

Integrating MicroRNA Expression Profiling Studies to Systematically Evaluate the Diagnostic Value of MicroRNAs in Pancreatic Cancer and Validate Their Prognostic Significance with the Cancer Genome Atlas Data

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Supplemental table 1. 693 microRNAs were measured in more than 9 profiling studies. SN-sample number, SEN-Sensitivity, SPE-Specificity, PLR-positive likelihood ratio, NLR-negative likelihood ratio, DOR-Diagnostic odds ratio, AUC-area under curve, PN-profiling number.

MicroRNAs ID	Gene Expression Omnibus accession number	SN	SEN	SPE	PLR	NLR	DOR	AUC	PN
hsa-miR-130b-3p	GSE24279,GSE71533,GSE60980,GSE31568,GSE85589,E-MTAB-753,GSE29352,E-TABM-664,GSE59856	925	0.732 72878	0.80 169 67	3.87	0.353	13.8	0.8385 23	9
hsa-miR-148a-3p	GSE71533,GSE24279,E-TABM-664,GSE60980,GSE31568,GSE29352,GSE59856,E-MTAB-753,GSE85589	924	0.749 09788	0.79 076 29	3.74	0.345	13.3	0.8383 31	9
hsa-miR-200c-3p	GSE24279,GSE71533,GSE60980,GSE85589,GSE31568,GSE29352,E-TABM-664,GSE59856,E-MTAB-753	925	0.556 80021	0.85 120 74	3.77	0.526	7.78	0.8319 92	9
hsa-miR-217	GSE24279,GSE71533,GSE60980,E-MTAB-753,E-TABM-664,GSE31568,GSE85589,GSE59856,GSE29352	925	0.863 81845	0.52 009 43	2	0.297	8.5	0.8254 33	9
hsa-miR-30c-5p	GSE24279,GSE71533,GSE60980,GSE29352,E-MTAB-753,E-TABM-664,GSE31568,GSE85589,GSE59856	925	0.829 73191	0.64 707 36	2.74	0.302	11.3	0.8244 44	9
hsa-miR-129-1-3p	GSE71533,GSE32678,GSE85589,E-MTAB-753,GSE60980,GSE24279,GSE31568,GSE59856,GSE29352	839	0.573 24663	0.84 603 35	3.7	0.506	7.68	0.8244 27	9
hsa-miR-23b-3p	GSE24279,GSE29352,GSE60980,GSE31568,GSE71533,E-TABM-664,GSE32678,E-MTAB-753,GSE85589,GSE59856	957	0.659 88357	0.80 897 53	3.46	0.425	8.34	0.8179 02	10
hsa-miR-30b-5p	GSE71533,GSE24279,GSE60980,GSE29352,E-MTAB-753,GSE31568,GSE85589,E-TABM-664,GSE59856	925	0.841 07225	0.60 803 39	2.4	0.288	9.71	0.8178 01	9
hsa-miR-222-3p	GSE71533,GSE24279,E-TABM-664,E-MTAB-753,GSE29352,GSE60980,GSE85589,GSE59856,GSE31568	925	0.762 83367	0.74 069 29	3.09	0.344	11.1	0.8141 09	9
hsa-miR-155-5p	GSE24279,GSE85589,GSE32678,E-TABM-664,E-MTAB-753,GSE29352,GSE71533,GSE60980,GSE31568,GSE59856	957	0.723 70394	0.76 522 44	3.08	0.37	8.87	0.8098 77	10
hsa-miR-575	GSE59856,GSE71533,GSE60980,GSE32678,E-MTAB-753,GSE85589,GSE29352,GSE31568,GSE24279	839	0.808 19324	0.69 439 51	2.74	0.293	10.8	0.8088 63	9
hsa-miR-1290	GSE71533,GSE59856,GSE24279,GSE85589,GSE29352,E-MTAB-753,GSE60980,GSE31568,GSE32678	839	0.670 60464	0.80 187 86	3.52	0.414	8.97	0.8074 04	9
hsa-miR-30e-5p	GSE71533,GSE24279,GSE60980,GSE29352,E-MTAB-753,GSE31568,E-TABM-664,GSE85589,GSE59856	925	0.560 71231	0.83 513 45	3.44	0.523	7.52	0.8067 18	9

hsa-miR-107	GSE71533,GSE24279,GSE60980,E-MTAB-753,E-TABM-664,GSE29352,GSE31568,GSE59856,GSE32678,GSE85589	957	0.743 90374	0.74 362 98	3.01	0.356	9.18	0.8065 31	10
hsa-miR-181b-5p	GSE24279,GSE71533,E-TABM-664,GSE60980,E-MTAB-753,GSE85589,GSE31568,GSE32678,GSE59856,GSE29352	957	0.617 51715	0.80 862 04	3.33	0.477	7.84	0.8059 72	10
hsa-miR-663a	GSE59856,GSE32678,GSE71533,GSE60980,GSE31568,E-MTAB-753,GSE85589,GSE29352,GSE24279	839	0.639 71826	0.80 614 35	3.37	0.454	8.07	0.8043 21	9
hsa-miR-93-5p	GSE24279,GSE59856,GSE71533,GSE29352,E-TABM-664,GSE85589,GSE60980,E-MTAB-753,GSE31568	925	0.737 93612	0.74 799 29	3.01	0.359	8.84	0.8038 05	9
hsa-miR-628-5p	GSE85589,GSE60980,GSE29352,GSE24279,GSE32678,GSE31568,GSE59856,GSE71533,E-MTAB-753	839	0.580 75354	0.81 960 72	3.22	0.527	6.4	0.7974 68	9
hsa-miR-27b-3p	GSE24279,GSE71533,GSE29352,GSE60980,E-TABM-664,E-MTAB-753,GSE31568,GSE59856,GSE85589	925	0.727 77893	0.74 285 27	2.9	0.376	8.51	0.7973 28	9
hsa-miR-132-5p	GSE71533,GSE60980,GSE31568,GSE32678,GSE29352,E-MTAB-753,GSE85589,E-TABM-664,GSE24279,GSE59856	957	0.469 43445	0.83 687 39	2.91	0.631	4.72	0.7968 12	10
hsa-miR-15b-5p	GSE24279,GSE71533,E-MTAB-753,GSE29352,GSE60980,GSE31568,E-TABM-664,GSE85589,GSE59856	925	0.664 97547	0.77 964 45	3	0.436	7.36	0.7962 97	9
hsa-miR-30a-5p	GSE24279,GSE71533,GSE60980,E-MTAB-753,GSE29352,GSE59856,GSE31568,GSE85589,E-TABM-664	925	0.765 33917	0.72 160 44	2.82	0.334	9.64	0.7960 85	9
hsa-miR-423-5p	GSE59856,GSE71533,GSE31568,GSE60980,GSE29352,E-TABM-664,GSE85589,GSE24279,E-MTAB-753	925	0.732 91044	0.73 376 39	2.81	0.369	7.83	0.7946 8	9
hsa-miR-1301-3p	GSE71533,GSE24279,GSE60980,GSE31568,GSE32678,GSE29352,E-MTAB-753,GSE85589,GSE59856	839	0.569 56017	0.80 949 53	2.94	0.54	5.84	0.7932 85	9
hsa-miR-647	GSE71533,GSE31568,E-MTAB-753,GSE29352,GSE24279,GSE32678,GSE85589,GSE60980,GSE59856	839	0.556 18042	0.81 616 02	3	0.548	5.81	0.7919 91	9
hsa-miR-29c-3p	GSE71533,GSE24279,GSE60980,GSE29352,GSE31568,E-MTAB-753,GSE85589,E-TABM-664,GSE59856	925	0.555 77023	0.80 930 44	2.9	0.558	6	0.7898 56	9
hsa-miR-221-5p	GSE71533,GSE60980,GSE85589,GSE32678,GSE31568,GSE29352,E-MTAB-753,GSE24279,GSE59856	839	0.570 56502	0.80 673 51	2.91	0.545	5.91	0.7880 1	9
hsa-miR-455-3p	GSE71533,GSE85589,GSE31568,E-MTAB-753,GSE29352,GSE60980,GSE32678,GSE59856,GSE24279,E-TABM-664	957	0.764 08353	0.69 916 81	2.57	0.359	8.13	0.7875 11	10

hsa-miR-26a-5p	GSE71533,GSE59856,GSE29352,GSE60980,GSE24279,GSE31568,E-MTAB-753,GSE85589,E-TABM-664	925	0.687 94975	0.75 840 43	2.89	0.414	7.12	0.7871 03	9
hsa-miR-335-5p	GSE71533,GSE24279,GSE85589,GSE60980,E-MTAB-753,GSE29352,GSE31568,E-TABM-664,GSE59856	924	0.502 62431	0.86 638 62	3.87	0.579	7.02	0.7860 32	9
hsa-miR-1225-5p	GSE60980,GSE59856,GSE85589,E-MTAB-753,GSE32678,GSE24279,GSE29352,GSE31568,GSE71533	839	0.471 66884	0.80 514 73	2.44	0.655	3.81	0.7835 36	9
hsa-miR-103a-3p	GSE71533,GSE24279,GSE29352,E-MTAB-753,E-TABM-664,GSE60980,GSE59856,GSE31568,GSE85589	925	0.702 8185	0.75 163 32	2.96	0.399	7.82	0.7828 69	9
hsa-miR-125b-5p	GSE24279,GSE71533,GSE60980,GSE29352,E-MTAB-753,E-TABM-664,GSE85589,GSE31568,GSE59856	925	0.565 48983	0.81 308 41	3.05	0.546	5.85	0.7794 29	9
hsa-miR-92a-3p	GSE71533,GSE24279,GSE29352,E-MTAB-753,E-TABM-664,GSE85589,GSE60980,GSE59856,GSE31568	925	0.645 33727	0.78 425 27	3.04	0.459	6.82	0.7794 05	9
hsa-miR-320a	GSE71533,GSE31568,E-TABM-664,GSE24279,GSE29352,GSE59856,GSE60980,E-MTAB-753,GSE85589	925	0.727 31428	0.70 854 8	2.56	0.392	6.73	0.7769 25	9
hsa-miR-30a-3p	GSE24279,GSE71533,GSE60980,E-MTAB-753,GSE29352,GSE31568,GSE59856,E-TABM-664,GSE85589	925	0.602 18553	0.79 524 57	3.08	0.517	6.66	0.7761 02	9
hsa-miR-10a-5p	GSE71533,GSE24279,E-TABM-664,GSE60980,GSE31568,GSE85589,E-MTAB-753,GSE59856,GSE29352	925	0.610 7817	0.79 466 99	2.99	0.494	6.16	0.7755 81	9
hsa-miR-203a-3p	E-TABM-664,GSE29352,E-MTAB-753,GSE71533,GSE31568,GSE60980,GSE59856,GSE24279,GSE85589	925	0.523 49562	0.84 567 43	3.46	0.565	6.27	0.7751 7	9
hsa-miR-32-5p	GSE71533,GSE85589,GSE29352,GSE60980,E-TABM-664,GSE24279,E-MTAB-753,GSE31568,GSE59856	925	0.484 25611	0.84 255 92	3.11	0.604	5.49	0.7750 27	9
hsa-miR-125a-3p	GSE59856,GSE71533,GSE60980,E-MTAB-753,GSE85589,E-TABM-664,GSE29352,GSE31568,GSE24279	925	0.714 75133	0.73 407 56	2.94	0.411	8.52	0.7735 89	9
hsa-miR-23a-3p	GSE24279,GSE60980,GSE71533,E-MTAB-753,GSE29352,E-TABM-664,GSE85589,GSE59856,GSE31568	925	0.633 45357	0.80 330 82	3.32	0.462	7.68	0.7722 55	9
hsa-miR-16-5p	GSE60980,GSE31568,GSE24279,GSE59856,GSE29352,GSE71533,E-MTAB-753,E-TABM-664,GSE85589	925	0.715 28064	0.71 288 94	2.54	0.405	6.34	0.7717 32	9
hsa-miR-891a-5p	GSE71533,GSE29352,GSE32678,E-MTAB-753,GSE31568,GSE24279,GSE60980,GSE59856,GSE85589	839	0.766 36471	0.66 296 02	2.33	0.367	6.76	0.7711 71	9

hsa-miR-17-5p	GSE59856,GSE29352,GSE71533,GSE31568,E-MTAB-753,E-TABM-664,GSE60980,GSE85589,GSE24279,GSE32678	957	0.749 76727	0.66 041 54	2.26	0.385	6.07	0.7710 84	10
hsa-miR-1246	GSE59856,GSE60980,GSE24279,E-MTAB-753,GSE85589,GSE32678,GSE71533,GSE29352,GSE31568	839	0.746 23272	0.67 307 53	2.36	0.384	6.42	0.7691 89	9
hsa-miR-320c	GSE71533,GSE31568,GSE24279,GSE59856,GSE32678,GSE60980,GSE85589,GSE29352,E-MTAB-753	839	0.724 20129	0.69 801 08	2.43	0.403	6.13	0.7689 36	9
hsa-miR-1275	GSE71533,GSE32678,GSE60980,GSE85589,GSE24279,GSE29352,GSE31568,GSE59856,E-MTAB-753	839	0.633 85116	0.76 129 8	2.69	0.482	5.92	0.7685 85	9
hsa-miR-375	GSE71533,GSE24279,GSE60980,E-TABM-664,GSE85589,GSE59856,E-MTAB-753,GSE29352,GSE31568	925	0.686 02133	0.73 376 02	2.68	0.435	6.63	0.7683 89	9
hsa-miR-1307-3p	GSE71533,GSE32678,GSE24279,GSE59856,GSE85589,GSE29352,E-MTAB-753,GSE60980,GSE31568	839	0.626 04802	0.78 008 85	2.93	0.486	6.36	0.7683 24	9
hsa-miR-106a-5p	GSE59856,GSE71533,GSE29352,GSE32678,GSE85589,GSE31568,E-TABM-664,E-MTAB-753,GSE24279	900	0.698 02948	0.72 246 47	2.52	0.424	6.24	0.7680 8	9
hsa-miR-138-2-3p	GSE71533,GSE60980,GSE24279,GSE32678,GSE29352,GSE31568,GSE85589,GSE59856,E-MTAB-753	839	0.834 51916	0.46 118 05	1.64	0.387	4.71	0.7675 55	9
hsa-miR-584-5p	GSE71533,GSE60980,GSE85589,GSE32678,GSE29352,GSE31568,E-MTAB-753,GSE24279,GSE59856	839	0.786 44261	0.58 151 51	1.95	0.38	5.26	0.7648 97	9
hsa-miR-564	GSE71533,GSE59856,GSE60980,GSE31568,GSE32678,GSE29352,GSE85589,GSE24279,E-MTAB-753	839	0.738 90214	0.67 860 99	2.4	0.395	6.29	0.7644 19	9
hsa-miR-551a	GSE71533,GSE31568,GSE24279,GSE32678,GSE29352,E-MTAB-753,GSE60980,GSE59856,GSE85589	839	0.720 49022	0.69 538 39	2.37	0.412	5.96	0.7621 99	9
hsa-miR-20b-5p	GSE29352,GSE59856,GSE71533,GSE60980,E-MTAB-753,E-TABM-664,GSE85589,GSE31568,GSE24279	925	0.787 84905	0.61 213 01	2.08	0.362	6.2	0.7620 02	9
hsa-miR-604	GSE71533,GSE24279,GSE60980,GSE31568,GSE32678,E-MTAB-753,GSE85589,GSE29352,GSE59856	839	0.723 40876	0.68 495 44	2.65	0.429	6.38	0.7619 64	9
hsa-miR-181a-5p	GSE24279,GSE71533,E-TABM-664,GSE31568,E-MTAB-753,GSE29352,GSE85589,GSE59856,GSE60980	925	0.756 53724	0.71 805 89	2.66	0.35	8.09	0.7615 78	9
hsa-miR-484	GSE71533,GSE24279,GSE29352,GSE31568,E-MTAB-753,GSE59856,E-TABM-664,GSE60980,GSE85589	925	0.606 26267	0.75 678 05	2.52	0.52	5.05	0.7606 34	9

hsa-miR-141-3p	GSE71533,GSE60980,GSE24279,GSE29352,GSE85589,E-TABM-664,GSE31568,E-MTAB-753,GSE59856	925	0.543 7346	0.79 541 87	2.69	0.575	5.22	0.7601 99	9
hsa-miR-603	GSE85589,GSE32678,GSE29352,GSE60980,E-MTAB-753,GSE71533,GSE31568,GSE24279,GSE59856	839	0.584 9596	0.78 918 7	2.82	0.534	5.52	0.7599 66	9
hsa-miR-127-3p	GSE71533,GSE24279,GSE85589,GSE60980,GSE31568,E-MTAB-753,E-TABM-664,GSE59856,GSE29352	925	0.515 53085	0.84 626 06	3.42	0.58	6.03	0.7598 29	9
hsa-miR-1262	GSE32678,GSE29352,GSE24279,GSE71533,GSE31568,GSE60980,E-MTAB-753,GSE85589,GSE59856	839	0.861 09364	0.45 221 85	1.69	0.332	5.26	0.7597 39	9
hsa-miR-155-3p	GSE71533,GSE85589,E-MTAB-753,GSE32678,GSE60980,GSE29352,GSE24279,GSE31568,GSE59856	839	0.507 79733	0.81 001 52	2.67	0.61	4.66	0.7591 24	9
hsa-miR-550a-3p	GSE71533,GSE31568,GSE85589,GSE29352,GSE32678,GSE60980,E-MTAB-753,GSE24279,GSE59856	839	0.673 77991	0.72 586 15	2.43	0.464	5.8	0.7585 13	9
hsa-miR-26b-5p	GSE71533,GSE24279,GSE29352,GSE31568,GSE85589,GSE60980,E-MTAB-753,E-TABM-664,GSE59856	925	0.759 33909	0.63 715 77	2.2	0.394	5.68	0.7582 74	9
hsa-miR-101-3p	GSE71533,GSE60980,GSE29352,GSE85589,E-MTAB-753,GSE31568,E-TABM-664,GSE24279,GSE59856	925	0.743 36673	0.65 446 09	2.25	0.415	5.82	0.7566 72	9
hsa-miR-146a-3p	GSE32678,GSE24279,GSE71533,E-MTAB-753,GSE60980,GSE85589,GSE29352,GSE31568,GSE59856	839	0.480 37763	0.80 849 68	2.51	0.639	4.12	0.7557 93	9
hsa-miR-362-5p	GSE29352,GSE71533,GSE24279,GSE60980,E-MTAB-753,E-TABM-664,GSE31568,GSE59856,GSE32678,GSE85589	957	0.462 76147	0.83 190 78	2.76	0.644	4.38	0.7556 87	10
hsa-miR-1181	GSE60980,GSE29352,GSE24279,GSE59856,GSE71533,GSE31568,GSE85589,GSE32678,E-MTAB-753	839	0.614 55886	0.77 442 94	2.77	0.509	5.52	0.7553 41	9
hsa-miR-1258	GSE71533,GSE32678,GSE29352,GSE60980,GSE85589,GSE31568,E-MTAB-753,GSE59856,GSE24279	839	0.832 62293	0.48 950 27	1.77	0.372	5.3	0.7547 81	9
hsa-miR-192-3p	GSE32678,GSE31568,GSE24279,GSE29352,E-MTAB-753,GSE71533,GSE85589,GSE59856,GSE60980	839	0.566 21144	0.79 017 11	2.72	0.552	5.03	0.7547 31	9
hsa-miR-210-3p	GSE71533,GSE24279,E-TABM-664,GSE31568,GSE29352,GSE60980,GSE85589,GSE59856,E-MTAB-753	925	0.621 99847	0.74 973 6	2.5	0.507	5.21	0.7545 27	9
hsa-miR-10a-3p	GSE71533,E-TABM-664,GSE85589,E-MTAB-753,GSE60980,GSE24279,GSE29352,GSE31568,GSE59856	925	0.662 35931	0.73 345 82	2.58	0.472	5.7	0.7544 17	9

