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Segmentation Given Partial Grouping Constraints

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Segmentation Given Partial Grouping Constraints

Stella X. Yu, Member, IEEE Computer Society, and Jianbo Shi, Member, IEEE Computer Society

Abstract—We consider data clustering problems where partial grouping is known a priori. We formulate such biased grouping problems as a constrained optimization problem, where structural properties of the data define the goodness of a grouping and partial grouping cues define the feasibility of a grouping. We enforce grouping smoothness and fairness on labeled data points so that sparse partial grouping information can be effectively propagated to the unlabeled data. Considering the normalized cuts criterion in particular, our formulation leads to a constrained eigenvalue problem. By generalizing the Rayleigh-Ritz theorem to projected matrices, we find the global optimum in the relaxed continuous domain by eigendecomposition, from which a near-global optimum to the discrete labeling problem can be obtained effectively. We apply our method to real image segmentation problems, where partial grouping priors can often be derived based on a crude spatial attentional map that binds places with common salient features or focuses on expected object locations. We demonstrate not only that it is possible to integrate both image structures and priors in a single grouping process, but also that objects can be segregated from the background without specific object knowledge.

Index Terms—Grouping, image segmentation, graph partitioning, bias, spatial attention, semisupervised clustering, partially labeled classification.

1 INTRODUCTION

A good image segmentation respects not only the structural properties of the image [1] but also the needs of later visual processing such as object recognition [2]. In this paper, we will develop a method that integrates both data-driven and task-driven knowledge for making a global decision on segmentation.

The approach where task-driven knowledge is used to constrain the segmentation at the very beginning contrasts with the sequential processing theory popularized by Marr [3]. According to his theory, visual processing starts with what can be computed directly from an image and ends with the information required to support goals such as navigation or object recognition. Intermediate representations are derived to turn the available information at one level to the required information at the succeeding level. Accordingly, most current image segmentation algorithms adopt a bottom-up approach. They start with an oversegmentation based on low-level cues such as feature similarity and boundary continuity and then build up larger perceptual units (e.g., surface, foreground, and background) by adding high-level knowledge such as statistical properties of regions into the grouping process [4].

Although a sequential system can relieve computational burden from later stages of perceptual processing, such a feed-forward system is vulnerable to mistakes made at each step: the low-level processing alone often produces an unreliable representation, e.g., missing object boundaries of weak contrast caused by lighting and background clutter, which may not be remediable by the later high-level processing.

We demonstrate that it is possible to integrate both bottom-up and top-down information in a single grouping process. We consider the type of task-driven knowledge presented as partial grouping information. For example, in Fig. 1, based on intensity distribution and viewers’ expectation, for the image with the tiger, a set of bright pixels are likely to be foreground and a set of dark pixels are likely to be background; for the image with the fashion-model, pixels near image boundaries are probably background. Such information provides bias to a natural grouping process that is based solely on data themselves.

Our work is concerned with the following issue: What is a simple and principled approach for incorporating these often sparse partial grouping cues directly into low-level image segmentation?

A straightforward approach that we adopt in this work is to formulate the problem as a constrained optimization problem, where the goodness of a segmentation is based on low-level data coherence and the feasibility of a segmentation is based on partial grouping constraints. For the normalized cuts criterion under the spectral graph-theoretic framework [5], we show that this straightforward formulation leads to a constrained eigenvalue problem. By generalizing the standard Rayleigh-Ritz theorem, we can compute a near-global optimum efficiently.

We then show through a simple point set example that segmentation performance breaks down especially when partial grouping cues are sparse. This observation leads to a new formulation with smoothed constraints. In the spectral graph framework, the smoothing operator is readily derived from the existing pairwise relationships between grouping elements. We present numerous image segmentation examples to demonstrate the efficacy of the new formulation.
they are not required to be different across the \( n \) groups. For a unique representation of \( U_t \)'s, we assume there is no common pixel between any two sets: \( U_t \cap U_s = \emptyset, s \neq t \). In other words, if there is a common pixel, then the two sets should be merged into one.

