miRNA targets

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May 19, 2011

Abstract

This document is generated by Anduril [1] (Engine 1.2.0) and it provides a human readable summary for the network outputs.

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1 hsa-mir-106b down-regulate ENSG00000141503

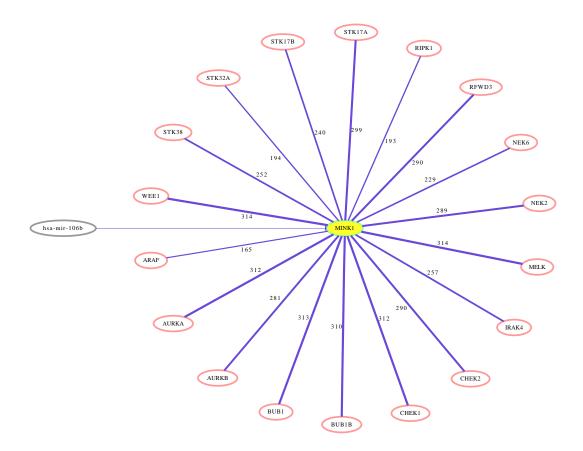


Figure 1: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

2 hsa-mir-106b down-regulate ENSG00000153233

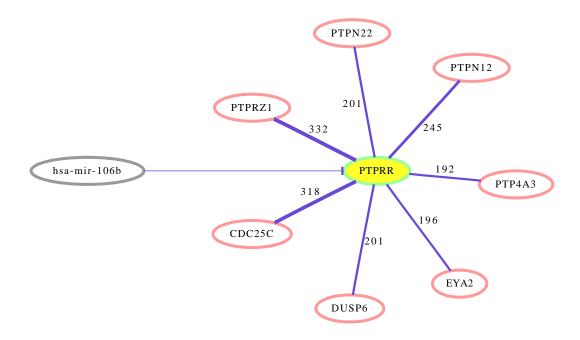


Figure 2: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

3 hsa-mir-142-5p down-regulate ENSG00000088179

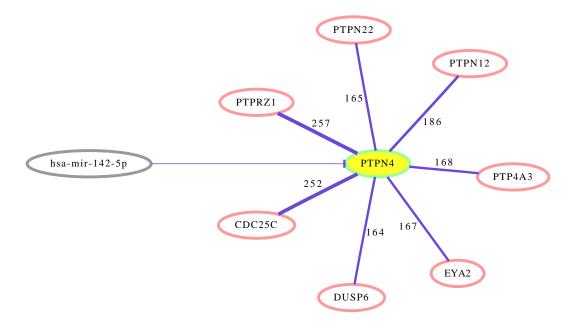


Figure 3: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

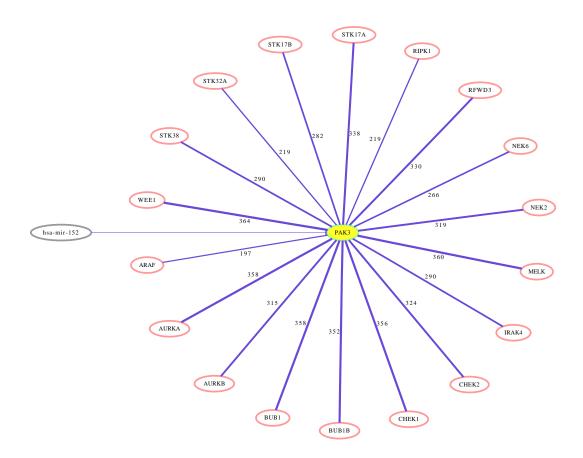


Figure 4: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

5 hsa-mir-155 down-regulate ENSG00000002746

