

# Graph-Sparse Decomposition

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## 1 Graph-Sparse Decomposition

We would like to find a low-rank decomposition of an input matrix  $X \in \mathbb{R}^{L \times N}$  with  $N$  gene measurements for  $L$  samples. We'd also like the components of the decomposition to be informed by a prior knowledge graph,  $\mathcal{G}$ , such that the support of each component forms a tree with low-cost edges. We formalize this as the following optimization problem:

$$\arg \min_{D, Z} \|X - ZD\|_2^2 + \lambda \sum_{i=1}^K c(D_i), \quad (1)$$

$$\arg \min_{U, V} \|X - UV^T\|_F^2 + \lambda \sum_{i=1}^K c(U_i), \quad (2)$$

$$f(\mathbf{u}) \xrightarrow{\text{PCST}} \mathbf{u}' \quad (3)$$

$$\arg \min_{T=(V', E')} \pi(\overline{V'}) + c(E') \quad (4)$$

where  $D = \{D_i\} \in \mathbb{R}^{K \times N}$  is a dictionary of components such that the support of each  $D_i$  forms a tree when projected onto  $\mathcal{G}$ , and  $Z \in \mathbb{R}^{L \times K}$  is the corresponding scores matrix that weights each component. The second term adds a sparsity constraint to the optimization in the form  $c(D_i)$ , which is the minimum sum of edges used to connect the support of  $D_i$  in  $\mathcal{G}$  (aka minimum spanning tree). Finally, the  $\lambda$  parameter weights the edge costs and effectively modulates the sparsity of  $D$ .

## 2 Method

To solve this minimization problem, we alternate updates to the score matrix,  $Z$ , and the dictionary,  $D$ .

First, we fix  $D$  and optimize  $Z$ . Because  $D$  is constant, the sparsity term in (5) can be ignored so that

$$Z^* = \arg \min_Z \|X - ZD\|_2^2 = (DD^T)^{-1}DX, \quad (5)$$

where  $Z^*$  is given by the ordinary least squared (OLS) estimator. Next, we update one dictionary component  $D_u$  while keeping the remaining components  $\{D_i\}_{i \neq u}$  and  $Z$  fixed. Breaking the product  $ZD$  into  $K$  individual outer products, (5) becomes

$$D_u^* = \arg \min_{D_u} \left\| X - \sum_{i=1}^K Z_i D_i^T \right\|_2^2 + \lambda \sum_{i=1}^K c(D_i), \quad (6)$$

$$= \arg \min_{D_u} \|X' - Z_u D_u^T\|_2^2 + \lambda c(D_u), \quad (7)$$

where  $X' = X - \sum_{i \neq u} Z_i D_i^T$ . In (7), we have ignored all sparsity terms for components that are not  $D_u$  because they are fixed. Next we note that, because the edge cost function is invariant given the support of  $D_u$ , each element of  $D_u = \{D_u(v)\}_N$  must be either 0 or  $D'_u(v)$ , where

$$D'_u = \arg \min_{D_u} \|X' - Z_u D_u^T\|_2^2. \quad (8)$$

Again,  $D'_u$  can be computed directly via OLS. If we denote  $X'_i$  to be the vector of values for gene  $i$  across all  $L$  samples, we can rewrite our objective as

$$D_u^* = \arg \min_{D_u \in \{0, D'_u(\cdot)\}^N} \sum_{i=1}^N \|X'_i - Z_u D_u(i)\|^2 + \lambda c(D_u) \quad (9)$$

$$= \arg \min_{D_u \in \{0, D'_u(\cdot)\}^N} \sum_{i=1}^N (\|X'_i - Z_u D_u(i)\|^2 - \|X'_i\|^2) + \lambda c(D_u) \quad (10)$$

where we have subtracted the constant value  $\sum_N \|X'_i\|^2$  to obtain (10). Now, we set the prize of node  $i$  to be the amount of error reduced if  $D_u(i)$  is  $D'_u(i)$  instead of 0:

$$p(i) = \|X'_i\|^2 - \|X'_i - Z_u D_u(i)\|^2. \quad (11)$$

Note that  $p(i) = 0$  if  $D_u(i) = 0$ , namely if  $i$  is not in the support (tree) of  $D_u$ . Then we have

$$D_u^* = \arg \min_{D_u} - \sum_{i=1}^N p(i) + \lambda c(D_u) \quad (12)$$

$$= \arg \min_{D_u} - \sum_{i \in \text{supp}(D_u)} p(i) + \lambda c(D_u) \quad (13)$$

$$= \arg \min_{D_u} \sum_{i \notin \text{supp}(D_u)} p(i) + \lambda c(D_u) \quad (14)$$

$$(15)$$

So the solution to (6) can be found using a PCST solver.