Generally, the Laplacian matrix of a graph network is defined as:

$$\mathbf{L} = \mathbf{\Delta} - \mathbf{A} \tag{S1}$$

where A is the adjacency matrix. Δ is given by:

$$\Delta_{ij} = \delta_{ij} \sum_{k=1}^{n} A_{ik} \qquad \qquad \delta_{ij} = \begin{cases} 1 & \text{if } i = j \\ 0 & \text{if } i \neq j \end{cases}$$
(S2)

When use of the transition matrix of a random walk $D^{-1}A$ is favored, the Laplacian matrix is changed into:

$$\mathbf{L}^* = \mathbf{\Delta}^* - \mathbf{D}^{-1}\mathbf{A} \tag{S3}$$

$$\Delta_{ij}^{*} = \delta_{ij} \sum_{k=1}^{n} (\mathbf{D}^{-1} \mathbf{A})_{ik}$$
(S4)

$$\delta_{ij} = \begin{cases} 1 \ if \ i = j \\ 0 \ if \ i \neq j \end{cases}$$

Here, the set of all the eigenvalues λ and the corresponding eigenvectors v for L^{*}, called spectrum, is given by:

$$\mathbf{L}^* \cdot \mathbf{v}_i = \lambda \cdot \mathbf{v}_i \tag{S5}$$

where $\{v_i\}$ provides the node coordinates in the Euclidean space to perform network clustering. Since ADMSC corresponds to the spectral clustering that employs a modified transition matrix $\mathbf{D}^{-\beta}\mathbf{A}$, it can be regarded as the diffusion model-based probabilistic interpretation of spectral analysis.