The most straightforward way to incorporate the partial grouping information is to encode it as constraints. With a little abuse of notation, we use \( \Gamma^K_V(i, l) \) to denote a Boolean function that returns 1 if \( i \in V_l \). Among the segmentations partially determined by \( U_t \)'s, we seek one that optimizes the goodness of grouping measured by \( \varepsilon \):

\[
\begin{align*}
\text{maximize} & \quad \varepsilon(\Gamma^K_V; f) \\
\text{subject to} & \quad \Gamma^K_V(i, l) = \Gamma^K_V(j, l), i, j \in U_t, l \in [K], t \in [n].
\end{align*}
\]

Since partial grouping cues are encoded as hard constraints, they have to be reliable enough to be enforced. Fig. 1 illustrates two basic scenarios where we can derive such cues. The first type is feature-driven, where pixels conforming to a particular generative model are biased together. For example, we probably perceive a white object against a dark background before we realize that it is a tiger in a river. In this case, \( U_1 \) contains pixels of the brightest intensities and \( U_2 \) the darkest. The second type is solely location-driven, it reflects our expectation as to where an object is going to appear. For example, pictures taken in a fashion show often have fashion models at the center. To segment out the fashion models, we consider pixels at image boundaries as the background group \( U_1 \). Such seemingly insignificant information provides long-range binding cues that are often lacking in low-level grouping.

For some particular forms of \( \varepsilon \), such as the above mentioned probability criteria using generative models and the minimum cuts criteria in discriminative approaches [7], [8], [9], the constraints in (3) can be trivially incorporated in an algorithm that optimizes the objective. For the former, Markov Chain Monte Carlo (MCMC) is a general solution technique and the constraints can be realized by generating legitimate samples [10]. For the latter, assuming that \( U_1 \) and \( U_2 \) take distinct labels, we can solve (3) using maximum-flow algorithms, in which two special nodes called source and sink are introduced, with infinite weights between the source and \( U_1 \), and between the sink and \( U_2 \) [7]. For others such as the normalized cuts criterion [5], it is not clear whether the solution can be obtained using the same technique that was used for the unconstrained problem. We will explore this criterion further.

### 3 Constrained Normalized Cuts Criterion

A weighted graph is specified by \( G = (V, E, W) \), where \( V \) is the set of all nodes, \( E \) is the set of edges connecting nodes, and \( W \) is an affinity matrix, with weights characterizing the likelihood that two nodes belong to the same group. We assume that \( W \) is nonnegative and symmetric.

In graph-theoretic methods for image segmentation, an image is first transcribed into a weighted graph, where each node represents a pixel and weights on edges connecting two nodes describe the pairwise feature similarity between the pixels. Segmentation then becomes a node partitioning problem. A good segmentation desires a partitioning that
has tight connections within partitions and loose connections across partitions. These two goals can both be achieved in the normalized cuts criterion [5], a brief self-contained account of which is given below.

### 3.1 Representation

Given weight matrix \( W \), the multiclass normalized cuts criterion tries to maximize the average of all \( K \) linkratios [11]:

\[
\epsilon_{NC}(\Gamma^k) = \frac{1}{K} \sum_{i=1}^{K} \text{linkratio}(\Gamma^k_i, \Gamma^k) \tag{4}
\]

\[
\text{linkratio}(\Gamma^k_i, \Gamma^k) = \frac{\sum_{i \neq j} W(i,j)}{\sum_{i \neq j} W(i,j)}.
\]

\( \text{linkratio}(\Gamma^k_i, \Gamma^k) \) is the fraction of the total weights within a group to the total weights all the member nodes have. Its complement \( \text{linkratio}(\Gamma^k_i, \Gamma^k) \) is the fraction of the weights between nodes in one group and the rest nodes in the graph. Since these two quantities sum up to one, maximizing the within-group weights and minimizing the between-group weights can both be achieved in the continuous domain and it can be solved by linear algebra.

In principle, we can solve \( \text{PNCZ} \) by applying the standard Rayleigh-Ritz theorem to its unconstrained version. That is, we first find a basis in the feasible solution space defined by \( U^T Z = 0 \). Let \( U^\perp \) denote an orthonormal basis in this space. Any solution that satisfies the partial grouping constraints can be represented by an \( (N - n) \times K \) coefficient matrix \( Y \) using this basis:

\[
Z = U^\perp Y, \quad U^T U^\perp = 0. \tag{14}
\]

We thus reduce \( \text{PNCZ} \) to a program in \( Y \):

\[
\begin{aligned}
\text{maximize} & \quad \epsilon_{NC}(Y) = \frac{1}{K} \text{tr}(Y^T W^p Y) \tag{15} \\
\text{subject to} & \quad Y^T D^p Y = I, \tag{16}
\end{aligned}
\]

where \( W^p = (U^\perp)^T W U^\perp \) and \( D^p = (U^\perp)^T D U^\perp \) are the equivalent weight and degree matrices for \( Y \). This is a standard Rayleigh quotient optimization problem. If \( (V^p, S^p) \) is the eigendecomposition of the matrix pair \( (W^p, D^p) \), where \( S^p = \text{Diag}(s^p) \) with nonincreasingly ordered eigenvalues in \( s^p \), then the global optimum is given by the eigenvectors corresponding to the first \( K \) largest eigenvalues and

\[
\epsilon_{NC}([V^p_1, \ldots, V^p_K]) = \frac{1}{K} \sum_{k=1}^{K} s^p_k \max_{V^p Y = I} \epsilon_{NC}(Y). \tag{17}
\]

From (14), we recover the global optimum in the original \( Z \)-space as \( Z^* = U^\perp [V^p_1, \ldots, V^p_K] \).