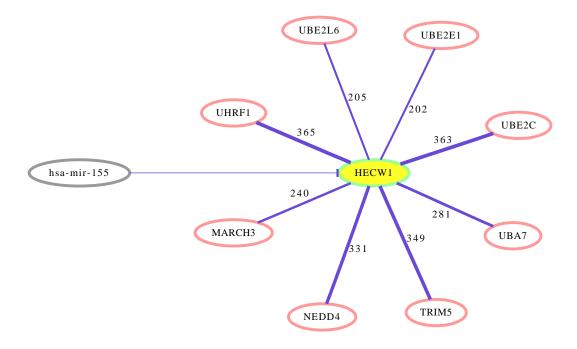


Figure 5: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

6 hsa-mir-155 down-regulate ENSG00000067606

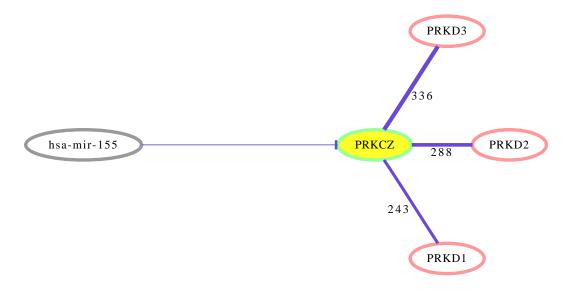


Figure 6: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

7 hsa-mir-155 down-regulate ENSG00000077264

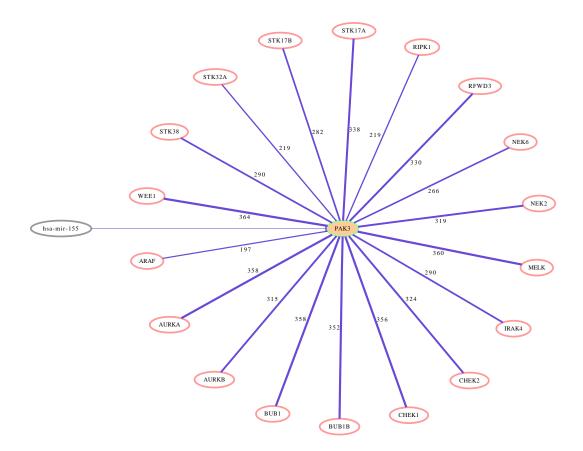


Figure 7: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

8 hsa-mir-155 down-regulate ENSG00000101349

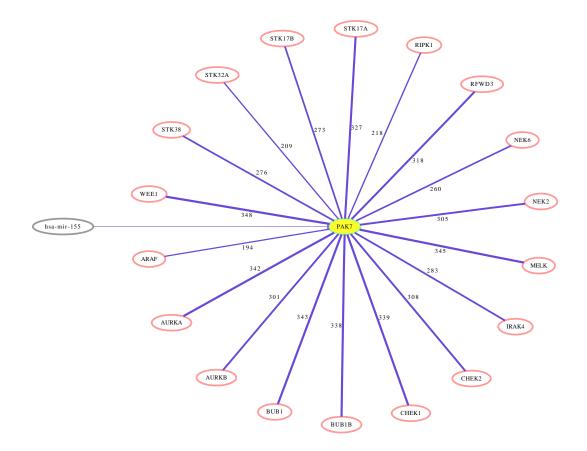


Figure 8: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

9 hsa-mir-155 down-regulate ENSG00000154914

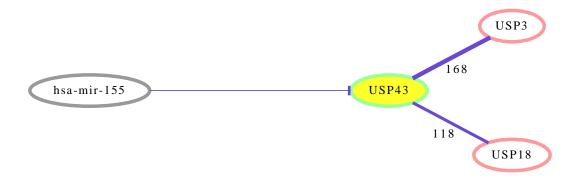


Figure 9: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

$10 \quad \text{hsa-mir-15b down-regulate ENSG00000153233}$

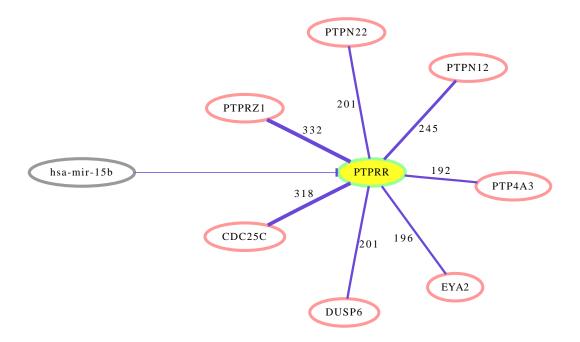


Figure 10: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

