

Performance of Matching Algorithms for Signal Approximation

¹U.L Jyothirmayee, ²K.V.V.Kumar
¹M.Tech, Student, ²Asst. Professor

^{1,2}Rise Krishna Sai Prakasam Group of Institutions

Abstract: One of the important classes of sparse signals is the non-negative signals. Many algorithms have already been proposed to recover such non-negative representations, where various and convex relaxed algorithms are among the most popular methods. The conventional techniques have been changed to incorporate the non-negativity of the depictions. One such modification has been proposed for Extrinsic Equivalent Detection (EED), which first chooses positive constants and uses a non-negative optimization technique as a replacement for the orthogonal projection onto the nominated support. Which gradually builds the sparse representation of a signal by iteratively adding the most correlated element of the lexicon, which is called an atom, to the set of selected elements. A disadvantage of ED is that the depiction found by the algorithm is not the best representation using selected atoms. It may also reselect already selected atoms in the later iterations, which slows down the junction of the algorithm. As a result, we present a novel fast implementation of the Nonnegative EED, which is based on the QR decomposition and an iterative constants apprise. We will empirically show that such a modification can easily accelerate the implementation by a factor of ten in a reasonable size problem. We explain how the non-negativity constraint of the coefficients stops us of using the canonical EED and how we can modify the algorithm to not only have a more intuitive atom selection step, but have a lower computational complexity.

Keywords: Detection, decomposition, iteration, non-negative

I. Introduction

The phrase compressed sensing refers to the problem of realizing a sparse input using few linear measurements that possess some incoherence properties. The field originated recently from an unfavorable opinion about the current signal compression methodology. The conventional scheme in signal processing, acquiring the entire signal and then compressing it, was questioned by Donoho [2]. Indeed, this technique uses tremendous resources to acquire often very large signals, just to throw away information during compression. The natural question then is whether we can combine these two processes, and directly sense the signal or its essential parts using few linear measurements. Recent work in compressed sensing has answered this question in positive, and the field continues to rapidly produce encouraging results.

The key objective in compressed sensing (also referred to as sparse signal recovery or compressive sampling) is to reconstruct a signal accurately and efficiently from a set of few non-adaptive linear measurements. Signals in this context are vectors, many of which in the applications will represent images. Of course, linear algebra easily shows that in general it is not possible to reconstruct an arbitrary signal from an incomplete set of linear measurements. Thus one must restrict the domain in which the signals belong. To this end, we consider sparse signals, those with few non-zero coordinates. It is now known that many signals such as real-world images or audio signals are sparse either in this sense, or with respect to a different basis. Since sparse signals lie in a lower dimensional space, one would think intuitively that they may be represented by few linear measurements.

This is indeed correct, but the difficulty is determining in which lower dimensional subspace such a signal lies. Multicarrier modulation has regained interest over the last decade. Several all-digital variants have been proposed: discrete multitone (DMT) is adopted as transmission format for asymmetric digital subscriber line (ADSL) and presented as a candidate for very high bit rate digital subscriber line (VDSL); orthogonal frequency division multiplexing (OFDM) is proposed for wireless local area applications, e.g. HiperLAN. DMT schemes divide the bandwidth into parallel subbands or tones. The incoming bitstream is split into parallel streams that are used to QAM-modulate the different tones. The modulation is done by means of an inverse fast Fourier transform (IFFT). Before transmission of a DMT symbol, a cyclic prefix of samples is added. If the channel impulse response order is less than or equal to the cyclic prefix length v , demodulation can be implemented by means of an FFT, followed by a (complex) 1-tap frequency domain equalizer (FEQ) per tone to compensate for channel amplitude and phase effects. The challenges of building a classifier that can distinguish between high-dimensional members of various classes based on their shape differences, involves devising a reliable dissimilarity measure that can perform shape-based comparisons of very high-dimensional signals.

