



Supplementary Figure 1: The results of TPDS on 4 data-sets when SOM was used with hexagonal geometry. (a) Parkinson's data-set (b) protein expression data-set (c) SCADI data (d) single-cell expression.

Supplementary Table-1 : Parameters and time taken by TPDS for optimal solution for different data-sets

Data-sets	SOM grid size	Lambda	SVD decomposition	Time taken
Parkinson's	2 X 2	1	True	5.6 second
Protein expression	5 X 5	1	True	2.86 minutes
SCADI	7 X 71	1	FALSE	1.26 second
Single-cell expression	3 X 3	1	True	14 minutes