting [6] and super-resolution [26], just to name a few. The TV model originated in [33] is isotropic, and later an anisotropic formulation has been addressed in the literature [10, 30] among others. We give mathematical definition for both the isotropic and anisotropic TV in the discrete setting. Denoting  $u$  as the column vector by a lexicographical ordering of a 2D image, we have

$$J_{iso}(u) := \|\nabla u\|_2 = \|\sqrt{|D_x u|^2 + |D_y u|^2}\|_1, \quad (1.2)$$

$$J_{ani}(u) := \|D_x u\|_1 + \|D_y u\|_1, \quad (1.3)$$

where  $D_x, D_y$  denote the horizontal and vertical partial derivative operators. Throughout this paper, we shall use notations  $\|\nabla u\|_2$  and  $\|\sqrt{|D_x u|^2 + |D_y u|^2}\|_1$  interchangeably.

Another interpretation of TV can be given from the perspective of compressive sensing (CS) [3, 12], which is to reconstruct a signal from an under-determined system provided that the signal is

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## Report Documentation Page

*Form Approved*  
*OMB No. 0704-0188*

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1. REPORT DATE <b>SEP 2014</b>	2. REPORT TYPE	3. DATES COVERED <b>00-00-2014 to 00-00-2014</b>	
4. TITLE AND SUBTITLE <b>A Weighted Difference of Anisotropic and Isotropic Total Variation Model for Image Processing</b>		5a. CONTRACT NUMBER	
		5b. GRANT NUMBER	
		5c. PROGRAM ELEMENT NUMBER	
6. AUTHOR(S)		5d. PROJECT NUMBER	
		5e. TASK NUMBER	
		5f. WORK UNIT NUMBER	
7. PERFORMING ORGANIZATION NAME(S) AND ADDRESS(ES) <b>University of California, Los Angeles, Department of Mathematics, Los Angeles, CA, 90095</b>		8. PERFORMING ORGANIZATION REPORT NUMBER <b>CAM14-69</b>	
9. SPONSORING/MONITORING AGENCY NAME(S) AND ADDRESS(ES)		10. SPONSOR/MONITOR'S ACRONYM(S)	
		11. SPONSOR/MONITOR'S REPORT NUMBER(S)	
12. DISTRIBUTION/AVAILABILITY STATEMENT <b>Approved for public release; distribution unlimited</b>			
13. SUPPLEMENTARY NOTES			
14. ABSTRACT <b>We propose a weighted difference of anisotropic and isotropic total variation (TV) as a regularization for image processing tasks, based on the well-known TV model and natural image statistics. Due to the difference form of our model, it is natural to compute via a difference of convex algorithm (DCA). We draw its connection to the Bregman iteration for convex problems, and prove that the iteration generated from our algorithm converges to a stationary point with the objective function values decreasing monotonically. A stopping strategy based on the stable oscillatory pattern of the iteration error from the ground truth is introduced. In numerical experiments on image denoising, image deblurring, and magnetic resonance imaging (MRI) reconstruction, our method improves on the classical TV model consistently, and is on par with representative start-of-the-art methods.</b>			
15. SUBJECT TERMS			
16. SECURITY CLASSIFICATION OF:			17. LIMITATION OF ABSTRACT
a. REPORT <b>unclassified</b>	b. ABSTRACT <b>unclassified</b>	c. THIS PAGE <b>unclassified</b>	<b>Same as Report (SAR)</b>
			18. NUMBER OF PAGES <b>17</b>
			19a. NAME OF RESPONSIBLE PERSON

sufficiently sparse or sparse in a transform domain. For example, a natural image is mostly sparse after taking gradient. Mathematically, it amounts to minimizing the  $L_0$  norm of the image gradient, *i.e.*,  $J(u) = \|\nabla u\|_0$ . To bypass the NP-hard  $L_0$  norm, the convex relaxation approach in CS is to replace  $L_0$  by  $L_1$ , and  $L_1$  on the gradient is the total variation. The restricted isometry property (RIP) condition [3] theoretically guarantees the exact recovery of sparse solutions by  $L_1$ . The RIP regime is where the sensing matrix is *incoherent*, such as a random Gaussian matrix. Several non-convex penalties have been proposed and studied as alternatives to  $L_1$ , [19]. A few notable examples are  $L^p$  for  $p \in (0, 1)$  [8, 21, 39],  $L_1/L_2$  (scale invariant  $L_1$ ) and  $L_1 - L_2$  [13, 22, 23, 40, 41]. In particular,  $L_1 - L_2$  penalty is found to be the best among existing methods for recovering sparse solutions when the sensing matrix is highly coherent or significantly violating the RIP condition [23, 41].

The TV-regularization has been a very active research topic in the past two decades. Though a gradient descent approach in the original paper can be slow to converge, a projection algorithm is later proposed by Chambolle [5] to speed up convergence based on duality. More recently, the Bregman and split Bregman methodology [9, 15, 29] offers another line of fast algorithms equivalent to the role of alternating direction method of multipliers (ADMM) and Douglas-Rachford splitting algorithm in the optimization literature dating back to the 1970's. There are also a few approaches to solve the  $L_0$  minimization directly. In [38], a special alternating minimization strategy with half-quadratic splitting is adopted for image smoothing. Image restoration via  $L_0$  is considered in [31], which uses hard shrinkage for  $L_0$  as opposed to soft shrinkage for  $L_1$ . In addition, the  $L_0$  on the gradient can be interpreted as the length of the partition boundaries, which leads to the classical Potts model [32] or piece-wise constant Mumford-Shah model [28] for image segmentation or partition. Recently, Storath *et. al.* [34] propose a hybrid ADMM and dynamic programming method to solve the Potts model.

Motivated from  $L_1 - L_2$  minimization of coherent CS [23, 41], we propose the following weighted difference of convex regularization,

$$J(u) := J_{ani} - \alpha J_{iso} = \|D_x u\|_1 + \|D_y u\|_1 - \alpha \|\sqrt{|D_x u|^2 + |D_y u|^2}\|_1, \quad (1.4)$$

where  $\alpha \in [0, 1]$  is a parameter for a more general model. When  $\alpha = 1$ ,  $J(u)$  is to apply  $L_1 - L_2$  on the gradient vector. Two advantages of  $L_1 - L_2$  over other nonconvex measures are its Lipschitz regularity, and guaranteed convergence via the difference of convex algorithm (DCA) [35, 36]. We find that the DCA requires solving the  $L_1$  type of minimization as a subproblem, which can be handled efficiently by utilizing the split Bregman technique. We prove that the DCA approach converges to stationary points, a typical situation for nonconvex problems. In practice, the DCA iterations, when properly stopped, are often close to global minima and produce excellent results. The stopping issue is discussed later based on the oscillatory pattern of the iteration errors.

The rest of the paper is organized as follows. Section 2 describes our model in detail including numerical algorithms and convergence analysis. Section 3 is devoted to numerical experiments, where three image processing applications (denoising, deblurring and MRI reconstruction) are examined. Finally, discussions and conclusions are given in Section 4 and Section 5 respectively.