The introduction of \( Y \) gets rid of the constraint in (13) and turns program \( \text{PNCZ} \) into an unconstrained eigenvalue problem. However, it requires finding an orthonormal basis for the feasible space first. Given that \( n \ll N \), this process has a space and time complexity of \( O(N^2) \) and \( O(N^3) \), respectively, which is prohibitively expensive for a large \( N \). We have to find another way out.

There is such an alternative through the use of matrix projectors. \( Q \) is called a projector if it is idempotent, i.e., \( Q^2 = Q \). If \( Q \) is a projector onto the space of feasible solutions of \( \text{PNCZ} \), then \( Q Z \) is the projection of \( Z \) on the feasible space. The key property of \( Q Z \) is that \( Q Z = Z \) if and only if \( Z \) is feasible. Therefore, we can guarantee the feasibility of a solution by projecting it to the feasible set in the original space without resorting to any reparameterization in a reduced space.

We introduce a few symbols to simplify notation. Let \( \pi \) be a vector of \( K \) distinct integers from \( [N] \). For any eigenvector matrix \( V \) and its corresponding eigenvalue matrix \( S = \text{Diag}(s) \), let \( V_z = [V_{z_1}, \ldots, V_{z_K}] \) and \( S_z = \text{Diag}([s_{z_1}, \ldots, s_{z_K}]) \).
Theorem 1 (Generalized Rayleigh-Ritz Theorem). Let 
\((V, S)\) be the following eigendecomposition of matrix \(QPQ\):
\[
\begin{align*}
Q P Q V &= V S \quad (18) \\
V ^T D V &= I. \quad (19)
\end{align*}
\]
where \(P\) is the row-normalized weight matrix and \(Q\) is a projector onto the feasible solution space:
\[
\begin{align*}
P &= D ^{-1} W \quad (20) \\
Q &= I - D ^{-1} U (U ^T D^{-1} U) ^{-1} U ^T. \quad (21)
\end{align*}
\]
For any local optimum candidate \(Z^*\) to program PNCZ, there exists an index vector \(\pi\) and an orthonormal matrix \(R\) such that:
\[
Z^* = V _\pi R, \quad R ^T R = I \quad (22)
\]
\[
\varepsilon _{NC}(Z^*) = \frac{1}{K} \text{tr}(S _\pi). \quad (23)
\]
Assuming that the eigenvectors are ordered according to their eigenvalues, where \(s_1 \geq \ldots \geq s_N\), any global optimum of PNCZ can thus be specified by the first \(K\) largest eigenvectors and any orthonormal matrix:
\[
\begin{align*}
Z^* &= V _{[K]} , \quad R ^T R = I \quad (24) \\
\varepsilon _{NC}(Z^*) &= \frac{1}{K} \text{tr}(S _{[K]}) = \max _{\pi} \varepsilon _{NC}(Z). \quad (25)
\end{align*}
\]
Proof. We define a Lagrangian for PNCZ:
\[
L(Z, \Lambda, \Theta) = \frac{1}{2} \text{tr}(Z ^T W Z) - \frac{1}{2} \text{tr}(\Lambda ^T (Z ^T D Z - I)) - \Theta ^T U ^T Z, \]
where \(\Lambda\) is a \(K \times K\) symmetric matrix and \(\Theta\) is an \(n \times K\) matrix. An optimal solution \((Z^*, \Lambda^*, \Theta^*)\) must satisfy:
\[
\begin{align*}
L _Z(Z, \Lambda, \Theta) &= W Z - D \Lambda Z - U \Theta = 0, \quad (26) \\
L _\Lambda(Z, \Lambda, \Theta) &= Z ^T D Z - I = 0, \quad (27) \\
L _\Theta(Z, \Lambda, \Theta) &= U ^T Z = 0. \quad (28)
\end{align*}
\]
Multiplying (26) with \(U ^T D^{-1}\) leads to:
\[
\Theta^* = (U ^T D^{-1} U) ^{-1} U ^T D^{-1} W Z^*, \quad (29)
\]
where \(D\) and \(U ^T D^{-1} U\) are invertible since both \(D\) and \(U\) assume full rank. Eliminating \(\Theta\) in (26) by (29), we obtain
\[
QPZ^* = Z^* \Lambda^*. \quad (30)
\]
From (28), we also have \(QZ^* = Z^*\). Substituting it into the above equation, we obtain \(QPQZ^* = Z^* \Lambda^*\). Therefore, there are three necessary conditions for the optimality: \(\Lambda^*\) is symmetric and
\[
QPQZ^* = Z^* \Lambda^*, \quad Z ^T D Z^* = I. \quad (31)
\]
Next, we show that there exists an eigendecomposition \((V, S)\) of \(QPQ\) that not only meets these conditions but can also generate all such solutions through orthonormal matrices.
