









miR-199-5p	CCCAGUGUUCAGACUACCUGUU				2
miR-199-3p	UACAGUAGUCUGCACAUUGGUU		1		
miR-200a	UACACUGUCUGGUAACGAUG		1		
miR-200b	UAAUACUGCCUGGUAUGAUGAC	2		1	
miR-201 <sup>c</sup>	UACUCAGUAAGGCAUUGUUCU			1	
miR-202	AGAGGUAUAGCGCAUGGGAAGA			1	
miR-203	UGAAAUGUUUAGGACCACUAGA		2	1	
miR-204	UUCCCUUUGUCAUCCUAUGCCUG				1
miR-205	UCCUUCAUCCACCGGAGUCUG		1		
miR-206 <sup>d</sup>	UGGAAUGUAAGGAAGUGUGUGG		2		
miR-207	GCUUCUCCUGGCUCUCCUCCUC				1
miR-208	AUAAGACGAGCAAAAAGCUUGU	1			

Mouse and human miRNA sequences. The mouse miRNA sequences are shown by default. If differences between mouse and human orthologous sequences are found, the human miRNA sequence is listed on a separate line with *H. sap.* as prefix. The number of clones identified from the indicated mouse tissues or the human osteoblast sarcoma cell line Saos-2 and cervical cancer derived HeLa cells are presented. Abbreviations: ht, heart; ln, lung; lv, liver; sp, spleen; si, small intestine; co, colon; kd, kidney; sk, skin; ts, testis; ov, ovary; thy, thymus; sc, spinal cord; cx, cortex; cb, cerebellum; mb, midbrain; S, Saos-2; H1, HeLa SS3 (this study and (Lagos-Quintana et al., 2001)); H2, HeLa published by (Mourelatos et al., 2002).

Footnotes:

<sup>a</sup>Cloned sequence not found in the mouse genome database.

<sup>b</sup>Cloned sequence not found in any genome database.

<sup>c</sup>Cloned sequence not found in the human genome database.

<sup>d</sup>miR-206 is homologous to miR-1 variants but has three A to G single-nucleotide changes.