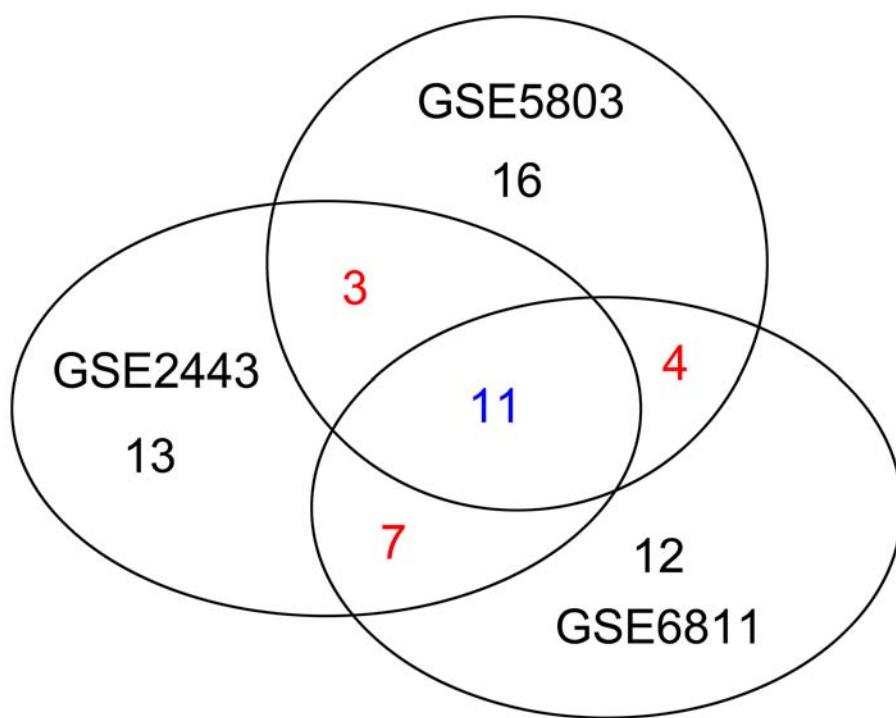


SUPPLEMENTARY FIGURE AND TABLES



Supplementary Figure S1: Venn diagram illustrating the overlap in microRNA detection across the three data sets.

**Supplementary Table S1: The predicted outlier miRNAs in each dataset**

| miRNA ID         | GSE2443 | GSE5803 | GSE6811 |
|------------------|---------|---------|---------|
| hsa-miR-1        | ✓       | ✓       | ✓       |
| hsa-miR-210      | ✓       | ✓       | ✓       |
| hsa-miR-218(-5p) | ✓       | ✓       | ✓       |
| hsa-miR-155      | ✓       | ✓       | ✓       |
| hsa-miR-145(-5p) | ✓       | ✓       | ✓       |
| hsa-miR-122(-5p) | ✓       | ✓       | ✓       |
| hsa-miR-197(-3p) | ✓       | ✓       | ✓       |
| hsa-miR-346      | ✓       | ✓       | ✓       |
| hsa-let-7b(-5p)  | ✓       | ✓       | ✓       |
| hsa-miR-125b     | ✓       | ✓       | ✓       |
| hsa-miR-124      | ✓       | ✓       | ✓       |
| hsa-miR-149(-5p) | ✓       |         | ✓       |
| hsa-miR-585      |         | ✓       | ✓       |
| hsa-miR-146a     |         | ✓       | ✓       |
| hsa-miR-326      | ✓       | ✓       |         |
| hsa-miR-593      | ✓       | ✓       |         |
| hsa-miR-626      | ✓       | ✓       |         |
| hsa-miR-339(-3p) | ✓       |         | ✓       |
| hsa-miR-375      | ✓       |         | ✓       |
| hsa-miR-192      | ✓       |         | ✓       |
| hsa-miR-126      | ✓       |         | ✓       |
| hsa-miR-627      | ✓       |         | ✓       |
| hsa-miR-938      | ✓       |         | ✓       |
| hsa-miR-490(-3p) |         | ✓       | ✓       |
| hsa-miR-378      |         | ✓       | ✓       |

**Supplementary Table S2: Highly enriched (P-value< 0.05) KEGG pathways by target genes of 8 experimentally validated key miRNAs**

