Graph-based Methods

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Image and Video Understanding a.y. 2018/19

Images as graphs





- Node for every pixel
- Edge between every pair of pixels (or every pair of "sufficiently close" pixels)
- Each edge is weighted by the *affinity* or similarity of the two nodes

Graph-theoretic segmentation

- Represent tokens using a weighted graph.
 - affinity matrix
- Cut up this graph to get subgraphs with strong interior links



Graphs and matrices









Measuring affinity

- Suppose we represent each pixel by a feature vector x, and define a distance function appropriate for this feature representation
- Then we can convert the distance between two feature vectors into an affinity with the help of a Gaussian kernel:

$$\exp\left(-\frac{1}{2\sigma^2}\operatorname{dist}(\mathbf{x}_i,\mathbf{x}_j)^2\right)$$

Scale affects affinity

- Small σ : group only nearby points
- Large σ : group far-away points



Eigenvector-based clustering

Let us represent a cluster using a vector \mathbf{x} whose k-th entry captures the participation of node k in that cluster. If a node does not participate in a cluster, the corresponding entry is zero.

We also impose the restriction that $\mathbf{x}^T \mathbf{x} = 1$

We want to maximize:

$$\sum_{i=1}^n \sum_{j=1}^n w_{ij} x_i x_j = \mathbf{x}^T \mathbf{A} \mathbf{x}$$

which is a measure for the cluster's cohesiveness.

This is an **eigenvalue problem**! *Choose the eigenvector of A with largest eigenvalue*

Example eigenvector





points



0.45



matrix

More than two segments

- Two options
 - Recursively split each side to get a tree, continuing till the eigenvalues are too small
 - Use the other eigenvectors

Segmentation by eigenvectors: Algorithm

- 1. Construct (or take as input) the affinity matrix A
- 2. Compute the eigenvalues and eigenvectors of *A*
- 3. Repeat
- 4. Take the eigenvector corresponding to the largest unprocessed eigenvalue
- 5. Zero all components corresponding to elements that have already been clustered
- 6. Threshold the remaining components to determine which elements belong to this cluster
- 7. If all elements have been accounted for, there are sufficient clusters
- 8. Until there are sufficient clusters

Segmentation as graph partitioning

Let G=(V, E, w) a weighted graph.

Given a "cut" (A, B), with $B = V \setminus A$, define:

$$cut(A,B) = \sum_{i \in A} \sum_{j \in B} w(i,j)$$



Minimum Cut Problem

Among all possible cuts (A, B), find the one which minimizes cut(A, B)

Segmentation as graph partitioning



MinCut clustering

Good news Solvable in polynomial time

Bad news

Favors highly unbalanced clusters (often with isolated vertices)



Graph terminology

Degree of nodes

$$d_i = \sum_j w_{i,j}$$





Volume of a set







Normalized Cut

$$Ncut(A,B) = cut(A,B) \begin{pmatrix} 1 & 1 \\ vol(A) & vol(B) \end{pmatrix}$$



Adapted from D. Sontag

Graph Laplacian (unnormalized)

Defined as

L = D - W

Example:



Assume the weights of edges are 1

Key fact

For all vectors f in \mathbf{R}^n , we have:

$$f^{\top}Lf = \frac{1}{2} \sum_{ij=1}^{n} w_{ij} (f_i - f_j)^2$$

Indeed:

$$\begin{split} f^{\top}Lf &= f^{\top}Df - f^{\top}Wf \\ &= \sum_{i} d_{i}f_{i}^{2} - \sum_{i,j} f_{i}f_{j}w_{ij} \\ &= \frac{1}{2} \left(\sum_{i} (\sum_{j} w_{ij})f_{i}^{2} - 2\sum_{ij} f_{i}f_{j}w_{ij} + \sum_{j} (\sum_{i} w_{ij})f_{j}^{2} \right) \\ &= \frac{1}{2} \sum_{ij} w_{ij}(f_{i} - f_{j})^{2} \end{split}$$

Properties

• *L* is symmetric (by assumption) and positive semi-definite:

```
f'Lf \geq 0
```

for all vectors f (by "key fact")

- Smallest eigenvalue of *L* is 0; corresponding eigenvector is **1**
- Thus eigenvalues are: $0 = \lambda_1 \le \lambda_2 \le \dots \le \lambda_n$

