

Supplemental Table 4 (S4): All identified miR groups based on “seed” sequence (2-7nt)

Group	Hsa-miRs
Let-7	let-7a, let-7b, let-7c, let-7d, let-7e, let-7f, let-7g, let-7i; 98, 202
miR-146	146a, 146b
miR-26	26a, 26b
miR-17	20a, 20b, 106a, 106b, 17-5p, 93
miR-29	29a, 29b, 29c
miR-18	18a, 18b
miR-34	34a, 34c
miR-19	19a, 19b
miR-15/16	503, 15a, 16, 195, 424
miR-125	125a, 125b
miR-23	23a, 23b
miR-373/520	302d, 373, 520b
miR-130	130a, 130b
miR-30	30a-5p, 30b, 30c, 30d, 30e
miR-18a*/324-3p	18a*, 324-3p
miR-518	520c, 526a, 518f*
miR-342/608	342, 608
miR-181	181a, 181b
miR-378/422a	378, 422a
miR-200	429, 200b, 200c
miR-221/222	221, 222

Note: Bolded miR families were analyzed using functional class scoring
 MiRBase release 17.0 (<http://microrna.sanger.ac.uk/>) was used to determine miR groups