**2. Our model.** Let  $(u_{jx}, u_{jy})$  be gradient vector at pixel  $j$ . Then equation (1.4) can be rewritten as

$$J(u) = \sum_j \left( |u_{jx}| + |u_{jy}| - \alpha \sqrt{u_{jx}^2 + u_{jy}^2} \right). \quad (2.1)$$

This point-wise formulation suggests that sparsity is enforced on every gradient vector. More specifically, we encourage the gradient to be 1-sparse at every pixel, which implies that horizontal or vertical edges are more preferable in this model. In order to understand the image gradient and 1-sparsity, we plot the histogram of gradient angles over the range of  $[0, 90]$  degree in Figure 2.1 for a large number of natural images. The angle distribution in other quadrants is similar. As shown in Figure 2.1, the

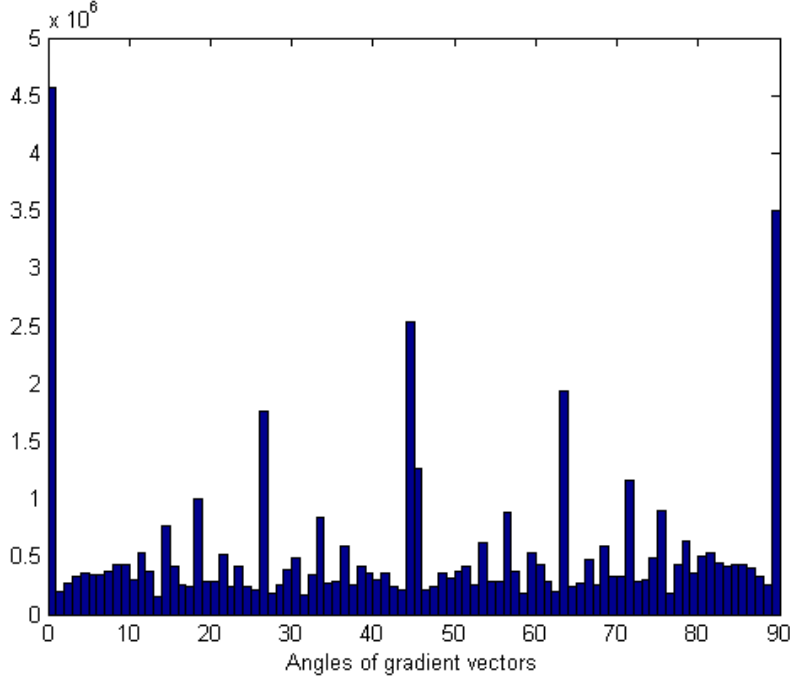


FIG. 2.1. The histogram of gradient angles over 300 images from Berkeley segmentation dataset [27]. Two largest peaks are at 0 and 90 degrees, indicating that gradient vectors are mostly 1-sparse.

two largest peaks are at 0 and 90 degrees, which implies that gradient vectors are 1-sparse at a fairly good chance, with non-sparse occurrences also at positive probability. Hence we insert a constant  $\alpha$  in (1.4) to reflect such behavior in the histogram. In Figure 2.2, we plot the level lines of  $L_0$  norm on the gradient, whose value is 0 at origin, 1 at axes, and 2 elsewhere. The level lines corresponding to  $\alpha < 1$  in (1.4) is closer to  $L_0$  than that of  $\alpha = 1$  in the sense that the latter yields 0 at both axes.

Let us derive the value of  $\alpha$  based on the gradient distribution. Suppose that the gradient value  $D_x u$  follows the distribution [21],  $\frac{p}{2\Gamma(\frac{1}{p})}e^{-|x|^p}$  where  $\Gamma(t) = \int_0^{+\infty} x^{t-1}e^{-x}$ . It is Gaussian distribution for  $p = 2$ , Laplacian distribution for  $p = 1$ , and hyper-Laplacian for  $0 < p < 1$ . We have

$$E_1 = E|D_x u| = \frac{p}{2\Gamma(\frac{1}{p})} \int_{-\infty}^{+\infty} e^{-|x|^p} |x| dx = \frac{1}{\Gamma(\frac{1}{p})} \int_0^{+\infty} e^{-t} t^{\frac{2}{p}-1} dt = \frac{\Gamma(\frac{2}{p})}{\Gamma(\frac{1}{p})}, \quad (2.2)$$

$$E_2 = E|D_x u|^2 = \frac{p}{2\Gamma(\frac{1}{p})} \int_{-\infty}^{+\infty} e^{-|x|^p} |x|^2 dx = \frac{1}{\Gamma(\frac{1}{p})} \int_0^{+\infty} e^{-t} t^{\frac{3}{p}-1} dt = \frac{\Gamma(\frac{3}{p})}{\Gamma(\frac{1}{p})}. \quad (2.3)$$

The value of  $\alpha$  corresponds to the ratio of  $L_1$  and  $L_2$ , *i.e.*,

$$\alpha = \frac{E_1}{\sqrt{E_2}} = \frac{\Gamma(2/p)}{\sqrt{\Gamma(3/p)\Gamma(1/p)}}. \quad (2.4)$$

Table 2.1 lists the values of  $\alpha$  based on gradient distributions for  $p = 0.5, 1, 2$ . We analyze the gradient distribution in Figure 2.3 which shows that the distribution of image gradient data matches the  $p = 1/2$  distribution better than classical Gaussian ( $p = 2$ ) or Laplacian ( $p = 1$ ) distribution. This observation is consistent with the choice of hyper-Laplacian [4, 21] for image processing ( $p \in [0.5, 0.8]$ ). In the rest of the paper, we shall fix the weighting coefficient  $\alpha = 1/2$  to approximate the desired value in Table 2.1.

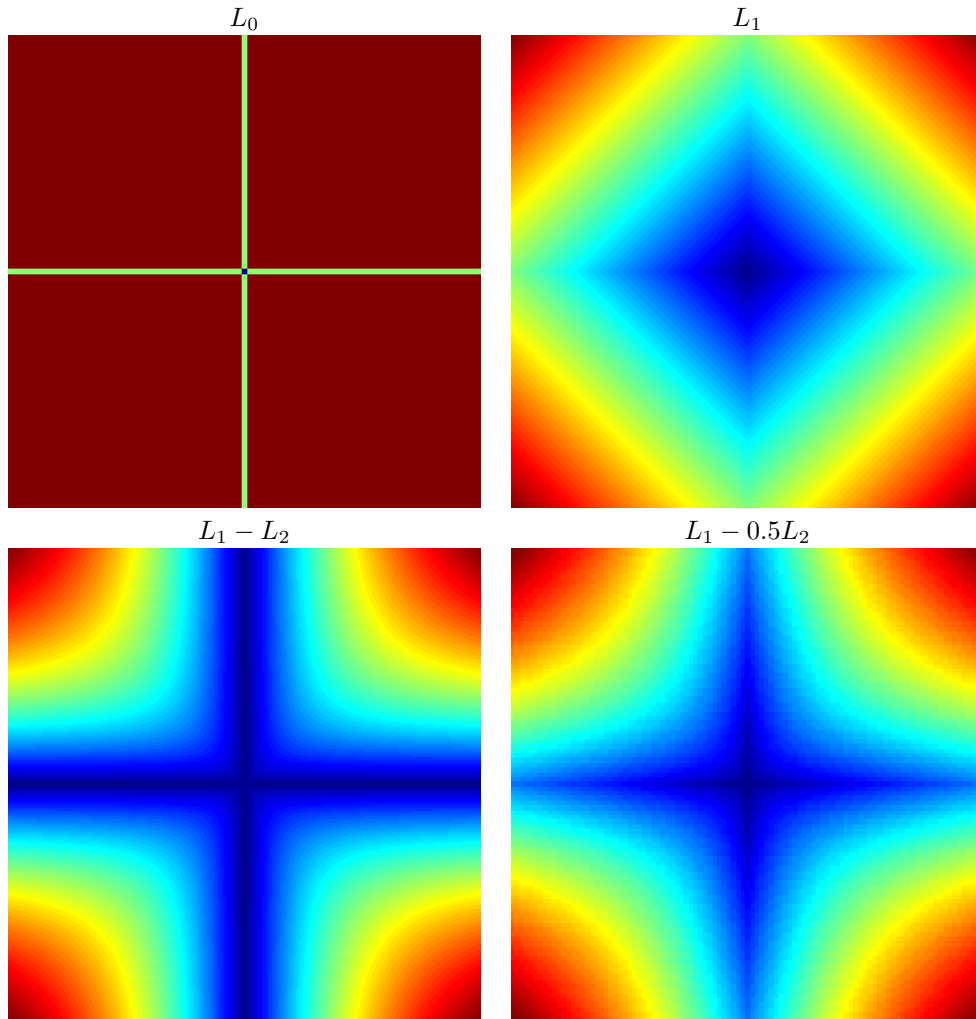


FIG. 2.2. Level curves of different metrics. The level lines corresponding to  $\alpha < 1$  in (1.4) is closer to  $L_0$  than that of  $\alpha = 1$  in the sense that the latter yields 0 at both axes.

TABLE 2.1  
The value of  $\alpha$  based on the gradient distribution.