First relation between spectrum and clusters:

- Multiplicity of eigenvalue $\lambda_1 = 0$ is the number of connected components of the graph
- eigenspace is spanned by the characteristic functions of these components (so all eigenvectors are piecewise constant)

Normalized graph Laplacians

• Row sum (random walk) normalization:

$$L_{\rm rw} = D^{-1}L$$
$$= I - D^{-1}W$$

• Symmetric normalization:

$$L_{\text{sym}} = D^{-1/2} L D^{-1/2}$$
$$= I - D^{-1} W D^{-1/2}$$

Spectral properties of both matrices similar to the ones of *L*.

Solving Ncut

Any cut (*A*, *B*) can be represented by a binary indicator vector *x*:

$$x_i = \begin{cases} +1 & \text{if } i \in A \\ -1 & \text{if } i \in B \end{cases}$$

It can be shown that:

$$\min_{x} Ncut(x) = \min_{y} \frac{y'(D-W)y}{y'Dy}$$

Rayleigh quotient

subject to the constraint that $y'D\mathbf{1} = \sum_i y_i d_i = 0$ (with $y_i \in \{1, -b\}$).

This is NP-hard!

Ncut as an eigensystem

If we *relax* the constraint that *y* be a discrete-valued vector and allow it to take on real values, the problem

$$\min_{y} \frac{y'(D-W)y}{y'Dy}$$

is equivalent to:

$$\min_{y} y'(D-W)y \quad \text{subject to } y'Dy = 1$$

This amounts to solving a *generalized* eigenvalue problem:

Laplacian
$$\langle (D-W) y = \lambda D y$$

Note: Equivalent to a *standard* eigenvalue problem using the normalized Laplacian: $L_{rw} = D^{-1}L = I - D^{-1}W$.

2-way Ncut

- 1. Compute the affinity matrix *W*, compute the degree matrix *D*
- 2. Solve the generalized eigenvalue problem $(D W)y = \lambda Dy$
- 3. Use the eigenvector associated to the second smallest eigenvalue to bipartition the graph into two parts.

Why the *second* smallest eigenvalue?

Remember, the smallest eigenvalue of Laplacians is always 0 (corresponds to the trivial partition $A = V, B = \{\}$)



Fig. 5. (a) Point set generated by two Poisson processes, with densities of 2.5 and 1.0 on the left and right clusters respectively, (b) \triangle and \times indicate the partition of point set in (a). Parameter settings: $\sigma_X = 5$, r = 3.

The effect of relaxation





How to choose the splitting point?

- Pick a constant value (0 or 0.5)
- Pick the median value as splitting point
- Look for the splitting point that has minimum Ncut value:
 - 1. Choose *n* possible splitting points
 - 2. Compute Ncut value
 - 3. Pick minimum

Random walk intepretation

Problem: Finding a cut (A, B) in a graph G such that a random walk does not have many opportunities to jump between the two clusters.

This is equivalent to the *Ncut* problem due to the following relation:

Ncut(A, B) = P(A | B) + P(B | A)

(Meila and Shi, 2001)

Ncut: More than 2 clusters

Approach #1: Recursive two-way cuts

- 1. Given a weighted graph G = (V, E, w), summarize the information into matrices W and D
- 2. Solve $(D W)y = \lambda Dy$ for eigenvectors with the smallest eigenvalues
- 3. Use the eigenvector with the second smallest eigenvalue to bipartition the graph by finding the splitting point such that Ncut is minimized
- 4. Decide if the current partition should be subdivided by checking the stability of the cut, and make sure Ncut is below the prespecified value
- 5. Recursively repartition the segmented parts if necessary

Note. The approach is computationally wasteful; only the second eigenvector is used, whereas the next few small eigenvectors also contain useful partitioning information.

Ncut: More than 2 clusters

Approach #2: Using first *k* eigenvectors

- 1. Construct a similarity graph and compute the unnormalized graph Laplacian L.
- 2. Compute the k smallest **generalized** eigenvectors u_1, u_2, \dots, u_k of the generalized eigenproblem $Lu = \lambda Du$.