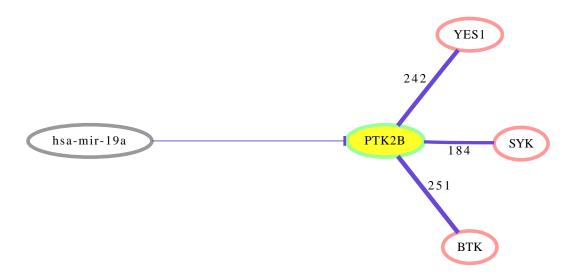


Figure 11: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

12 hsa-mir-19b down-regulate ENSG00000120899

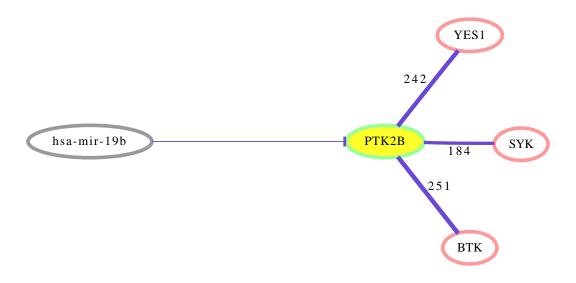


Figure 12: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

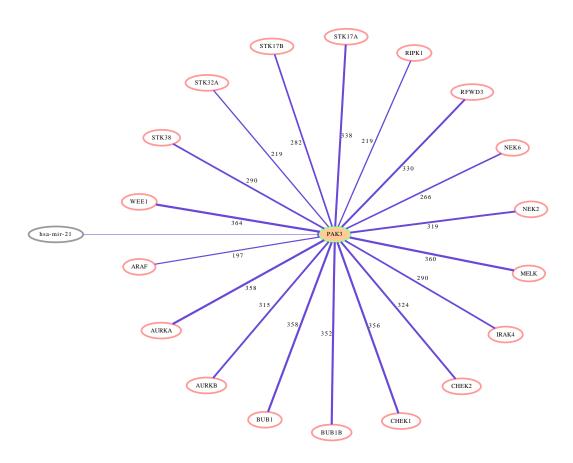


Figure 13: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

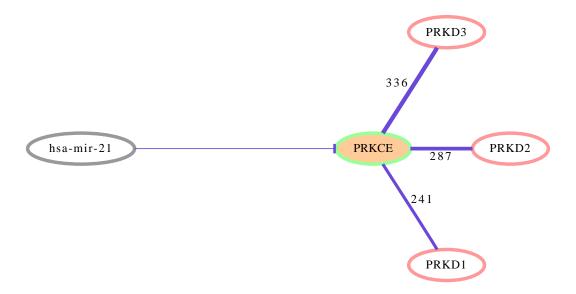


Figure 14: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

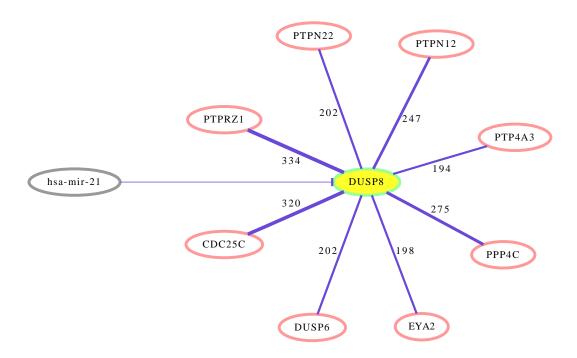


Figure 15: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

$16 \quad \text{hsa-mir-} 210 \ \text{down-regulate ENSG} 00000068383$



Figure 16: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

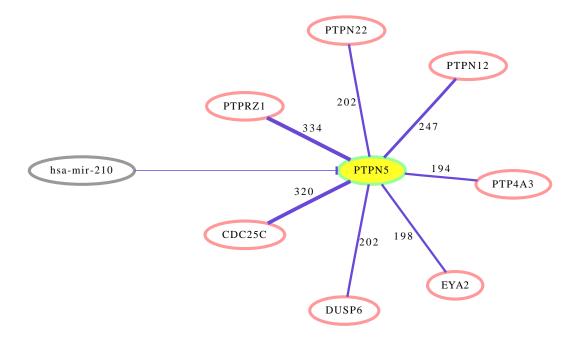


Figure 17: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