$p$	$\alpha$
0.5	0.5477
1	0.7071
2	0.7979

**2.1. Numerical algorithms.** To solve (1.1) with  $J(u)$  defined in (1.4), we apply the technique of difference of convex algorithm (DCA) by linearizing the isotropic term

$$u^{n+1} = \arg \min_u \|D_x u\|_1 + \|D_y u\|_1 - \alpha \langle \nabla u, q^n \rangle + \frac{\mu}{2} \|Au - f\|_2^2, \quad (2.5)$$

for  $q^n = (q_x^n, q_y^n) = (D_x u^n, D_y u^n) / \sqrt{|D_x u^n|^2 + |D_y u^n|^2}$  at step  $u^n$ . Note that  $q^n$  is a point-wise calculation; and if the denominator is zero at some point, the corresponding  $q^n$  value is set to be zero. Each DCA subproblem, eq. (2.5), amounts to solving a TV type of minimization. We employ the split

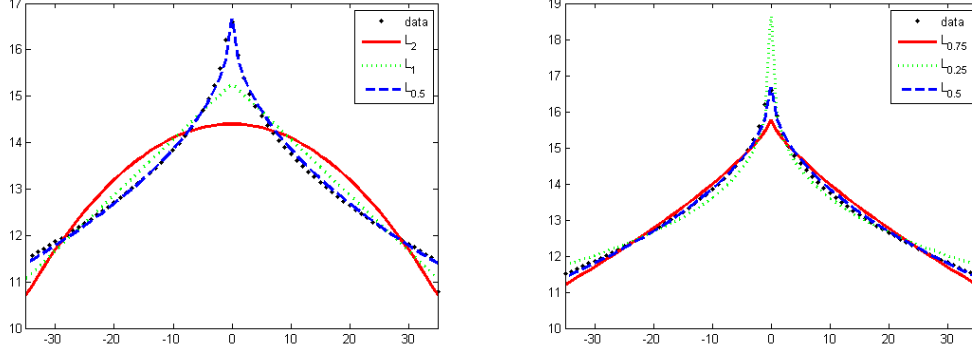


FIG. 2.3. The plot of log probability v.s. gradient in comparison with different distributions, indicating that the gradient distribution of a large natural image dataset matches  $L_{1/2}$  or the  $p = 1/2$  hyper-Laplacian distribution better than classical Gaussian or Laplacian distribution.

Bregman technique [15] to do the job. Specifically, we introduce two auxiliary variables and split the anisotropic term in the following way,

$$\begin{aligned}
 u^{k+1} = \arg \min_{u, d_x, d_y} & \|d_x\|_1 + \|d_y\|_1 - \alpha(d_x^T \cdot q_x^n + d_y^T \cdot q_y^n) \\
 & + \frac{\mu}{2} \|Au - f\|_2^2 + \frac{\lambda}{2} \|d_x - D_x u\|_2^2 + \frac{\lambda}{2} \|d_y - D_y u\|_2^2,
 \end{aligned} \tag{2.6}$$

in which  $d_x, d_y$  can be updated via soft shrinkage, defined as

$$\text{shrink}(s, \gamma) = \text{sgn}(s) \max\{|s| - \gamma, 0\} . \tag{2.7}$$

The pseudo-code is summarized in Algorithm 1. The algorithm is efficient for many applications where the matrix to be inverted is diagonal or can be diagonalized by Fourier transform, which is true for image denoising, deconvolution, and MRI reconstruction.

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**Algorithm 1** for solving unconstrained problem (2.5)

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Define  $u = q_x = q_y = 0, z = f$  and  $MaxDCA, MaxBregman$ 
for 1 to  $MAXDCA$  do
   $b_x = b_y = 0$ 
  for 1 to  $MAXBregman$  do
     $u = (\mu A^T A - \lambda \Delta)^{-1} (\mu A z + \lambda D_x^T (d_x - b_x) + \lambda D_y^T (d_y - b_y))$ 
     $d_x = \text{shrink}(D_x u + b_x + \alpha q_x / \lambda, 1 / \lambda)$ 
     $d_y = \text{shrink}(D_y u + b_y + \alpha q_y / \lambda, 1 / \lambda)$ 
     $b_x = b_x + D_x u + d_x$ 
     $b_y = b_y + D_y u + d_y$ 
  end for
   $(q_x, q_y) = (D_x u, D_y u) / \sqrt{|D_x u|^2 + |D_y u|^2}$ 
end for

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For the corresponding constrained problem,

$$\min \|D_x u\|_1 + \|D_y u\|_1 - \alpha \|\nabla u\|_2 \quad \text{s.t.} \quad Au = f, \tag{2.8}$$

the DCA is expressed as

$$u^{n+1} = \arg \min_u \{ \|D_x u\|_1 + \|D_y u\|_1 - \alpha \langle \nabla u, q^n \rangle \quad \text{s.t.} \quad Au = f \} . \tag{2.9}$$

Each DCA subproblem could be reduced to a sequence of unconstrained problems of the form

$$u_{k+1} = \arg \min_u \|D_x u\|_1 + \|D_y u\|_1 - \alpha \langle \nabla u, q^n \rangle + \frac{\mu}{2} \|Au - z_k\|_2^2, \quad (2.10)$$

$$z_{k+1} = z_k + f - Au_{k+1}. \quad (2.11)$$

Again the first equation can be solved by the split Bregman method. Algorithm 2 for solving the constrained problem (2.9) is almost the same as Algorithm 1, except for an additional update on  $z$ .

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**Algorithm 2** for solving constrained problem (2.9)

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Define  $u = q_x = q_y = 0, z = f$  and *MaxDCA*, *MaxBregmanInner*, *MaxBregmanOuter*

**for** 1 **to** *MAXDCA* **do**

$b_x = b_y = 0$

**for** 1 **to** *MaxBregmanOuter* **do**

**for** 1 **to** *MAXBregmanInner* **do**

$u = (\mu A^T A - \lambda \Delta)^{-1} (\mu Az + \lambda D_x^T (d_x - b_x) + \lambda D_y^T (d_y - b_y))$

$d_x = \text{shrink}(D_x u + b_x + \alpha q_x / \lambda, 1 / \lambda)$

$d_y = \text{shrink}(D_y u + b_y + \alpha q_y / \lambda, 1 / \lambda)$

$b_x = b_x + D_x u + d_x$

$b_y = b_y + D_y u + d_y$

**end for**

$z = z + f - Au$

**end for**

$(q_x, q_y) = (D_x u, D_y u) / \sqrt{|D_x u|^2 + |D_y u|^2}$

**end for**

---

**2.2. Convergence analysis.** We want to show that the sequence of  $\{u^n\}$  obtained from the DCA iterations, *i.e.*, eq. (2.5), converges to a stationary point. The standard DCA requires strong convexity to prove convergence [35], here we can get rid of this requirement using the fact that  $L_1$  is convex (not strictly though), and its subgradient<sup>1</sup> is a close set. We first prove two lemmas saying that the objective function is coercive and monotonically non-increasing for the minimizing sequence; and then complete the convergence proof.

LEMMA 2.1. *Suppose  $\mu > 0, 0 < \alpha < 1$ , and  $\ker(A) \cap \ker(D) = \{\mathbf{0}\}$ , where  $D = [D_x; D_y]$ . Then the objective function*

$$F(u) := \|D_x u\|_1 + \|D_y u\|_1 - \alpha \sqrt{|D_x u|^2 + |D_y u|^2} + \frac{\mu}{2} \|Au - f\|_2^2,$$

*is coercive.*

*Proof.* It suffices to show that for any fixed  $u \in \mathbb{R}^N \setminus \{\mathbf{0}\}$ ,  $F(\gamma u) \rightarrow \infty$  as  $\gamma \rightarrow \infty$ . We discuss two cases separately.

- If  $u \notin \ker(D)$ , then we have

$$F(\gamma u) > (1 - \alpha)\gamma(\|D_x u\|_1 + \|D_y u\|_1) \rightarrow \infty \quad \text{as } \gamma \rightarrow \infty,$$

since  $\sqrt{|D_x u|^2 + |D_y u|^2} < \|D_x u\|_1 + \|D_y u\|_1$ .