3. Let
$$U = [u_1 u_2 \cdots u_k] \in \mathbb{R}^{n \times k}$$
.

4. Let $y_i \in \mathbb{R}^k$ be the vector corresponding to the *i*th row of U.

$$U = \begin{bmatrix} u_{11} & u_{12} & \cdots & u_{1k} \\ u_{21} & u_{22} & \cdots & u_{2k} \\ \vdots & \vdots & \ddots & \vdots \\ u_{n1} & u_{n2} & \cdots & u_{nk} \end{bmatrix} = \begin{bmatrix} y_1^T \\ y_2^T \\ \vdots \\ y_n^T \end{bmatrix}$$

5. Thinking of y_i 's as points in \mathbb{R}^k , cluster them with k-means algorithms.



Fig. 2. A gray level image of a baseball game.















(e)



(f)



(g)





(i)

Fig. 3. Subplot (a) plots the smallest eigenvectors of the generalized eigenvalue system (11). Subplots (b)-(i) show the eigenvectors corresponding the second smallest to the ninth smallest eigenvalues of the system. The eigenvectors are reshaped to be the size of the image.

(h)



Fig. 4. (a) shows the original image of size 80×100 . Image intensity is normalized to lie within 0 and 1. Subplots (b)-(h) show the components of the partition with *Ncut* value less than 0.04. Parameter setting: $\sigma_I = 0.1$, $\sigma_X = 4.0$, r = 5.



Fig. 8. (a) shows a 126×106 weather radar image. (b)-(g) show the components of the partition with *Ncut* value less than 0.08. Parameter setting: $\sigma_I = 0.007$, $\sigma_x = 15.0$, r = 10.



Fig. 10. (a) shows an image of a zebra. The remaining images show the major components of the partition. The texture features used correspond to convolutions with DOOG filters [16] at six orientations and five scales.

Spectral clustering

- 1. Construct a similarity graph and compute the normalized graph Laplacian L_{sym} .
- 2. Compute the k smallest eigenvectors u_1, u_2, \cdots, u_k of L_{sym} .
- 3. Let $U = [u_1 u_2 \cdots u_k] \in \mathbb{R}^{n \times k}$.
- 4. Normalized the rows of U to norm 1.

$$U_{ij} \leftarrow \frac{U_{ij}}{(\sum_k U_{ik}^2)^{1/2}}$$

- 5. Let $y_i \in \mathbb{R}^k$ be the vector corresponding to the *i*th row of U.
- 6. Thinking of y_i 's as points in \mathbb{R}^k , cluster them with k-means algorithms.

Ng, Jordan and Weiss (2002)

K-means vs Spectral clustering

Applying k-means to Laplacian eigenvectors allows us to find cluster with non-convex boundaries.


K-means vs Spectral clustering

Applying k-means to Laplacian eigenvectors allows us to find cluster with non-convex boundaries.



K-means vs Spectral clustering

Applying k-means to Laplacian eigenvectors allows us to find cluster with non-convex boundaries.





Second eigenvector of graph Laplacian



Examples



Examples (choice of *k*)



Choosing *k*

The eigengap heuristic: Choose *k* such that all eigenvalues $\lambda_1, \dots, \lambda_k$ are very small, but λ_{k+1} is relatively large



Four 1D Gaussian clusters with increasing variance and corresponding eigevalues of L_{rw} (von Luxburg, 2007).

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Graph-based Methods in Computer Vision: Recent Advances

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Clustering on Graphs

Given:

- an *n* × *n* matrix A of pairwise similarities

Goal: Group the the input objects (the vertices of the graph) into maximally homogeneous classes (i.e., clusters).



What is a Cluster?

No universally accepted (formal) definition of a "cluster" but, informally, a cluster should satisfy two criteria:

Internal criterion

all "objects" inside a cluster should be highly similar to each other

External criterion

all "objects" outside a cluster should be highly dissimilar to the ones inside

How to formalize these criteria?

Basic Definitions

Let $S \subseteq V$ be a non-empty subset of vertices, and $i \in S$.

The (average) weighted degree of *i* w.r.t. S is defined as: $\operatorname{awdeg}_{S}(i) = \frac{1}{|S|} \sum_{j \in S} a_{ij}$

Moreover, if $j \notin S$, we define:

 $\phi_{S}(i,j) = a_{ij} - \operatorname{awdeg}_{S}(i)$



Intuitively, $\Phi_{S}(i,j)$ measures the similarity between vertices *j* and *i*, with respect to the (average) similarity between vertex *i* and its neighbors in *S*

Basic Definitions

Let $S \subseteq V$ be a non-empty subset of vertices, and $i \in S$.