18 hsa-mir-23a down-regulate ENSG00000008086

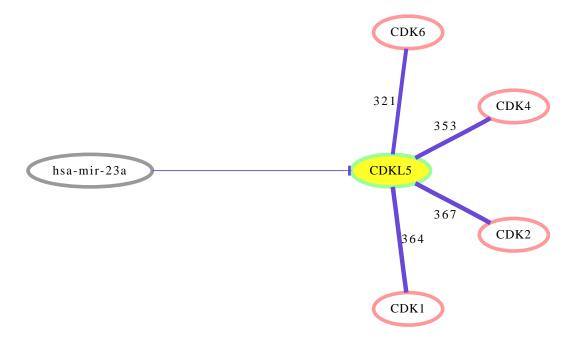


Figure 18: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

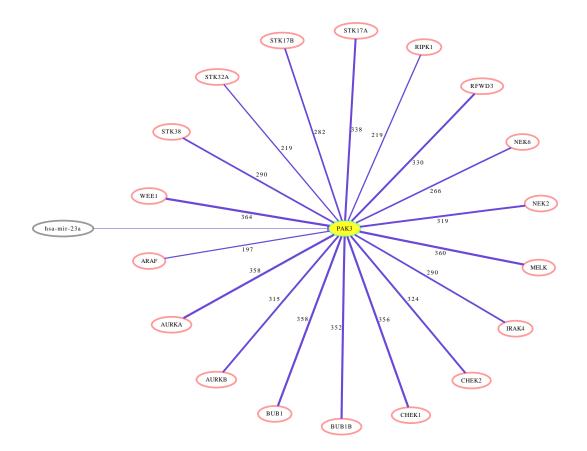


Figure 19: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

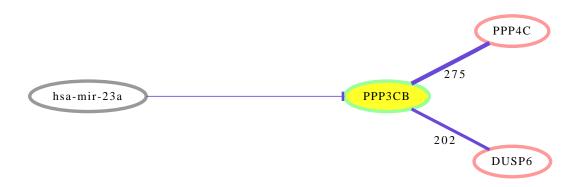


Figure 20: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

$21 \quad \text{hsa-mir-} 23 \text{a down-regulate ENSG} 00000126583$

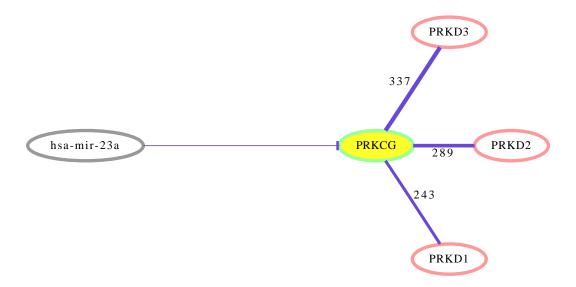


Figure 21: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

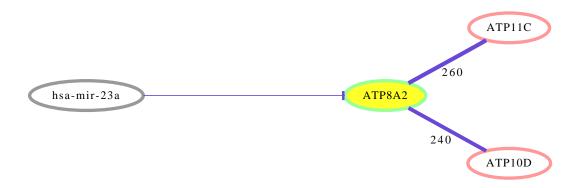


Figure 22: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

$23 \quad \text{hsa-mir-} 23 \text{a down-regulate ENSG} 00000138769$

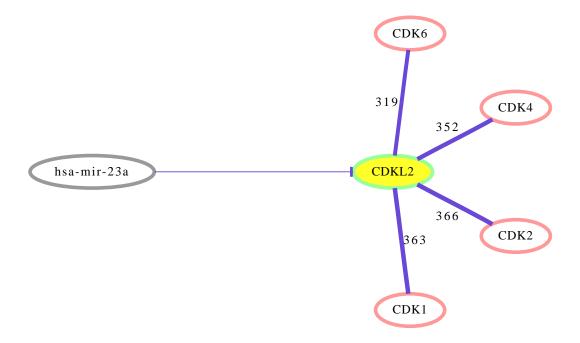


Figure 23: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

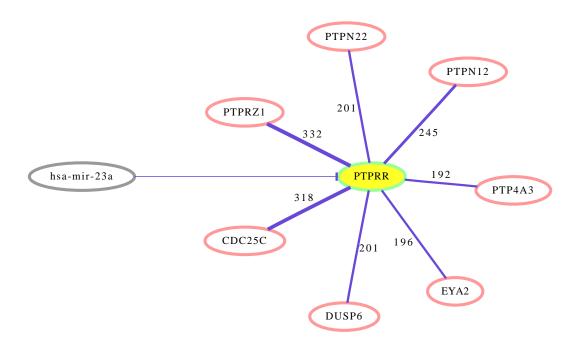