- If  $u \in \ker(D)$ , then  $Au \neq 0$ , since  $\ker(A) \cap \ker(D) = \{\mathbf{0}\}$  and  $u \neq 0$ . Therefore, we have

$$F(\gamma u) \geq \frac{\mu}{2} \|\gamma Au - f\|_2^2 \geq \frac{\mu}{2} (\gamma \|Au\|_2^2 - \|f\|_2^2) \rightarrow \infty \quad \text{as } \gamma \rightarrow \infty.$$

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<sup>1</sup>We say a vector  $g$  is a *subgradient* of  $f$  at  $x \in \mathbf{dom}(f)$  if  $f(z) \geq f(x) + g^T(z - x)$  for all  $z \in \mathbf{dom}(f)$ .

□

LEMMA 2.2. *If the sequence  $\{u^n\}$  is generated by the DCA algorithm (2.5), i.e.,*

$$u^{n+1} = \arg \min \|D_x u\|_1 + \|D_y u\|_1 - \alpha \left\langle \frac{\nabla u^n}{\|\nabla u^n\|}, \nabla u \right\rangle + \frac{\mu}{2} \|Au - f\|_2^2,$$

then

$$F(u^n) - F(u^{n+1}) \geq 0. \quad (2.12)$$

*Proof.* It follows from the first-order optimality condition at  $u^{n+1}$  that there exist  $p^{n+1} \in \partial \|Du^{n+1}\|_1$  such that

$$p^{n+1} - \alpha q^n + \mu A^T(Au^{n+1} - f) = 0. \quad (2.13)$$

A simple calculation shows that

$$\begin{aligned} & F(u^n) - F(u^{n+1}) \\ &= \frac{\mu}{2} \|A(u^n - u^{n+1})\|_2^2 + \mu \langle A(u^n - u^{n+1}), Au^{n+1} - f \rangle + \|Du^n\|_1 - \|Du^{n+1}\|_1 - \alpha (\|\nabla u^n\|_2 - \|\nabla u^{n+1}\|_2) \\ &= \frac{\mu}{2} \|A(u^n - u^{n+1})\|_2^2 - \langle p^{n+1} - \alpha q^n, u^n - u^{n+1} \rangle + \|Du^n\|_1 - \|Du^{n+1}\|_1 - \alpha (\|\nabla u^n\|_2 - \|\nabla u^{n+1}\|_2) \\ &= \frac{\mu}{2} \|A(u^n - u^{n+1})\|_2^2 + (\|Du^n\|_1 - \langle p^{n+1}, u^n \rangle) + \alpha (\|\nabla u^{n+1}\|_2 - \langle q^n, \nabla u^{n+1} \rangle). \end{aligned}$$

The second equality above is obtained from left multiplying (2.13) by  $(u^n - u^{n+1})^T$ , and third one uses the fact that  $\langle p^{n+1}, u^{n+1} \rangle = \|Du^{n+1}\|_1$ .

The chain rule of subgradient [18] suggests that  $\partial \|Du\|_1 = D^T \partial \|Du\|_1$ , where

$$\partial |r|_1 = \begin{cases} [-1, 1] & r = 0, \\ \text{sign}(r) & \text{otherwise}. \end{cases}$$

It implies that  $\|Du^n\|_1 - \langle p^{n+1}, u^n \rangle = \|Du^n\|_1 - \langle p^{n+1}, Du^n \rangle \geq 0$ , since  $p^{n+1} \leq 1$  for each component. Using the definition of subgradient and the fact that  $q^n \in \partial \|\nabla u\|_2$ , we have  $\|\nabla u^{n+1}\|_2 - \langle q^n, \nabla u^{n+1} \rangle \geq 0$ , which concludes the proof. □

THEOREM 2.3. *Under the assumptions in Lemma 2.1, any non-zero limit point  $u^*$  of  $\{u^n\}$  satisfies the first-order optimality condition, which means  $u^*$  is a stationary point.*

*Proof.* It follows from Lemma 2.1 and Lemma 2.2 that the objective function  $F$  is coercive, monotonically decreasing, and convergent. As a result, the sequence  $\{u^n\}$  is bounded, and

$$\|A(u^n - u^{n+1})\|_2 \rightarrow 0, \quad (2.14)$$

$$\|\nabla u^{n+1}\|_2 - \langle q^n, \nabla u^{n+1} \rangle \rightarrow 0. \quad (2.15)$$

We start our algorithm from  $u^0 = \mathbf{0}$ , and  $u^1$  is obtained from

$$u^1 = \arg \min \|Du\|_1 + \frac{\mu}{2} \|Au - f\|_2^2.$$

If  $u^1$  is a constant vector, we stop our algorithm; otherwise  $F(u^1) \leq F(v)$  for any constant vector  $v$ . Since  $F$  monotonically decreases, any subsequent solution  $u^n$  for  $n \geq 1$  is not constant. Then we can properly define

$$c^n := \frac{\langle \nabla u^n, \nabla u^{n+1} \rangle}{\|\nabla u^n\|_2 \|\nabla u^{n+1}\|_2}.$$



Eq. (2.15) suggests that  $(1 - c^n)\|\nabla u^{n+1}\| \rightarrow 0$ . Therefore,  $c^n \rightarrow 1$ . It follows from the minimizing sequence that  $\|\nabla u^n\|_1 - \|\nabla u^{n+1}\|_1 \rightarrow 0$ , and so  $\nabla(u^n - u^{n+1}) \rightarrow 0$ . Combining (2.14) and  $\ker(A) \cap \ker(D) = \{\mathbf{0}\}$ , we get  $u^n - u^{n+1} \rightarrow 0$ .

This implies that there exists a subsequence of  $\{u^n\}$  converging to  $u^*$ , denoted as  $\{u^{n_k}\}$ . The optimality condition at the  $n_k$ -th step of DCA reads

$$\mathbf{0} \in \partial\|Du^{n_k}\|_1 + \alpha \nabla \cdot \frac{\nabla u^{n_k-1}}{\|\nabla u^{n_k-1}\|_2} + \mu A^T(Au^{n_k} - f), \quad (2.16)$$

or

$$-\alpha \nabla \cdot \frac{\nabla u^{n_k-1}}{\|\nabla u^{n_k-1}\|_2} - \mu A^T(Au^{n_k} - f) \in \partial\|Du^{n_k}\|_1. \quad (2.17)$$

We can show  $Du^{n_k}$  converges to  $Du^*$ , as

$$\|Du^{n_k} - Du^*\| \leq \|D\| \cdot \|u^{n_k} - u^*\| \rightarrow 0 \quad \text{as } n_k \rightarrow \infty.$$

When  $n_k$  is sufficiently large,  $\text{supp}(Du^*) \subseteq \text{supp}(Du^{n_k})$  and  $\text{sign}(Du^{n_k}) = \text{sign}(Du^*)$ . Using the chain rule of subgradient, we have  $\partial\|Du^{n_k}\|_1 \subseteq \partial\|Du^*\|_1$ , and  $D^T \partial\|Du^{n_k}\|_1 \subseteq D^T \partial\|Du^*\|_1$ . Consequently, it follows from (2.17) that

$$-\nabla \cdot \frac{\nabla u^{n_k-1}}{\|\nabla u^{n_k-1}\|} - \mu A^T(Au^{n_k} - f) \in D\|Du^*\|_1.$$

We assume  $\frac{\nabla u}{\|\nabla u\|} \doteq 0$  if  $\|\nabla u\| = 0$  at some points. Letting  $n_k \rightarrow \infty$ , we obtain

$$-\nabla \cdot \frac{\nabla u^*}{\|\nabla u^*\|} - \mu A^T(Au^* - f) \in D\|Du^*\|_1,$$

which means that  $u^*$  satisfies the first-order optimality condition.  $\square$