hsa-miR-181d-5p	GSE24279,GSE71533,E-TABM-664,E-MTAB-753,GSE85589,GSE29352,GSE60980,GSE31568,GSE59856	925	0.714 40058	0.68 243 89	2.47	0.444	5.91	0.7543 11	9
hsa-miR-610	GSE71533,GSE60980,GSE32678,GSE85589,GSE29352,GSE31568,E-MTAB-753,GSE59856,GSE24279	839	0.678 52848	0.71 661 97	2.44	0.461	5.62	0.7542 5	9
hsa-miR-199a-5p	GSE29352,GSE24279,GSE71533,GSE60980,E-MTAB-753,E-TABM-664,GSE32678,GSE85589,GSE31568,GSE59856	957	0.662 98064	0.72 633 86	2.45	0.467	5.48	0.7534 93	10
hsa-miR-920	GSE71533,GSE60980,GSE32678,GSE31568,GSE59856,E-MTAB-753,GSE29352,GSE85589,GSE24279	839	0.474 96976	0.81 191 5	2.56	0.652	4.06	0.7530 34	9
hsa-miR-374b-3p	GSE29352,GSE32678,GSE60980,GSE71533,E-MTAB-753,GSE31568,GSE85589,GSE24279,GSE59856	839	0.502 22094	0.81 563 23	2.7	0.605	4.63	0.7529 08	9
hsa-miR-150-5p	GSE71533,GSE24279,E-MTAB-753,GSE60980,GSE29352,E-TABM-664,GSE59856,GSE31568,GSE85589	925	0.731 98016	0.67 494 51	2.29	0.417	6.3	0.7522 73	9
hsa-miR-145-5p	GSE24279,E-MTAB-753,GSE60980,E-TABM-664,GSE29352,GSE71533,GSE85589,GSE59856,GSE31568	925	0.657 03909	0.72 569 02	2.43	0.476	5.3	0.7520 55	9
hsa-miR-100-5p	GSE71533,GSE24279,GSE29352,E-TABM-664,GSE60980,E-MTAB-753,GSE85589,GSE31568,GSE59856	925	0.595 5206	0.76 551 04	2.53	0.54	4.96	0.7518 72	9
hsa-miR-513a-5p	GSE71533,GSE60980,GSE59856,GSE32678,GSE85589,GSE24279,E-TABM-664,GSE29352,GSE31568,E-MTAB-753	957	0.673 82903	0.71 605 6	2.4	0.461	5.53	0.7518 3	10
hsa-miR-1180-3p	GSE60980,GSE32678,GSE29352,GSE85589,GSE71533,GSE24279,GSE59856,GSE31568,E-MTAB-753	839	0.760 94441	0.61 323 97	2.03	0.397	5.22	0.7512 43	9
hsa-miR-892b	GSE60980,GSE29352,GSE71533,GSE31568,GSE32678,E-MTAB-753,GSE24279,GSE85589,GSE59856	838	0.442 41899	0.84 985 23	3.01	0.655	4.8	0.7511 34	9
hsa-miR-206	GSE71533,GSE60980,GSE24279,E-TABM-664,GSE31568,GSE85589,E-MTAB-753,GSE29352,GSE59856	925	0.583 69904	0.77 024 47	2.51	0.541	4.86	0.7511 33	9
hsa-miR-195-5p	GSE71533,GSE24279,GSE29352,E-MTAB-753,GSE60980,GSE31568,E-TABM-664,GSE85589,GSE59856	925	0.750 70115	0.63 029 4	2.13	0.415	5.24	0.7507 98	9
hsa-miR-346	GSE71533,GSE85589,GSE32678,GSE59856,GSE24279,GSE60980,E-MTAB-753,E-TABM-664,GSE31568,GSE29352	957	0.729 00425	0.66 499 67	2.24	0.417	5.75	0.7499 9	10
hsa-miR-518a-3p	GSE71533,GSE29352,GSE32678,GSE31568,GSE24279,E-MTAB-753,GSE60980,E-TABM-664,GSE85589,GSE59856	957	0.531 56804	0.78 209 47	2.42	0.602	4.15	0.7498 88	10

hsa-miR-361-3p	GSE24279,GSE71533,GSE60980,GSE32678,GSE31568,GSE59856,GSE29352,GSE85589,E-TABM-664,E-MTAB-753	957	0.779 32609	0.60 671 72	2.03	0.373	5.77	0.7497 35	10
hsa-miR-19b-1-5p	GSE71533,E-MTAB-753,GSE85589,GSE31568,GSE29352,GSE32678,GSE24279,GSE59856,GSE60980	839	0.496 4856	0.81 682 45	2.73	0.614	4.54	0.7491 99	9
hsa-miR-20a-3p	GSE29352,GSE85589,GSE71533,GSE60980,GSE32678,GSE31568,GSE24279,GSE59856,E-MTAB-753	839	0.536 21708	0.79 354 43	2.56	0.583	4.67	0.7490 91	9
hsa-miR-1233-3p	GSE59856,GSE71533,GSE29352,GSE32678,GSE85589,GSE60980,GSE31568,GSE24279,E-MTAB-753	839	0.816 73406	0.54 692 97	1.85	0.345	5.52	0.7490 69	9
hsa-miR-125b-2-3p	GSE71533,GSE60980,E-MTAB-753,E-TABM-664,GSE29352,GSE24279,GSE85589,GSE31568,GSE59856	925	0.531 64949	0.78 963 14	2.53	0.6	4.49	0.7487 39	9
hsa-miR-1184	GSE71533,GSE32678,GSE85589,GSE60980,GSE31568,GSE29352,E-MTAB-753,GSE24279,GSE59856	839	0.551 28416	0.79 143 29	2.72	0.57	5.13	0.7483 98	9
hsa-miR-641	E-MTAB-753,GSE60980,GSE71533,GSE29352,GSE31568,GSE24279,GSE32678,GSE85589,GSE59856	839	0.445 71074	0.85 165 01	3.02	0.65	4.71	0.7482 02	9
hsa-miR-186-5p	GSE71533,GSE60980,E-MTAB-753,GSE85589,GSE24279,E-TABM-664,GSE29352,GSE59856,GSE31568	925	0.658 3298	0.72 601 93	2.53	0.484	5.61	0.7475 32	9
hsa-miR-1260a	GSE71533,GSE60980,GSE29352,GSE59856,E-MTAB-753,GSE85589,GSE24279,GSE31568,GSE32678	839	0.750 64837	0.65 372 11	2.17	0.399	5.88	0.7474 13	9
hsa-miR-28-5p	GSE71533,GSE32678,GSE60980,GSE85589,E-MTAB-753,E-TABM-664,GSE24279,GSE29352,GSE31568,GSE59856	957	0.560 47228	0.78 035	2.54	0.572	4.67	0.7469 44	10
hsa-miR-574-3p	GSE60980,GSE71533,E-MTAB-753,GSE24279,GSE29352,GSE32678,GSE59856,GSE31568,GSE85589	839	0.658 49686	0.72 350 6	2.42	0.481	5.22	0.7468 87	9
hsa-miR-143-5p	GSE71533,GSE24279,GSE60980,GSE85589,GSE29352,GSE31568,E-MTAB-753,GSE32678,GSE59856	839	0.681 71306	0.70 271 48	2.31	0.455	5.33	0.7465 72	9
hsa-miR-20a-5p	GSE59856,GSE29352,E-MTAB-753,GSE71533,GSE31568,E-TABM-664,GSE32678,GSE60980,GSE24279,GSE85589	957	0.732 2798	0.66 178 9	2.17	0.415	5.49	0.7464 73	10
hsa-miR-181c-5p	GSE24279,E-TABM-664,GSE31568,E-MTAB-753,GSE60980,GSE71533,GSE29352,GSE85589,GSE59856	925	0.614 11632	0.74 756 08	2.41	0.518	4.85	0.7462 25	9
hsa-miR-548a-3p	GSE32678,GSE85589,E-MTAB-753,GSE31568,GSE29352,GSE60980,GSE24279,GSE59856,GSE71533	839	0.565 6798	0.77 627 23	2.55	0.566	4.71	0.7461 16	9

hsa-miR-135b-5p	GSE71533,GSE29352,GSE60980,GSE31568,GSE85589,GSE24279,GSE59856,E-MTAB-753,E-TABM-664	925	0.721 68264	0.66 446 76	2.18	0.435	5.34	0.7457 03	9
hsa-miR-34a-5p	GSE24279,E-MTAB-753,GSE71533,GSE31568,E-TABM-664,GSE60980,GSE59856,GSE29352,GSE85589	925	0.492 68845	0.80 862 83	2.55	0.627	4.2	0.7456 1	9
hsa-miR-578	GSE29352,GSE60980,GSE31568,GSE32678,E-MTAB-753,GSE85589,GSE71533,GSE24279,GSE59856	839	0.501 82778	0.81 842 75	2.77	0.603	4.78	0.7455 09	9
hsa-miR-1231	GSE59856,GSE85589,GSE71533,GSE32678,GSE24279,GSE29352,GSE31568,E-MTAB-753,GSE60980	839	0.533 37944	0.83 808 47	3.36	0.562	6.12	0.7447 93	9
hsa-miR-143-3p	GSE71533,GSE60980,GSE29352,E-MTAB-753,E-TABM-664,GSE24279,GSE85589,GSE31568,GSE59856	925	0.711 44458	0.67 223 68	2.21	0.435	5.22	0.7445 97	9
hsa-miR-509-3-5p	GSE32678,GSE60980,GSE71533,GSE29352,GSE85589,E-MTAB-753,GSE31568,GSE24279,GSE59856	839	0.486 69396	0.82 848 63	2.94	0.624	4.89	0.7440 43	9
hsa-miR-340-5p	GSE71533,GSE60980,E-TABM-664,E-MTAB-753,GSE85589,GSE29352,GSE24279,GSE31568,GSE32678,GSE59856	957	0.802 66628	0.53 966 3	1.84	0.38	4.97	0.7439 52	10
hsa-miR-1825	GSE32678,GSE85589,GSE60980,GSE71533,GSE24279,GSE29352,GSE31568,GSE59856,E-MTAB-753	839	0.695 21279	0.68 927 03	2.23	0.453	5.22	0.7438 71	9
hsa-miR-519e-5p	GSE71533,GSE31568,E-MTAB-753,GSE32678,GSE60980,GSE24279,GSE85589,GSE29352,GSE59856	839	0.595 8883	0.76 252 46	2.55	0.529	5.1	0.7438 67	9
hsa-miR-1207-3p	GSE71533,E-MTAB-753,GSE31568,GSE85589,GSE59856,GSE60980,GSE24279,GSE32678,GSE29352	839	0.731 54963	0.65 068 8	2.12	0.432	5.29	0.7438 16	9
hsa-miR-489-3p	GSE71533,GSE32678,GSE29352,E-MTAB-753,GSE85589,GSE60980,E-TABM-664,GSE59856,GSE24279,GSE31568	957	0.572 49255	0.77 924 59	2.71	0.557	5.19	0.7432 99	10
hsa-miR-141-5p	GSE71533,GSE32678,GSE60980,E-MTAB-753,GSE31568,GSE29352,GSE59856,GSE24279,GSE85589	839	0.459 21534	0.82 732 47	2.73	0.649	4.58	0.7432 88	9
hsa-miR-34b-5p	GSE32678,GSE71533,GSE24279,GSE60980,GSE29352,E-TABM-664,GSE31568,GSE59856,E-MTAB-753,GSE85589	957	0.588 51659	0.78 332 12	2.84	0.535	5.51	0.7432 27	10
hsa-miR-1252-5p	GSE71533,GSE32678,E-MTAB-753,GSE60980,GSE85589,GSE31568,GSE59856,GSE29352,GSE24279	839	0.569 29421	0.80 007 13	3.02	0.547	5.75	0.7427 42	9
hsa-miR-505-5p	GSE71533,GSE60980,GSE32678,E-MTAB-753,GSE29352,GSE31568,GSE59856,GSE24279,GSE85589	839	0.683 07184	0.69 633 58	2.31	0.465	5.3	0.7426 95	9

hsa-miR-214-3p	GSE24279,GSE60980,GSE29352,E-TABM-664,E-MTAB-753,GSE59856,GSE85589,GSE31568,GSE71533	925	0.681 75759	0.70 012 35	2.31	0.47	5.56	0.7421 38	9
hsa-miR-195-3p	GSE71533,GSE32678,GSE60980,GSE24279,GSE85589,GSE29352,E-MTAB-753,GSE59856,GSE31568	839	0.729 58092	0.64 754 51	2.15	0.432	5.29	0.7415 41	9
hsa-miR-215-5p	GSE24279,GSE32678,GSE60980,GSE85589,GSE29352,E-MTAB-753,GSE59856,GSE31568,E-TABM-664,GSE71533	957	0.467 39097	0.83 122 2	2.81	0.644	4.7	0.7413 9	10
hsa-miR-320b	GSE71533,GSE31568,GSE24279,GSE29352,GSE59856,GSE32678,GSE60980,E-MTAB-753,GSE85589	839	0.760 76959	0.60 996 75	1.98	0.402	5.05	0.7409 86	9
hsa-miR-191-3p	GSE24279,GSE29352,GSE85589,E-MTAB-753,GSE60980,GSE32678,GSE71533,GSE31568,GSE59856	839	0.517 377	0.78 741 72	2.42	0.612	4.1	0.7402 96	9
hsa-miR-548b-5p	GSE29352,E-MTAB-753,GSE60980,GSE71533,GSE31568,GSE24279,GSE85589,GSE59856,GSE32678	839	0.467 76786	0.79 721 42	2.3	0.657	3.7	0.7398 08	9
hsa-miR-21-3p	GSE71533,GSE24279,GSE32678,E-TABM-664,GSE31568,GSE29352,E-MTAB-753,GSE60980,GSE85589,GSE59856	957	0.625 66008	0.73 870 33	2.45	0.512	4.98	0.7397 27	10
hsa-miR-223-3p	GSE60980,GSE59856,E-TABM-664,GSE71533,GSE29352,GSE85589,GSE24279,GSE31568,E-MTAB-753	925	0.664 21003	0.71 107 83	2.36	0.48	5.12	0.7396 19	9
hsa-miR-597-5p	GSE71533,GSE32678,GSE29352,GSE85589,GSE31568,GSE60980,GSE24279,E-MTAB-753,GSE59856	839	0.534 95164	0.80 845 83	2.89	0.586	5.12	0.7392 71	9
hsa-miR-640	GSE71533,GSE32678,GSE85589,E-MTAB-753,GSE29352,GSE24279,GSE31568,GSE59856,GSE60980	839	0.532 92302	0.79 482 53	2.65	0.597	4.69	0.7389 42	9
hsa-miR-1263	GSE85589,GSE71533,GSE24279,GSE29352,GSE60980,GSE32678,GSE31568,E-MTAB-753,GSE59856	839	0.472 25518	0.83 812 4	2.93	0.626	4.81	0.7384 78	9
hsa-miR-607	GSE32678,GSE24279,GSE60980,GSE71533,GSE29352,GSE31568,E-MTAB-753,GSE85589,GSE59856	839	0.538 32711	0.83 534 69	3.57	0.564	6.27	0.7382 74	9
hsa-miR-1321	GSE71533,GSE60980,GSE32678,E-MTAB-753,GSE24279,GSE29352,GSE31568,GSE85589,GSE59856	839	0.487 55549	0.82 107 61	2.77	0.622	4.62	0.7377 55	9
hsa-miR-425-3p	GSE71533,GSE24279,GSE60980,GSE85589,GSE31568,GSE59856,E-MTAB-753,GSE29352,E-TABM-664	925	0.644 86659	0.72 129 2	2.37	0.495	4.93	0.7375 55	9
hsa-miR-22-3p	GSE85589,GSE60980,GSE29352,GSE71533,GSE59856,GSE24279,E-MTAB-753,E-TABM-664,GSE31568	925	0.610 81317	0.76 560 7	2.7	0.512	5.37	0.7374 63	9