Also, the automation of the feature selection process to minimize human intervention and reliance on domain knowledge. Finally, a robust prototype-based classifier that can detect outliers in test data. In order to achieve the above objectives, a Matching Pursuits Dissimilarity Measure is presented. The EDDM extends the well-known signal approximation technique Equivalent Detections (ED) for signal comparison purposes [1]. MP is a greedy algorithm that approximates a signal x as a linear combination of signals from a pre-defined dictionary. MP is commonly used for signal representation and compression, particularly image and video compression [5, 6]. The dictionary and coefficients information produced by the ED algorithm has been previously used in some classification applications. However, most of these applications work on some underlying assumptions about the data and the MP dictionary (section 2.3). The EDDM is the first MP based comparison measure that does not require any assumptions about the problem domain. It is versatile enough to perform shape-based comparisons of very high-dimensional signals and it can also be adopted to perform magnitude-based comparisons, similar to the Euclidean Distance. Since the EDDM is a differentiable measure, it can be seamlessly used with existing clustering or discrimination algorithms.

Therefore, the EDDM may find application in a variety of classification and approximation problems of very high-dimensional signals, including image and video signals. The experimental results show that EDDM is more useful than the Euclidean distance for shape-based comparison between signals in high dimensions.

The potential usefulness of the MPDM for a variety of problems is demonstrated by devising two important EDDM-based algorithms. The first algorithm, called CAMP, deals with the prototype-based classification of high-dimensional signals. The second algorithm is called the EK-SVD algorithm and it automates the dictionary learning process for the MP approximation of signals. In the CAMP algorithm, EDDM is used with the Competitive Agglomeration (CA) clustering algorithm by Frigui and Krishnapuram to propose a probabilistic classification model [2]. The CA algorithm is a fuzzy clustering algorithm that learns the optimal number of clusters during training. Therefore, it eliminates the need for manually specifying the number of clusters beforehand. This algorithm has been named as CAMP as an abbreviation of CA and ED algorithms.

II. Non-Negative Least Squares Algorithm

Let A be an $m \times n$ matrix and b be a vector of dimension m . Consider the following feasibility problem:

$$Ax = b \quad (1)$$

$$x \geq 0 \quad (2)$$

A straightforward way of solving the above problem through linear programming is by solving the following LP problem:

$$(LP) : \min \sum_{j=1}^n |s_j|$$

$$Ax + s = bx \geq 0$$

Observe that this norm 1 minimization can be carried out very efficiently by the simplex in most cases. However, there are some constraint matrices for which the Simplex method performs a large number of degenerate pivots, not improving the solution for many iterations, leading to a poor performance. Our approach to solve the feasibility problem posed by relations 1 and 2 will also be the minimization of a p -norm, but different from the norm 1 considered in (LP), we will consider the norm 2, i.e., we will solve the following problem:

At first glance, problem (PLS) seems much harder than problem (LP). However, in cases where (LP) is highly degenerate (as pointed out before), it is usually simpler to solve (PLS). E. Barnes et al. in [6] showed that the normalized direction obtained by (PLS) is the direction of the steepest ascent at $\pi/2$ on the dual polyhedron (D). This suggests that the dual direction obtained by (PLS) may be much better in practice than the one obtained by the linear update in (LP). E. Barnes et al. in [6] showed empirically that this is indeed true for some classes of problems. Since (PLS) is a convex program, KKT conditions are necessary and sufficient for optimality. Thus, the vector (x, s) is a solution for P if and only if there exists π such that:

The NNLS algorithm starts with a primal feasible solution, i.e., one that is feasible for (PLS), and tries to find a solution for the problem (DLS). The NNLS algorithm is similar to the simplex method in the sense that we have a subset of the columns of A that is a primal feasible basis, and then we move from one primal feasible basis to another. Unlike the simplex method, our 'basis' is not required to be square. The only requirement is that it is composed of linearly independent columns. Let B be a basis, i.e., a linearly independent subset of the columns of A . Then one crucial step of the nonnegative least squares algorithm is to solve the following problem:

$$\min \|Bx - b\|_2$$

Since the columns of B are linearly independent, the solution will be:

$$x = B^+b, \text{ where } B^+ = (B^T B)^{-1} B^T$$

The matrix B^+ is called the generalized inverse or pseudo inverse. If B is a basis, we say that it is feasible for (PLS) if we have:

$$x = B^+b > 0$$

Theorem 1. Algorithm 1 terminates with a solution of problem (PLS).