**3. Experiments.** We apply the proposed method to three applications: image denoising, deconvolution, and the MRI construction. The matrix  $A$  in these examples can be diagonalized by Fourier transform, and hence Algorithm 1 or Algorithm 2 can be efficiently implemented. We compare  $L_1$  and  $L_1 - \alpha L_2$  for  $\alpha = 0.5$  or  $1$  with some existing methods, such as  $L_0$  for image smoothing in [38],  $L_0$  in [31],  $L_p$  for  $p = 2/3$  in [21], and  $L_1 + L_2^2$  in [2] for image deblurring. We use structural similarity (SSIM) index [37] as a quantitative measure for image quality. Let us first define *local* similarity index computed on windows  $x$  and  $y$ ,

$$\text{ssim}(x, y) := \frac{(2\mu_x\mu_y + c_1)(2\sigma_{xy} + c_2)}{(\mu_x^2 + \mu_y^2 + c_1)(\sigma_x^2 + \sigma_y^2 + c_2)}, \quad (3.1)$$

where  $\mu_x, \mu_y$  are the average of  $x, y$ ,  $\sigma_x^2, \sigma_y^2$  are the variance,  $\sigma_{xy}$  is covariance of  $x, y$ , and  $c_1, c_2$  are two variables to stabilize the division with weak denominator. The overall SSIM is the mean of local similarity indexes, *i.e.*,

$$\text{SSIM}(X, Y) := \frac{1}{N} \sum_{i=1}^N \text{ssim}(x_i, y_i), \quad (3.2)$$

where  $X$  is a reference image,  $Y$  is a distorted one,  $x_i, y_i$  are corresponding windows indexed by  $i$ , and  $N$  is the number of windows. Here we consider windows of size  $8 \times 8$ .

**Image denoising.** We examine the problem of image denoising using an artificial piece-wise constant image in Figure 3.1 and a Lena image in Figure 3.2. We assume zero-mean additive Gaussian noise with standard deviations being 0.2 and 0.05 respectively. Not only does our method work

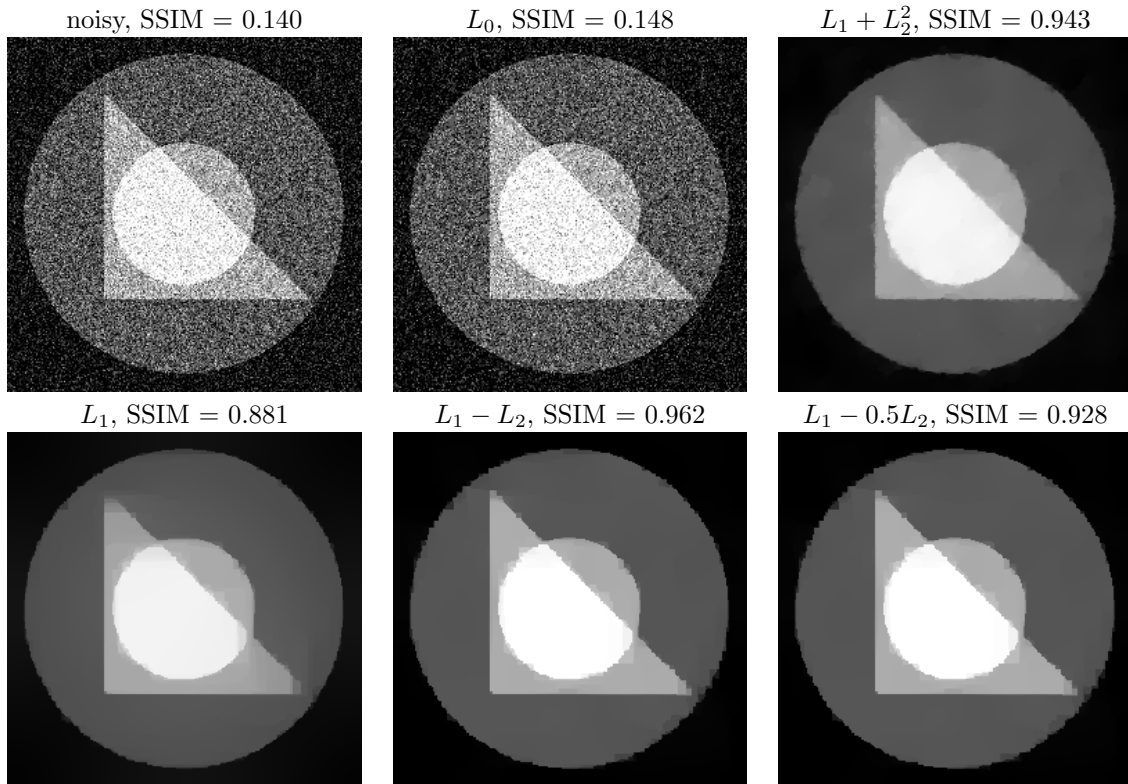


FIG. 3.1. Denoising results with comparison to  $L_0$  in [38] and  $L_1 + L_2^2$  in [2].

particularly well on horizontal or vertical edges by design, it can deal with natural images as well. To verify convergence analysis, the difference of  $u^n$  and  $u^{n-1}$  versus iterations is plotted in logarithm scale for both denoising examples shown in Figure 3.3, which shows that  $L_1 - 0.5L_2$  converges faster than  $L_1 - L_2$ . As the ground-truth is available, we plot the relative errors versus cpu runtime for  $L_1, L_1 - L_2, L_1 - 0.5L_2$  in Figure 3.4. This figure implies that our solutions oscillate around the ground truth due to the nonconvex nature of our model. Additionally we observe that the larger  $\alpha$  is (say approaching 1), the less well-behaved DCA becomes due to more weight on the nonconvex term. On the other hand,  $L_1 - L_2$  yields better results than  $L_1 - 0.5L_2$  for the first few DCA iterates. The denoising results presented in Figure 3.1 and Figure 3.2 are from stopping DCA after 2 iterations.

**Image deblurring.** In Figure 3.5, a binary image is vertically blurred by motion blur of 15 pixels plus Gaussian additive noise with zero mean and standard deviation 0.1. Our method outperforms  $L_0$  in [31],  $L_p$  for  $p = 2/3$  in [21],  $L_1 + L_2$  in [2], and the state-of-the-art deblurring method BM3D [11]. In Figure 3.6, we present deblurring results for a natural image: Cameraman. The original image is blurred by  $15 \times 15$  Gaussian blur whose standard deviation is 1.5 plus Gaussian additive noise with zero mean and standard deviation 0.05. Although  $L_0, L_{2/3}$  and BM3D are better than ours in terms of SSIM, their results have some ringing artifacts. In both deblurring examples, our method is better than the classical  $L_1$  approach. The relative errors versus computational time is plotted in Figure 3.7 for both examples. It shows similar behavior as in the denoising problem that  $L_1 - L_2$  tends to worsen beyond certain iterations while  $L_1 - 0.5L_2$  is more stable. The deblurring results presented in Figure 3.5 and Figure 3.6 are from stopping DCA after 2 and 10 iterations for  $L_1 - 0.5L_2$  and  $L_1 - L_2$  respectively. A discussion on stopping criterion is given later.

**MRI reconstruction.** In Figure 3.8, we investigate the MRI reconstruction problem using a Shepp-Logan phantom from 7 and 8 radial projections. There is no noise when we synthesize the data. Consequently we adopt the constrained formulation, *i.e.*, Algorithm 2 for solving eq.(2.9). Due to the