The weight of *i* w.r.t. *S* is defined as:

$$w_{S}(i) = \begin{cases} I & \text{w.r.t. S} \text{ is defined as:} \\ \sum_{j \in S - \{i\}} \phi_{S - \{i\}}(j,i) w_{S - \{i\}}(j) & \text{otherwise} \end{cases}$$

Further, the **total weight** of *S* is defined as:

$$W(S) = \sum_{i \in S} w_S(i)$$



Interpretation

Intuitively, $w_{S}(i)$ gives us a measure of the overall (relative) similarity between vertex *i* and the vertices of S \ {*i*} with respect to the overall similarity among the vertices in S \ {*i*}.



Dominant Sets

Let $S \subseteq V$ be a subset of vertices of a graph G and $i \in S$.

Define a measure for the similarity between vertex i and the vertices of $S \setminus \{i\}$ with respect to the overall internal similarity of $S \setminus \{i\}$.

Call it $w_{s}(i)$.

S is said to be a **dominant set** if:

1. $w_{s}(i) > 0$, for all $i \in S$ (internal homogeneity

2. $w_{S \cup \{i\}}(i) < 0$, for all $i \notin S$ (external homogeneity

S

Dominant Sets

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The Many Facets of Dominant Sets

Dominant sets have intriguing connections wth:

Game theory

Nash equilibria of "clustering games"

Optimization theory

Local maximizers of (continuous) quadratic problems

Graph theory

Maximal cliques

Dynamical systems theory

Stable attractors of evolutionary game dynamics

Using Symmetric Affinities

Given a symmetric affinity matrix *A*, consider the following continuous quadratic optimization problem (QP):

 $\begin{array}{ll} \text{maximize} & f(\mathbf{x}) = \mathbf{x}' A \mathbf{x} \\ \text{subject to} & \mathbf{x} \in \Delta \end{array}$

where Δ is the standard simplex (probability space).

The function f(x) provides a measure of cohesiveness of a cluster.

Dominant sets are in one-to-one correspondence to (strict) local solutions of QP

Note. In the 0/1 case, dominant sets correspond to maximal cliques

Finding Dominant Sets

Replicator dynamics from evolutionary game theory are a popular and principled way to find DS's.

$$x_i(t+1) = x_i(t) \frac{A(x(t))_i}{x(t)^T A x(t)}$$

MATLAB

distance implementation

while distance>epsilon

old_x=x;

```
x = x.*(A*x);
```

```
x = x./sum(x);
```

```
distance=pdist([x,old_x]');
```

Faster dynamics available! (See Rota Bulò and Pelillo, 2017)

end

Measuring Cluster Membership

The components of the converged vector \mathbf{x} give us a measure of the participation of the corresponding vertices in the cluster, while the value of the objective function measures the cluster's cohe





In a Nutshell

The dominant-set approach to clustering:

- does not require a priori knowledge on the number of clusters
- ✓ is robust against outliers
- allows to rank the cluster's elements according to "centrality"
- ✓ allows extracting overlapping clusters (ICPR'08)
- generalizes naturally to hypergraph clustering problems (PAMI'13)
- makes no assumption on the structure of the similarity matrix, (works also with asymmetric and even negative

Some Computer Vision Applications

- Image and video segmentation
- Anomaly detection
- Video summarization
- Feature selection
- Image matching and registration
- 3D reconstruction
- Human action recognition
- Content-based image retrieval
- ...

But also in neuroscience, bioinformatics, medical image analysis, etc.