hsa-miR-212-3p	GSE71533,GSE60980,GSE31568,E-MTAB-753,GSE24279,E-TABM-664,GSE85589,GSE29352,GSE59856	925	0.608 84358	0.74 909 85	2.46	0.524	4.83	0.7368 03	9
hsa-miR-1281	GSE71533,GSE32678,GSE85589,GSE31568,GSE29352,E-MTAB-753,GSE59856,GSE24279,GSE60980	839	0.810 89241	0.59 196 61	1.99	0.327	6.41	0.7364 51	9
hsa-miR-1272	GSE71533,GSE85589,GSE32678,GSE60980,E-MTAB-753,GSE24279,GSE59856,GSE31568,GSE29352	839	0.665 46327	0.70 152 86	2.28	0.489	4.93	0.7362 33	9
hsa-miR-1200	GSE71533,GSE32678,GSE29352,GSE60980,GSE31568,GSE85589,GSE24279,GSE59856,E-MTAB-753	839	0.552 78596	0.76 631 86	2.36	0.586	4.22	0.7358 15	9
hsa-miR-624-3p	GSE71533,GSE32678,GSE29352,GSE85589,GSE31568,GSE60980,GSE59856,E-MTAB-753,GSE24279	839	0.623 52452	0.73 171 4	2.36	0.516	4.75	0.7355 19	9
hsa-miR-146a-5p	GSE24279,GSE71533,E-MTAB-753,GSE60980,E-TABM-664,GSE31568,GSE59856,GSE85589,GSE29352	925	0.668 83867	0.69 656 04	2.21	0.477	4.75	0.7354 38	9
hsa-miR-20b-3p	GSE71533,E-MTAB-753,GSE85589,GSE60980,GSE31568,GSE32678,E-TABM-664,GSE24279,GSE29352,GSE59856	957	0.487 29152	0.77 399 16	2.12	0.664	3.42	0.7346 72	10
hsa-miR-125a-5p	GSE71533,GSE24279,GSE60980,E-MTAB-753,E-TABM-664,GSE85589,GSE31568,GSE29352,GSE59856	925	0.615 32718	0.73 124 43	2.32	0.531	4.61	0.7345 23	9
hsa-miR-645	GSE71533,GSE60980,GSE32678,GSE85589,GSE24279,GSE29352,GSE59856,GSE31568,E-MTAB-753	839	0.771 55045	0.58 229 91	1.89	0.405	4.79	0.7338 03	9
hsa-miR-891b	GSE60980,GSE32678,GSE31568,GSE71533,GSE29352,GSE85589,E-MTAB-753,GSE24279,GSE59856	839	0.515 66337	0.82 774 17	3.1	0.585	5.3	0.7335 68	9
hsa-miR-1225-3p	GSE32678,GSE59856,GSE29352,GSE31568,GSE71533,E-MTAB-753,GSE24279,GSE85589,GSE60980	839	0.716 53265	0.64 052 68	2.03	0.451	4.56	0.7334 68	9
hsa-miR-132-3p	GSE24279,GSE71533,E-MTAB-753,GSE60980,E-TABM-664,GSE29352,GSE31568,GSE85589,GSE59856	925	0.669 45536	0.69 359 46	2.18	0.496	4.79	0.7333 28	9
hsa-miR-501-3p	GSE60980,E-MTAB-753,GSE31568,GSE24279,GSE71533,E-TABM-664,GSE29352,GSE59856,GSE85589	925	0.588 81552	0.74 693 68	2.32	0.554	4.37	0.7327 55	9
hsa-miR-223-5p	GSE60980,GSE31568,GSE24279,GSE71533,GSE29352,GSE32678,E-MTAB-753,GSE85589,GSE59856	839	0.764 88798	0.54 266 81	1.84	0.456	4.1	0.7324 9	9
hsa-miR-1277-3p	GSE32678,GSE29352,GSE60980,GSE31568,GSE85589,GSE24279,E-MTAB-753,GSE59856,GSE71533	839	0.538 16442	0.77 635 37	2.4	0.604	4.12	0.7316 04	9

hsa-miR-200b-3p	GSE24279,GSE71533,GSE29352,GSE60980,E-MTAB-753,E-TABM-664,GSE85589,GSE59856,GSE31568	925	0.541 88472	0.78 962 17	2.65	0.588	4.71	0.7315 87	9
hsa-miR-324-5p	GSE24279,GSE60980,GSE29352,GSE85589,GSE71533,E-TABM-664,GSE31568,E-MTAB-753,GSE59856	925	0.680 55626	0.68 465 87	2.17	0.477	4.87	0.7311 18	9
hsa-miR-1278	GSE85589,GSE32678,GSE31568,GSE24279,E-MTAB-753,GSE29352,GSE71533,GSE59856,GSE60980	839	0.625 37614	0.73 134 37	2.37	0.521	4.63	0.7308 93	9
hsa-miR-24-3p	GSE29352,GSE71533,GSE59856,GSE60980,GSE24279,E-TABM-664,E-MTAB-753,GSE85589,GSE31568	925	0.606 30561	0.79 085 63	3.01	0.502	6.15	0.7308 24	9
hsa-miR-1-3p	GSE60980,GSE71533,GSE29352,GSE85589,E-TABM-664,E-MTAB-753,GSE24279,GSE31568,GSE59856	925	0.534 17655	0.77 687 52	2.38	0.601	4.1	0.7307 79	9
hsa-miR-32-3p	GSE71533,GSE32678,GSE31568,GSE60980,E-TABM-664,GSE85589,GSE24279,GSE59856,GSE29352,E-MTAB-753	957	0.465 44116	0.83 285 67	2.85	0.64	4.56	0.7306 09	10
hsa-miR-1228-3p	GSE59856,GSE32678,GSE71533,GSE60980,GSE31568,GSE29352,GSE85589,GSE24279,E-MTAB-753	839	0.685 33877	0.67 584 95	2.18	0.471	4.7	0.7305 9	9
hsa-miR-630	GSE71533,GSE32678,GSE60980,GSE31568,GSE29352,E-MTAB-753,GSE59856,GSE24279,GSE85589	839	0.575 08633	0.77 162 28	2.62	0.558	4.84	0.7303 93	9
hsa-miR-429	GSE32678,GSE71533,GSE29352,GSE24279,E-MTAB-753,GSE31568,GSE85589,GSE60980,E-TABM-664,GSE59856	957	0.732 83651	0.62 340 82	2.02	0.441	4.74	0.7302 34	10
hsa-miR-486-5p	GSE71533,GSE85589,GSE24279,GSE60980,GSE32678,E-MTAB-753,GSE59856,GSE31568,GSE29352,E-TABM-664	957	0.750 03254	0.60 596 12	1.94	0.425	4.88	0.7301 94	10
hsa-miR-92b-5p	GSE32678,GSE60980,GSE85589,GSE71533,GSE59856,GSE24279,E-MTAB-753,GSE31568,GSE29352	839	0.697 65161	0.65 878 75	2.09	0.461	4.64	0.7298 05	9
hsa-miR-1267	GSE71533,GSE24279,E-MTAB-753,GSE29352,GSE60980,GSE85589,GSE59856,GSE31568,GSE32678	839	0.448 4762	0.83 595 74	2.82	0.661	4.42	0.7297 93	9
hsa-miR-493-5p	GSE71533,GSE60980,GSE29352,GSE32678,GSE24279,E-MTAB-753,GSE85589,E-TABM-664,GSE59856,GSE31568	957	0.699 49508	0.66 163 25	2.08	0.465	4.75	0.7294 02	10
hsa-miR-133a-3p	GSE31568,GSE71533,GSE24279,GSE60980,GSE85589,E-TABM-664,E-MTAB-753,GSE29352,GSE59856	925	0.453 6297	0.81 118 56	2.42	0.672	3.65	0.7292 26	9
hsa-miR-16-2-3p	GSE85589,GSE71533,E-TABM-664,GSE32678,GSE31568,GSE24279,GSE60980,E-MTAB-753,GSE29352,GSE59856	957	0.664 54284	0.69 048 82	2.16	0.496	4.59	0.7289 05	10

hsa-miR-450b-5p	GSE85589,GSE32678,E-MTAB-753,GSE31568,GSE29352,GSE71533,GSE24279,GSE59856,GSE60980	839	0.728 50513	0.62 545 36	2.03	0.454	4.6	0.7287 88	9
hsa-miR-1257	GSE71533,GSE32678,GSE24279,GSE31568,E-MTAB-753,GSE29352,GSE85589,GSE59856,GSE60980	839	0.567 68087	0.76 897 89	2.5	0.567	4.49	0.7286 73	9
hsa-miR-29b-3p	GSE71533,GSE29352,GSE60980,GSE24279,GSE31568,GSE85589,E-MTAB-753,E-TABM-664,GSE59856	925	0.776 76948	0.62 848 98	2.05	0.376	6.33	0.7286 24	9
hsa-miR-449b-5p	GSE71533,GSE31568,GSE32678,GSE24279,E-MTAB-753,GSE29352,GSE85589,GSE59856,GSE60980	839	0.591 67182	0.74 966 98	2.42	0.552	4.52	0.7284 21	9
hsa-miR-100-3p	GSE71533,GSE32678,GSE31568,GSE24279,GSE60980,GSE29352,GSE85589,E-MTAB-753,E-TABM-664,GSE59856	957	0.840 30288	0.45 674 7	1.59	0.368	4.6	0.7281 53	10
hsa-miR-1248	GSE71533,GSE32678,GSE29352,GSE24279,GSE60980,GSE31568,E-MTAB-753,GSE59856,GSE85589	839	0.540 03661	0.78 394 18	2.59	0.594	4.61	0.7278 43	9
hsa-miR-1273a	GSE71533,GSE32678,GSE59856,GSE85589,E-MTAB-753,GSE29352,GSE24279,GSE31568,GSE60980	839	0.734 7055	0.61 537 86	1.94	0.437	4.57	0.7275 84	9
hsa-miR-424-5p	GSE71533,GSE32678,GSE60980,GSE29352,GSE85589,E-MTAB-753,E-TABM-664,GSE59856,GSE31568,GSE24279	957	0.622 44735	0.72 744 25	2.34	0.523	4.55	0.7266 71	10
hsa-miR-570-3p	GSE71533,GSE32678,GSE85589,GSE29352,E-MTAB-753,GSE24279,GSE59856,GSE31568,GSE60980	839	0.542 91181	0.79 207 72	2.69	0.583	4.68	0.7265 34	9
hsa-miR-513b-5p	GSE60980,GSE71533,GSE32678,GSE24279,GSE31568,E-MTAB-753,GSE29352,GSE85589,GSE59856	839	0.782 38948	0.56 951 47	1.88	0.402	4.95	0.7265 25	9
hsa-miR-92a-2-5p	GSE32678,GSE71533,GSE59856,GSE31568,GSE24279,GSE60980,GSE85589,GSE29352,E-MTAB-753	839	0.667 87044	0.69 282 49	2.25	0.493	4.6	0.7264 39	9
hsa-miR-338-5p	GSE71533,GSE60980,GSE24279,GSE31568,GSE29352,E-TABM-664,GSE59856,GSE85589,E-MTAB-753	925	0.656 78948	0.69 264 77	2.15	0.505	4.57	0.7262	9
hsa-miR-320d	GSE71533,GSE31568,GSE60980,GSE24279,GSE59856,GSE29352,GSE32678,GSE85589,E-MTAB-753	839	0.757 81229	0.59 001 3	1.88	0.419	4.66	0.7261 35	9
hsa-miR-331-3p	GSE24279,E-MTAB-753,GSE60980,GSE71533,GSE85589,GSE31568,GSE29352,GSE59856,E-TABM-664	924	0.628 78108	0.71 013 89	2.17	0.526	4.3	0.7260 29	9
hsa-miR-634	GSE71533,GSE29352,GSE60980,GSE24279,E-MTAB-753,GSE31568,GSE85589,GSE32678,GSE59856	839	0.666 70576	0.68 464 02	2.13	0.497	4.79	0.7259 57	9

hsa-miR-221-3p	GSE71533,GSE24279,E-TABM-664,GSE60980,E-MTAB-753,GSE59856,GSE85589,GSE29352,GSE31568	925	0.786 92047	0.70 696 9	2.68	0.314	9.66	0.7251 93	9
hsa-miR-126-5p	GSE71533,GSE29352,GSE60980,GSE31568,GSE85589,GSE24279,E-TABM-664,E-MTAB-753,GSE59856	925	0.610 29255	0.72 745 86	2.28	0.542	4.47	0.7251 52	9
hsa-miR-1288-3p	GSE60980,GSE24279,GSE32678,GSE29352,E-MTAB-753,GSE31568,GSE71533,GSE85589,GSE59856	839	0.612 9212	0.71 916 63	2.2	0.542	4.14	0.7249 58	9
hsa-miR-500a-5p	GSE71533,GSE60980,GSE29352,GSE31568,GSE24279,E-MTAB-753,GSE32678,GSE59856,GSE85589	839	0.577 66783	0.74 793 76	2.32	0.572	4.25	0.7249 13	9
hsa-miR-660-5p	GSE71533,GSE29352,GSE60980,GSE32678,GSE85589,GSE31568,GSE24279,E-MTAB-753,GSE59856	839	0.578 39155	0.75 374 42	2.42	0.564	4.46	0.7242 82	9
hsa-miR-1204	GSE71533,GSE85589,GSE29352,GSE32678,E-MTAB-753,GSE60980,GSE31568,GSE24279,GSE59856	839	0.593 21534	0.73 453 64	2.28	0.562	4.26	0.7242 13	9
hsa-miR-128-3p	GSE60980,GSE31568,E-MTAB-753,GSE71533,E-TABM-664,GSE85589,GSE24279,GSE29352,GSE59856	925	0.479 29389	0.82 609 6	2.77	0.627	4.46	0.7239 09	9
hsa-miR-658	GSE59856,GSE60980,GSE71533,GSE31568,GSE85589,GSE32678,GSE24279,E-MTAB-753,GSE29352	839	0.673 16223	0.68 088 42	2.25	0.494	4.69	0.7235 95	9
hsa-miR-628-3p	GSE71533,GSE32678,GSE60980,GSE24279,GSE31568,GSE85589,GSE59856,GSE29352,E-MTAB-753	839	0.593 32277	0.75 621 39	2.5	0.546	4.62	0.7233 04	9
hsa-miR-15a-5p	GSE71533,GSE29352,GSE60980,GSE24279,E-MTAB-753,GSE31568,E-TABM-664,GSE59856,GSE85589	925	0.698 8401	0.64 812 13	2.01	0.467	4.41	0.7227 72	9
hsa-miR-500a-3p	E-MTAB-753,GSE60980,GSE24279,GSE29352,GSE31568,GSE71533,E-TABM-664,GSE85589,GSE32678,GSE59856	957	0.554 21224	0.75 794 49	2.29	0.593	3.94	0.7226 63	10
hsa-miR-1244	E-MTAB-753,GSE24279,GSE71533,GSE32678,GSE85589,GSE29352,GSE31568,GSE59856,GSE60980	839	0.733 05698	0.62 481 69	1.95	0.446	4.66	0.7224 4	9
hsa-miR-548n	GSE32678,GSE71533,GSE24279,E-MTAB-753,GSE60980,GSE31568,GSE85589,GSE59856,GSE29352	839	0.734 53192	0.59 376 8	1.91	0.466	4.16	0.7223 64	9
hsa-miR-525-5p	GSE71533,GSE32678,GSE29352,E-MTAB-753,GSE24279,GSE85589,GSE60980,GSE31568,GSE59856	839	0.600 84615	0.71 950 58	2.13	0.561	3.92	0.7217 98	9
hsa-miR-940	GSE71533,GSE24279,GSE85589,GSE59856,GSE32678,GSE31568,E-MTAB-753,GSE29352,GSE60980	839	0.538 18734	0.74 280 18	2.1	0.621	3.45	0.7210 69	9

hsa-miR-30d-5p	GSE71533,GSE24279,GSE60980,GSE29352,E-MTAB-753,GSE85589,GSE31568,GSE59856,E-TABM-664	925	0.802 50509	0.64 048 51	2.21	0.323	7.76	0.7210 38	9
hsa-miR-216a-5p	GSE24279,GSE71533,GSE60980,E-TABM-664,E-MTAB-753,GSE85589,GSE31568,GSE59856,GSE29352	925	0.842 69424	0.60 262 71	2.11	0.277	8.59	0.7204 91	9
hsa-miR-1285-3p	GSE59856,GSE71533,GSE60980,GSE31568,GSE24279,GSE85589,GSE29352,E-MTAB-753,GSE32678	839	0.602 25921	0.72 935 5	2.24	0.554	4.2	0.7199 72	9
hsa-miR-555	GSE71533,GSE60980,GSE29352,GSE32678,E-MTAB-753,GSE31568,GSE85589,GSE24279,GSE59856	839	0.688 93225	0.65 069 69	2.18	0.51	4.43	0.7197 59	9
hsa-miR-766-3p	GSE71533,GSE29352,GSE60980,GSE24279,GSE59856,GSE32678,E-MTAB-753,GSE31568,GSE85589	839	0.696 39396	0.65 563 93	2.04	0.472	4.63	0.7197 13	9
hsa-miR-218-5p	GSE60980,GSE85589,GSE71533,GSE24279,E-TABM-664,GSE29352,GSE31568,E-MTAB-753,GSE59856	925	0.659 83803	0.68 096 29	2.08	0.53	4.22	0.7196 63	9
hsa-miR-498	GSE71533,E-MTAB-753,GSE29352,GSE60980,GSE32678,GSE85589,GSE31568,E-TABM-664,GSE24279,GSE59856	957	0.669 35268	0.67 269 81	2.08	0.498	4.4	0.7195 8	10
hsa-miR-1243	GSE71533,GSE60980,GSE29352,GSE31568,GSE32678,E-MTAB-753,GSE85589,GSE24279,GSE59856	839	0.669 64032	0.67 233 5	2.14	0.507	4.32	0.7194 41	9
hsa-miR-1298-5p	GSE71533,GSE60980,GSE29352,GSE24279,E-MTAB-753,GSE59856,GSE85589,GSE31568,GSE32678	839	0.683 63148	0.65 733 44	2.07	0.494	4.41	0.7192 09	9
hsa-miR-106b-3p	GSE71533,GSE24279,GSE85589,E-MTAB-753,E-TABM-664,GSE31568,GSE29352,GSE60980,GSE59856	925	0.560 77699	0.75 077 63	2.28	0.585	3.97	0.7188 21	9
hsa-miR-548h-5p	GSE32678,GSE29352,E-MTAB-753,GSE85589,GSE31568,GSE24279,GSE60980,GSE59856,GSE71533	839	0.464 19384	0.81 482 27	2.53	0.662	3.91	0.7185 58	9
hsa-miR-494-3p	GSE59856,GSE60980,E-TABM-664,GSE85589,E-MTAB-753,GSE71533,GSE29352,GSE31568,GSE24279	925	0.732 74823	0.62 253 54	1.96	0.443	4.87	0.7183 11	9
hsa-miR-135b-3p	GSE32678,E-TABM-664,GSE60980,GSE71533,GSE29352,GSE31568,GSE85589,E-MTAB-753,GSE24279,GSE59856	957	0.581 5706	0.74 446 52	2.32	0.57	4.16	0.7181 41	10
hsa-miR-122-5p	GSE29352,E-TABM-664,GSE31568,E-MTAB-753,GSE60980,GSE85589,GSE24279,GSE59856,GSE71533	925	0.650 03357	0.68 632 04	2.08	0.511	4.16	0.7180 9	9
hsa-miR-324-3p	GSE24279,GSE60980,E-TABM-664,GSE71533,GSE31568,GSE59856,E-MTAB-753,GSE85589,GSE29352	925	0.740 59935	0.63 724 68	2.05	0.411	5.19	0.7180 02	9

