Elastic Graph Data Structure

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January 31, 2020

>> EG{1}

ans =

struct with fields:

Abeta: [2×101×5×5 double]
beta: [2×101×4 double]
nodeXY: [2×5 double]
A: [5*5 double]
G: [5*8 double]
H: [5×8 double]

Fields in EG structure:

- Abeta: Generalized adjacency matrix whose entry is a curve. size(Abeta,3)=size(Abeta,4)= number of nodes. Abeta(:,:,i,j) is an edge curve between node i and j. All edges are original curves rather than SRVF
- beta: Flattened version of upper triangular of Abeta. size(beta,3) is the number of edges. Refer to .././src/adj2beta.m
- nodeXY: Matrix of node coordinate. nodeXY(:,i) is the coordinate of node i
- A: Binary version of adjacency matrix
- G: Graph incidence matrix. size(G,1) is the number of nodes. size(G,2) is 2 times of number of edges. Row index is the nodes while column index is the edges. Refer to [1] for more details.
- H: Graph incidence matrix. size(H,1) is the number of nodes. size(H,2) is 2 times of number of edges. Row index is the nodes while column index is the edges. Refer to [1] for more details.

Note G and H are only used for graph matching algorithm. Specifically, we are using the algorithm for directed graph matching from [1]. The edge between i and j is treated as 2 virtual directed edges: one from i to j and one from j to i. By doing so, we are able to optimize over a flipped reparametrization. Accordingly, Abeta(:,:,i,j) = fliplr(Abeta(:,:,j,i)). G and H's columns are doubled.

References

[1] Feng Zhou and Fernando De la Torre. Factorized graph matching. *IEEE transactions on pattern analysis and machine intelligence*, 38(9):1774–1789, 2015.