

Supplementary data

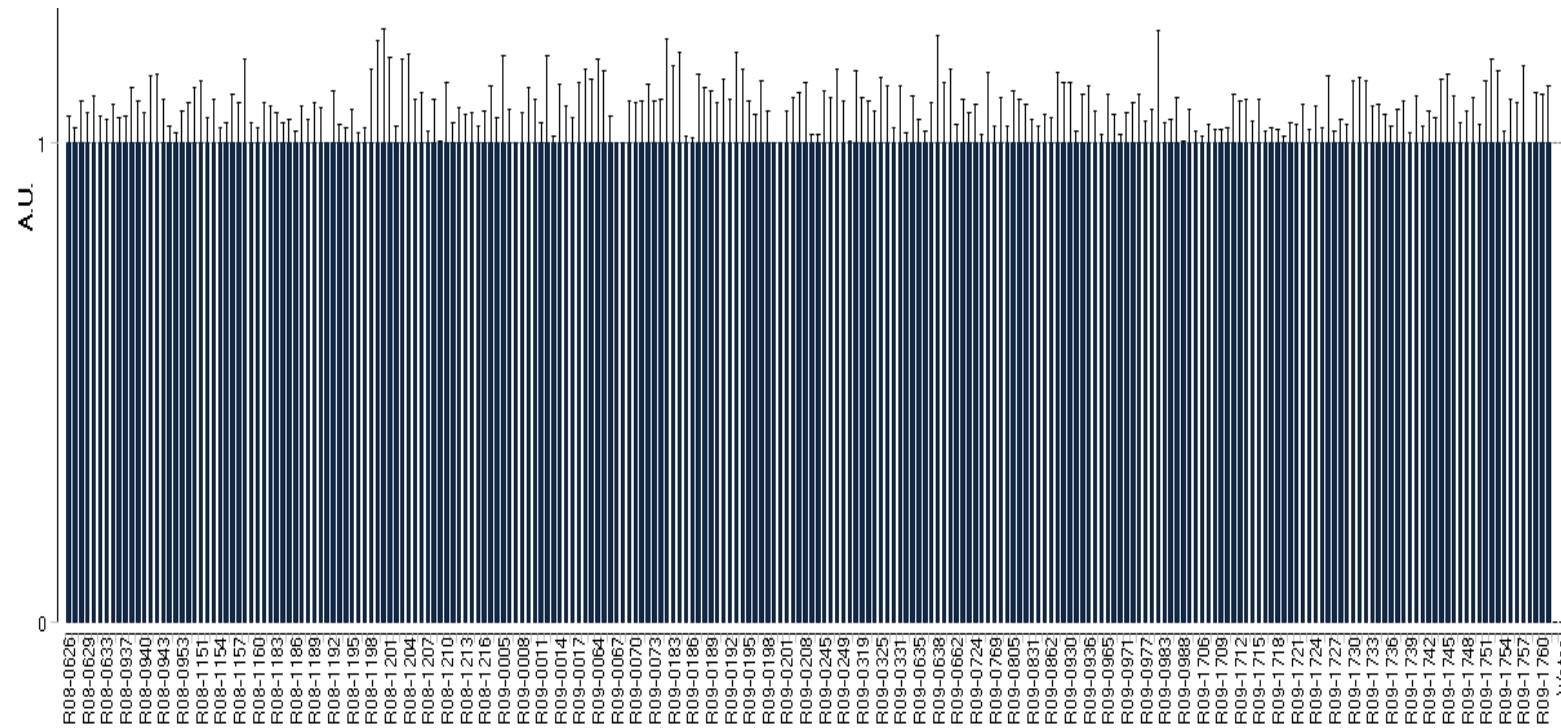
Table 1. MiRNAs with altered expression in malignancy

Tissue/tumor type	Increased expression	Decreased expression
Breast ^{21, 22}	miR-21, miR-29b-2	miR-125b, miR-145 miR-10b, miR-155, miR-17-5p, miR-27b
Ovarian ^{23, 24}	miR-141, miR-200(a-c), miR-221	let-7f, miR-140, miR-145, miR199a, miR-424
Endometrial ²⁵⁻²⁷	miR-103, miR-107, miR-185, miR-205, miR-210, miR-449	miR-99b, miR-152, miR-193, miR-204, miR-221, let-7i
Glioblastoma ^{28, 29}	miR-221, miR-21	miR-181a, miR-181b, miR-181c
Chronic lymphocytic leukemia ⁸		miR-15, miR-16
Lymphoma ^{28, 30}	miR-155, miR-17-92cluster	miR-15a
Colorectal ^{9, 28-31}	miR-10a, miR-17-92 cluster, miR-20a, miR-24-1, miR-29b-2, miR-31	miR-143, miR-145, let-7
Thyroid ^{28, 29}	miR-221, miR-222, miR-146, miR-181b, miR-197, miR-346	
Hepatocellular ^{28, 29}	miR-18, miR-224	miR-199a, miR-195, miR-200a, miR-125a
Testicular ³⁰	miR-372, miR-373	
Pancreatic ²⁸⁻³⁰	miR-221, miR-376a, miR301, miR-21, miR-24-2, miR-100, miR-103-1,2, miR-107, miR-125b-1	miR-375
Cholangiocarcinoma ²⁹	miR-21, miR-141, miR-200b	
Prostate ^{30, 32-34}	let-7d, miR-195, miR-203	miR-128a
Gastric ²⁸⁻³⁰	miR-223, miR-21, miR-103-2	miR-218-2
Lung ^{13, 28-30}	mir-17-92 cluster, miR-17-5p	let-7 family

Supplementary Data Figure 1.

miR-16 expression

(226 samples: breast cancer cases and controls)



MiR-16 expression in whole blood samples of all 226 participants. Systemic miR-16 is stably expressed across all samples and thus is a suitable reference gene for normalising RQ-PCR data in this study