hsa-miR-194-5p	GSE24279,GSE71533,GSE31568,E-MTAB-753,GSE29352,E-TABM-664,GSE59856,GSE60980,GSE85589	924	0.555 63233	0.74 768 74	2.23	0.591	3.88	0.7179 75	9
hsa-miR-345-5p	GSE32678,GSE24279,GSE31568,GSE71533,E-TABM-664,GSE29352,GSE60980,GSE85589,GSE59856,E-MTAB-753	957	0.638 36947	0.69 463 44	2.1	0.529	4.1	0.7178 81	10
hsa-miR-632	GSE29352,GSE32678,GSE24279,E-MTAB-753,GSE71533,GSE85589,GSE59856,GSE31568,GSE60980	839	0.440 89223	0.80 130 94	2.24	0.695	3.3	0.7175 59	9
hsa-miR-541-3p	GSE31568,GSE29352,GSE60980,GSE85589,GSE32678,GSE24279,E-MTAB-753,GSE59856,GSE71533	839	0.491 36352	0.82 307 85	2.86	0.622	4.63	0.7174 46	9
hsa-miR-503-5p	GSE24279,GSE32678,GSE60980,GSE85589,E-MTAB-753,GSE31568,E-TABM-664,GSE71533,GSE59856,GSE29352	957	0.650 89713	0.68 768 44	2.15	0.515	4.31	0.7172 12	10
hsa-miR-454-3p	GSE60980,GSE32678,GSE85589,GSE31568,E-MTAB-753,GSE24279,GSE71533,GSE29352,GSE59856	839	0.498 23396	0.79 989 9	2.53	0.629	4.09	0.7170 95	9
hsa-miR-576-5p	GSE71533,GSE31568,E-MTAB-753,GSE29352,GSE32678,GSE85589,GSE60980,GSE24279,GSE59856	839	0.592 05559	0.72 634 83	2.15	0.565	3.99	0.7167 65	9
hsa-miR-19b-3p	GSE71533,GSE29352,GSE59856,GSE85589,GSE24279,E-TABM-664,GSE60980,GSE31568,E-MTAB-753	925	0.732 95296	0.60 253 33	1.87	0.454	4.2	0.7166 26	9
hsa-miR-363-5p	GSE24279,GSE31568,GSE29352,E-TABM-664,E-MTAB-753,GSE60980,GSE59856,GSE85589,GSE71533	925	0.706 52182	0.63 022 4	1.93	0.472	4.16	0.7162 84	9
hsa-miR-518f-5p	GSE60980,GSE71533,GSE32678,E-MTAB-753,GSE29352,GSE31568,GSE24279,GSE85589,GSE59856	839	0.450 6491	0.82 624 31	2.62	0.66	4.06	0.7161 46	9
hsa-miR-602	GSE59856,GSE32678,GSE71533,GSE29352,GSE31568,E-MTAB-753,GSE24279,GSE60980,GSE85589	839	0.735 75007	0.59 656 23	1.86	0.452	4.28	0.7160 5	9
hsa-miR-519e-3p	GSE71533,GSE60980,E-MTAB-753,GSE29352,E-TABM-664,GSE32678,GSE59856,GSE31568,GSE24279,GSE85589	957	0.352 62197	0.87 072 42	2.86	0.739	3.97	0.7160 11	10
hsa-miR-890	GSE71533,GSE85589,GSE29352,GSE24279,E-MTAB-753,GSE32678,GSE31568,GSE59856,GSE60980	839	0.620 46594	0.70 455 24	2.11	0.541	4.06	0.7159 6	9
hsa-miR-140-3p	GSE60980,GSE29352,GSE24279,E-MTAB-753,E-TABM-664,GSE85589,GSE71533,GSE31568,GSE59856	925	0.553 03619	0.74 337 57	2.15	0.601	3.63	0.7158 43	9
hsa-miR-98-5p	GSE71533,GSE60980,GSE31568,GSE29352,GSE24279,E-TABM-664,E-MTAB-753,GSE85589,GSE59856	925	0.539 28415	0.76 012 92	2.27	0.609	3.84	0.7155 92	9

hsa-miR-191-5p	GSE85589,E-TABM-664,GSE24279,GSE29352,GSE59856,GSE71533,E-MTAB-753,GSE31568,GSE60980	924	0.624 43972	0.72 309 14	2.28	0.524	4.38	0.7149 87	9
hsa-miR-595	GSE31568,GSE32678,GSE71533,GSE85589,GSE29352,E-MTAB-753,GSE60980,GSE24279,GSE59856	839	0.545 59322	0.76 060 2	2.28	0.599	3.87	0.7149 08	9
hsa-miR-616-5p	GSE29352,GSE60980,GSE24279,GSE31568,GSE32678,E-MTAB-753,GSE71533,GSE59856,GSE85589	839	0.539 58875	0.77 908 68	2.48	0.596	4.25	0.7146 88	9
hsa-miR-30c-1-3p	GSE71533,GSE60980,GSE29352,E-TABM-664,GSE24279,GSE85589,E-MTAB-753,GSE59856,GSE31568	925	0.631 9603	0.69 426 4	2.09	0.533	4.08	0.7146 62	9
hsa-miR-452-5p	GSE60980,GSE71533,GSE32678,GSE24279,GSE31568,GSE85589,E-TABM-664,GSE59856,GSE29352,E-MTAB-753	957	0.545 73115	0.78 646 59	2.58	0.58	4.55	0.7145 62	10
hsa-miR-505-3p	GSE60980,GSE24279,GSE71533,E-TABM-664,GSE85589,GSE31568,GSE29352,GSE59856,E-MTAB-753	925	0.588 01057	0.72 946 55	2.18	0.574	3.93	0.7144 01	9
hsa-miR-379-3p	GSE71533,E-MTAB-753,GSE32678,GSE24279,GSE31568,GSE29352,GSE85589,E-TABM-664,GSE59856,GSE60980	957	0.432 20191	0.81 403 58	2.32	0.698	3.41	0.7141 26	10
hsa-miR-371a-5p	GSE71533,GSE32678,GSE59856,E-MTAB-753,GSE31568,GSE85589,GSE29352,E-TABM-664,GSE24279,GSE60980	957	0.741 49156	0.57 638 54	1.75	0.459	3.93	0.7137 1	10
hsa-miR-99a-5p	GSE71533,GSE29352,GSE60980,E-TABM-664,GSE85589,E-MTAB-753,GSE24279,GSE59856,GSE31568	925	0.650 63886	0.68 049 56	2.03	0.518	4.07	0.7137 05	9
hsa-miR-222-5p	GSE60980,GSE24279,GSE31568,E-MTAB-753,GSE71533,GSE32678,GSE59856,GSE29352,GSE85589	839	0.609 49249	0.71 358 22	2.14	0.552	4.03	0.7135 1	9
hsa-miR-656-3p	GSE71533,GSE32678,GSE60980,E-MTAB-753,GSE29352,GSE31568,GSE85589,GSE59856,GSE24279	839	0.463 90505	0.81 202 63	2.49	0.659	3.85	0.7134 16	9
hsa-miR-515-3p	GSE29352,GSE85589,GSE32678,E-MTAB-753,GSE71533,E-TABM-664,GSE31568,GSE24279,GSE59856,GSE60980	957	0.369 53536	0.86 725 04	2.85	0.726	3.96	0.7133 31	10
hsa-miR-10b-5p	GSE71533,GSE60980,E-TABM-664,GSE29352,GSE24279,GSE85589,E-MTAB-753,GSE31568,GSE59856	925	0.594 20639	0.71 138 19	2.05	0.572	3.74	0.7132 77	9
hsa-miR-450a-5p	GSE60980,GSE32678,GSE71533,GSE31568,GSE24279,GSE29352,E-TABM-664,GSE85589,GSE59856,E-MTAB-753	957	0.515 58782	0.78 123 55	2.36	0.624	3.84	0.7131 73	10
hsa-miR-1255a	GSE71533,GSE60980,E-MTAB-753,GSE32678,GSE85589,GSE59856,GSE29352,GSE24279,GSE31568	839	0.754 26496	0.59 72	1.88	0.434	5.1	0.7131 38	9

hsa-miR-552-3p	GSE71533,GSE32678,GSE24279,GSE60980,E-MTAB-753,GSE31568,GSE29352,GSE59856,GSE85589	839	0.661 79616	0.66 961 62	2.05	0.521	4.16	0.7129 47	9
hsa-miR-181c-3p	E-MTAB-753,GSE29352,GSE71533,GSE32678,GSE85589,GSE24279,GSE60980,GSE59856,GSE31568	839	0.575 48596	0.73 373 41	2.21	0.59	3.94	0.7126 98	9
hsa-miR-425-5p	GSE71533,GSE24279,GSE85589,GSE60980,E-MTAB-753,E-TABM-664,GSE59856,GSE29352,GSE31568	925	0.683 87801	0.64 398 39	1.97	0.495	4.01	0.7126 66	9
hsa-miR-25-3p	GSE71533,GSE59856,E-TABM-664,GSE29352,E-MTAB-753,GSE31568,GSE85589,GSE60980,GSE24279	925	0.585 63522	0.74 135 02	2.32	0.561	4.2	0.7123 82	9
hsa-miR-202-5p	GSE32678,GSE85589,E-MTAB-753,GSE71533,GSE24279,GSE60980,GSE29352,GSE59856,GSE31568	839	0.758 75065	0.53 840 13	1.72	0.471	3.88	0.7119 45	9
hsa-miR-1324	GSE71533,GSE32678,GSE60980,GSE24279,GSE29352,GSE85589,GSE31568,E-MTAB-753,GSE59856	839	0.473 27829	0.82 970 99	2.82	0.633	4.48	0.7116 68	9
hsa-miR-148a-5p	GSE71533,GSE24279,GSE60980,E-MTAB-753,GSE29352,GSE31568,GSE85589,E-TABM-664,GSE59856	925	0.537 36783	0.74 980 14	2.15	0.618	3.66	0.7114 89	9
hsa-miR-1304-5p	GSE71533,GSE32678,GSE29352,GSE60980,GSE31568,GSE59856,E-MTAB-753,GSE85589,GSE24279	839	0.598 56038	0.72 565 75	2.21	0.561	4.06	0.7112 13	9
hsa-miR-548a-5p	GSE71533,GSE32678,GSE85589,E-MTAB-753,GSE31568,GSE24279,GSE29352,GSE59856,GSE60980	839	0.786 40827	0.46 524 63	1.54	0.477	3.31	0.7111 63	9
hsa-miR-146b-5p	GSE60980,E-TABM-664,GSE85589,GSE71533,E-MTAB-753,GSE31568,GSE24279,GSE59856,GSE29352	925	0.683 28851	0.65 239 4	1.97	0.489	4.12	0.7111 62	9
hsa-miR-1183	GSE71533,GSE60980,GSE31568,GSE85589,GSE32678,E-MTAB-753,GSE24279,GSE59856,GSE29352	839	0.527 81654	0.75 157 24	2.11	0.631	3.48	0.7107 93	9
hsa-miR-770-5p	GSE85589,GSE60980,GSE32678,GSE24279,E-MTAB-753,GSE59856,GSE31568,GSE29352,GSE71533	839	0.564 00975	0.72 611 09	2.04	0.598	3.58	0.7100 23	9
hsa-miR-376a-3p	GSE71533,GSE24279,GSE60980,GSE29352,GSE85589,GSE31568,GSE59856,E-MTAB-753,E-TABM-664,GSE32678	957	0.532 78308	0.76 349 89	2.28	0.617	3.78	0.7096 28	10
hsa-miR-200c-5p	GSE71533,GSE60980,E-TABM-664,GSE31568,GSE24279,GSE29352,E-MTAB-753,GSE85589,GSE59856	925	0.461 46648	0.82 428 53	2.71	0.658	4.15	0.7095 52	9
hsa-miR-149-3p	GSE71533,GSE60980,GSE85589,GSE24279,GSE31568,E-MTAB-753,GSE59856,GSE29352,GSE32678	839	0.719 83349	0.60 933 78	1.86	0.463	4.1	0.7094 7	9

hsa-miR-302e	GSE71533,E-MTAB-753,GSE29352,GSE31568,GSE85589,GSE24279,GSE32678,GSE60980,GSE59856	839	0.624 14978	0.69 545 83	2.08	0.549	3.93	0.7092 18	9
hsa-miR-631	GSE32678,GSE29352,GSE60980,GSE71533,GSE85589,E-MTAB-753,GSE31568,GSE59856,GSE24279	839	0.693 29842	0.64 545 74	1.95	0.484	4.15	0.7090 24	9
hsa-miR-135a-5p	GSE29352,GSE31568,GSE85589,GSE24279,GSE71533,E-TABM-664,GSE60980,GSE59856,E-MTAB-753	925	0.529 47305	0.77 354 48	2.36	0.607	3.94	0.7089 35	9
hsa-miR-187-3p	GSE71533,E-TABM-664,GSE31568,GSE29352,E-MTAB-753,GSE85589,GSE60980,GSE59856,GSE24279	925	0.415 06488	0.85 498 06	2.94	0.68	4.36	0.7089 32	9
hsa-miR-376c-3p	GSE71533,GSE24279,GSE29352,GSE60980,E-TABM-664,GSE85589,E-MTAB-753,GSE59856,GSE31568,GSE32678	957	0.482 59264	0.79 709 49	2.39	0.649	3.76	0.7086 53	10
hsa-miR-889-3p	GSE71533,GSE60980,E-MTAB-753,GSE29352,GSE32678,GSE31568,GSE24279,GSE85589,GSE59856	839	0.509 10603	0.76 810 46	2.21	0.634	3.61	0.7085 05	9
hsa-miR-409-3p	GSE71533,GSE24279,GSE60980,GSE29352,E-MTAB-753,GSE31568,GSE85589,E-TABM-664,GSE59856	925	0.634 66333	0.68 355 55	1.98	0.548	3.81	0.7084 38	9
hsa-miR-29a-3p	GSE29352,GSE60980,GSE24279,GSE31568,E-MTAB-753,GSE71533,GSE85589,E-TABM-664,GSE59856	925	0.622 14689	0.69 864 66	2.1	0.544	4.05	0.7083 95	9
hsa-miR-1291	GSE71533,E-MTAB-753,GSE32678,GSE29352,GSE31568,GSE24279,GSE60980,GSE85589,GSE59856	839	0.646 10567	0.67 568 95	1.97	0.52	4.06	0.7082 83	9
hsa-miR-487a-3p	GSE71533,GSE32678,GSE60980,GSE24279,E-TABM-664,GSE29352,GSE85589,GSE31568,E-MTAB-753,GSE59856	957	0.528 13196	0.76 106 74	2.22	0.624	3.63	0.7082 47	10
hsa-miR-1256	GSE31568,GSE32678,E-MTAB-753,GSE71533,GSE60980,GSE85589,GSE24279,GSE59856,GSE29352	839	0.491 22603	0.77 941 51	2.24	0.659	3.48	0.7080 62	9
hsa-miR-619-3p	GSE60980,E-MTAB-753,GSE32678,GSE71533,GSE29352,GSE31568,GSE24279,GSE85589,GSE59856	839	0.587 16562	0.71 394 99	2.02	0.578	3.66	0.7078 32	9
hsa-miR-589-5p	GSE71533,GSE32678,GSE31568,GSE24279,GSE60980,GSE85589,E-MTAB-753,GSE29352,GSE59856	839	0.588 08088	0.72 085 59	2.12	0.578	3.79	0.7076 58	9
hsa-miR-1255b-5p	GSE60980,GSE32678,GSE24279,E-MTAB-753,GSE31568,GSE85589,GSE71533,GSE59856,GSE29352	839	0.428 08385	0.82 304 03	2.41	0.691	3.56	0.7073 76	9
hsa-miR-643	GSE32678,GSE29352,GSE85589,GSE60980,GSE71533,GSE31568,GSE59856,GSE24279,E-MTAB-753	839	0.600 68521	0.71 251 39	2.09	0.563	3.82	0.7072 63	9

hsa-miR-548i	GSE24279,GSE32678,GSE71533,GSE29352,GSE85589,E-MTAB-753,GSE31568,GSE59856,GSE60980	839	0.610 22519	0.70 782 68	2.16	0.561	3.99	0.7070 59	9
hsa-miR-122-3p	GSE71533,GSE32678,GSE85589,GSE60980,GSE31568,E-MTAB-753,GSE59856,GSE29352,GSE24279	838	0.576 14333	0.72 322 26	2.13	0.588	3.82	0.7069 08	9
hsa-miR-548g-3p	GSE71533,GSE60980,GSE31568,GSE24279,GSE85589,E-MTAB-753,GSE32678,GSE29352,GSE59856	839	0.613 45139	0.70 573 39	2.13	0.557	3.9	0.7068 59	9
hsa-miR-888-5p	E-MTAB-753,GSE85589,GSE24279,GSE31568,GSE29352,GSE71533,GSE32678,GSE59856,GSE60980	839	0.628 98317	0.68 935 32	2.07	0.547	3.89	0.7067 35	9
hsa-miR-508-3p	GSE60980,GSE29352,GSE31568,GSE32678,GSE71533,GSE24279,E-MTAB-753,GSE85589,E-TABM-664,GSE59856	957	0.566 04018	0.72 434 61	2.05	0.601	3.53	0.7067 29	10
hsa-miR-151a-3p	GSE71533,GSE24279,GSE60980,GSE31568,E-TABM-664,GSE59856,GSE32678,GSE85589,GSE29352,E-MTAB-753	957	0.598 45473	0.70 131 16	2	0.575	3.58	0.7063 91	10
hsa-miR-451a	GSE59856,GSE29352,GSE71533,GSE24279,GSE85589,GSE60980,E-MTAB-753,E-TABM-664,GSE31568	925	0.655 24477	0.69 048 54	2.19	0.504	4.43	0.7062 89	9
hsa-miR-1276	GSE71533,GSE29352,GSE32678,GSE60980,GSE31568,GSE85589,GSE59856,GSE24279,E-MTAB-753	839	0.588 87567	0.72 053 97	2.15	0.574	3.91	0.7062 03	9
hsa-miR-302a-3p	GSE71533,GSE29352,GSE24279,GSE32678,GSE85589,GSE31568,E-MTAB-753,GSE59856,GSE60980	839	0.709 79963	0.59 805 2	1.86	0.504	3.74	0.7058 26	9
hsa-miR-33b-5p	GSE71533,E-TABM-664,GSE24279,GSE60980,GSE29352,E-MTAB-753,GSE31568,GSE85589,GSE59856,GSE32678	957	0.628 34743	0.69 010 51	2.07	0.545	3.85	0.7058 03	10
hsa-miR-136-5p	GSE71533,GSE31568,GSE24279,GSE29352,GSE85589,GSE60980,E-TABM-664,GSE59856,E-MTAB-753	925	0.466 79509	0.81 807 07	2.64	0.653	4.14	0.7056 53	9
hsa-miR-30b-3p	GSE32678,GSE59856,GSE71533,E-MTAB-753,GSE24279,E-TABM-664,GSE31568,GSE60980,GSE29352,GSE85589	957	0.655 18005	0.68 557 17	2.16	0.509	4.37	0.7056 44	10
hsa-miR-1208	GSE29352,GSE32678,GSE60980,GSE31568,GSE24279,GSE85589,GSE59856,E-MTAB-753,GSE71533	839	0.504 82407	0.75 501 28	2.07	0.655	3.23	0.7055 83	9
hsa-miR-620	GSE71533,GSE24279,GSE32678,E-MTAB-753,GSE85589,GSE60980,GSE31568,GSE59856,GSE29352	839	0.593 63764	0.71 835 42	2.16	0.578	3.99	0.7055 21	9
hsa-miR-338-3p	GSE71533,GSE60980,GSE24279,E-TABM-664,GSE85589,GSE31568,GSE29352,E-MTAB-753,GSE59856	925	0.522 20946	0.77 056 32	2.29	0.623	3.75	0.7054 98	9