1. Let B be the feasible basis for problem P
2. Let I_B be the index set of the columns in B
3. $\bar{x} \leftarrow B^+ b$
4. $\bar{\pi} \leftarrow b - B \bar{x}$
5. $S \leftarrow \{j : A_j > 0\}$
6. if $S = \emptyset$ the
7. stop: optimal solution found
8. end if
9. Let $k \in S$
10. $d \leftarrow B^+ A_k$
11. $\theta \leftarrow \min_{d_j > 0} \frac{x_j}{d_j}$
12. $P \leftarrow I - BB^+$
13. $\bar{\theta} \leftarrow \frac{\bar{\pi}^t A_j}{\|PA_j\|^2}$
14. $\theta \leftarrow \min\{\theta, \bar{\theta}\}$
15. if $\theta = \bar{\theta}$ then
16. $I_B \leftarrow I_B \cup \{j\}$
17. if $\theta = \bar{\theta}$ then
18. $\bar{x}(\theta) \leftarrow \bar{x} - \theta d$
19. $I_B \leftarrow I_B - \{j : \bar{x}(\theta) = 0\}$
20. end if
21. $B \leftarrow [A_j] \forall j \in I_B$
22. Return to 3
23. else
24. $I_B \leftarrow I_B \left\{ J : \theta = \frac{x_j}{d_j} \right\}$
25. $B \leftarrow [A_j] \forall j \in I_B$
26. $\bar{x} \leftarrow B^+ b$
27. Return to 10
28. end if

Proof. We will show that the algorithm terminates by showing that no basis can be repeated. Since the number of basis is finite, the result follows. In order to prove that no basis can be repeated, we will show that, if a basis B is updated to B^* , then we must have:

$$\min_{x \geq 0} \|\widehat{B}x - b\|^2 < \min_{x \geq 0} \|Bx - b\|^2$$

Let us suppose first that $\theta = \bar{\theta} \leq \theta$

Let B be the current basis and A_j be the entering column. If x is the current primal solution, then

$$x = (B^+ b) - 1 B^+ b$$

$$\min_{x \geq 0} \|\hat{B}x - b\|^2 = \min_{x, t \geq 0} \|Bx + A_j t - b\|^2 \leq \min_{x \geq 0} \|Bx - (b - A_j \theta)\|^2$$

Let \hat{B} be the new basis. Then the solution of the last minimization problem is:

$$\bar{x} = (B^T \hat{B})^{-1} B^T (b - A_j \theta) = \bar{x} - \theta d$$

III. Extranious Equivalent Detection

One such greedy algorithm is Orthogonal Matching Pursuit (OMP), put forth by Mallat and his collaborators (see e.g. [47]) and analyzed by Gilbert and Tropp [62]. OMP uses sub-Gaussian measurement matrices to reconstruct sparse signals. If Φ is such a measurement matrix, then $\Phi^* \Phi$ is in a loose sense close to the identity. Therefore one would expect the largest coordinate of the observation vector $y = \Phi^* \Phi x$ to correspond to a non-zero entry of x . Thus one coordinate for the support of the signal x is estimated. Subtracting off that contribution from the observation vector and repeating eventually yields the entire support of the signal x . OMP is quite fast, both in theory and in practice, but its guarantees are not as strong as those of Basis Pursuit.

The OMP algorithm can thus be described as follows:

Orthogonal Matching Pursuit (OMP)

Input: Measurement matrix Φ , measurement vector $u = \Phi x$, sparsity level s

Output: Index set $I \subset \{1, \dots, d\}$

Procedure:

Initialize Let the index set $I = \emptyset$ and the residual $r = u$.

Repeat the following s times:

Identify Select the largest coordinate λ of $y = \Phi^* r$ in absolute value. Break ties lexicographically.