Noting that \(QPQZ^* = Z^* \Lambda^*\) is equivalent to:
\[
D ^\dagger Q D ^\dagger \cdot D ^\dagger P D ^\dagger \cdot D ^\dagger Q D ^\dagger \cdot D ^\dagger Z^* = D ^\dagger Z^* \Lambda^*, \quad (32)
\]
we rewrite (31) using a transformed variable \(Z\):
\[
\begin{align*}
QPQZ &= Z \Lambda^*, \quad Z ^T Z = I, \quad (33) \\
Z &= D ^\dagger Z^*, \quad (34) \\
P &= D ^\dagger P D ^\dagger = D ^\dagger W D ^\dagger, \quad (35) \\
Q &= D ^\dagger Q D ^\dagger = I - D ^\dagger U (U ^T D^{-1} U) ^{-1} U ^T D ^\dagger. \quad (36)
\end{align*}
\]
Since both \(P\) and \(Q\) are symmetric, \(QPQ\) is symmetric, which means that all its eigenvectors are real and orthogonal. Therefore, if \((V, S)\) is an orthonormal eigendecomposition of \(QPQ\), then any \(K\) distinct eigenvectors and their eigenvalues, i.e., \((V _\pi, S _\pi)\), form a solution to (33).
If \((Z, \Lambda^*)\) is a solution that satisfies (33) with \(\Lambda^*\) orthonormal and \(\Lambda^*\) symmetric, since \(V\) is a complete basis in the \(N\)-dimensional space, there exists an index vector \(\pi\) and an orthonormal matrix \(R\) such that
\[
Z = V _\pi R, \quad R ^T R = I \quad (37) \\
\Lambda^* = R ^T S _\pi R. \quad (38)
\]
Multiplying (26) with \(Z ^T\) and using \(\text{tr}(AB) = \text{tr}(BA)\), we derive:
\[
K \varepsilon _{NC}(Z^*) = \text{tr}(Z ^T W Z^*) = \text{tr}(\Lambda^*) = \text{tr}(S _\pi). \quad (39)
\]
Therefore, \(\{V _\pi, S _\pi : \pi\}\) produce all possible local optimal values. The global optimal value is thus given by the average of the first \(K\) largest eigenvalues. Transforming \(Z\) back to the \(Z\) space based on (34), we have \(V = D ^\dagger V\) and \((V, S)\) as an eigendecomposition of \(QPQ\). This completes the proof.

When there is no constraint, \(Q = I\), then \(QPQ = P\) can be considered as a transition probability matrix of random walks, and the normalized cuts criterion is equivalent to a maximum conductance problem where subsets of states only occasionally visit each other [13]. When there are constraints, \(Q \neq I\), \(QPQ\) usually has negative entries and it no longer has a transition probability interpretation. In other words, the solution to constrained grouping can no longer be cast as the equilibrium of a natural diffusion process.

To summarize, the optimal solution to PNCZ is not unique. It is a subspace spanned by the first \(K\) largest eigenvectors of \(QPQ\) by orthonormal matrices:
\[
Z^* \in \{V _{[K]} R : Q P Q V _{[K]} = V _{[K]} S _{[K]}, R ^T R = I\}. \quad (40)
\]
Unless all \(K\) eigenvalues are the same, \(V _{[K]} R\) are no longer the eigenvectors of \(QPQ\). Yet, all these solutions have the optimal objective value.
After we compute \(V _{[K]} S _{[K]}\) from \(QPQ\), the same procedure for the unconstrained normalized cuts can be applied to find a near global-optimal discrete solution to PNCX. The only difference is that now the eigenvectors are from \(QPQ\) rather than \(P\). In the discretization procedure, we honor the constraints that were ignored when we relaxed the program PNCX into the program PNCZ. That is, we find a discrete solution that satisfies the binary and exclusion constraints in (7), yet is closest to the continuous optima given in (40). This is another optimization problem which can be solved efficiently, since the objective function is bilinear in the discrete solution \(X\) and the orthonormal transform \(R\). The details can be found in [11].
3.3 Algorithm
To summarize, given data \( f \) defined over \( V \), and \( n \) partial grouping node sets \( \{ U_t : t \in [n] \} \), we use the following constrained normalized cuts algorithm to find an optimal \( K \)-way grouping:

1. Compute the affinity matrix \( W \) from the data \( f \), e.g.:
   \[
   W(i, j) = e^{-\frac{(d(i, j))^2}{\sigma^2}}, \quad i, j \in V.
   \]

2. Derive the constraint matrix \( U \) from \( \{ U_t : t \in [n] \} \):
   \[
   k = 0 \\
   \text{For } t = 1:n, \\
   \quad \text{For } s = 1:|U_t| - 1, \\
   \quad \quad k = k + 1 \\
   \quad U(U_t(s), k) = 1 \\
   \quad U(U_t(s + 1), k) = 1
   \]

3. Compute the degree matrix \( D = \text{Diag}(W1_N) \).

4. Compute \( P, U \), and \( H \) as:
   \[
   P = D^{-\frac{1}{2}}WD^{-\frac{1}{2}} \\
   U = D^{-\frac{1}{2}}U \\
   H = (U^TU)^{-1}
   \]

5. Compute the first \( K \) eigenvectors of \( QPQ \) by solving:
   \[
   (I - UHU^T)P(I - UHU^T)V_{[K]} = V_{[K]}S_{[K]} \\
   V_{[K]}V_{[K]}^T = I
   \]

6. Compute the first \( K \) eigenvectors of \( QPQ \) by
   \[
   V_{[K]} = D^{-\frac{1}{2}}V_{[K]}. 
   \]

7. Obtain a discrete segmentation \( X^* \) closest to \( V_{[K]} \) [11].

In Step 5, we avoid directly computing \( QPQ \) since it can become a dense matrix even when \( U \) and \( P \) are sparse. Specifically, we modify the innermost iteration in an eigensolver. For that, we only need to precompute \( U = D^{-\frac{1}{2}}U \), which is as sparse as \( U \), and \( H = (U^TU)^{-1} \), which is an \( n \times n \) matrix. \( U \) and \( H \) are the only two other matrices apart from those already used for unconstrained cuts. During each iteration of \( x := QPQx \), we compute:

\[
\begin{align*}
   z & := Qx = x - UHU^T \\ & x \\
   y & := Px \\
   x & := Qy = y - UHU^T \\ & x
\end{align*}
\]

If \( P \) has an average of \( k \) nonzeros per row, then (42) has \( O(Nk) \) multiplications. Equations (41) and (43) each requires \( O(2NN + n^2) \) multiplications, which are the only extra computation needed for constrained cuts. Given that \( n \ll N \) but comparable to \( k \), the increase in time complexity is linear. However, since the solution space is reduced; fewer iterations are needed to converge to the leading eigenvectors. Therefore, the net increase in the computational space and time is negligible if the number of constraints \( n \) is small. We can further reduce the complexity by sampling the constraints.

We can also avoid the matrix inversion in computing \( H \). To see this, let \( (A, \Sigma, B) \) be the singular value decomposition (SVD) of \( U \). Since

\[
U = A_{N \times N} \Sigma_{N \times n} B_{n \times N}^T, \quad A^T A = I, \quad B^T B = I, \quad (44)
\]

we have

\[
H = (U^TU)^{-1} = (B \Sigma^T \Sigma B^T)^{-1} = B(\Sigma^T \Sigma)^{-1} B^T. \quad (45)
\]

Therefore, we can eliminate \( H \) altogether since we only need \( UHU^T \) and it becomes:

\[
UHU^T = A \Sigma (\Sigma^T \Sigma)^{-1} \Sigma^T A = A_{[n]} A_{[n]}^T. \quad (46)
\]

That is, instead of keeping both \( U \) and \( H \), we only need to compute the \( n \) right eigenvectors of \( U \) and replace \( UHU^T \) with \( A_{[n]} A_{[n]}^T \) in Step 5.

Whether to use \( A_{[n]} \) or both \( U \) and \( H \) depends on the conditions of the constraint matrix. When the number of constraints is small, \( H \) is small and \( U \) is very sparse, whereas \( A_{[n]} \) is a full \( N \times n \) matrix. With the additional cost of computing \( A \) from \( U \), using \( A_{[n]} \) might not be a good choice. However, when the number of constraints is large, the matrix inversion involved in computing \( H \) could be costly and unstable. Later when we smooth the constraints, the columns of \( U \) can become dense and correlated. In these cases, we can use the SVD of \( U \) to find a small set of significant constraints, i.e., the first few columns of \( A \), making the computation stable and manageable.