hsa-miR-423-3p	GSE24279,GSE60980,E-MTAB-753,E-TABM-664,GSE31568,GSE29352,GSE59856,GSE85589,GSE71533	925	0.552 32813	0.73 404 51	2.08	0.618	3.47	0.7053 77	9
hsa-miR-627-5p	GSE60980,GSE32678,GSE29352,GSE85589,GSE71533,GSE24279,E-MTAB-753,GSE31568,GSE59856	839	0.554 35754	0.75 140 97	2.26	0.603	3.82	0.7052 85	9
hsa-miR-1827	GSE71533,GSE60980,GSE29352,GSE31568,E-MTAB-753,GSE24279,GSE32678,GSE85589,GSE59856	839	0.531 12904	0.75 061 37	2.12	0.619	3.54	0.7050 56	9
hsa-miR-106b-5p	GSE29352,GSE71533,GSE60980,E-TABM-664,E-MTAB-753,GSE24279,GSE59856,GSE85589,GSE31568	925	0.769 06187	0.53 452 09	1.65	0.439	3.84	0.7050 47	9
hsa-miR-497-5p	GSE29352,GSE60980,GSE71533,GSE24279,GSE85589,GSE31568,E-MTAB-753,E-TABM-664,GSE59856	925	0.595 90661	0.72 743 31	2.27	0.568	4.04	0.7050 36	9
hsa-miR-184	GSE71533,GSE24279,GSE31568,GSE29352,GSE59856,E-TABM-664,E-MTAB-753,GSE85589,GSE60980	925	0.664 7036	0.65 671 31	1.96	0.514	3.91	0.7049 59	9
hsa-miR-1265	GSE29352,GSE32678,E-MTAB-753,GSE31568,GSE71533,GSE24279,GSE59856,GSE85589,GSE60980	839	0.755 48097	0.53 308 34	1.67	0.475	3.62	0.7049 59	9
hsa-miR-1245a	GSE31568,GSE24279,E-MTAB-753,GSE29352,GSE32678,GSE85589,GSE60980,GSE71533,GSE59856	839	0.542 39771	0.75 936	2.29	0.605	3.85	0.7047 81	9
hsa-miR-516b-5p	GSE32678,GSE71533,GSE60980,E-TABM-664,GSE85589,E-MTAB-753,GSE29352,GSE24279,GSE31568,GSE59856	957	0.478 70396	0.81 016 28	2.57	0.649	4.01	0.7047 21	10
hsa-miR-518d-3p	GSE71533,GSE85589,GSE60980,GSE31568,GSE24279,GSE29352,GSE32678,E-MTAB-753,E-TABM-664,GSE59856	957	0.444 87688	0.81 385 49	2.42	0.683	3.59	0.7046 18	10
hsa-miR-301b-3p	GSE32678,GSE71533,GSE60980,GSE85589,GSE24279,GSE29352,GSE59856,E-MTAB-753,GSE31568	839	0.565 6773	0.73 046 92	2.11	0.591	3.66	0.7045 57	9
hsa-miR-591	GSE60980,GSE71533,GSE31568,GSE85589,GSE24279,GSE29352,GSE59856,GSE32678,E-MTAB-753	839	0.534 25354	0.76 443 93	2.32	0.61	3.87	0.7043 76	9
hsa-miR-330-3p	GSE71533,GSE60980,GSE29352,E-TABM-664,GSE32678,GSE31568,E-MTAB-753,GSE24279,GSE85589,GSE59856	957	0.584 43669	0.71 450 87	2.04	0.583	3.63	0.7043 06	10
hsa-miR-625-5p	GSE32678,GSE71533,GSE29352,GSE24279,E-MTAB-753,GSE85589,GSE60980,GSE31568,GSE59856	839	0.637 11516	0.67 579 25	1.98	0.54	3.89	0.7041 76	9
hsa-miR-611	GSE32678,GSE71533,E-MTAB-753,GSE29352,GSE85589,GSE60980,GSE24279,GSE31568,GSE59856	839	0.537 17731	0.76 694 47	2.38	0.611	3.95	0.7041 54	9

hsa-miR-1303	GSE71533,GSE32678,GSE31568,GSE59856,E-MTAB-753,GSE60980,GSE24279,GSE85589,GSE29352	839	0.665 07267	0.65 036 71	1.93	0.519	3.83	0.7031 18	9
hsa-miR-885-5p	GSE32678,E-MTAB-753,GSE31568,GSE71533,GSE24279,GSE29352,GSE60980,GSE85589,GSE59856	839	0.730 38833	0.57 775 36	1.75	0.478	3.82	0.7013 78	9
hsa-miR-377-5p	GSE32678,GSE85589,GSE71533,GSE29352,E-MTAB-753,E-TABM-664,GSE60980,GSE31568,GSE24279,GSE59856	957	0.338 38118	0.85 476 44	2.34	0.772	3.06	0.7013 48	10
hsa-miR-937-3p	GSE71533,GSE32678,GSE29352,GSE31568,GSE59856,GSE24279,GSE85589,E-MTAB-753,GSE60980	839	0.685 88016	0.62 303 66	1.86	0.513	3.75	0.7012 69	9
hsa-miR-369-3p	GSE71533,GSE29352,GSE60980,GSE31568,GSE85589,GSE24279,E-TABM-664,GSE59856,E-MTAB-753	925	0.826 52665	0.49 198 16	1.65	0.374	4.87	0.7012 25	9
hsa-miR-34c-5p	GSE60980,GSE32678,GSE71533,GSE24279,GSE85589,E-MTAB-753,GSE31568,GSE29352,E-TABM-664,GSE59856	957	0.445 92675	0.79 699 17	2.2	0.689	3.25	0.7011 97	10
hsa-miR-136-3p	GSE71533,GSE60980,GSE24279,GSE85589,E-TABM-664,E-MTAB-753,GSE29352,GSE59856,GSE31568	925	0.485 45063	0.76 163 05	2.04	0.673	3.16	0.7007 71	9
hsa-miR-557	GSE32678,GSE71533,GSE59856,GSE60980,GSE24279,GSE85589,GSE31568,E-MTAB-753,GSE29352	839	0.571 95579	0.71 745 3	2.05	0.6	3.57	0.7005 47	9
hsa-miR-551b-5p	GSE32678,GSE24279,GSE60980,GSE29352,GSE59856,GSE31568,GSE71533,GSE85589,E-MTAB-753	839	0.607 76603	0.68 979	1.97	0.573	3.49	0.7005 46	9
hsa-miR-16-1-3p	GSE71533,GSE29352,E-MTAB-753,GSE31568,E-TABM-664,GSE32678,GSE24279,GSE85589,GSE60980,GSE59856	957	0.729 97123	0.56 703 12	1.78	0.499	3.66	0.7005 11	10
hsa-miR-147a	GSE31568,GSE32678,E-TABM-664,GSE29352,E-MTAB-753,GSE71533,GSE24279,GSE85589,GSE59856,GSE60980	957	0.747 31647	0.55 097 57	1.71	0.469	3.71	0.7004 34	10
hsa-miR-1237-3p	GSE32678,GSE59856,GSE24279,GSE29352,E-MTAB-753,GSE60980,GSE71533,GSE31568,GSE85589	839	0.692 59757	0.61 463 32	1.84	0.507	3.75	0.7003 75	9
hsa-miR-665	GSE59856,GSE71533,GSE85589,GSE32678,GSE29352,GSE31568,GSE60980,GSE24279,E-MTAB-753	839	0.691 79641	0.61 263 38	1.83	0.511	3.79	0.7000 71	9
hsa-miR-512-3p	GSE60980,GSE85589,E-MTAB-753,GSE31568,GSE29352,GSE24279,GSE71533,GSE59856,E-TABM-664,GSE32678	957	0.481 94842	0.78 304 12	2.25	0.66	3.48	0.6998 56	10
hsa-miR-654-5p	GSE59856,GSE71533,GSE31568,GSE29352,E-MTAB-753,GSE24279,GSE60980,GSE85589,GSE32678	839	0.559 82839	0.76 076 04	2.39	0.583	4.15	0.6996 73	9

hsa-let-7a-5p	GSE32678,GSE60980,GSE59856,GSE71533,GSE29352,E-TABM-664,GSE85589,GSE31568,GSE24279	923	0.606 86742	0.69 093 31	1.97	0.569	3.53	0.6994 78	9
hsa-miR-1297	GSE71533,GSE29352,GSE60980,GSE24279,GSE85589,E-MTAB-753,GSE31568,GSE59856,GSE32678	839	0.666 76517	0.64 494 1	1.89	0.534	3.7	0.6993 78	9
hsa-miR-454-5p	GSE29352,GSE32678,GSE31568,GSE24279,E-MTAB-753,GSE60980,GSE71533,GSE59856,GSE85589	839	0.630 11708	0.67 452 56	1.92	0.561	3.59	0.6993 09	9
hsa-miR-21-5p	GSE29352,GSE60980,E-TABM-664,GSE71533,E-MTAB-753,GSE31568,GSE24279,GSE59856,GSE85589	925	0.650 48883	0.68 754 09	2.08	0.514	4.26	0.6990 76	9
hsa-miR-302a-5p	GSE29352,E-TABM-664,GSE71533,GSE32678,E-MTAB-753,GSE31568,GSE85589,GSE24279,GSE59856,GSE60980	957	0.486 55566	0.78 529 92	2.29	0.657	3.54	0.6988 97	10
hsa-miR-548l	GSE71533,E-MTAB-753,GSE85589,GSE24279,GSE29352,GSE59856,GSE32678,GSE31568,GSE60980	839	0.475 88772	0.76 546 67	2.05	0.686	3.07	0.6986 38	9
hsa-miR-561-3p	GSE71533,GSE31568,GSE32678,E-MTAB-753,GSE29352,GSE60980,GSE85589,GSE24279,GSE59856	839	0.497 45694	0.75 208 42	2	0.665	3.14	0.6985 92	9
hsa-miR-541-5p	GSE29352,GSE60980,GSE71533,GSE32678,GSE85589,GSE59856,E-MTAB-753,GSE24279,GSE31568	839	0.626 6421	0.67 732 86	1.94	0.569	3.64	0.6985 81	9
hsa-miR-380-5p	GSE71533,GSE60980,GSE32678,GSE24279,GSE31568,GSE29352,E-TABM-664,GSE59856,GSE85589,E-MTAB-753	957	0.486 21712	0.79 371 6	2.39	0.65	3.74	0.6983 71	10
hsa-miR-135a-3p	GSE71533,GSE60980,GSE31568,E-MTAB-753,GSE29352,GSE59856,GSE24279,E-TABM-664,GSE85589	925	0.550 49213	0.74 334 3	2.18	0.612	3.74	0.6981 35	9
hsa-miR-767-3p	GSE71533,GSE60980,GSE32678,GSE31568,GSE59856,E-MTAB-753,GSE24279,GSE29352,GSE85589	839	0.569 63389	0.71 800 47	2.02	0.602	3.43	0.6981 08	9
hsa-miR-513c-5p	GSE60980,GSE24279,E-MTAB-753,GSE29352,GSE31568,GSE71533,GSE85589,GSE59856,GSE32678	839	0.612 3536	0.68 883 55	2	0.573	3.59	0.6980 64	9
hsa-miR-659-3p	GSE60980,GSE32678,GSE24279,GSE29352,GSE71533,GSE85589,GSE59856,GSE31568,E-MTAB-753	839	0.759 8113	0.54 517 34	1.68	0.446	3.89	0.6979 65	9
hsa-miR-192-5p	GSE24279,GSE60980,GSE71533,GSE29352,E-MTAB-753,GSE31568,E-TABM-664,GSE59856,GSE85589	925	0.587 17528	0.69 384 73	1.95	0.594	3.48	0.6979 34	9
hsa-miR-613	GSE71533,GSE32678,GSE29352,GSE60980,E-MTAB-753,GSE24279,GSE85589,GSE59856,GSE31568	839	0.468 22963	0.81 363 29	2.62	0.657	4.09	0.6977 45	9

hsa-miR-190a-5p	GSE29352,GSE71533,GSE60980,E-MTAB-753,GSE31568,GSE24279,E-TABM-664,GSE59856,GSE85589	925	0.566 55391	0.72 659 48	2.1	0.605	3.59	0.6975 27	9
hsa-miR-633	GSE71533,GSE32678,GSE60980,GSE85589,E-MTAB-753,GSE24279,GSE29352,GSE59856,GSE31568	839	0.747 05477	0.57 048 56	1.74	0.455	4.07	0.6972 5	9
hsa-miR-144-3p	GSE31568,E-MTAB-753,GSE85589,GSE60980,GSE71533,GSE24279,E-TABM-664,GSE29352,GSE59856	925	0.729 68886	0.56 028 08	1.75	0.501	3.65	0.6972 27	9
hsa-miR-1294	GSE71533,GSE24279,GSE32678,E-MTAB-753,GSE85589,GSE31568,GSE59856,GSE29352,GSE60980	839	0.574 70508	0.71 797 04	2.07	0.598	3.66	0.6971 11	9
hsa-miR-377-3p	GSE71533,GSE29352,GSE60980,GSE85589,E-MTAB-753,GSE59856,GSE24279,GSE31568,E-TABM-664	925	0.548 0498	0.73 763 23	2.1	0.619	3.45	0.6970 31	9
hsa-miR-1250-5p	GSE31568,GSE85589,E-MTAB-753,GSE60980,GSE32678,GSE24279,GSE29352,GSE71533,GSE59856	839	0.556 10629	0.71 958 28	2	0.616	3.33	0.6965 26	9
hsa-miR-515-5p	GSE24279,E-MTAB-753,GSE32678,GSE31568,GSE60980,E-TABM-664,GSE59856,GSE85589,GSE29352,GSE71533	957	0.559 96443	0.72 583 87	2.05	0.611	3.42	0.6963 4	10
hsa-miR-513a-3p	GSE29352,GSE71533,GSE31568,GSE85589,GSE32678,GSE24279,GSE60980,E-TABM-664,E-MTAB-753,GSE59856	957	0.547 43985	0.73 682 47	2.09	0.61	3.49	0.6962 51	10
hsa-miR-224-5p	GSE60980,GSE24279,GSE29352,GSE71533,GSE85589,E-TABM-664,GSE31568,E-MTAB-753,GSE59856	925	0.751 03607	0.52 608 53	1.63	0.487	3.42	0.6962 48	9
hsa-miR-501-5p	GSE24279,GSE60980,GSE32678,GSE29352,GSE71533,GSE31568,E-MTAB-753,GSE85589,E-TABM-664,GSE59856	957	0.589 43544	0.70 481 77	2.02	0.588	3.58	0.6961 6	10
hsa-miR-342-3p	GSE71533,GSE29352,GSE60980,E-TABM-664,GSE24279,GSE31568,E-MTAB-753,GSE85589,GSE59856	925	0.622 75243	0.68 389 56	2	0.557	3.67	0.6961	9
hsa-miR-624-5p	GSE71533,GSE32678,GSE85589,GSE24279,GSE31568,GSE29352,E-MTAB-753,GSE59856,GSE60980	839	0.471 32001	0.80 406 33	2.45	0.66	3.75	0.6960 94	9
hsa-miR-648	GSE60980,GSE31568,GSE32678,E-MTAB-753,GSE24279,GSE85589,GSE71533,GSE29352,GSE59856	839	0.455 47333	0.80 577 89	2.35	0.679	3.52	0.6959 63	9
hsa-miR-490-3p	GSE71533,GSE29352,GSE60980,GSE31568,E-MTAB-753,GSE24279,GSE59856,E-TABM-664,GSE85589	925	0.448 32014	0.76 059	1.86	0.724	2.65	0.6958 87	9
hsa-miR-342-5p	GSE24279,GSE60980,GSE71533,E-TABM-664,GSE29352,E-MTAB-753,GSE59856,GSE85589,GSE31568	925	0.727 27895	0.54 465 01	1.63	0.512	3.36	0.6958 36	9