Update Add the coordinate λ to the index set: $I \leftarrow I \cup \{\lambda\}$, and update the residual:

$x^{\wedge} = \operatorname{argmin}_z$

$\|u - \Phi|_I z\|_2$; $r = u - \Phi^{\wedge} x$.

Once the support I of the signal x is found, the estimate can be reconstructed as $x^{\wedge} = \Phi^{\dagger} I u$, where recall we define the pseudoinverse by $\Phi^{\dagger} I \stackrel{\text{def}}{=} (\Phi^* I \Phi I)^{-1} \Phi^* I$. The algorithm's simplicity enables a fast runtime. The algorithm iterates s times, and each iteration does a selection through d elements, multiplies by Φ^* , and solves a least squares problem. The selection can easily be done in $O(d)$ time, and the multiplication of Φ^* in the general case takes $O(md)$. When Φ is an unstructured matrix, the cost of solving the least squares problem is $O(s^2 d)$. However, maintaining a QR-Factorization of $\Phi|_I$ and using the modified Gram-Schmidt algorithm reduces this time to $O(|I| d)$ at each iteration. Using this method, the overall cost of OMP becomes $O(sm d)$. In the case where the measurement matrix Φ is structured with a fast-multiply, this can clearly be improved.

IV. Stagewise Extranious Equivalent Detection

An alternative greedy approach, Stagewise Orthogonal Matching Pursuit (StOMP) developed and analyzed by Donoho and his collaborators [23], uses ideas inspired by wireless communications. As in OMP, StOMP utilizes the observation vector $y = \Phi^* u$ where $u = \Phi x$ is the measurement vector. However, instead of simply selecting the largest component of the vector y , it selects all of the coordinates whose values are above a specified threshold. It then solves a least-squares problem to update the residual. The algorithm iterates through only a fixed number of stages and then terminates, whereas OMP requires s iterations where s is the sparsity level.

The pseudo-code for StOMP can thus be described by the following.

Input: Measurement matrix Φ , measurement vector $u = \Phi x$,

Output: Estimate \hat{x} to the signal x

Procedure: Initialize Let the index set $I = \emptyset$, the estimate $\hat{x} = 0$, and the residual $r = u$. Repeat the following until stopping condition holds:

Identify Using the observation vector $y = \Phi^* r$, set $J = \{j : |y_j| > t_k \sigma_k\}$, where σ_k is a formal noise level and t_k is a threshold parameter for iteration k .

Update Add the set J to the index set: $I \leftarrow I \cup J$, and update the residual and estimate: $x^{\wedge} | I = (\Phi^* I \Phi I)^{-1} \Phi^* I u$, $r = u - \Phi^{\wedge} x$.

The thresholding strategy is designed so that many terms enter at each stage, and so that algorithm halts after a fixed number of iterations. The formal noise level σ_k is proportional the Euclidean norm of the residual at that iteration.

Syphilis antigen in a bloodsample. Since this test was expensive, the method was to sample a group of mentogether and test the entire pool of blood samples. If the pool did not contain theantigen, then one test replaced many. If it was found, then the process could eitherbe repeated with that group, or each individual in the group could then be tested.These sublinear algorithms in compressed sensing use this same idea to test for elements of the support of the signal x . Chaining pursuit, for example, uses a measurement matrixconsisting of a row tensor product of a bit test matrix and anisolation matrix, both of which are 0-1 matrices. Chaining pursuit first uses bit teststo locate the positions of the large components of the signal x and estimate thosevalues. Then the algorithm retains a portion of the coordinates that are largestmagnitude and repeats. In the end, those coordinates which appeared throughouta large portion of the iterations are kept, and the signal is estimated using these.Pseudo-code is available in [3], where the following result is proved.

