**Table S7:** Robustness analysis results for selected network motifs <sup>a,b</sup>

Motif	Core network	5% removal	5% addition
miR Gene	Z score = 9.22	Z score = $8.79 \pm 0.41$ (p-value = $0.22$ )	$Z \text{ score} = 7.23 \pm 1.16$ (p-value = 0.10)
Gene	Z score = 9.22	Z score = $7.06 \pm 3.02$ (p-value = $0.61$ )	$Z \text{ score} = 5.18 \pm 3.06$ (p-value = 0.19)
↑↓	Z score = 7.02	Z score = $7.05 \pm 3.05$ (p-value = $0.34$ )	Z score = $6.03 \pm 2.22$ (p-value = $0.43$ )
miR TA	Z score = 2.52	Z score = $2.67 \pm 0.36$ (p-value = $0.43$ )	$Z \text{ score} = 2.57 \pm 0.43$ (p-value = 0.85)
miR Gene	Z score = 11.02	Z score = $8.50 \pm 4.17$ (p-value = $0.32$ )	$Z \text{ score} = 10.10 \pm 2.27$ (p-value = 0.48)
Gene	Z score = 8.22	$Z \text{ score} = 8.02 \pm 0.60$ (p-value = 0.56)	$Z \text{ score} = 6.83 \pm 1.14$ (p-value = 0.07)

<sup>&</sup>lt;sup>a</sup> Z-scores were represented by mean  $\pm$  standard error of mean (SEM).

<sup>&</sup>lt;sup>b</sup> The statistical significance of z-scores (p-value) was exhibited using two-tailed Student's T-test.