hsa-miR-204-5p	GSE24279,GSE31568,GSE71533,E-MTAB-753,GSE29352,E-TABM-664,GSE60980,GSE85589,GSE59856	925	0.507 02815	0.76 397 07	2.17	0.647	3.4	0.6957 11	9
hsa-miR-362-3p	GSE71533,GSE60980,GSE29352,GSE32678,GSE24279,E-TABM-664,GSE59856,GSE31568,E-MTAB-753,GSE85589	957	0.532 80331	0.74 145 09	2.07	0.644	3.38	0.6957	10
hsa-miR-943	GSE71533,GSE32678,GSE31568,GSE85589,GSE59856,GSE29352,E-MTAB-753,GSE24279,GSE60980	839	0.589 72237	0.70 344 32	2.01	0.585	3.52	0.6956 75	9
hsa-miR-1284	GSE31568,GSE71533,GSE29352,E-MTAB-753,GSE24279,GSE60980,GSE32678,GSE59856,GSE85589	839	0.547 76133	0.74 247 68	2.17	0.611	3.61	0.6955 61	9
hsa-miR-523-3p	GSE32678,E-MTAB-753,GSE60980,GSE24279,GSE31568,GSE29352,GSE59856,GSE85589,GSE71533	839	0.492 43875	0.75 327 66	1.99	0.675	3.04	0.6955 51	9
hsa-miR-635	GSE71533,GSE32678,GSE85589,GSE31568,GSE29352,E-MTAB-753,GSE60980,GSE24279,GSE59856	839	0.602 42195	0.71 079 28	2.28	0.577	3.98	0.6954 49	9
hsa-miR-944	GSE71533,GSE32678,GSE60980,GSE85589,GSE31568,GSE29352,GSE24279,GSE59856,E-MTAB-753	839	0.561 02581	0.71 343 12	1.96	0.618	3.27	0.6954 18	9
hsa-miR-520a-5p	GSE71533,GSE32678,GSE24279,GSE31568,GSE85589,E-MTAB-753,GSE60980,GSE59856,GSE29352	839	0.522 04819	0.74 938 27	2.1	0.641	3.39	0.6953 22	9
hsa-miR-300	GSE31568,GSE32678,E-MTAB-753,GSE71533,GSE85589,GSE60980,GSE29352,GSE24279,GSE59856	839	0.481 03465	0.76 326 44	2.02	0.673	3.11	0.6951 76	9
hsa-miR-7-5p	GSE71533,GSE24279,E-MTAB-753,GSE60980,GSE85589,E-TABM-664,GSE29352,GSE31568,GSE59856	925	0.493 11304	0.79 979 3	2.55	0.644	4.02	0.6949 2	9
hsa-miR-1264	GSE71533,GSE24279,GSE32678,GSE29352,GSE85589,GSE60980,E-MTAB-753,GSE59856,GSE31568	839	0.768 38477	0.51 936 61	1.65	0.46	3.73	0.6949 17	9
hsa-miR-148b-3p	GSE60980,E-TABM-664,GSE71533,GSE29352,GSE24279,GSE59856,E-MTAB-753,GSE31568,GSE85589	925	0.632 84672	0.67 282 59	2.02	0.552	3.76	0.6939 94	9
hsa-miR-1295a	GSE60980,GSE31568,GSE29352,E-MTAB-753,GSE32678,GSE24279,GSE71533,GSE59856,GSE85589	839	0.656 50804	0.64 695 72	1.86	0.539	3.6	0.6939 25	9
hsa-miR-934	GSE71533,GSE60980,GSE24279,GSE31568,GSE85589,E-MTAB-753,GSE32678,GSE29352,GSE59856	839	0.459 61659	0.80 264 29	2.42	0.678	3.69	0.6936 95	9
hsa-miR-577	GSE71533,GSE29352,GSE85589,GSE32678,GSE24279,GSE60980,GSE31568,E-MTAB-753,GSE59856	838	0.812 26055	0.49 025 08	1.61	0.396	4.33	0.6936 06	9

hsa-miR-1305	GSE60980,E-MTAB-753,GSE31568,GSE85589,GSE71533,GSE29352,GSE24279,GSE32678,GSE59856	839	0.705 7958	0.57 819 45	1.76	0.519	3.46	0.6935 72	9
hsa-miR-1266-5p	GSE71533,GSE31568,GSE29352,GSE24279,E-MTAB-753,GSE60980,GSE59856,GSE85589,GSE32678	839	0.451 58115	0.80 769 4	2.4	0.683	3.55	0.6933 93	9
hsa-miR-518c-5p	GSE71533,GSE60980,GSE32678,GSE59856,GSE29352,GSE85589,GSE31568,E-MTAB-753,GSE24279	839	0.637 70494	0.66 426 38	1.92	0.552	3.52	0.6932 69	9
hsa-miR-550a-5p	GSE59856,GSE85589,GSE31568,E-MTAB-753,GSE24279,GSE29352,GSE71533,GSE60980,GSE32678	839	0.680 97505	0.62 190 3	1.8	0.515	3.55	0.6928 99	9
hsa-miR-509-5p	GSE60980,GSE71533,GSE85589,GSE32678,E-MTAB-753,GSE29352,GSE31568,E-TABM-664,GSE24279,GSE59856	957	0.485 45397	0.82 828 63	2.98	0.625	4.83	0.6927 84	10
hsa-miR-522-3p	GSE29352,GSE32678,GSE24279,GSE60980,GSE31568,GSE71533,E-MTAB-753,GSE59856,GSE85589	839	0.694 48259	0.60 205 65	1.8	0.518	3.54	0.6927 51	9
hsa-miR-769-3p	GSE71533,E-MTAB-753,GSE32678,GSE59856,GSE60980,GSE31568,GSE85589,GSE24279,GSE29352	838	0.739 67051	0.57 449 63	1.74	0.461	3.94	0.6927 25	9
hsa-miR-580-3p	GSE71533,GSE60980,GSE24279,GSE31568,E-MTAB-753,GSE85589,GSE29352,GSE32678,GSE59856	839	0.425 90818	0.81 088 38	2.25	0.706	3.24	0.6926 33	9
hsa-miR-488-5p	GSE60980,GSE71533,GSE31568,E-TABM-664,GSE85589,E-MTAB-753,GSE59856,GSE24279,GSE29352,GSE32678	957	0.508 46498	0.74 668 14	2.03	0.658	3.22	0.6920 94	10
hsa-miR-1224-3p	GSE29352,GSE31568,GSE85589,E-MTAB-753,GSE71533,GSE32678,GSE24279,GSE60980,GSE59856	839	0.661 00616	0.64 169 5	1.85	0.534	3.59	0.6915 22	9
hsa-miR-299-5p	GSE71533,E-MTAB-753,E-TABM-664,GSE60980,GSE24279,GSE29352,GSE31568,GSE85589,GSE59856	925	0.573 38249	0.70 898 65	1.99	0.611	3.34	0.6913 86	9
hsa-miR-301a-3p	GSE60980,GSE29352,GSE24279,GSE31568,GSE85589,E-MTAB-753,E-TABM-664,GSE71533,GSE59856	925	0.551 65279	0.72 415 61	2.02	0.624	3.35	0.6913 36	9
hsa-miR-520g-3p	GSE71533,GSE29352,GSE32678,GSE60980,E-TABM-664,GSE24279,GSE85589,E-MTAB-753,GSE31568,GSE59856	957	0.585 99567	0.71 213 88	2.06	0.59	3.57	0.6912 68	10
hsa-miR-939-5p	E-MTAB-753,GSE60980,GSE71533,GSE85589,GSE32678,GSE29352,GSE59856,GSE31568,GSE24279	839	0.669 26501	0.64 627 81	1.86	0.522	3.84	0.6912 3	9
hsa-miR-1234-3p	GSE60980,E-MTAB-753,GSE71533,GSE29352,GSE24279,GSE59856,GSE32678,GSE85589,GSE31568	839	0.678 92151	0.62 729 45	1.83	0.517	3.64	0.6908 51	9

hsa-miR-553	GSE71533,GSE32678,GSE31568,E-MTAB-753,GSE85589,GSE29352,GSE60980,GSE24279,GSE59856	839	0.695 20968	0.57 901 66	1.81	0.559	3.44	0.6904 65	9
hsa-miR-765	GSE71533,GSE60980,GSE32678,GSE24279,GSE29352,E-MTAB-753,GSE31568,GSE59856,GSE85589	839	0.582 44386	0.69 739 16	1.93	0.599	3.27	0.6904 2	9
hsa-miR-1286	GSE71533,GSE32678,GSE60980,GSE85589,E-MTAB-753,GSE29352,GSE24279,GSE31568,GSE59856	839	0.446 15514	0.77 095 09	1.96	0.717	2.79	0.6900 46	9
hsa-miR-193a-5p	GSE24279,GSE60980,E-TABM-664,GSE85589,GSE31568,GSE29352,E-MTAB-753,GSE59856,GSE71533,GSE32678	957	0.492 31979	0.74 225 75	1.92	0.684	2.84	0.6900 37	10
hsa-miR-101-5p	GSE60980,E-MTAB-753,E-TABM-664,GSE31568,GSE71533,GSE24279,GSE85589,GSE59856,GSE29352	925	0.464 86745	0.80 829 15	2.46	0.661	3.73	0.6899 76	9
hsa-miR-639	GSE71533,E-MTAB-753,GSE85589,GSE31568,GSE24279,GSE32678,GSE59856,GSE60980,GSE29352	839	0.482 22343	0.77 393 59	2.14	0.671	3.25	0.6898 81	9
hsa-miR-23b-5p	GSE60980,GSE29352,GSE71533,GSE85589,E-MTAB-753,GSE31568,GSE59856,GSE32678,GSE24279	839	0.460 90484	0.79 317 59	2.24	0.68	3.35	0.6898 34	9
hsa-miR-491-3p	GSE71533,E-TABM-664,GSE32678,GSE31568,GSE85589,E-MTAB-753,GSE60980,GSE29352,GSE59856,GSE24279	957	0.724 9569	0.55 022 02	1.68	0.513	3.38	0.6896 91	10
hsa-miR-1279	GSE29352,GSE60980,GSE31568,GSE32678,GSE24279,GSE85589,GSE59856,GSE71533,E-MTAB-753	839	0.492 23524	0.77 760 73	2.26	0.658	3.47	0.6895 44	9
hsa-miR-663b	GSE71533,E-MTAB-753,GSE32678,GSE60980,GSE59856,GSE24279,GSE31568,GSE29352,GSE85589	839	0.598 81357	0.68 162 51	1.89	0.589	3.28	0.6895 32	9
hsa-miR-589-3p	GSE32678,GSE60980,GSE85589,E-MTAB-753,GSE31568,GSE29352,GSE71533,GSE59856,GSE24279	839	0.644 20436	0.65 095 14	1.83	0.558	3.58	0.6892 54	9
hsa-miR-376a-5p	GSE60980,GSE29352,GSE32678,E-MTAB-753,GSE31568,GSE24279,GSE71533,GSE59856,GSE85589	839	0.581 02872	0.69 806 2	1.92	0.607	3.27	0.6892 52	9
hsa-miR-892a	GSE29352,GSE32678,GSE60980,GSE24279,GSE31568,GSE85589,E-MTAB-753,GSE59856,GSE71533	839	0.546 64916	0.71 104 07	1.88	0.633	3.11	0.6890 84	9
hsa-miR-1224-5p	GSE71533,GSE32678,GSE60980,GSE59856,GSE85589,GSE24279,GSE29352,GSE31568,E-MTAB-753	839	0.594 36522	0.68 950 76	1.91	0.586	3.33	0.6889 81	9
hsa-miR-211-5p	GSE85589,GSE71533,GSE32678,GSE31568,GSE29352,GSE60980,GSE24279,E-MTAB-753,E-TABM-664,GSE59856	957	0.503 52173	0.75 890 02	2.12	0.654	3.33	0.6886 42	10

hsa-miR-326	GSE71533,GSE60980,E-MTAB-753,GSE29352,GSE24279,GSE59856,GSE85589,GSE31568,E-TABM-664	925	0.606 09365	0.68 28	1.95	0.584	3.46	0.6886 23	9
hsa-miR-548c-3p	E-MTAB-753,GSE29352,GSE85589,GSE71533,GSE32678,GSE60980,GSE24279,GSE59856,GSE31568	838	0.477 24229	0.77 082 06	2.11	0.681	3.16	0.6884 64	9
hsa-miR-218-1-3p	GSE71533,GSE31568,E-MTAB-753,GSE29352,GSE85589,GSE60980,GSE32678,GSE59856,GSE24279	839	0.573 47448	0.72 930 4	2.17	0.594	3.68	0.6883 77	9
hsa-miR-96-5p	GSE71533,GSE60980,GSE24279,E-TABM-664,E-MTAB-753,GSE29352,GSE85589,GSE31568,GSE59856	925	0.601 55837	0.68 835 38	1.97	0.584	3.45	0.6881 82	9
hsa-miR-183-3p	GSE71533,GSE85589,GSE24279,E-MTAB-753,E-TABM-664,GSE31568,GSE59856,GSE60980,GSE29352	925	0.625 1434	0.66 260 11	1.85	0.575	3.39	0.6880 74	9
hsa-miR-9-5p	GSE24279,GSE60980,E-TABM-664,GSE31568,GSE71533,E-MTAB-753,GSE85589,GSE29352,GSE59856	925	0.425 53497	0.84 562 11	2.83	0.68	4.17	0.6878 78	9
hsa-miR-520e	GSE85589,GSE71533,GSE24279,GSE29352,E-TABM-664,GSE59856,GSE32678,GSE60980,GSE31568,E-MTAB-753	957	0.474 50784	0.78 775 7	2.29	0.67	3.45	0.6877 35	10
hsa-miR-559	E-MTAB-753,GSE32678,GSE24279,GSE71533,GSE60980,GSE85589,GSE29352,GSE31568,GSE59856	839	0.622 28563	0.66 374 44	1.8	0.579	3.38	0.6877 25	9
hsa-miR-521	GSE29352,GSE71533,GSE31568,GSE85589,GSE24279,E-TABM-664,GSE60980,E-MTAB-753,GSE59856,GSE32678	957	0.508 32825	0.75 773 05	2.1	0.646	3.31	0.6876 35	10
hsa-miR-26a-1-3p	GSE71533,GSE29352,GSE32678,GSE60980,E-MTAB-753,GSE85589,GSE24279,GSE31568,GSE59856,E-TABM-664	957	0.504 61937	0.78 490 94	2.38	0.633	3.8	0.6875 86	10
hsa-miR-197-3p	GSE71533,GSE29352,GSE60980,E-MTAB-753,E-TABM-664,GSE31568,GSE85589,GSE59856,GSE24279	925	0.802 46632	0.50 025 93	1.61	0.408	4.14	0.6875 63	9
hsa-miR-374a-3p	GSE71533,E-MTAB-753,GSE29352,GSE32678,GSE31568,GSE24279,E-TABM-664,GSE60980,GSE85589,GSE59856	957	0.749 72555	0.47 253 21	1.48	0.552	2.77	0.6872 18	10
hsa-miR-601	GSE32678,GSE31568,GSE85589,E-MTAB-753,GSE60980,GSE71533,GSE24279,GSE29352,GSE59856	839	0.551 37181	0.71 254 6	1.93	0.629	3.18	0.6868 19	9
hsa-miR-193a-3p	GSE71533,GSE60980,GSE24279,GSE31568,E-TABM-664,GSE85589,GSE59856,GSE29352,E-MTAB-753	925	0.601 83727	0.69 184 99	2	0.587	3.48	0.6868 11	9
hsa-miR-1293	GSE29352,GSE31568,GSE60980,GSE24279,GSE71533,GSE85589,GSE59856,E-MTAB-753,GSE32678	838	0.575 75608	0.69 989 78	1.92	0.61	3.21	0.6868 09	9

hsa-miR-146b-3p	GSE31568,GSE24279,E-MTAB-753,GSE32678,GSE71533,GSE60980,GSE85589,GSE29352,GSE59856	839	0.640 79162	0.65 683 44	2.03	0.571	3.59	0.6865 46	9
hsa-miR-556-5p	GSE31568,E-MTAB-753,GSE85589,GSE32678,GSE71533,GSE24279,GSE59856,GSE29352,GSE60980	839	0.528 82735	0.74 413 36	2.09	0.63	3.43	0.6863 74	9
hsa-miR-938	GSE60980,GSE24279,GSE85589,GSE71533,GSE29352,GSE59856,GSE31568,E-MTAB-753,GSE32678	839	0.582 64069	0.69 492 71	1.91	0.602	3.31	0.6860 15	9
hsa-miR-497-3p	GSE71533,GSE24279,GSE60980,GSE29352,GSE85589,E-MTAB-753,GSE32678,GSE31568,GSE59856	839	0.588 90936	0.69 851 78	2.01	0.596	3.42	0.6858 23	9
hsa-miR-1296-5p	GSE71533,GSE32678,GSE85589,GSE60980,GSE59856,E-MTAB-753,GSE31568,GSE29352,GSE24279	839	0.645 09917	0.64 327 2	1.83	0.552	3.39	0.6856 97	9
hsa-miR-637	GSE29352,E-MTAB-753,GSE32678,GSE31568,GSE71533,GSE59856,GSE24279,GSE85589,GSE60980	839	0.581 3377	0.71 211 16	2.07	0.594	3.52	0.6856 53	9
hsa-miR-1238-3p	GSE60980,GSE85589,GSE71533,GSE32678,GSE29352,E-MTAB-753,GSE59856,GSE31568,GSE24279	839	0.637 14274	0.65 046 6	1.83	0.563	3.36	0.6856 1	9
hsa-miR-202-3p	GSE32678,E-TABM-664,GSE85589,GSE71533,GSE31568,GSE24279,E-MTAB-753,GSE60980,GSE29352,GSE59856	957	0.691 52358	0.58 006 6	1.71	0.541	3.19	0.6851 44	10
hsa-miR-942-5p	GSE32678,GSE31568,GSE29352,GSE24279,GSE85589,GSE71533,GSE60980,GSE59856,E-MTAB-753	839	0.646 15879	0.64 226 68	1.8	0.556	3.31	0.6850 99	9
hsa-miR-382-5p	GSE71533,GSE60980,E-MTAB-753,GSE24279,GSE85589,GSE31568,E-TABM-664,GSE29352,GSE59856	925	0.607 05177	0.67 241 96	1.84	0.586	3.25	0.6849 57	9
hsa-miR-664a-3p	GSE60980,GSE71533,GSE29352,GSE24279,GSE32678,GSE31568,E-MTAB-753,GSE85589,GSE59856	839	0.610 52533	0.67 345 72	1.89	0.587	3.3	0.6849 03	9
hsa-miR-492	E-TABM-664,GSE71533,GSE85589,GSE32678,GSE31568,GSE29352,E-MTAB-753,GSE24279,GSE60980,GSE59856	957	0.617 83749	0.66 266 59	1.81	0.58	3.26	0.6847 36	10
hsa-miR-608	GSE32678,GSE31568,GSE71533,GSE29352,GSE60980,GSE24279,E-MTAB-753,GSE59856,GSE85589	839	0.654 15354	0.63 142 81	1.8	0.562	3.3	0.6844 15	9
hsa-miR-185-5p	GSE71533,GSE60980,GSE85589,E-TABM-664,GSE29352,GSE24279,E-MTAB-753,GSE59856,GSE31568	925	0.697 26138	0.58 124 1	1.69	0.53	3.32	0.6842 77	9
hsa-miR-605-5p	GSE71533,GSE31568,GSE32678,GSE24279,GSE85589,E-MTAB-753,GSE29352,GSE60980,GSE59856	839	0.632 18578	0.65 754 34	1.96	0.582	3.39	0.6842 34	9