Detecting conversational groups in images and sequences: A robust game-theoretic approach^{**}



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F-formations

"Whenever two or more individuals in close proximity orient their bodies in such a way that each of them has an easy, direct and equal access to every other participant's transactional segment"



System Architecture



Frustrum of visual attention

- A person in a scene is described by his/her position (x,y) and the head orientation θ
- The frustum represents the area in which a person can sustain a conversation and is defined by an aperture and by a length

-		CoffeeBre		eak (S1+S2)		PosterData		Gdet		
	Method *	Prec	Rec	F1	Prec	Rec	F1	Prec	Rec	F1
	IRPM [61],[22]	0.60	0.41	0.49	-		-	-	10	-
	HFF [22]	0.82	0.83	0.82	0.93	0.96	0.94	0.67	0.57	0.62
	DS ([6], [22])*	0.68	0.65	0.66	0.93	0.92	0.92	<u>115</u>		3 <u>—</u> 3
	MULTISCALE [46]	0.82	0.77	0.80	1000			18-91	0	10-00
	GTCG [47] KL	0.80	0.84	0.82	0.90	0.94	0.92	0.76	0.75	0.75
	GTCG [47] JS	0.83	0.89	0.86	0.92	0.96	0.94	0.76	0.76	0.76
	R-GTCG SC	0.52	0.59	0.55	0.26	0.27	0.26	0.75	0.75	0.75
	R-GTCG	0.86	0.88	0.87	0.92	0.96	0.94	0.76	0.76	0.76
Spectral		$\sigma = 0.2$, $l = 145$		$\sigma = 0.25$, $l = 115$			$\sigma = 0.7 l = 180$			
Clustering		cocktain raity		Synth						
Clustering	Method \star	Prec	Rec	F1	Prec	Rec	F1			
	IRPM [22,61]	-			0.71	0.54	0.61			
	HFF ([7], [46])	0.59	0.74	0.66	0.73	0.83	0.78			
	MULTISCALE [46]	0.69	0.74	0.71	0.86	0.94	0.90			
	GTCG [47] KL	0.85	0.81	0.83	1.00	1.00	1.00			
	GTCG [47] JS	0.86	0.82	0.84	1.00	1.00	1.00			
	R-GTCG SC	0.77	0.72	0.74	0.40	0.90	0.56			
		$\sigma = 0.6$, $l = 170$		$\sigma = 0.1$	$\sigma = 0.1$, $l = 75$					
	R-GTCG	0.87	0.82	0.84	1.00	1.00	1.00			



Qualitative results on the CoffeeBreak dataset compared with the state of the art HFF.

Yellow = ground truth Green = our method Red = HFF.

Dominant Sets for "Constrained" Image Segmentation

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Abstract—Image segmentation has come a long way since the early days of computer vision, and still remains a challenging task. Modern variations of the classical (purely bottom-up) approach, involve, e.g., some form of user assistance (interactive segmentation) or ask for the simultaneous segmentation of two or more images (co-segmentation). At an abstract level, all these variants can be thought of as "constrained" versions of the original formulation, whereby the segmentation process is guided by some external source of information. In this paper, we propose a new approach to tackle this kind of problems in a unified way. Our work is based on some properties of a family of quadratic optimization problems related to *dominant sets*, a graph-theoretic notion of a cluster which generalizes the concept of a maximal clique to edge-weighted graphs. In particular, we show that by properly controlling a regularization parameter which determines the structure and the scale of the underlying problem, we are in a position to extract groups of dominant-set clusters that are constrained to contain predefined elements. In particular, we shall focus on interactive segmentation and co-segmentation (in both the unsupervised and the interactive versions). The proposed algorithm can deal naturally with several types of constraints and input modalities, including scribbles, sloppy contours and bounding boxes, and is able to robustly handle noisy annotations on the part of the user. Experiments on standard benchmark datasets show the effectiveness of our approach as compared to state-of-the-art algorithms on a variety of natural images under several input conditions and constraints.

Index Terms-Interactive segmentation, co-segmentation, dominant sets, quadratic optimization, game dynamics.

Constrained Dominant Sets

Given $S \subseteq V$ and a parameter $\alpha > 0$, define the following parameterized family of quadratic programs:

maximize $f_S^{\alpha}(\mathbf{x}) = \mathbf{x}'(A - \alpha \hat{I}_S)\mathbf{x}$ subject to $\mathbf{x} \in \Delta$

where I_S is the diagonal matrix whose elements are set to 1 in correspondence to the vertices outside S, and to zero otherwise:

$$\hat{I}_S = \left(\begin{array}{cc} 0 & 0\\ 0 & I_{n-k} \end{array}\right)$$

Property. By setting:

 $\alpha > \lambda_{\max}(A_{V \setminus S})$

all local solutions will have a support containing elements of S

Interactive Image Segmentation

Given an image and some information provided by a user, in the form of a scribble or of a bounding box, to provide as output a foreground object that best reflects the user's intent



System Overview



Left: Over-segmented image with a user scribble (blue label).