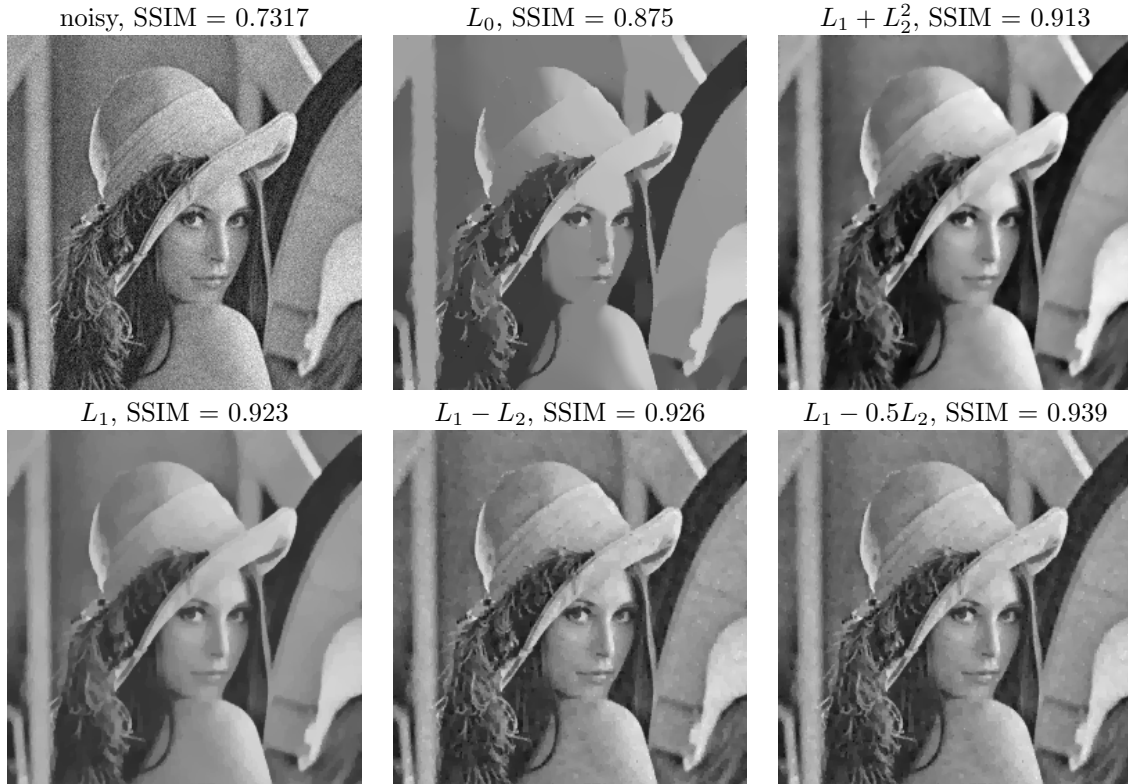


FIG. 3.2. Denoising results with comparison to  $L_0$  in [38] and  $L_1 + L_2^2$  in [2].

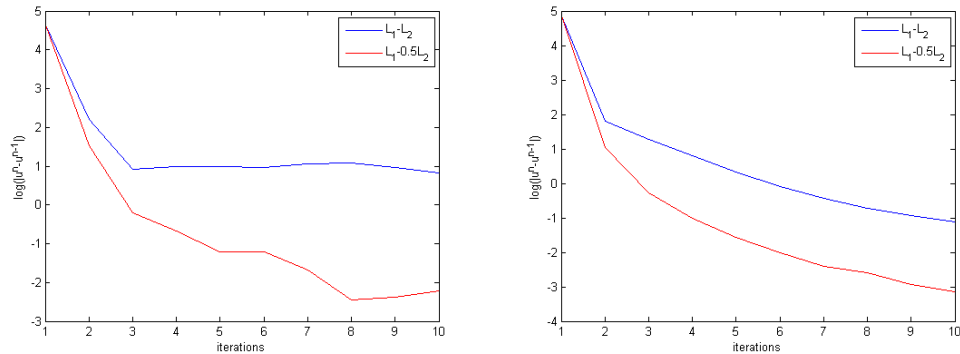


FIG. 3.3. The difference of  $u^n$  and  $u^{n-1}$  versus iterations is plotted in logarithm scale for denoising examples in Figure 3.1 (left) and Figure 3.2 (right).  $L_1 - 0.5L_2$  converges faster than  $L_1 - L_2$ .

presence of complex values in MRI reconstruction problem, SSIM is no longer applicable; instead we use root-mean-square (RMS) error to measure the performance quantitatively. RMS between reference and distorted images  $X, Y$  is defined as  $\text{RMS}(X, Y) = \frac{1}{\sqrt{M}} \|X - Y\|_2$  where  $M$  is the number of pixels in images  $X, Y$ . Figure 3.8 shows that our method can get a perfect reconstruction using only 8 projections, while a similar work [8] reports that 10 projections are required. When the number of projections is down to 7,  $L_1 - 0.5L_2$  is much better than  $L_1$  and  $L_1 - L_2$  visually as well as in terms of RMS. The relative errors versus cpu time is plotted in Figure 3.9. The relative errors of  $L_1 - L_2$  iterations in the constrained formulation appear as stable oscillations in contrast to the unstable

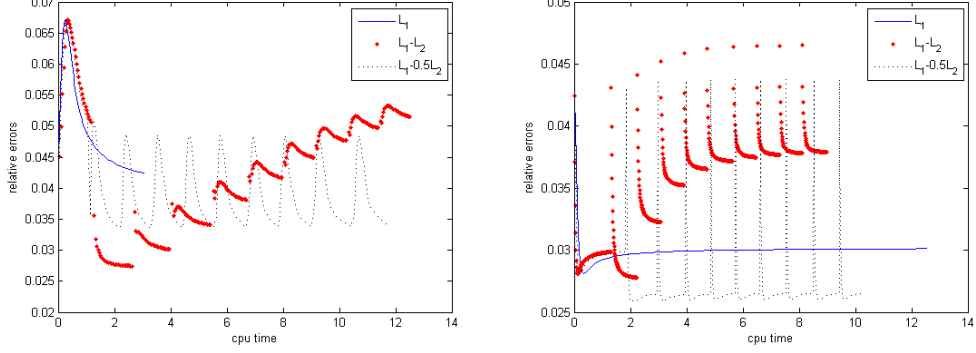


FIG. 3.4. The relative errors versus runtime for methods  $L_1$ ,  $L_1-L_2$ ,  $L_1-0.5L_2$  for denoising examples in Figure 3.1 (left) and Figure 3.2 (right). Our model solutions are seen to oscillate around the ground truth due to nonconvexity.

oscillations in the unconstrained problems.

**4. Discussions.** Let us draw some connections of this work to two existing methods, Lysaker-Osher-Tai (LOT) model [24] and Bregman iterations [29]. Additionally, we comment on the stopping criterion.

**4.1. Relation to existing methods.** At first, the iterative scheme (2.6) for  $\alpha = 1$  resembles the work of denoising the normals, proposed by Lysaker-Osher-Tai [24],

$$u^{n+1} = \operatorname{argmin}_u \|\nabla u\|_2 - q^n \cdot \nabla u + \frac{\mu}{2} \|Au - f\|_2^2, \quad (4.1)$$

where  $q^n = \frac{\nabla u^n}{|\nabla u^n|}$  is the surface normal. Notice that the TV norm in (4.1) is isotropic, while the first term in our model is the anisotropic TV; and hence  $L_1 - L_2$  applied to the gradient with linearized  $L_2$  term is different from the LOT.

On the other hand, the LOT model leads to the discovery of Bregman iterations [29], which relates to the DCA as well. Specifically, the Bregman distance [1] based on a convex functional  $J(\cdot)$  between two points  $u$  and  $v$  is defined as

$$D_J^p(u, v) := J(u) - J(v) - \langle p, u - v \rangle, \quad (4.2)$$

where  $p \in \partial J(v)$  is the subgradient of  $J$  at the point  $v$ . Osher *et. al.* [29] suggest an iterative refinement procedure to update  $u$  as follows,

$$u^{n+1} = \operatorname{argmin} D_J^{p^n}(u, u^n) + \frac{\mu}{2} \|Au - f\|_2^2, \quad (4.3)$$

$$= \operatorname{argmin} J(u) - \langle p^n, u \rangle + \frac{\mu}{2} \|Au - f\|_2^2, \quad (4.4)$$

which is referred to as the Bregman iterations. Let  $J(u) = \|\nabla u\|_2$  be the isotropic TV as in the LOT model, and its subgradient has the form  $-\nabla \cdot \frac{\nabla u}{|\nabla u|}$ . Consequently, we rewrite the second term in Eq. (4.4) as

$$\langle p^n, u \rangle = \langle -\nabla \cdot \frac{\nabla u^n}{|\nabla u^n|}, u \rangle = \langle \frac{\nabla u^n}{|\nabla u^n|}, \nabla u \rangle, \quad (4.5)$$

which coincides with the second term in the LOT model (4.1).