Figure 24: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

$25 \quad \text{hsa-mir-} 25 \ \text{down-regulate ENSG} 00000099308$

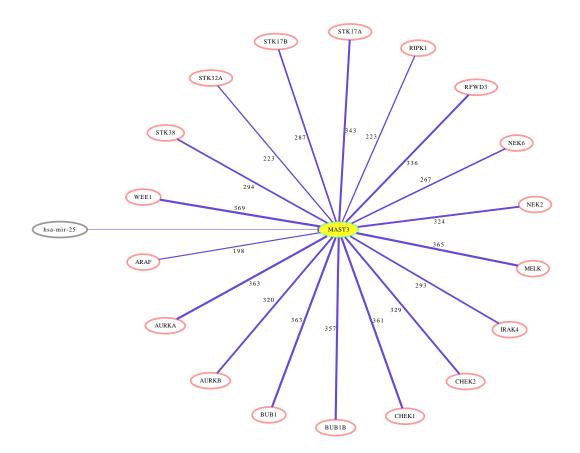


Figure 25: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

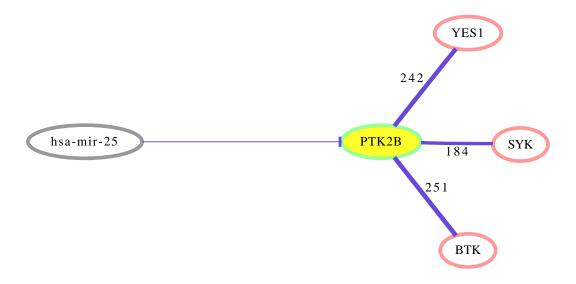


Figure 26: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

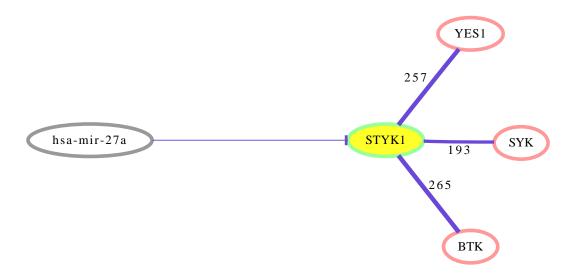


Figure 27: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

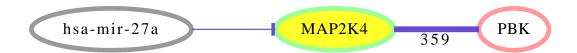


Figure 28: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

$29 \quad \text{hsa-mir-} 27 \text{a down-regulate ENSG} 00000077264$

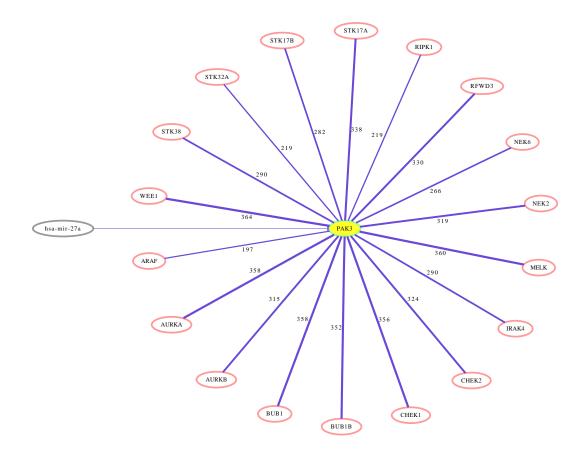


Figure 29: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

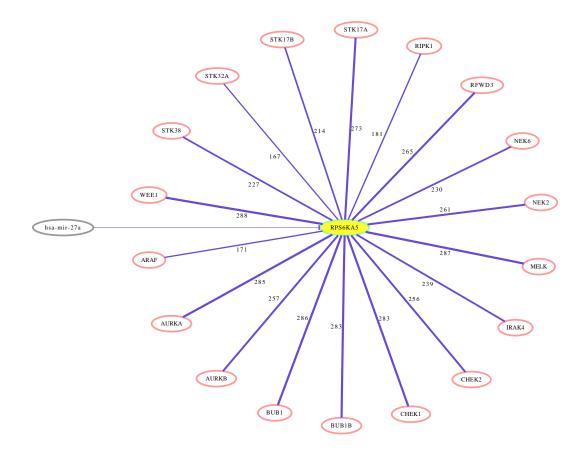


Figure 30: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.