| Pathway Term   | Pathway Class                        | Subclass                 | P-Value  |
|--|--------------------------------------|--------------------------|----------|
| 1. Pancreatic cancer   | Human Diseases                       | Cancers                  | 6.44E-06 |
| 2. Pathways in cancer  | Human Diseases                       | Cancers                  | 9.18E-06 |
| 3. Small cell lung cancer                                      | Human Diseases                       | Cancers                  | 3.27E-05 |
| 4. Chronic myeloid leukemia                                    | Human Diseases                       | Cancers                  | 5.47E-05 |
| 5. Glioma  | Human Diseases                       | Cancers                  | 6.02E-05 |
| 6. Colorectal cancer   | Human Diseases                       | Cancers                  | 1.57E-04 |
| 7. Non-small cell lung cancer                                  | Human Diseases                       | Cancers                  | 5.44E-04 |
| 8. Bladder cancer  | Human Diseases                       | Cancers                  | 5.85E-04 |
| 9. Melanoma  | Human Diseases                       | Cancers                  | 8.00E-04 |
| 10. Prostate cancer  | Human Diseases                       | Cancers                  | 1.08E-03 |
| 11. Adherens junction  | Cellular Processes                   | Cell Communication       | 1.44E-03 |
| 12. Endometrial cancer   | Human Diseases                       | Cancers                  | 2.15E-03 |
| 13. Apoptosis  | Cellular Processes                   | Cell Growth and Death    | 3.39E-03 |
| 14. Cell cycle   | Cellular Processes                   | Cell Growth and Death    | 4.45E-03 |
| 15. MAPK signaling pathway                                     | Environmental Information Processing | Signal Transduction      | 5.95E-03 |
| 16. p53 signaling pathway                                      | Cellular Processes                   | Cell Growth and Death    | 9.71E-03 |
| 17. Epithelial cell signaling in Helicobacter pylori infection | Human Diseases                       | Cardiovascular Diseases  | 1.60E-02 |
| 18. Acute myeloid leukemia                                     | Human Diseases                       | Cancers                  | 1.60E-02 |
| 19. Dilated cardiomyopathy                                     | Human Diseases                       | Cardiovascular Diseases  | 1.73E-02 |
| 20. Arrhythmogenic right ventricular cardiomyopathy (ARVC)     | Human Diseases                       | Cardiovascular Diseases  | 2.51E-02 |
| 21. Focal adhesion   | Cellular Processes                   | Cell Communication       | 2.91E-02 |
| 22. Endocytosis  | Cellular Processes                   | Transport and Catabolism | 3.00E-02 |
| 23. Hypertrophic cardiomyopathy (HCM)                          | Human Diseases                       | Cardiovascular Diseases  | 3.35E-02 |
| 24. TGF-beta signaling pathway                                 | Environmental Information Processing | Signal Transduction      | 3.85E-02 |
| 25. Viral myocarditis  | Human Diseases                       | Cardiovascular Diseases  | 3.88E-02 |
| 26. Regulation of actin cytoskeleton                           | Cellular Processes                   | Cell Communication       | 4.71E-02 |
| 27. Tight junction   | Cellular Processes                   | Cell Communication       | 4.71E-02 |
| 28. Jak-STAT signaling pathway                                 | Environmental Information Processing | Signal Transduction      | 4.86E-02 |
| 29. B cell receptor signaling pathway                          | Organismal Systems                   | Immune System            | 4.89E-02 |

**Supplementary Table S3: Top 50 GeneGo pathways enriched by target genes of 8 experimentally validated key miRNAs**

| Pathway Term   | Pathway Map Folder      | P-Value  |
|--|-------------------------|----------|
| 1. Influence of Ras and Rho proteins on G1/S Transition                        | Cell cycle              | 6.68E-09 |
| 2. ESR1 regulation of G1/S transition  | Cell cycle              | 9.59E-08 |
| 3. Regulation of G1/S transition (part 1)                                      | Cell cycle              | 3.62E-07 |
| 4. Cell cycle (generic schema)   | Cell cycle              | 5.87E-07 |
| 5. TGF, WNT and cytoskeletal remodeling  | Cytoskeleton remodeling | 8.68E-07 |
| 6. AKT signaling   | Signal transduction     | 1.12E-06 |
| 7. Regulation of G1/S transition (part 2)                                      | Cell cycle              | 2.97E-06 |
| 8. Role of SCF complex in cell cycle regulation                                | Cell cycle              | 6.61E-06 |
| 9. Brca1 as a transcription regulator  | DNA damag               | 8.43E-06 |
| 10. PIP3 signaling in cardiac myocytes   | Development             | 2.26E-05 |
| 11. Endothelial cell contacts by junctional mechanisms                         | Cell adhesion           | 4.09E-05 |
| 12. Role of Activin A in cell differentiation and proliferation                | Development             | 6.20E-05 |
| 13. CD16 signaling in NK cells   | Immune response         | 6.30E-05 |
| 14. RalA regulation pathway  | Cytoskeleton remodeling | 9.68E-05 |
| 15. FGF2-dependent induction of EMT  | Development             | 1.22E-04 |
| 16. Thrombopoietin-regulated cell processes                                    | Development             | 1.35E-04 |
| 17. Chromosome condensation in prometaphase                                    | Cell cycle              | 1.57E-04 |
| 18. Caspase cascade  | Apoptosis and survival  | 1.69E-04 |
| 19. Normal and pathological TGF-beta-mediated regulation of cell proliferation | Development             | 1.69E-04 |
| 20. CREB pathway   | Transcription           | 1.79E-04 |
| 21. Macropinocytosis regulation by growth factors                              | Transport               | 1.96E-04 |
| 22. LRRK2 and immune function in Parkinson's disease                           | Disease                 | 2.00E-04 |
| 23. Chemokines and adhesion  | Cell adhesion           | 2.40E-04 |
| 24. Regulation of actin cytoskeleton by Rho GTPases                            | Cytoskeleton remodeling | 2.50E-04 |
| 25. Cross-talk between Ras-family GTPases                                      | G-protein signaling     | 2.50E-04 |
| 26. Cytoskeleton remodeling  | Cytoskeleton remodeling | 2.83E-04 |
| 27. VEGF signaling via VEGFR2 - generic cascades                               | Development             | 2.93E-04 |
| 28. Endothelial cell contacts by non-junctional mechanisms                     | Cell adhesion           | 3.09E-04 |
| 29. IGF-1 receptor signaling   | Development             | 3.42E-04 |
| 30. Nucleocytoplasmic transport of CDK/Cyclins                                 | Cell cycle              | 3.50E-04 |
| 31. K-RAS regulation pathway   | G-protein signaling     | 3.78E-04 |