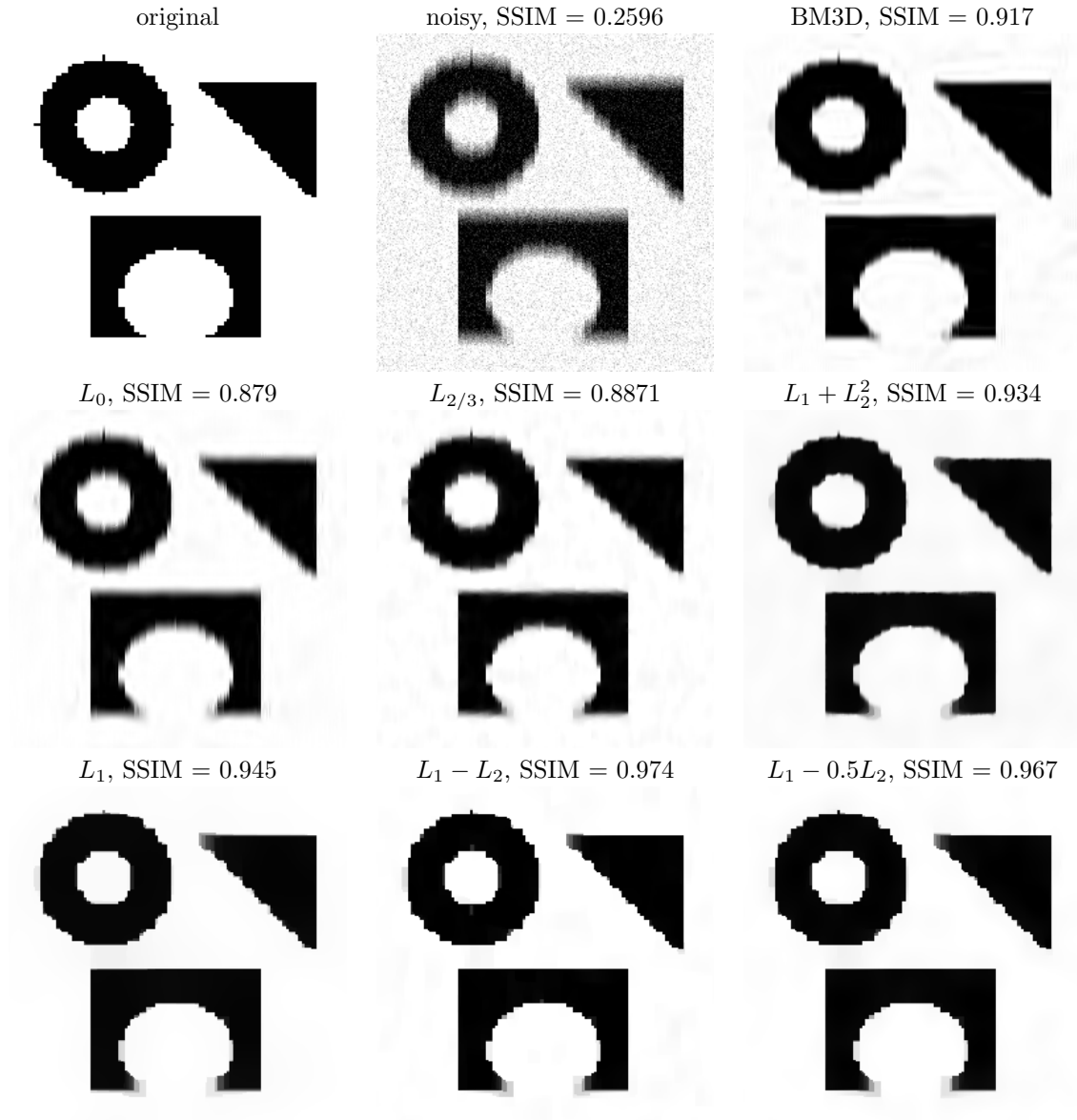


FIG. 3.5. Deblurring results with comparison to  $L_0$  in [31],  $L_p$  for  $p = 2/3$  in [21],  $L_1 + L_2^2$  in [2] and the state-of-the-art deblurring method BM3D [11].

Bregman iterations can be viewed as an optimization technique. Computing the optimality condition for each subproblem (4.4), we obtain

$$p^{n+1} - p^n + \mu A^T (A u^{n+1} - f) = 0. \quad (4.6)$$

Summing up to  $n + 1$ , we have  $p^{n+1} - \mu A^T (u^{n+1} - z^n)$  for  $p^0 = 0$  and  $z^{n+1} = z^n + (f - A u^n)$ . It is the optimality condition for solving  $u^{n+1}$  from  $\operatorname{argmin} J(u) + \frac{\mu}{2} \|A u - z^n\|_2^2$ . In short, the Bregman iterations can be rewritten as

$$u^{n+1} = \operatorname{argmin} J(u) + \frac{\mu}{2} \|A u - z^n\|_2^2, \quad (4.7)$$

$$z^{n+1} = z^n + (f - A u^n). \quad (4.8)$$



FIG. 3.6. Deblurring results with comparison to  $L_0$  in [31],  $L_p$  for  $p = 2/3$  in [21],  $L_1 + L_2^2$  in [2] and the state-of-the-art deblurring method BM3D [11].

The DCA for solving  $L_1 - L_2$  minimization can be derived from a similar way of the Bregman iterations. Let  $p$  and  $q$  be the subgradient of anisotropic  $J_{ani}$  and isotropic  $J_{iso}$  respectively. Lagging the isotropic term gives us

$$p^{n+1} - p^n - \alpha(q^n - q^{n-1}) + \mu A^T(Au^{n+1} - f) = 0. \quad (4.9)$$

We apply the same summation technique as in (4.6) and obtain

$$p^{n+1} - \alpha q^n + \mu A^T(Au^{n+1} - z^{n+1}) = 0, \quad (4.10)$$

$$z^{n+1} = z^n + (f - Au^n). \quad (4.11)$$

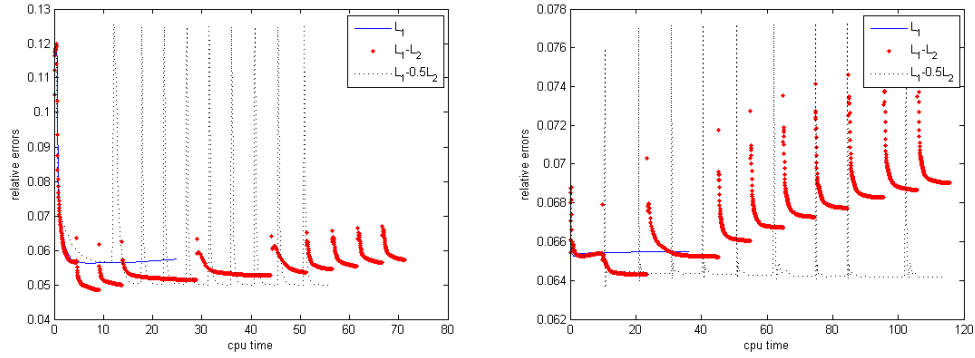


FIG. 3.7. The relative errors versus runtime for methods  $L_1, L_1 - L_2, L_1 - 0.5L_2$  for deblurring examples in Figure 3.5 (left) and Figure 3.6 (right).