Figure 31: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

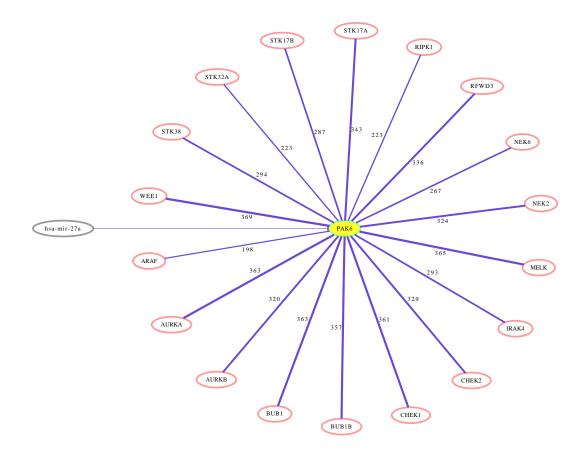


Figure 32: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

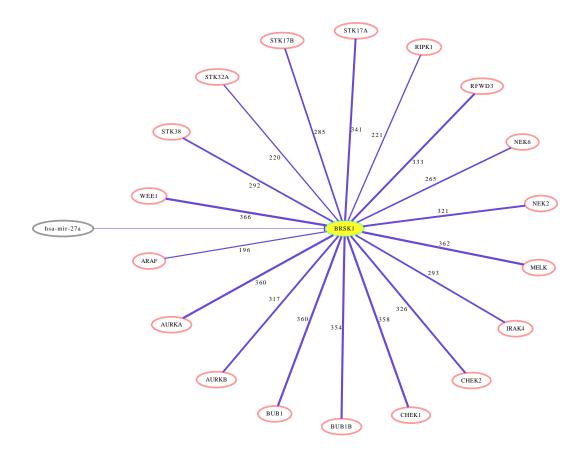


Figure 33: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

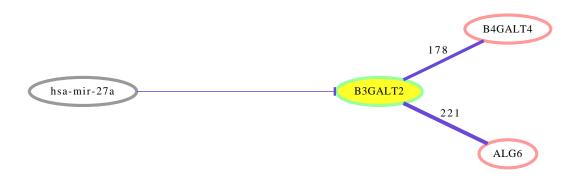


Figure 34: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

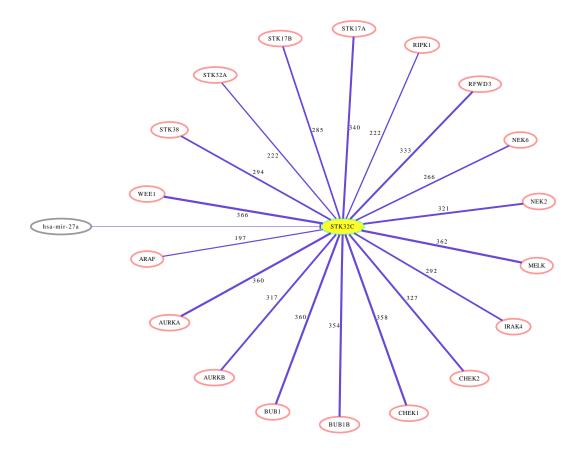


Figure 35: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

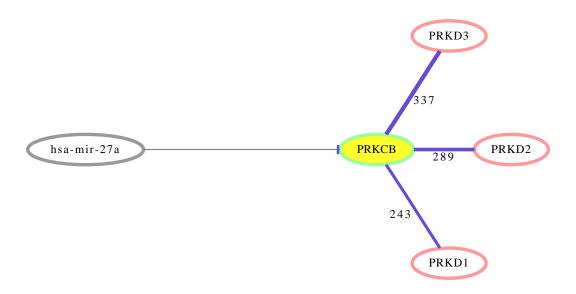


Figure 36: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

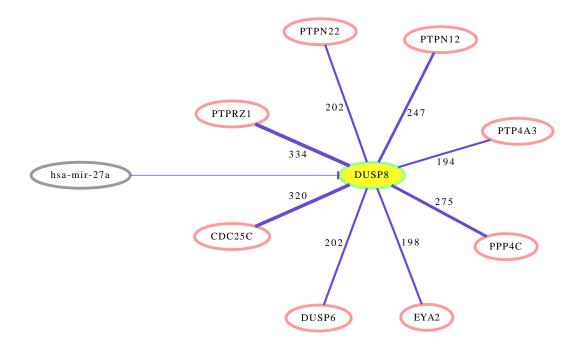


Figure 37: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

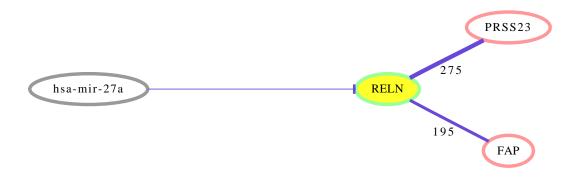


Figure 38: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

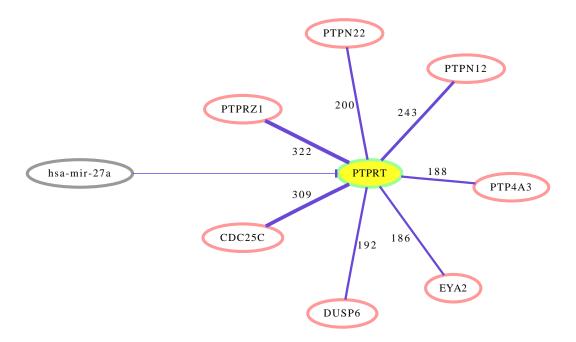


Figure 39: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

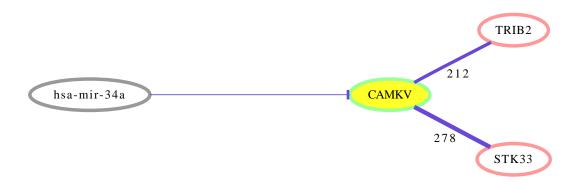


Figure 40: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

