

Supplementary Information (Tables S1 to S4) for Computing persistent homology by spanning trees and critical simplices

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Table S1: Torus triangulation network

This table lists the Morse function values of all simplices, the spanning trees of the boundary matrixes B_1 and B_2 , and the simplices composed of 1- and 2-cavities for the network.

Table S2: C. elegans neural network³³

This table lists the Morse function values of all simplices, the simplices composed of 1-, 2- and 3-cavities, the iterative process of 2-cavities, and an iterative example for the network.

Table S3: BA scale-free model network³⁰

This table lists the Morse function values of all simplices, the simplices composed of 1-order cavities, and the results obtained by Kannan's³⁰ method for the network which is simulated here.

Variables in Kannan's method are as follows:

DFM==Discrete Morse function values;

Flag==To keep track with the size of the set U_α for each simplex α ;

IsCritical==To indicate if a given simplex is critical;

FiltrationWeight==To store the filtration weight corresponding to each simplex.

Table S4: Stanford dragon graphic network³⁴

This table lists the points in the (x, y, z) -coordinates, the present thresholds of all simplices, the persistence barcodes of 1- and 2-cavities calculated by **javaplex** in Ref. [34], the representative cycles with two lengths of the only 2-cavity, and the Morse function values of all simplices obtained by the new method for the network.

Data of **Tables S1 to S4** are available

<https://github.com/ChuangMa1900/Supplementary-Information-Tables-S1-to-S4.git>