4 PROPAGATING CONSTRAINTS
The basic formulation works reasonably well if there are enough partial grouping cues. This is not very useful since in reality only a few such cues are given. Sparse cues expose an inherent flaw in the formulation; however, it can be remedied.

4.1 Point Set Example
In Fig. 2, points are naturally organized into four clusters based on proximity. Since the vertical gap is larger than the horizontal gap, an ideal 2-class clustering is obtained by a horizontal cut that divides the four clusters into top and bottom groups. Now, if a few points at the horizontal boundary are grouped together a priori, the horizontal cut violates the partial grouping constraints and the vertical cut becomes optimal. However, when the number of grouping cues is reduced, the formulation in (3) fails to produce the desired vertical cut that divides the four clusters into left and right groups. In particular, the labeled points tend to stand out, while having little impact on the grouping of the rest of the points.

4.2 Why Simple Constraints Are Insufficient
When we preassign points from top and bottom clusters together, we do not just want a group to lose its labeled points to the other group (Fig. 2c), but rather we desire a grouping process that explores their neighboring connections and discovers the left-right division instead.

The formulation in (3), however, does not entail the desire of propagating grouping information on the constrained data points to their neighbors. Often, a slightly perturbed version of the optimal unbiased segmentation becomes the legitimate optimum (Fig. 3). This observation is made from a general optimization point of view and, thus, holds for all choices of \( \varepsilon \). The basic formulation in (3), although straightforward, is flawed.
The more similar a data point is to the biased ones, the average is taken based on the coherence among data points. We desire an average of their labels to be the same. The enforcing exactly the same labels on biased data points, remedy to our basic formulation. Rather than strictly from the perturbation of the unbiased optimal grouping.

Local segmentations are dissimilar in any solution resulting similar affinity patterns to their nearby points, yet their friends. In Fig. 3, the two points in a labeled pair have feasible space.

Optimal unbiased grouping solutions are ruled out from the condition a grouping to the extent that many trivial near-

Such smoothed constraints on the biased data points can condition a grouping to the extent that many trivial near-

The second reason is that such a biased grouping lacks fairness with regard to labeled points. Intuitively, if two labeled points, \( i \) and \( j \), have similar connections to their neighbors, we desire a fair segmentation so that if \( i \) gets grouped with \( i \)'s friends, \( j \) also gets grouped with \( j \)'s friends. In Fig. 3, the two points in a labeled pair have similar affinity patterns to their nearby points, yet their local segmentations are dissimilar in any solution resulting from the perturbation of the unbiased optimal grouping.

These two conditions, smoothness and fairness of the local segmentations on biased data points, provide a remedy to our basic formulation. Rather than strictly enforcing exactly the same labels on biased data points, we desire an average of their labels to be the same. The average is taken based on the coherence among data points. The more similar a data point is to the biased ones, the heavier the weight is on the label that it takes. Formally, let \( g_1 \circ g_2 \) be the compound function of \( g_1 \) and \( g_2 \). Let \( S_f \) denote a smoothing function contingent on the data \( f \). We modify the formulation in (3) to be:

\[
\begin{align*}
\text{maximize} & \quad \varepsilon(\Gamma V, f), \\
\text{subject to} & \quad S_f \circ \Gamma V(i, l) = S_f \circ \Gamma V(j, l), \\
& \quad i, j \in U, l \in K, t \in [n].
\end{align*}
\]

Such smoothed constraints on the biased data points can condition a grouping to the extent that many trivial near-optimal unbiased grouping solutions are ruled out from the feasible space.

Our new formulation is not equivalent to the introduction of smoothness priors in a generative approach. There, prior knowledge such as piecewise constancy is usually imposed on the solution independently of the goodness of fit [6], whereas ours is closely coupled with the coherence of the data. Our essential message, in this regard, is that an effective propagation of priors requires an intimate interaction with the data themselves.