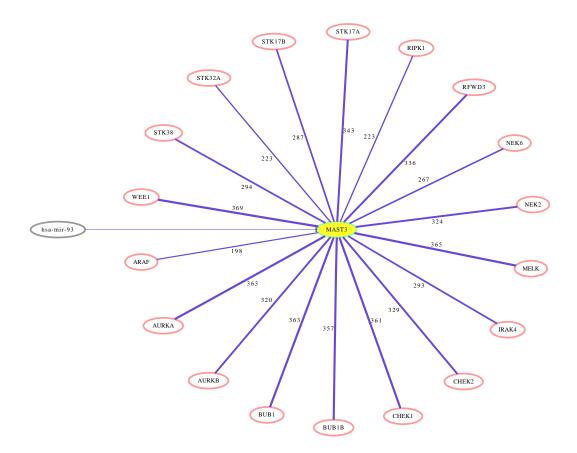


Figure 41: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

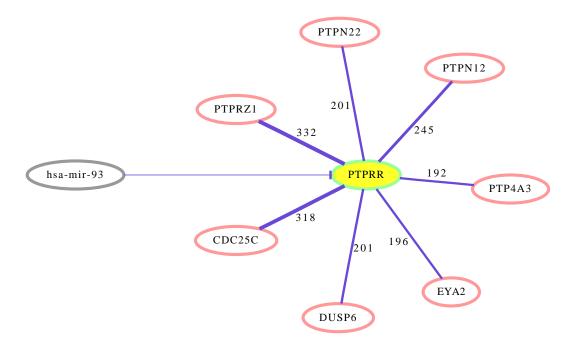


Figure 42: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

$43 \quad \text{hsa-mir-93 down-regulate ENSG00000169032}$



Figure 43: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

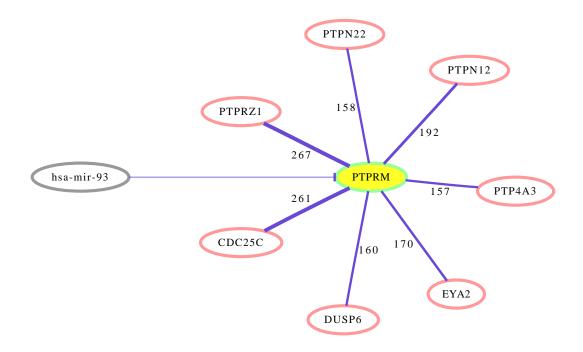


Figure 44: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.



Figure 45: Enzyme pairs. Up or down regulated genes are in red or green. Enzyme pair frequencies are displayed on edges. Nodes filled with orange or yellow are validated or predicted miRNA targets in databases.

References

[1] K. Ovaska, M. Laakso, S. Haapa-Paananen, R. Louhimo, P. Chen, V. Aittomäki, E. Valo, J. Núñez-Fontarnau, V. Rantanen, S. Karinen, K. Nousiainen, A.-M. Lahesmaa-Korpinen, M. Miettinen, L. Saarinen, P. Kohonen, J. Wu, J. Westermarck, and S. Hautaniemi. Large-scale data integration framework provides a comprehensive view on glioblastoma multiforme. *Genome Medicine*, 2(9):65, September 2010.