hsa-miR-876-3p	GSE32678,GSE60980,GSE71533,GSE29352,E-MTAB-753,GSE85589,GSE31568,GSE24279,GSE59856	839	0.580 49953	0.70 830 85	2.1	0.603	3.53	0.6841 92	9
hsa-miR-548d-5p	GSE60980,GSE29352,GSE24279,GSE31568,GSE32678,E-MTAB-753,GSE85589,GSE59856,GSE71533	838	0.745 07947	0.49 252 46	1.53	0.536	2.91	0.6840 36	9
hsa-miR-378a-3p	GSE71533,GSE31568,GSE85589,E-TABM-664,GSE24279,GSE60980,GSE59856,GSE29352,E-MTAB-753	925	0.633 368	0.65 145 14	1.82	0.565	3.28	0.6840 32	9
hsa-miR-374a-5p	GSE71533,GSE60980,GSE31568,GSE29352,E-TABM-664,GSE85589,GSE59856,GSE24279,E-MTAB-753,GSE32678	957	0.691 32129	0.57 681 96	1.71	0.563	3.18	0.6839 21	10
hsa-miR-27a-3p	GSE24279,GSE60980,GSE71533,GSE29352,E-MTAB-753,GSE31568,E-TABM-664,GSE59856,GSE85589	925	0.524 20301	0.80 316 04	2.73	0.598	4.67	0.6839 14	9
hsa-miR-488-3p	GSE32678,GSE29352,GSE24279,GSE85589,GSE31568,GSE71533,GSE60980,E-MTAB-753,GSE59856,E-TABM-664	957	0.539 53055	0.74 423 98	2.15	0.628	3.48	0.6838 1	10
hsa-miR-181a-3p	E-MTAB-753,GSE60980,GSE71533,GSE24279,GSE31568,GSE85589,GSE59856,GSE29352,E-TABM-664	925	0.567 59112	0.72 681 88	2.12	0.599	3.56	0.6835 02	9
hsa-miR-302c-3p	GSE31568,E-MTAB-753,GSE71533,GSE24279,GSE29352,GSE60980,E-TABM-664,GSE59856,GSE85589	925	0.751 44655	0.51 798 12	1.61	0.495	3.39	0.6828 89	9
hsa-miR-329-3p	GSE60980,GSE71533,GSE29352,GSE32678,GSE85589,E-MTAB-753,GSE31568,E-TABM-664,GSE59856,GSE24279	957	0.654 63766	0.63 325 08	1.78	0.555	3.33	0.6827 19	10
hsa-miR-130b-5p	GSE71533,GSE24279,GSE31568,GSE60980,E-TABM-664,GSE29352,GSE85589,GSE59856,E-MTAB-753	925	0.760 38881	0.51 897 82	1.6	0.466	3.53	0.6824 38	9
hsa-miR-372-3p	GSE60980,GSE71533,GSE85589,GSE29352,E-MTAB-753,GSE24279,GSE32678,GSE59856,GSE31568	839	0.554 26695	0.73 120 21	2.11	0.616	3.46	0.6821 89	9
hsa-miR-370-3p	GSE71533,GSE59856,E-TABM-664,GSE85589,GSE60980,GSE24279,GSE31568,GSE29352,E-MTAB-753	925	0.557 0383	0.71 660 52	1.98	0.618	3.24	0.6821 71	9
hsa-miR-941	GSE71533,GSE24279,E-MTAB-753,GSE29352,GSE59856,GSE32678,GSE85589,GSE60980,GSE31568	839	0.617 82375	0.66 114 5	1.84	0.584	3.25	0.6821 4	9
hsa-miR-1205	GSE31568,GSE71533,GSE29352,GSE24279,E-MTAB-753,GSE60980,GSE32678,GSE85589,GSE59856	839	0.807 41107	0.46 400 18	1.53	0.428	3.74	0.6820 85	9
hsa-miR-18a-5p	GSE71533,GSE32678,GSE60980,E-TABM-664,E-MTAB-753,GSE29352,GSE85589,GSE31568,GSE24279,GSE59856	957	0.665 55745	0.62 593 45	1.75	0.54	3.47	0.6819 6	10

hsa-miR-96-3p	GSE32678,GSE24279,GSE31568,E-MTAB-753,E-TABM-664,GSE29352,GSE71533,GSE85589,GSE60980,GSE59856	957	0.668 73245	0.60 970 12	1.78	0.557	3.26	0.6816 4	10
hsa-miR-337-3p	GSE71533,E-MTAB-753,GSE31568,GSE60980,E-TABM-664,GSE32678,GSE85589,GSE29352,GSE24279,GSE59856	957	0.578 06609	0.70 115 95	1.98	0.605	3.31	0.6816 15	10
hsa-miR-182-3p	GSE32678,GSE29352,GSE71533,GSE60980,GSE85589,E-MTAB-753,GSE31568,GSE24279,GSE59856,E-TABM-664	957	0.616 0643	0.66 316 55	1.83	0.589	3.19	0.6815 92	10
hsa-miR-545-3p	GSE60980,GSE71533,E-MTAB-753,GSE31568,GSE85589,GSE24279,GSE29352,GSE59856,GSE32678	839	0.644 4683	0.63 837 97	1.77	0.558	3.29	0.6811 1	9
hsa-miR-675-5p	GSE32678,GSE71533,GSE59856,E-MTAB-753,GSE85589,GSE60980,GSE24279,GSE31568,GSE29352	839	0.744 47794	0.51 763 59	1.56	0.497	3.22	0.6810 83	9
hsa-miR-1261	GSE71533,GSE29352,GSE85589,GSE31568,E-MTAB-753,GSE60980,GSE32678,GSE24279,GSE59856	839	0.775 00928	0.45 662 27	1.49	0.51	3.03	0.6810 39	9
hsa-miR-598-3p	GSE71533,E-MTAB-753,GSE85589,GSE32678,GSE24279,GSE29352,GSE59856,GSE31568,GSE60980	839	0.560 79245	0.70 347 38	1.92	0.625	3.16	0.6807 59	9
hsa-miR-548d-3p	GSE60980,GSE31568,GSE71533,GSE24279,GSE32678,GSE85589,E-MTAB-753,GSE29352,GSE59856	839	0.593 1667	0.67 641 47	1.83	0.606	3.12	0.6805 76	9
hsa-miR-1282	GSE60980,GSE32678,GSE29352,GSE31568,GSE85589,E-MTAB-753,GSE24279,GSE59856,GSE71533	839	0.439 52546	0.80 842 61	2.36	0.695	3.46	0.6805 53	9
hsa-miR-548p	GSE32678,GSE85589,E-MTAB-753,GSE60980,GSE31568,GSE71533,GSE24279,GSE29352,GSE59856	839	0.686 51227	0.60 348 28	1.73	0.538	3.4	0.6802 22	9
hsa-miR-188-5p	GSE71533,GSE60980,GSE85589,GSE24279,E-TABM-664,GSE29352,GSE31568,E-MTAB-753,GSE59856	925	0.651 38469	0.62 929 06	1.81	0.568	3.33	0.6802 21	9
hsa-miR-142-5p	GSE60980,GSE29352,GSE71533,E-TABM-664,E-MTAB-753,GSE85589,GSE24279,GSE31568,GSE59856	925	0.510 04049	0.73 360 41	1.9	0.668	2.91	0.6801 12	9
hsa-miR-205-5p	GSE71533,GSE24279,E-TABM-664,GSE59856,GSE85589,GSE31568,GSE29352,GSE60980,E-MTAB-753	925	0.574 01246	0.73 821 01	2.25	0.583	3.88	0.6799 36	9
hsa-miR-384	GSE60980,E-MTAB-753,GSE71533,GSE31568,E-TABM-664,GSE32678,GSE85589,GSE29352,GSE24279,GSE59856	957	0.500 47325	0.76 149 14	2.13	0.659	3.27	0.6797 35	10

hsa-miR-378a-5p	GSE71533,E-MTAB-753,GSE60980,GSE29352,GSE31568,GSE59856,GSE85589,GSE24279,E-TABM-664	925	0.619 51726	0.66 251 49	1.87	0.579	3.26	0.6796 22	9
hsa-miR-181a-2-3p	GSE71533,GSE32678,GSE31568,E-TABM-664,GSE60980,GSE85589,GSE24279,GSE29352,GSE59856,E-MTAB-753	957	0.637 48869	0.64 252 75	1.76	0.565	3.26	0.6793 82	10
hsa-miR-563	GSE24279,GSE29352,GSE31568,GSE60980,GSE32678,GSE59856,E-MTAB-753,GSE85589,GSE71533	839	0.535 24529	0.73 856 39	2.1	0.636	3.34	0.6793 55	9
hsa-miR-545-5p	GSE32678,GSE29352,GSE60980,GSE31568,GSE24279,E-MTAB-753,GSE59856,GSE71533,GSE85589	839	0.329 83252	0.89 153 39	3.09	0.749	4.14	0.6793 39	9
hsa-miR-510-5p	GSE32678,GSE24279,GSE31568,E-MTAB-753,GSE29352,GSE71533,GSE60980,GSE85589,E-TABM-664,GSE59856	957	0.763 32473	0.50 669 29	1.59	0.486	3.4	0.6791 46	10
hsa-miR-194-3p	GSE32678,GSE31568,GSE71533,GSE60980,GSE29352,GSE85589,E-TABM-664,GSE59856,E-MTAB-753,GSE24279	957	0.509 87159	0.72 961 53	1.89	0.672	2.86	0.6785 76	10
hsa-miR-10b-3p	GSE71533,E-TABM-664,GSE24279,GSE29352,GSE60980,E-MTAB-753,GSE31568,GSE59856,GSE85589	925	0.568 45379	0.69 792 83	1.9	0.614	3.16	0.6782 67	9
hsa-miR-33a-5p	GSE71533,GSE60980,GSE31568,E-MTAB-753,E-TABM-664,GSE24279,GSE29352,GSE32678,GSE85589,GSE59856	956	0.677 05964	0.60 982 41	1.73	0.536	3.37	0.6782 03	10
hsa-miR-556-3p	GSE29352,GSE71533,GSE32678,GSE60980,GSE24279,E-MTAB-753,GSE31568,GSE85589,GSE59856	839	0.500 17626	0.73 778 4	1.9	0.678	2.88	0.6778 09	9
hsa-miR-140-5p	GSE60980,E-MTAB-753,GSE71533,GSE29352,GSE24279,E-TABM-664,GSE31568,GSE59856,GSE85589	925	0.601 25028	0.66 790 54	1.81	0.599	3.1	0.6776 6	9
hsa-miR-29b-1-5p	E-MTAB-753,GSE60980,GSE29352,E-TABM-664,GSE31568,GSE24279,GSE85589,GSE59856,GSE71533	925	0.520 81163	0.74 959 05	2.13	0.645	3.32	0.6774 78	9
hsa-miR-328-3p	GSE71533,GSE60980,GSE24279,E-MTAB-753,GSE29352,E-TABM-664,GSE59856,GSE31568,GSE85589	925	0.569 00888	0.69 722 65	1.89	0.618	3.09	0.6774 18	9
hsa-miR-518f-3p	GSE71533,GSE32678,GSE85589,GSE29352,GSE31568,GSE24279,E-TABM-664,GSE59856,GSE60980,E-MTAB-753	957	0.532 39469	0.72 105 24	1.92	0.649	3.01	0.6772	10
hsa-miR-583	GSE32678,GSE85589,GSE59856,GSE24279,GSE60980,GSE31568,GSE29352,GSE71533,E-MTAB-753	839	0.546 58934	0.71 341 84	1.91	0.636	3.05	0.6771 17	9
hsa-miR-296-3p	GSE71533,GSE60980,GSE31568,E-MTAB-753,E-TABM-664,GSE85589,GSE29352,GSE24279,GSE59856	925	0.666 44867	0.61 641 85	1.74	0.544	3.27	0.6769 21	9

hsa-miR-34c-3p	GSE29352,GSE71533,GSE24279,GSE59856,E-TABM-664,GSE85589,GSE32678,E-MTAB-753,GSE60980,GSE31568	957	0.624 01055	0.64 936 89	1.79	0.583	3.14	0.6768 62	10
hsa-miR-383-5p	GSE85589,GSE71533,E-TABM-664,GSE32678,GSE60980,GSE29352,GSE31568,GSE24279,GSE59856,E-MTAB-753	957	0.397 71969	0.83 805 76	2.5	0.718	3.51	0.6766 01	10
hsa-miR-18b-5p	GSE71533,GSE60980,E-MTAB-753,GSE85589,GSE32678,GSE31568,E-TABM-664,GSE24279,GSE29352,GSE59856	957	0.600 2772	0.66 710 65	1.82	0.601	3.07	0.6765 17	10
hsa-miR-1283	E-MTAB-753,GSE31568,GSE71533,GSE29352,GSE60980,GSE24279,GSE59856,GSE32678,GSE85589	839	0.415 98757	0.78 787 97	1.94	0.743	2.67	0.6762 6	9
hsa-miR-19a-3p	GSE71533,GSE60980,GSE29352,E-TABM-664,E-MTAB-753,GSE24279,GSE31568,GSE85589,GSE59856	925	0.586 17912	0.67 137 79	1.74	0.623	2.96	0.6759 84	9
hsa-miR-1202	GSE59856,GSE32678,E-MTAB-753,GSE29352,GSE31568,GSE24279,GSE71533,GSE85589,GSE60980	839	0.580 76087	0.67 083 37	1.77	0.63	2.91	0.6758 13	9
hsa-miR-767-5p	GSE32678,GSE71533,GSE24279,GSE60980,GSE31568,E-MTAB-753,GSE59856,GSE85589,GSE29352	839	0.620 24189	0.65 441 75	1.82	0.587	3.15	0.6756 49	9
hsa-miR-520a-3p	GSE32678,E-MTAB-753,GSE31568,GSE71533,E-TABM-664,GSE24279,GSE60980,GSE59856,GSE85589,GSE29352	957	0.519 50491	0.74 187 13	2.03	0.646	3.21	0.6754 81	10
hsa-miR-644a	GSE29352,GSE32678,GSE60980,E-MTAB-753,GSE85589,GSE31568,GSE71533,GSE59856,GSE24279	839	0.632 4919	0.63 996 34	1.75	0.588	3.1	0.6754 72	9
hsa-miR-99b-3p	GSE60980,E-MTAB-753,GSE24279,E-TABM-664,GSE59856,GSE31568,GSE71533,GSE85589,GSE29352	925	0.654 96623	0.61 858 26	1.72	0.564	3.13	0.6754 11	9
hsa-miR-623	GSE60980,GSE32678,GSE24279,GSE29352,GSE85589,GSE31568,GSE59856,E-MTAB-753,GSE71533	839	0.493 79936	0.78 640 58	2.39	0.649	3.7	0.6753 48	9
hsa-miR-208a-3p	GSE29352,GSE71533,GSE24279,GSE31568,E-MTAB-753,GSE85589,E-TABM-664,GSE59856,GSE60980	925	0.673 93346	0.59 894 12	1.71	0.558	3.18	0.6752 59	9
hsa-miR-340-3p	GSE71533,GSE24279,GSE29352,E-MTAB-753,GSE60980,GSE85589,GSE31568,E-TABM-664,GSE59856	925	0.497 17115	0.75 007 72	2.02	0.67	3.06	0.6748 1	9
hsa-miR-302d-5p	GSE24279,GSE31568,GSE29352,E-TABM-664,GSE60980,GSE32678,E-MTAB-753,GSE71533,GSE59856,GSE85589	957	0.363 28912	0.85 270 58	2.5	0.744	3.39	0.6745 37	10
hsa-miR-519b-3p	GSE32678,GSE29352,GSE71533,GSE85589,GSE24279,GSE31568,E-TABM-664,E-MTAB-753,GSE59856,GSE60980	957	0.485 16674	0.76 187 68	2.04	0.678	3.05	0.6745 02	10

hsa-miR-7-2-3p	GSE71533,GSE32678,GSE60980,GSE29352,GSE85589,GSE24279,E-MTAB-753,E-TABM-664,GSE31568,GSE59856	957	0.754 38475	0.52 284 66	1.58	0.476	3.47	0.6744 97	10
hsa-miR-506-3p	GSE71533,GSE60980,GSE31568,GSE24279,GSE59856,E-MTAB-753,GSE85589,E-TABM-664,GSE29352,GSE32678	957	0.428 25858	0.84 459 32	2.91	0.68	4.36	0.6743 77	10
hsa-miR-554	GSE71533,GSE29352,E-MTAB-753,GSE60980,GSE32678,GSE85589,GSE24279,GSE31568,GSE59856	839	0.549 42969	0.71 228 81	1.92	0.638	3.07	0.6743 51	9
hsa-miR-502-5p	GSE32678,E-MTAB-753,GSE60980,GSE71533,GSE29352,GSE31568,GSE85589,E-TABM-664,GSE59856,GSE24279	957	0.506 1828	0.74 301 35	1.98	0.67	3	0.6743 17	10
hsa-miR-339-5p	GSE71533,E-MTAB-753,GSE31568,GSE60980,GSE24279,GSE29352,GSE85589,GSE59856,E-TABM-664	925	0.701 46821	0.57 799 71	1.67	0.522	3.27	0.6740 23	9
hsa-miR-1269a	GSE29352,GSE60980,GSE85589,GSE24279,GSE59856,E-MTAB-753,GSE32678,GSE71533,GSE31568	839	0.535 34939	0.74 579 61	2.16	0.632	3.47	0.6739 91	9
hsa-miR-600	GSE71533,GSE32678,E-MTAB-753,GSE60980,GSE29352,GSE59856,GSE31568,GSE24279,GSE85589	839	0.555 14126	0.71 527 07	2.01	0.636	3.22	0.6736 93	9
hsa-miR-1206	GSE29352,GSE31568,GSE85589,GSE60980,GSE71533,E-MTAB-753,GSE24279,GSE32678,GSE59856	839	0.612 06886	0.66 072 34	1.89	0.604	3.16	0.6736 67	9
hsa-miR-650	GSE71533,GSE59856,GSE85589,GSE60980,E-MTAB-753,GSE24279,GSE29352,GSE31568,GSE32678	839	0.603 82619	0.65 876 89	1.77	0.614	3.04	0.6735 02	9
hsa-miR-586	GSE85589,GSE60980,GSE29352,GSE32678,GSE71533,E-MTAB-753,GSE31568,GSE59856,GSE24279	839	0.472 5715	0.75 830 01	1.96	0.694	2.88	0.6732 09	9
hsa-miR-504-5p	GSE71533,GSE24279,GSE29352,E-TABM-664,GSE31568,GSE85589,E-MTAB-753,GSE59856,GSE60980	924	0.510 33613	0.75 427 25	2.11	0.662	3.22	0.6731 34	9
hsa-miR-183-5p	GSE71533,GSE24279,GSE29352,E-TABM-664,E-MTAB-753,GSE31568,GSE60980,GSE59856,GSE85589	925	0.485 78219	0.74 119 61	1.88	0.691	2.83	0.6730 7	9
hsa-miR-629-5p	GSE32678,GSE60980,E-MTAB-753,GSE31568,GSE24279,GSE29352,GSE71533,GSE59856,GSE85589	839	0.733 2047	0.55 314	1.64	0.493	3.55	0.6729 36	9
hsa-miR-19b-2-5p	GSE71533,E-MTAB-753,GSE29352,GSE32678,GSE24279,GSE85589,GSE31568,GSE60980,GSE59856	839	0.846 58878	0.43 138 3	1.51	0.382	4.3	0.6729 26	9
hsa-miR-1271-5p	GSE60980,GSE71533,GSE24279,GSE31568,E-MTAB-753,GSE29352,GSE85589,GSE32678,GSE59856	839	0.589 69883	0.67 697 64	1.87	0.612	3.11	0.6728 83	9

