SUPPLEMENTARY NOTES
for
Determining the distance to monotonicity of a biological network: a graph-theoretical approach

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1   Performances of Algorithm 4

Table 1: Comparison between δ_{low} obtained by Algorithm 4 and δ_{low} obtained by the simple procedure of iteratively deleting the shortest negative cycles.

	δ_{low} (short cycle deletion)	δ_{low} (Algorithm 4)
EGFR	176	**186**
macrophage	288	**302**
yeast	40	**41**
E.coli	336	**365**

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