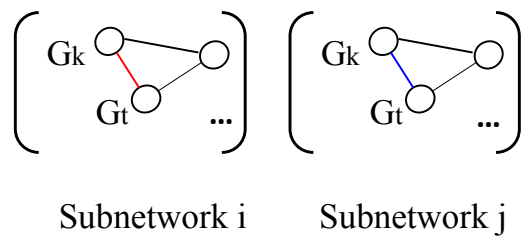


**A**

$$\left( \begin{array}{c} \text{Gk} \\ \text{Gt} \\ \dots \end{array} \right) = W_1 \left| \begin{array}{c} \text{Gk} \\ \text{Gt} \\ \dots \end{array} \right\rangle + W_2 \left| \begin{array}{c} \text{Gk} \\ \text{Gt} \\ \dots \end{array} \right\rangle$$

**B**



**S3 Fig. A diagram illustrating the process of gene co-expression subnetwork identification by COAC. (A)** Each co-expression sub-network can be treated as a superposition of two different gene expression state. **(B)** The gene co-expression relationship can be represented by different subnetworks.