Middle: The corresponding affinity matrix, using each over-segments as a node, showing its two parts: S, the constraint set which contains the user labels, and V n S, the part of the graph which takes the regularization parameter .

Right: RRp, starts from the barycenter and extracts the first dominant set and update x and M, for the next extraction till all the dominant sets which contain the user labeled regions are extracted.

Table 1. Error rates of differentscribble-based approaches on the Grab-Cut dataset

Methods	Error rate
Graph Cut [7]	6.7
Lazy Snapping [5]	6.7
Geodesic Segmentation [4]	6.8
Random Walker [33]	5.4
Transduction [34]	5.4
Geodesic Graph Cut [30]	4.8
Constrained Random Walker [31]	4.1
CDS_Self Tuning (Ours)	3.57
CDS_Single Sigma (Ours)	3.80
CDS_Best Sigma (Ours)	2.72

Table 2.Jaccard index of differentapproaches – first 5 bounding-box-based – on Berkeley dataset

Methods	Jaccard index
MILCut-Struct [3]	84
MILCut-Graph [3]	83
MILCut [3]	78
Graph Cut [1]	77
Binary Partition Trees [35]	71
Interactive Graph Cut [7]	64
Seeded Region Growing [36]	59
Simple Interactive O.E [37]	63
CDS_Self Tuning (Ours)	93
CDS_Single Sigma (Ours)	93
CDS_Best Sigma (Ours)	95



Bounding box

Result

Scribble

Result

Ground truth



Bounding box

Result

Scribble

Result

Ground truth

IEEE TRANSACTIONS ON PATTERN ANALYSIS AND MACHINE INTELLIGENCE

Large-scale Image Geo-Localization Using Dominant Sets

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Abstract—This paper presents a new approach for the challenging problem of geo-localization using image matching in a structured database of city-wide reference images with known GPS coordinates. We cast the geo-localization as a clustering problem of local image features. Akin to existing approaches to the problem, our framework builds on low-level features which allow local matching between images. For each local feature in the query image, we find its approximate nearest neighbors in the reference set. Next, we cluster the features from reference images using Dominant Set clustering, which affords several advantages over existing approaches. First, it permits variable number of nodes in the cluster, which we use to dynamically select the number of nearest neighbors for each query feature based on its discrimination value. Second, this approach is several orders of magnitude faster than existing approaches. Thus, we obtain multiple clusters (different local maximizers) and obtain a robust final solution to the problem using multiple weak solutions through constrained Dominant Set clustering also bypasses heuristic approaches to voting and selecting the reference image that matches to the query. We evaluate the proposed framework on an existing dataset of 102k street view images as well as a new larger dataset of 300k images, and show that it outperforms the state-of-the-art by 20% and 7%, respectively, on the two datasets.

Index Terms—Geo-localization, Dominant Set Clustering, Multiple Nearest Neighbor Feature Matching, Constrained Dominant Set

Image Geo-localization

A new approach for the problem of geo-localization using image matching in a structured database of city-wide reference images with known GPS coordinates.



200x time faster + 20% accuracy improvement w.r.t previous approach

Datasets:

- Datasets one:
 - Reference images:
 - 102K Google street view images from Pittsburgh,
 PA and Orlando, FL
 - Test Set:
 - 521 GPS-Tagged unconstrained images
 - Downloaded From Flickr, Panoramio, Picasa, ...
- WorldCities Datasets (NEW)*:
 - Reference images:
 - 300K Google street view images
 - 14 different cities from Europe, N. America and Australia
 - Test Set:
 - 500 GPS-Tagged unconstrained images
 - Downloaded From Flickr, Panoramio, Picasa, ...