Theorem (Chaining pursuit [31]). With probability at least $1 - O(d^{-3})$, the $O(s \log^2 d) \times d$ random measurement operator Φ has the following property. For $x \in \mathbb{R}^d$ and its measurements $u = \Phi x$, the Chaining Pursuit algorithm produces a signal \hat{x} with at most s nonzero entries. The output \hat{x} satisfies

$$\|x - \hat{x}\|_1 \leq C(1 + \log s)\|x - x_s\|_1.$$

The time cost of the algorithm is $O(s \log^2 s \log^2 d)$.HHS Pursuit, a similar algorithm but with improved guarantees, uses a measurement matrix that consists again of two parts. The first part is an identificationmatrix, and the second is an estimation matrix. As the names suggest, the identification matrix is used to identify the location of the large components of the signal,whereas the estimation matrix is used to estimate the values at those locations. Eachof these matrices consist of smaller parts, some deterministic and some random. Using this measurement matrix to locate large components and estimate their values, HHS Pursuit then adds the new estimate to the previous, and prunes it relative tothe sparsity level.

This estimation is itself then sampled, and the residual of thesignal is updated. Let $x \in \mathbb{R}^d$ and let $u = \Phi x$ be the measurement vector.The HHS Pursuit algorithm produces a signal approximation \hat{x} with $O(s/\epsilon^2)$ nonzeroentries. $\|x - \hat{x}\|_2 \leq \sqrt{s} \|x - x_s\|_1$, where again x_s denotes the vector consisting of the s largest entries in magnitudeof x . The number of measurements m is proportional to $(s/\epsilon^2) \text{polylog}(d/\epsilon)$, andHHS Pursuit runs in time $(s^2/\epsilon^4) \text{polylog}(d/\epsilon)$. The algorithm uses working space $(s/\epsilon^2) \text{polylog}(d/\epsilon)$, including storage of the matrix Φ .

There are other algorithms such as the Sudocodes algorithm that as of now onlywork in the noiseless, strictly sparse case. However, these are still interesting becauseof the simplicity of the algorithm. The Sudocodes algorithm is a simple two-phasealgorithm. In the first phase, an easily implemented avalanche bit testing schemes applied iteratively to recover most of the coordinates of the signal x . At thispoint, it remains to reconstruct an extremely low dimensional signal (one whosecoordinates are only those that remain). In the second phase, this part of the signalis reconstructed, which completes the reconstruction. Since the recovery is twophase, the measurement matrix is as well. For the first phase, it must contain asparse submatrix, one consisting of many zeros and few ones in each row. For thesecond phase, it also contains a matrix whose small submatrices are invertible. Thefollowing result for strictly sparse signals.Combinatorial algorithms such as HHS pursuit provide sublinear timerecovery withoptimal error bounds and optimal number of measurements. Some of these arestraightforward and easy to implement, and others require complicated structures.The majordisadvantage however is the structural requirement on the measurementmatrices. Not only do these methods only work with one particular kind of measurement matrix, but that matrix is highly structured which limits its use in practice.There are no known sublinear methods in compressed sensing that allow for unstructured or generic measurement matrices

V. Outputs

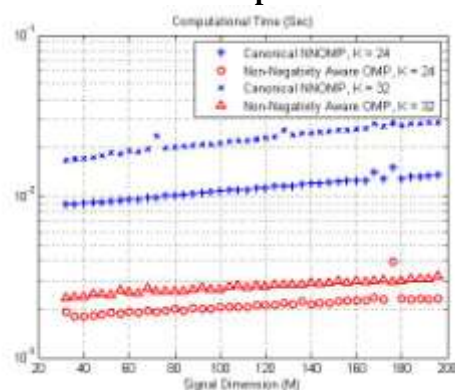


Fig 1 Computation time for the fixed $N = 256$ and $K = 24 \ \& \ 32$

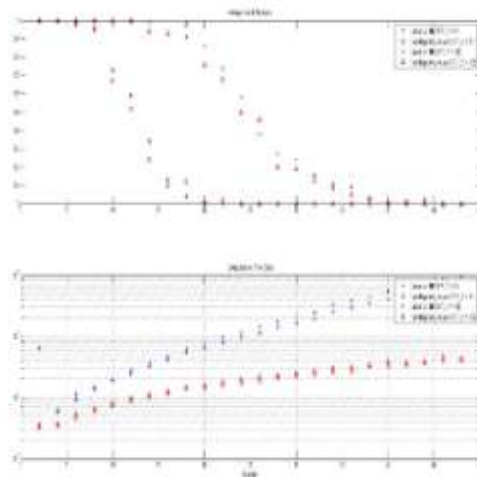


Fig 2: Average exact recovery and Computational time(sec)