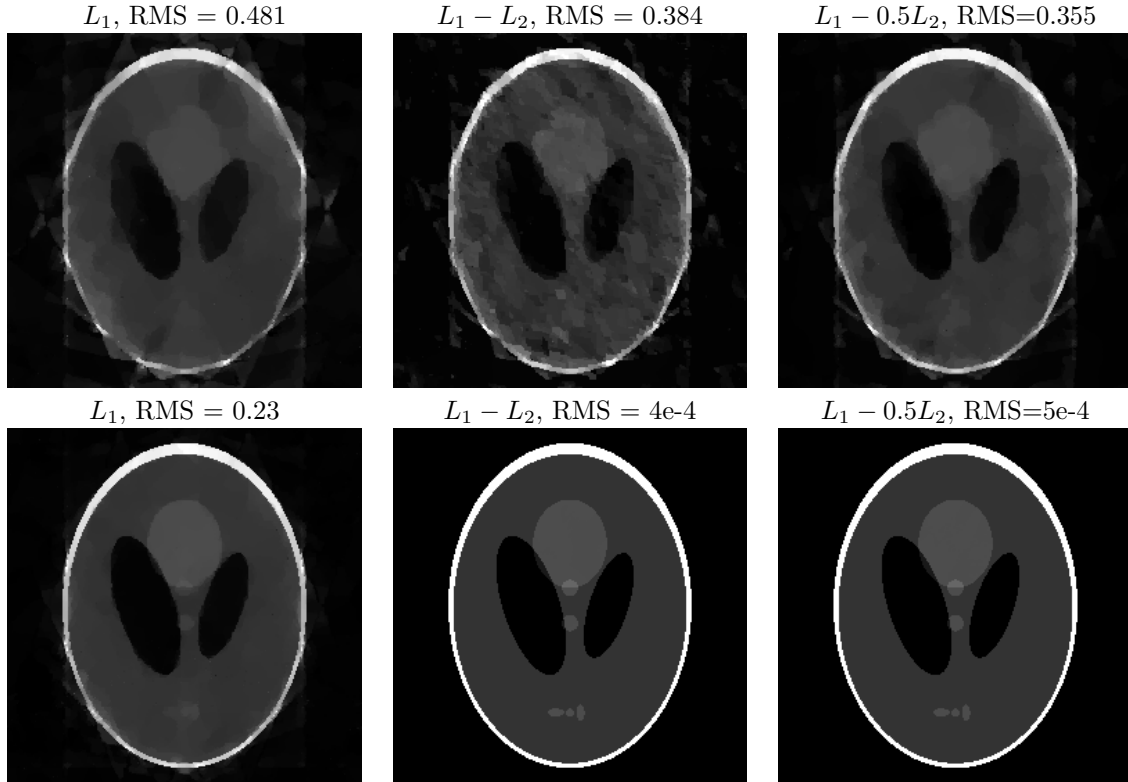


FIG. 3.8. MRI reconstruction using 7 (top) and 8 projections (bottom). The root-means-error (RMS) is provided for comparison.

for  $p^0 = q^0 = z^0 = 0$ . The subproblem (4.10) is equivalent to

$$u^{n+1} = \arg \min J_{ani}(u) - \alpha \langle q^n, u \rangle + \frac{\mu}{2} \|Au - z^n\|_2^2, \quad (4.12)$$

which looks very similar to applying the DCA for a constrained problem, eq. (2.5). The algorithm derived from the Bregman iterations is summarized in Algorithm 3. Its difference to Algorithm 2 lies in the update of  $z$  and  $q$ . For Algorithm 2,  $z$  is updated *MaxBregmanOuter* iterations and then  $q$  is updated, while Algorithm 3 is to update  $z$  and  $q$  simultaneously. The comparison between the

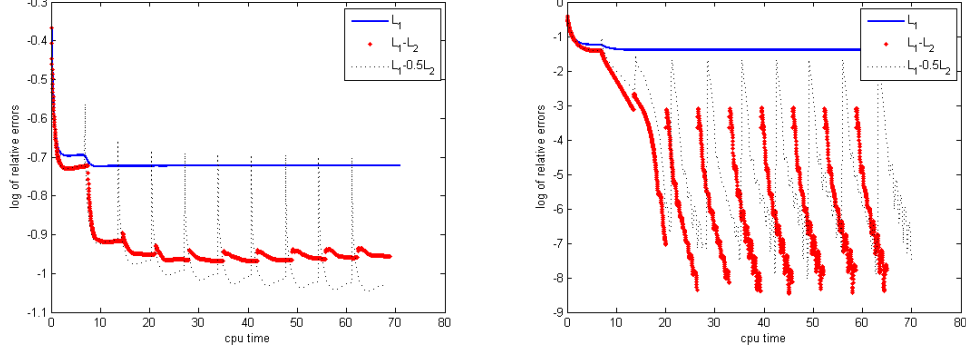


FIG. 3.9. The logarithm of relative errors versus runtime for methods  $L_1, L_1 - L_2, L_1 - 0.5L_2$  in MRI reconstruction problem using 7 (left) and 8 (right) projections. All are solved under constrained formulation.

Bregman and DCA iterations for solving such constrained nonconvex problems is a subject of further study.

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**Algorithm 3** for solving constrained problem (2.9) using Bregman method

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```

Define  $u = q_x = q_y = 0, z = f$  and MaxDCA, MaxBregman
for 1 to MAXDCA do
   $b_x = b_y = 0$ 
  for 1 to MaxBregman do
     $u = (\mu A^T A - \lambda \Delta)^{-1} (\mu A z + \lambda D_x^T (d_x - b_x) + \lambda D_y^T (d_y - b_y))$ 
     $d_x = \text{shrink}(D_x u + b_x + \alpha q_x / \lambda, 1 / \lambda)$ 
     $d_y = \text{shrink}(D_y u + b_y + \alpha q_y / \lambda, 1 / \lambda)$ 
     $b_x = b_x + D_x u + d_x$ 
     $b_y = b_y + D_y u + d_y$ 
  end for
   $z = z + f - Au$ 
   $(q_x, q_y) = (D_x u, D_y u) / \sqrt{|D_x u|^2 + |D_y u|^2}$ 
end for

```

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**4.2. Stopping criterion.** We discuss the stopping conditions of Algorithm 1 and Algorithm 2 for unconstrained and constrained problems respectively. Both algorithms have an outer DCA loop, which iteratively updates  $q$ , and inner iterations for updating  $u$ . We use  $u^n$  and  $u_k$  to specify the outer and inner outputs of  $u$ .

The inner loop is easier to impose a proper stopping criterion for, because the inner loop solves a convex subproblem. Some standard stopping criteria are either the relative error being small or objective function being stagnant or both *i.e.*,

$$\frac{\|u_{k+1} - u_k\|}{\|u_k\|} < \epsilon_u \quad \text{and/or} \quad \frac{|F(u_{k+1}) - F(u_k)|}{|F(u_k)|} < \epsilon_F \quad (4.13)$$

with pre-defined tolerance values  $\epsilon_u, \epsilon_F$ . In this paper, we choose to stop the inner iteration when the relative error is smaller than  $1e^{-6}$ .

As for the outer iterations, Figures 3.4, 3.7, and 3.9 show that the relative error develops an oscillatory pattern. One can estimate the onset time  $t_b$  of the oscillation stage of the error based on training images. In the denoising (deblurring) example,  $t_b = 2$  ( $= 10$ ). Hence, a good stopping time for the outer iteration is at the end of an inner loop when the cpu time exceeds  $t_b$ .



More generally, if the error does not follow a clear oscillatory pattern, one could inject random perturbations with slowly reduced magnitudes to steer away from unstable stationary points or directions to help convergence towards the ground truth [17]. This approach is closely related to simulated annealing [14, 20].

**5. Conclusion.** We proposed a weighted difference of anisotropic and isotropic total variation as a regularization term for image processing applications. We presented a difference of convex algorithm (DCA) for both the constrained and unconstrained formulations. We proved the convergence of the algorithm to ensure that each limiting point is a stationary point and the values of the objective function monotonically decrease. The behavior of the iterations was observed numerically to be oscillatory around the ground truth. The deviation occurs at the beginning of outer loops of DCA. A stopping criterion was introduced based on such oscillatory pattern of the errors.

In the numerical experiments, we examined three particular applications: image denoising, deblurring and MRI reconstruction. By design, our method works particularly well for piecewise constant images. For natural images, it improved the classical TV model, and is comparable to the state-of-the-art methods. In future work, we plan to carry out a detailed comparison between the DCA and Bregman methods, and further study the error pattern and the resulting stopping criterion for other imaging science problems.

**Acknowledgments.** YL would like to thank Dr. Ernie Esser at University of British Columbia for helpful discussions. JX would like to thank Profs. Krishna Nayak and Angel Pineda for their hospitality during a visit to USC in March 2014, and their suggestion to consider a weighted variant of  $L_1 - L_2$  for compressed sensing and the SSIM measure for image quality evaluation.

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