Google Maps Street View Datasets:

For each location: 4 side views and 1 top view is collected





Pittsburgh, PA Longitude = 40.44146° Latitude = 80.0037°
Overall Result

• Dataset 1: 102K Google street view images (Orlando and Pittsburg area)



Overall Result

• Dataset 2: WorldCities (14 different cities from Europa, North America, Australia)



Computational Time



Qualitative Results



Query

Match – Error: 70.01 m

```
Match – Error: 5.4 m
```





Query



Match – Error: 7.5 m





Match – Error: 62.7 m

Multi-Target Tracking in Multiple Non-Overlapping Cameras using Fast-Constrained Dominant Sets

Yonatan Tariku Tesfaye \cdot Eyasu Zemene \cdot Andrea Prati \cdot Marcello Pelillo \cdot Mubarak Shah

Submitted

Person Re-identification



- Recognize an individual over different non-overlapping cameras.
- Given a gallery of person images we want to recognize (between all of them) a new observed image, called probe.



Video-based Person Re-ID

Probe



Traditional methods focus on:

- Building better feature representation of objects
- Building a better distance metric
- Finally rank images from gallery based on the pairwise distances from the query

In our approach

- We use standard features and distance metric
- Extract constrained dominant sets for each query
- Perform ranking over shortlisted clips NOT over the whole set

We take into account **both** the relationship between query and elements in the gallery and elements in the gallery.

Re-ID with Constrained DS's

Probe

Constrained DS's

Final Rank





CNN features with XQDA metric used to compute the edge weights

Results on MARS Dataset

- Largest video Re-ID dataset (2016)
- 6 near-synchronized cameras
- 1,261 identities
- 3,248 distractors
- tracklets are of 25-30 frames long

Methods	rank 1
HLBP $[40] + XQDA$	18.60
BCov [24] + XQDA	9.20
LOMO [20] + XQDA	<u>30.70</u>
BoW $[49] + KISSME$	30.60
SDALF [8] + DVR	4.10
HOG3D [16] + KISSME	2.60
CNN + XQDA [48]	65.30
$\overline{\text{CNN}}$ + KISSME [48]	65.00
Ours	68.22

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Gallery **Probes** nna nnan As the set of the

The green and red boxes denote the same and different persons with the probes, respectively Gallery images are ordered based on their membership score (highest -> lowest).

Multi-target Multi-camera Tracking



Pipeline





- Appearance = CNN features
- Motion = Constant velocity



- Appearance = CNN features
- Motion = Constant velocity



- Appearance = CNN features
- Motion = Constant velocity



- Appearance = CNN features
- Motion = Constant velocity



- Appearance = CNN features
- Motion = Constant velocity



Another data association problem Nodes become tracklets CDSC is used to stitch tracklets







Another data association problem Nodes become tracklets CDSC is used to stitch tracklets



Another data association problem Nodes become tracklets CDSC is used to stitch tracklets

Within-Camera Tracking



Input: Human Detections







Layer 3: Cross-Camera Association



Tracks are nodes Cameras as constraints

Layer 3: Cross-Camera Association



Tracks are nodes Cameras as constraints

Layer 3: Cross-Camera Association



Tracks are nodes Cameras as constraints

Results on DukeMTMC

- Largest MTMC dataset (2016)
- 8 fixed synchronized cameras
- More than 2 million frames
- 0 to 54 persons per frame
- 2,700 Identities

	Methods		IDP	IDR
Multi-Camera Test-easy	[33]	47.3	59.6	39.2
	[26]	32.9	41.3	27.3
	Ours	50.9	63.2	42.6
				Ŕ
	Methods	IDF1↑	IDP↑	IDR↑
Multi-Camera Test-hard	[33]	56.2	67.0	<u>48.4</u>
	[26]	34.9	41.6	30.1
	Ours	60.0	68.3	53.5

MULL I IDDIA IDDA IDDA

- IDP = Fraction of computed detections that are correctly identified
- IDR = Fraction of ground-truth detections that are correctly identified
- IDF1 = Ratio of correctly identified detections over the average number of ground-truth and computed detections





Camera 1

Camera 2





Camera 5

Camera 6





Camera 6

Camera 8



Camera 1

Camera 7



Conclusions

Dominant sets and related concepts shown to be a powerful notion for attacking a variety of computer vision problems, e.g.,

- Interactive image segmentation and cosegmentation
- Geo-localization
- Group detection in image and videos
- Person re-identification
- Multi-target tracking

On-going work focuses on combining deep learning and DS's for improving performances.

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