

Supplementary Table 2: List of significantly ($p < .05$) differentially expressed probes on microarray comparison of NK-AML samples versus normal bone marrow. Only those microRNAs with a FC of ≥ 2 in either direction were included in this table. There were 26 overexpressed probes and 11 under expressed probes in NK-AML versus normal BM. The microRNAs depicted in bold represent proprietary miRPlus™ probe whose sequences have subsequently been annotated on miRBase. Those probes denoted as miRPlus™ have not been further annotated by the release of miRBase release 15.

Upregulated miRNAs		Downregulated miRNAs	
miR- probe	F C	miR- probe	F C
miR-10a	19.6	miR-451	-5.7
let-7b	6.0	miR-22	-3.8
miR-20a	4.8	miR-plus (17856)	-3.0
let-7c	4.7	miR-plus (17952)	-2.8
miR-222	4.6	miR-483	-2.5
miR-221	4.4	miR-125-3p	-2.5
miR-17-5p	4.3	miR-21	-2.3
miR-196b	4.3	miR-150	-2.2
miR-20a	3.8	miR-744	-2.1
let-7a	3.8	miR-plus (17881)	-2.1
miR-181a	3.7	miR-342	-2.0
miR-86-5p	3.5		
miR-363*	3.5		
miR-181b	3.3		
miR-155	2.8		
miR-plus (17848)	2.7		
miR-34a	2.7		
miR-20b	2.7		
miR-23*	2.6		
miR-19b	2.5		
miR-181a	2.5		
miR-21*	2.4		
miR-191	2.3		
miR-17-3p	2.1		
miR-345	2.1		
miR-plus (17818)	2.0		