(Continued)

|     | Pathway Term  | Pathway Map Folder         | P-Value  |
|-----|---|----------------------------|----------|
| 32. | Role of HDAC and calcium/calmodulin-dependent kinase (CaMK) in control of skeletal myogenesis | Development                | 4.34E-04 |
| 33. | Prolactin receptor signaling  | Development                | 6.75E-04 |
| 34. | Histamine H1 receptor signaling in the interruption of cell barrier integrity                 | Cell adhesion              | 9.62E-04 |
| 35. | Fibronectin-binding integrins in cell motility  | Cytoskeleton remodeling    | 1.06E-03 |
| 36. | ACM regulation of nerve impulse   | Neurophysiological process | 1.08E-03 |
| 37. | MIF - the neuroendocrine-macrophage connector   | Immune response            | 1.08E-03 |
| 38. | Alpha-1 adrenergic receptors signaling via cAMP   | Development                | 1.22E-03 |
| 39. | PDGF signaling via STATs and NF- $\kappa$ B   | Development                | 1.24E-03 |
| 40. | NF-AT signaling in Cardiac Hypertrophy  | Cardiac Hypertrophy        | 1.34E-03 |
| 41. | Histamine H1 receptor signaling in immune response  | Immune response            | 1.36E-03 |
| 42. | Integrin-mediated cell adhesion and migration   | Cell adhesion              | 1.36E-03 |
| 43. | Inhibition of telomerase activity and cellular senescence                                     | DNA damage                 | 1.50E-03 |
| 44. | Oncostatin M signaling via JAK-Stat in human cells  | Immune response            | 1.50E-03 |
| 45. | Integrin outside-in signaling   | Cytoskeleton remodeling    | 1.51E-03 |
| 46. | Erk Interactions: Inhibition of Erk   | Signal transduction        | 1.64E-03 |
| 47. | Cytoplasmic/mitochondrial transport of proapoptotic proteins Bid, Bmf and Bim                 | Apoptosis and survival     | 1.64E-03 |
| 48. | Some pathways of EMT in cancer cells  | Development                | 1.87E-03 |
| 49. | EPO-induced Jak-STAT pathway  | Development                | 1.87E-03 |
| 50. | Role of heterochromatin protein 1 (HP1) family in transcriptional silencing                   | Transcription              | 2.18E-03 |

**Supplementary Table S4: Literature validated GeneGO pathways from enrichment of genes targeted by 8 key miRNAs**

| Pathway Term   | Enrichment Genes   | Enrichment miRNAs                      | PMID                          |
|--|--|--|-------------------------------|
| Regulation of G1/S transition (part 1)                                     | CDKN1B;FBXW7;CDKN1A;ZFYVE9;CCNA2;CCND2;ANAPC1;CCND1;CDK6             | miR-218;miR-197;miR-145;let-7b         | 19646263;15216909             |
| Cell cycle (generic schema)  | E2F3;E2F1;CCNA2;CCND2;E2F5;E2F6;CCND1;CDK6                           | miR-210;miR-197;miR-145;let-7b         | 19632176;18097561             |
| AKT signaling  | IKBKB;CDKN1B;IGF1R;FOXO3;CDKN1A;IGF1R;IRS1;MYC;AKT3;GYS1;CCND2;CCND1 | miR-218;miR-197;miR-145;miR-122;let-7b | 14683476;21681193<br>21317204 |
| Regulation of G1/S transition (part 2)                                     | E2F1;CCNA2;AKT3;CCNA2;CCND2;CCND1;CDK6                               | miR-197;miR-145;miR-122;let-7b         | 19646263;15216909             |
| RalA regulation pathway  | ACTG1;RGL2;CALM1;HSF1;RALGDS;PGGTB;RAC1                              | miR-210;miR-149;miR-145;miR-122;let-7b | 16964283                      |
| Normal and pathological TGF-beta mediated regulation of cell proliferation | ZFYVE9;MYC;RAC1;PDGFRA;CCND1   | miR-145;miR-122;let-7b                 | 18163430                      |
| CREB pathway   | IGF1R;PRKAR2A;CACNA1D;CALM1;IRS1;RAC1;AKT3;CCND1                     | miR-197;miR-149;miR-145;miR-122;let-7b | 22710715                      |