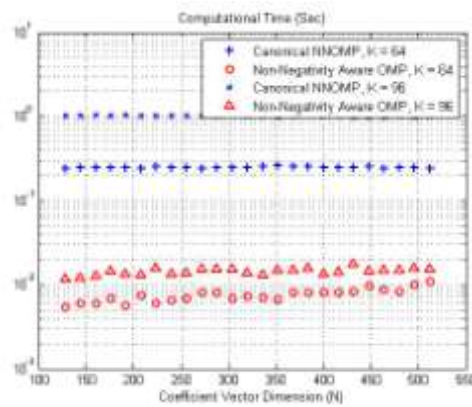


Fig. 3. Computation time for the fixed $M = 128$ and $K = 64$ & 96

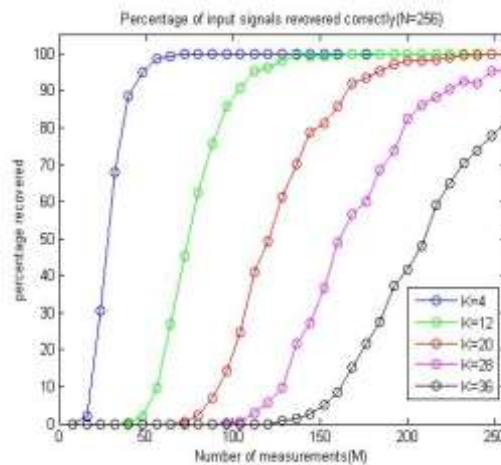


Fig 4: Percentage of recovered signals



Fig 5: Sparse AOMP data

VI. Conclusion

A matching pursuits dissimilarity measure has been presented, which is capable of performing accurate shape-based comparisons between high-dimensional data. It extends the matching pursuits signal approximation technique and uses its dictionary and coefficient information to compare two signals. AOMP is capable of performing shape-based comparisons of very high dimensional data and it can also be adapted to perform magnitude-based comparisons, similar to the Euclidean distance. Since AOMP is a differentiable measure, it can be seamlessly integrated with existing clustering or discrimination algorithms. Therefore, AOMP may find application in a variety of classification and approximation problems of very high dimensional data. The AOMP is used to develop an automated dictionary learning algorithm for MP approximation of signals, called Enhanced K-SVD. The EK-SVD algorithm uses the AOMP and the CA clustering algorithm to learn the required number of dictionary elements during training. Under-utilized and replicated dictionary elements are gradually pruned to produce a compact dictionary, without compromising its approximation capabilities. The experimental results show that the size of the dictionary learned by our method is 60% smaller but with same approximation capabilities as the existing dictionary learning algorithms. The AOMP is also used with the competitive agglomeration fuzzy clustering algorithm to build a prototype-based classifier called AMP. The AMP algorithm builds robust shape-based prototypes for each class and assigns a confidence to a test pattern based on its dissimilarity to the prototypes of all classes. If a test pattern is different from all the prototypes, it will be assigned a low confidence value. Therefore, our experimental results show that the CAMP algorithm is able to identify outliers in the given test data better than discrimination-based classifiers, like, multilayer and support vector machines. We presented a new greedy technique based on OMP, suitable for non-negative sparse representation, which is much faster than the state of the art algorithm. The new algorithm has a slightly different atom selection procedure, which guarantees the non-negativity of the signal approximations. Although this selection step is more involved, the overall algorithm has a much faster implementation. The reason is that with the new selection procedure, we can use fast QR implementation of the OMP. The computational complexity of two NNOMP's were derived and the differences were demonstrated.

The experimental results show that the size of the dictionary learned by our method is 60% smaller but with same approximation capabilities as the existing dictionary learning algorithms.

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