

Table S2

Positions of MassARRAY amplicons in hg18 and sequences of MassARRAY primers used for DNA methylation analysis

chr	start	end	amplicon name	10F primer sequence	T7R primer sequence
chr9	21,549,054	21,549,490	mir-31	aggaagagagTTTTTTTAAGAAGGGAAAGTTAGTTA	cagtaatacgactcactataggagaaggctCAAATAAACTAAAAAACCTTAATCCC
chr11	57,161,896	57,162,383	mir-130a	aggaagagagAGGTTTTTATTGGTTGTGGGTTAAT	cagtaatacgactcactataggagaaggctTTCTAAAACTACCTCTTCTAACCA
chr22	44,856,162	44,856,743	let-7a-3/let7b	aggaagagagTGGGTGGTATGAATTTTTAGTTTG	cagtaatacgactcactataggagaaggctAAAATCCCTCAAAAACCTTAAATC
chr1	98,284,168	98,284,631	mir-137	aggaagagagGTTTAGTGGTTAGTTGGTGGTTG	cagtaatacgactcactataggagaaggctATACCCAAAAACCTTACCACATCT
chr21	25,855,810	25,856,217	mir-155	aggaagagagTGTTTGTGTTTGGAAATTTATAGTTTGA	cagtaatacgactcactataggagaaggctCCATTTCTTCTCTTAAAAACCTA
chr16	68,516,150	68,516,613	mir-140	aggaagagagGGTTAAGTGAGTTTTGGAAAAGGT	cagtaatacgactcactataggagaaggctCAACTTCAAATTCAAACAAAAAAA
chr1	40,947,229	40,947,768	mir-30e/mir-30c-1	aggaagagagTAGGAGATGTAGAAATATGAGTTTGGAA	cagtaatacgactcactataggagaaggctAAACTCTAACCAACCAATAAAAAATCCC
chr7	25,956,149	25,956,538	mir-148a	aggaagagagGTTTTAGTTTTTTAGATGGGAAGGT	cagtaatacgactcactataggagaaggctATCTTCTCCAAAAATTTCTCTCTA
chr11	110,888,268	110,888,617	mir-34b/mir-34c	aggaagagagTTTTTTTGGAGTTTTTTAGGGAT	cagtaatacgactcactataggagaaggctAAAAATACCAAACCTCCCTTC
chr21	16,828,628	16,829,184	mir-99a/let-7c	aggaagagagTATTTTTGGTTGATGGGTAGAAAAA	cagtaatacgactcactataggagaaggctCCAATCAAAAAAACAAAAAAACA
chr11	121,476,136	121,476,612	mir-125b-1	aggaagagagTATAAATGATTGAAGAGAAAATAGGAGGT	cagtaatacgactcactataggagaaggctCTCTAAAAAAATTTACTACTCCCC
chr21	16,882,554	16,882,936	mir-125b-2	aggaagagagTTGTTTTTATTGATTGATTGTTT	cagtaatacgactcactataggagaaggctCCCTAAAAAAATAAAAAATTTCC