### 4.3 Smooth Constraints for Normalized Cuts

A natural choice of \( S_f \) for the normalized cuts criterion is the normalized weight matrix \( P \):

\[
S_f \circ \Gamma V V(i, l) = \sum_j P_{ij} X(j, l), \quad i \in V, l \in [K].
\]

This value measures the average density of \( V_i \) from node \( i \)'s point of view, with nodes of high affinity to it weighted more in the density. This discourages \( i \) to take a label different from those of its close neighbors. We may not know in advance what this density is for the optimal partitioning, but the fairness condition requires it to be the same for the labeled pair \((i, j)\): \( S_f \circ \Gamma V V(i, l) = S_f \circ \Gamma V V(j, l) \). The partial grouping constraints in (8) then become:

\[
U^T P X = (P^T U)X = 0.
\]

Since the only change here is that the constraint matrix \( U \) becomes \( P^T U \), the same solution technique applies. That is, the eigensolution to the program PNCZ is given by the eigenvectors of \( P^T Q \), where \( Q \) is a projector onto the solution space specified by \( (P^T U)^T X = 0 \) instead of \( U^T X = 0 \).

In Fig. 4, we show new results with the smoothed constraints. In addition to the basic results in Fig. 2, we also consider two other alternatives that directly utilize partial grouping cues. The simplest case of encoding the labeled pair \((i, j)\) is to modify their weights so that:

\[
W_{ij} = W_{ji} := 1,
\]

where an originally vanishingly small value increases to the maximum affinity. The influence of this change depends on the number of connections the biased nodes have. For example, if node \( i \) connects to 10 other nodes, this one more
connection would matter little after being normalized by the total connections. Unlike minimum cuts, where a change in one link can change the global optimum completely, normalized cuts are insensitive to perturbation in the weights. Another approach is to let $i$ and $j$ share each other's neighboring connections since $i$ and $j$ are indistinguishable in a desired grouping:

$$W_{ik} = W_{ki} = W_{kj} = W_{jk} := \max(W_{ik}, W_{jk}), \; k \in \mathcal{V}. \quad (51)$$

Short-circuiting labeled nodes as well as their neighbors produces a similar result as the simple biased grouping in Fig. 2. Their common problem is that only the labeled nodes expand their neighborhoods significantly, which make them distinct from the rest unlabeled data. If we extend (51) to modify the weights among the neighbors of labeled points, we can overcome the discontinuity of the segmentation. That's what (49) does, and in a principled way.

The inherent flaw in our basic formulation is also evident in the undesirable results from even dense grouping cues. Though it is unclear for this point set what the best 4-class clustering is with either dense or sparse partial grouping cues, as shown in Fig. 5, the labeled data points never stand out with smoothed constraints. In general, we don't know how many classes there are and whether the partial grouping cues are sufficient. Therefore, partial grouping constraints should always be smoothed with the coherence exhibited in the data in order to produce a meaningful segmentation.

5 Experiments

We calculate pixel affinity using a Gaussian function on the maximum magnitude of intensity edges separating two pixels. $W(i,j)$ is low if $i, j$ are on the opposite sides of a strong edge [14]. Using this simple feature, we will demonstrate how simple extra-image knowledge can improve low-level segmentation and how smoothed partial grouping constraints make a difference.

In Fig. 6, we derive partial groupings based on brightness values, e.g., the foreground is more likely to be lighter and the background is darker. We choose two thresholds to find the pixels at the two intensity extremes and then use morphological operations to further remove pixels appearing in the other set due to noise. As we have already seen in Fig. 2, with simple constraints, biased pixels stand out in segmentation, while with smoothed constraints, they bring...
their neighbors along and change the segmentation completely. This image has rich texture against a relatively simple background. Compared to segmentation using morphological operations on such images, our method can fill the holes caused by thresholding without losing thin structures or distorting region boundaries.

Partial grouping cues can also be derived from motion cues in a video sequence. In Fig. 7, for every image, we compute its difference with two preceding images in a video sequence, threshold and then apply morphological operations to the difference image to create a mask for the foreground. Our constrained segmentation can effectively shrink it to the head in motion.

Partial grouping cues can come not only from low-level cues, but also from high-level expectation. For fashion pictures featuring a fashion model at the center, we choose the background to be: 4-pixel wide at left and right sides, and 7-pixel high at top and bottom sides. Figs. 8 and 9 show the results with and without such background knowledge. Notice that all eigenvectors of $QPQ$ satisfy the constraints and pixels at the four image sides always have similar values in the eigensolutions. Through these constraints, the large uniform background is never broken up in a segmentation, which focuses on the more interesting foreground-background separation or a division within the foreground itself.

Using the same spatial mask and the same set of parameters for computing pixel affinity, we apply our
In our experiments, we also observe no significant improvement over the partial grouping further at the cost of more computation. However, higher-order smoothness constraints propagate to neighboring nodes that they have large affinity with. The extent of this detrimental effect depends on the connections of the constrained nodes, since partial grouping information is propagated to neighboring nodes that have large affinity with. Our formulation can neither spot nor correct mistakes in priors.

Technically, (49) can be replaced by an up-to $x$th order smoothness condition (or a subset of it): $S_I = [P^0, P^1, \ldots, P^x]$. However, higher-order smoothness constraints propagate the partial grouping further at the cost of more computation. In our experiments, we also observe no significant improvement over $S_I = P$ in the eigensolutions.

6 DISCUSSION OF RELATED WORK

Our work can be regarded as a small step toward bridging generative approaches and discriminative approaches for grouping. Generative models, including MRF [6] and variational formulations [16], [17], can be naturally cast in a Bayesian framework, where data fidelity and model specificity are treated at equal footing. However, they are sensitive to model mismatches and are usually solved by MCMC methods, which often find local optima with slow convergence.

Discriminative methods, for example graph approaches on image segmentation [5], [18], [19], [20], [21], [22], [23], achieve a global decision based on local pairwise relationships. These algorithms often have efficient computational solutions. These local pairwise comparisons can encode general grouping rules such as proximity and feature similarity. Promising segmentation results on a wide range of complex natural images were reported in [14]. Such pairwise comparisons, however, often have difficulty in deriving reliable long-range grouping information.

Attempts have been made to find MRF solutions by graph partitioning algorithms [7], [8], [24], [25], [26]. In particular, sufficient and necessary conditions on the properties of energy functions that can be solved by minimum cuts have been proven in [27], [28]. The work here shows that prior knowledge can be used to guide grouping for discriminative criteria such as normalized cuts [5] and that their global optima in the continuous domain can be solved algebraically with little extra cost.

Our work is also closely linked to the transduction problem, the goal of which is to complete the labeling of a partially labeled data set [29], [30], [31], [32]. If the labeled data set is rich enough to characterize both the structures of the data and the classification task, then using the induced classifier on the labeled set and interpolating it to the unlabeled set shall suffice, which is a supervised learning problem that has many efficient algorithms. However, usually the labeled set is small, so the problem becomes how to integrate the two types of information from both sets to reach a better solution. In [29], the classification problem is formulated in the support vector machine (SVM) framework and labeled data are treated similarly to the rest except that their labels have been instantiated. In [30], information about the labeled data is encoded in the prior distribution of the labeling and the goal is to find a projection of the best SVM discriminator onto the prior space. Through model averaging, partial labeling constraints are softly enforced. In [31], class-dependent data generation models are assumed and the labeled data can be used to estimate the parameters involved in the models. This might be the most effective way to propagate priors. However, these generative models are often too simple to be realistic. In [32], the class-dependent probability models are hidden in the pairwise affinity matrix of all the data points. Again, the labeled set is used to estimate the class-dependent label generation process.

Though our work was initially motivated by the gap between discriminative and generative approaches, we are aware of other works that put similar constraints into clustering algorithms such as $K$-means [33], [34]. Two types of constraints, must-link and cannot-link, are considered. Earlier versions of our work [35], [36] also considered cannot-link constraints, that is, two nodes cannot assume the same label. Such constraints are not transitive, which makes them difficult to propagate. In [35], repulsion weights...
are used to help enforcing such cues. It also involves approximation in the constraint formulation. For clarity, we choose not to include cannot-link constraints here. Our work is distinct from all these methods in two aspects. Rather than instantiating the labels or the constraints on labeled data points, we use them to regulate the form of a segmentation. We gave an intuitive computational account for the need of constraint propagation and provided a principled way to implement it. Secondly, we can solve near-global optima of our formulation, whereas most other works can only guarantee local optimality.

Our experimental results on image segmentation demonstrate that simple grouping bias can approach figure-ground segregation without knowing what the object is. Our spatial priors effectively take advantage of the asymmetry between figure and ground [37]. In other words, since the outcome of a grouping depends on global configurations, figure-ground segregation can be obtained not only by enhancing the saliency of object structures, but also by suppressing background structures, the latter of which is often easier than the former. Our next step is to explore the integration of more complicated priors in order to segment out only objects known a priori.

7 Summary

We developed a method that integrates both bottom-up and top-down information in a single grouping process. The former is based on low-level cues presented in the image, whereas the latter is based on partial grouping cues known a priori; the former defines the goodness of a segmentation, whereas the latter defines the feasibility of a segmentation. The two are unified in a constrained optimization problem. We showed that it is essential to propagate sparse partial grouping cues based on the coherence exhibited in the data. In particular, we developed an efficient solution for such constrained normalized cuts and applied the method successfully to segmenting a wide range of real images.

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References


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