hsa-miR-587	GSE71533,GSE60980,GSE31568,E-MTAB-753,GSE29352,GSE24279,GSE32678,GSE85589,GSE59856	839	0.653 25627	0.61 370 09	1.73	0.574	3.08	0.6726 15	9
hsa-miR-493-3p	GSE71533,GSE60980,GSE85589,E-MTAB-753,E-TABM-664,GSE29352,GSE24279,GSE31568,GSE59856	925	0.647 49315	0.62 418 21	1.7	0.571	3.18	0.6725 35	9
hsa-miR-134-5p	GSE71533,GSE31568,GSE60980,E-TABM-664,GSE24279,GSE29352,GSE59856,GSE85589,E-MTAB-753	925	0.676 06537	0.60 320 29	1.69	0.545	3.21	0.6724 19	9
hsa-miR-585-3p	GSE71533,GSE32678,GSE31568,GSE29352,GSE24279,GSE85589,GSE59856,E-MTAB-753,GSE60980	838	0.541 54958	0.70 471 59	1.86	0.654	2.92	0.6718 14	9
hsa-miR-337-5p	GSE60980,GSE71533,GSE24279,GSE31568,GSE29352,E-MTAB-753,GSE85589,GSE59856,GSE32678	839	0.590 99701	0.66 809 51	1.78	0.613	2.99	0.6715 4	9
hsa-miR-582-5p	GSE60980,GSE71533,GSE31568,GSE29352,GSE24279,GSE85589,E-MTAB-753,GSE32678,GSE59856	839	0.429 08773	0.82 287 7	2.49	0.697	3.6	0.6713 61	9
hsa-miR-507	GSE29352,GSE60980,GSE32678,E-MTAB-753,GSE71533,GSE31568,GSE24279,E-TABM-664,GSE59856,GSE85589	957	0.512 83851	0.74 758 97	2.09	0.656	3.24	0.6710 14	10
hsa-miR-582-3p	GSE32678,GSE24279,GSE71533,GSE29352,E-MTAB-753,GSE85589,GSE60980,GSE59856,GSE31568	839	0.703 88191	0.55 244 52	1.62	0.553	3.02	0.6709 73	9
hsa-miR-298	GSE32678,GSE85589,GSE29352,GSE59856,GSE31568,E-MTAB-753,GSE60980,GSE24279,GSE71533	839	0.602 25737	0.65 716 11	1.77	0.605	2.97	0.6706 07	9
hsa-miR-1287-5p	GSE32678,GSE60980,E-MTAB-753,GSE29352,GSE24279,GSE85589,GSE31568,GSE59856,GSE71533	839	0.390 47311	0.81 369 91	2.12	0.748	2.88	0.6698 81	9
hsa-miR-548e-3p	GSE71533,GSE32678,GSE29352,GSE85589,GSE59856,GSE24279,GSE31568,E-MTAB-753,GSE60980	839	0.505 9073	0.76 341 94	2.19	0.655	3.37	0.6696 37	9
hsa-miR-371a-3p	GSE71533,GSE60980,E-MTAB-753,GSE29352,GSE31568,GSE32678,GSE85589,GSE24279,GSE59856	839	0.493 71312	0.77 078 67	2.23	0.659	3.44	0.6696 16	9
hsa-miR-593-5p	GSE29352,GSE71533,GSE24279,GSE60980,GSE32678,GSE85589,GSE59856,GSE31568,E-MTAB-753	839	0.564 18383	0.69 178 26	1.85	0.632	2.96	0.6689 9	9
hsa-miR-636	GSE29352,GSE31568,GSE60980,GSE32678,GSE71533,E-MTAB-753,GSE59856,GSE24279,GSE85589	839	0.671 28083	0.61 248 21	1.74	0.542	3.3	0.6687 81	9
hsa-miR-581	GSE29352,GSE32678,GSE31568,E-MTAB-753,GSE24279,GSE85589,GSE59856,GSE71533,GSE60980	839	0.443 00172	0.78 878 85	2.13	0.7	3.09	0.6686 93	9

hsa-miR-599	GSE60980,E-MTAB-753,GSE31568,GSE24279,GSE29352,GSE32678,GSE59856,GSE85589,GSE71533	839	0.480 01264	0.77 288 79	2.15	0.675	3.21	0.6686 51	9
hsa-miR-1253	GSE71533,GSE32678,E-MTAB-753,GSE31568,GSE85589,GSE59856,GSE60980,GSE29352,GSE24279	839	0.461 38297	0.75 756 46	1.91	0.714	2.75	0.6685 85	9
hsa-miR-519d-3p	GSE32678,GSE29352,GSE24279,GSE60980,GSE31568,GSE85589,E-TABM-664,E-MTAB-753,GSE59856	905	0.604 48667	0.65 361 17	1.75	0.601	2.96	0.6683 57	9
hsa-miR-519c-3p	GSE32678,GSE24279,GSE60980,GSE31568,GSE71533,GSE29352,GSE85589,E-TABM-664,GSE59856,E-MTAB-753	957	0.478 57815	0.74 397 49	1.87	0.706	2.71	0.6682 75	10
hsa-miR-520c-3p	GSE29352,E-MTAB-753,GSE24279,GSE71533,GSE32678,GSE31568,GSE85589,GSE59856,GSE60980	839	0.565 12529	0.70 229 75	1.93	0.628	3.1	0.6682 59	9
hsa-miR-1236-3p	GSE71533,GSE32678,E-MTAB-753,GSE59856,GSE31568,GSE85589,GSE24279,GSE29352,GSE60980	839	0.644 55723	0.61 965 85	1.71	0.579	3.08	0.6682 44	9
hsa-miR-571	GSE29352,GSE71533,GSE32678,GSE31568,GSE60980,E-MTAB-753,GSE85589,GSE59856,GSE24279	839	0.759 05564	0.47 045 41	1.47	0.531	2.84	0.6681 54	9
hsa-miR-512-5p	GSE71533,GSE29352,GSE60980,GSE85589,GSE24279,E-TABM-664,GSE59856,GSE31568,E-MTAB-753,GSE32678	957	0.551 63659	0.68 962 98	1.77	0.647	2.81	0.6681 33	10
hsa-miR-566	GSE60980,GSE32678,GSE31568,E-MTAB-753,GSE71533,GSE29352,GSE59856,GSE85589,GSE24279	839	0.644 84492	0.61 507 13	1.69	0.591	2.93	0.6678 22	9
hsa-miR-612	GSE32678,GSE71533,GSE59856,E-MTAB-753,GSE60980,GSE24279,GSE85589,GSE29352,GSE31568	839	0.612 55147	0.64 787 09	1.76	0.605	2.99	0.6675 93	9
hsa-miR-199b-5p	GSE60980,GSE32678,E-MTAB-753,E-TABM-664,GSE71533,GSE85589,GSE31568,GSE59856,GSE29352,GSE24279	957	0.506 6802	0.74 961 94	2.07	0.662	3.15	0.6675 85	10
hsa-miR-138-5p	GSE71533,E-TABM-664,GSE31568,E-MTAB-753,GSE24279,GSE85589,GSE59856,GSE29352,GSE60980	925	0.515 9154	0.74 309 51	2.03	0.655	3.12	0.6674 62	9
hsa-miR-431-5p	GSE71533,E-MTAB-753,GSE60980,GSE29352,GSE31568,GSE85589,GSE59856,E-TABM-664,GSE24279	925	0.595 06674	0.66 143 25	1.78	0.62	2.95	0.6674 3	9
hsa-miR-92a-1-5p	GSE32678,GSE60980,GSE29352,GSE24279,E-TABM-664,GSE71533,GSE85589,GSE31568,E-MTAB-753,GSE59856	957	0.629 9721	0.63 692 99	1.91	0.607	3.18	0.6669 29	10
hsa-miR-144-5p	GSE85589,GSE60980,GSE24279,GSE32678,GSE31568,E-MTAB-753,GSE29352,GSE59856,E-TABM-664,GSE71533	957	0.520 54649	0.71 688 67	1.83	0.671	2.78	0.6666 5	10

hsa-miR-520h	GSE32678,GSE31568,GSE60980,GSE85589,E-TABM-664,E-MTAB-753,GSE29352,GSE59856,GSE24279	905	0.621 03985	0.63 630 79	1.7	0.602	2.95	0.6663 73	9
hsa-miR-651-5p	GSE29352,GSE85589,GSE71533,GSE24279,GSE31568,GSE60980,GSE32678,E-MTAB-753,GSE59856	839	0.429 14644	0.86 384 17	3.39	0.669	5.04	0.6662 06	9
hsa-miR-618	GSE29352,GSE60980,E-MTAB-753,GSE32678,GSE85589,GSE31568,GSE24279,GSE71533,GSE59856	839	0.592 00407	0.66 004 83	1.75	0.619	2.9	0.6660 45	9
hsa-miR-888-3p	GSE60980,GSE32678,E-MTAB-753,GSE71533,GSE24279,GSE85589,GSE29352,GSE31568,GSE59856	839	0.496 89036	0.75 042 72	2.02	0.674	3.04	0.6658 27	9
hsa-miR-299-3p	GSE71533,GSE32678,GSE60980,GSE24279,GSE29352,GSE85589,E-TABM-664,GSE31568,E-MTAB-753,GSE59856	957	0.589 65461	0.66 683 71	1.78	0.621	2.92	0.6658 15	10
hsa-miR-452-3p	GSE60980,GSE71533,GSE29352,E-MTAB-753,GSE32678,GSE24279,GSE59856,GSE31568,GSE85589	839	0.453 5874	0.76 557 13	1.94	0.714	2.77	0.6657 95	9
hsa-miR-1185-5p	GSE60980,GSE24279,GSE29352,GSE85589,E-MTAB-753,GSE31568,GSE32678,GSE59856,GSE71533	839	0.407 39923	0.82 597 74	2.4	0.719	3.37	0.6653 83	9
hsa-miR-409-5p	GSE71533,GSE60980,E-MTAB-753,E-TABM-664,GSE85589,GSE59856,GSE24279,GSE31568,GSE29352	925	0.539 90103	0.70 373 48	1.83	0.658	2.83	0.6652 78	9
hsa-miR-153-3p	GSE71533,GSE60980,GSE29352,E-MTAB-753,E-TABM-664,GSE85589,GSE59856,GSE31568,GSE24279	925	0.485 08473	0.74 338 12	1.92	0.696	2.82	0.6652 71	9
hsa-miR-129-5p	E-TABM-664,GSE29352,GSE60980,GSE24279,E-MTAB-753,GSE31568,GSE59856,GSE85589,GSE71533	925	0.499 52633	0.75 451 95	2.07	0.67	3.11	0.6651 89	9
hsa-miR-1254	E-MTAB-753,GSE29352,GSE60980,GSE59856,GSE32678,GSE71533,GSE24279,GSE85589,GSE31568	839	0.579 7488	0.67 246 51	1.78	0.627	2.88	0.6651 69	9
hsa-miR-924	E-MTAB-753,GSE71533,GSE29352,GSE85589,GSE32678,GSE24279,GSE60980,GSE59856,GSE31568	839	0.758 355	0.52 895 61	1.61	0.473	3.7	0.6649 27	9
hsa-miR-485-3p	GSE71533,GSE60980,GSE29352,GSE85589,GSE24279,E-TABM-664,GSE31568,E-MTAB-753,GSE59856	925	0.508 9434	0.71 568 56	1.79	0.691	2.64	0.6649 23	9
hsa-miR-638	GSE71533,GSE60980,GSE59856,GSE32678,GSE31568,GSE85589,GSE24279,E-MTAB-753,GSE29352	839	0.683 30842	0.56 512 84	1.59	0.564	2.86	0.6648 97	9
hsa-miR-1247-5p	GSE24279,GSE32678,GSE71533,GSE85589,GSE60980,GSE59856,E-MTAB-753,GSE31568,GSE29352	839	0.538 26091	0.72 419 5	1.98	0.644	3.11	0.6647 57	9

hsa-miR-1322	GSE60980,E-MTAB-753,GSE71533,GSE32678,GSE31568,GSE24279,GSE59856,GSE85589,GSE29352	839	0.401 38557	0.79 649 79	1.99	0.748	2.71	0.6642 78	9
hsa-miR-646	E-MTAB-753,GSE32678,GSE29352,GSE60980,GSE59856,GSE31568,GSE71533,GSE24279,GSE85589	839	0.672 26744	0.59 022 93	1.65	0.558	3.04	0.6639 1	9
hsa-miR-526b-3p	GSE32678,E-MTAB-753,GSE85589,GSE60980,GSE31568,GSE29352,GSE71533,GSE24279,GSE59856	839	0.535 82851	0.69 606 07	1.77	0.666	2.72	0.6638 86	9
hsa-miR-154-5p	GSE24279,GSE29352,GSE31568,GSE60980,E-MTAB-753,E-TABM-664,GSE85589,GSE59856,GSE71533	925	0.349 05126	0.86 226 93	2.59	0.755	3.45	0.6636 02	9
hsa-miR-1178-3p	GSE60980,GSE31568,E-MTAB-753,GSE85589,GSE71533,GSE59856,GSE24279,GSE29352,GSE32678	839	0.533 83417	0.73 665 14	2.1	0.638	3.32	0.6633 35	9
hsa-miR-769-5p	GSE71533,GSE60980,GSE24279,GSE31568,GSE29352,E-MTAB-753,GSE85589,GSE59856,GSE32678	839	0.562 57834	0.67 738 09	1.75	0.65	2.78	0.6630 76	9
hsa-miR-524-3p	GSE71533,GSE32678,GSE60980,E-TABM-664,GSE29352,GSE31568,GSE85589,GSE24279,E-MTAB-753,GSE59856	957	0.666 55562	0.59 418 08	1.64	0.565	3.02	0.6630 15	10
hsa-miR-219a-5p	GSE71533,GSE29352,GSE60980,E-TABM-664,GSE85589,GSE59856,GSE24279,E-MTAB-753,GSE31568	925	0.484 51597	0.75 579 49	2.02	0.684	2.98	0.6628 95	9
hsa-miR-935	GSE32678,GSE71533,GSE31568,GSE60980,GSE59856,GSE85589,GSE24279,E-MTAB-753,GSE29352	839	0.572 46896	0.67 521 63	1.78	0.635	2.83	0.6625 68	9
hsa-miR-34a-3p	GSE32678,GSE60980,E-TABM-664,GSE71533,GSE85589,E-MTAB-753,GSE31568,GSE29352,GSE59856,GSE24279	957	0.696 00136	0.55 247 22	1.57	0.561	2.86	0.6625 58	10
hsa-miR-410-3p	GSE24279,GSE71533,GSE60980,E-MTAB-753,GSE29352,E-TABM-664,GSE31568,GSE59856,GSE85589	925	0.652 00776	0.60 008 44	1.65	0.585	2.87	0.6618 65	9
hsa-miR-517-5p	GSE71533,GSE32678,GSE29352,E-MTAB-753,GSE60980,E-TABM-664,GSE85589,GSE31568,GSE24279,GSE59856	957	0.578 24408	0.65 890 83	1.68	0.641	2.75	0.6617 31	10
hsa-miR-323a-3p	GSE71533,E-MTAB-753,GSE60980,GSE24279,GSE31568,GSE85589,E-TABM-664,GSE29352,GSE59856	925	0.526 87826	0.69 639 73	1.73	0.677	2.64	0.6616 49	9
hsa-miR-516a-5p	GSE32678,GSE71533,GSE31568,E-MTAB-753,GSE60980,GSE29352,GSE85589,E-TABM-664,GSE59856,GSE24279	957	0.714 59839	0.54 620 39	1.57	0.53	3.05	0.6616 06	10
hsa-miR-485-5p	GSE29352,GSE60980,GSE31568,GSE24279,GSE71533,E-TABM-664,GSE85589,E-MTAB-753,GSE59856	925	0.547 51685	0.70 143 85	1.87	0.655	2.88	0.6615 06	9

hsa-miR-596	GSE32678,GSE29352,GSE71533,GSE60980,GSE31568,GSE24279,E-MTAB-753,GSE59856,GSE85589	839	0.598 15813	0.64 390 53	1.67	0.626	2.77	0.6605 47	9
hsa-miR-124-3p	GSE59856,GSE60980,E-TABM-664,GSE71533,GSE85589,E-MTAB-753,GSE31568,GSE24279,GSE29352	925	0.640 77192	0.60 470 54	1.67	0.606	2.84	0.6605 32	9
hsa-miR-662	GSE60980,GSE29352,GSE71533,GSE31568,GSE24279,E-MTAB-753,GSE85589,GSE59856,GSE32678	839	0.517 97087	0.71 142 49	1.79	0.682	2.68	0.6600 04	9
hsa-miR-593-3p	GSE29352,GSE60980,GSE71533,GSE85589,GSE59856,GSE24279,E-MTAB-753,GSE31568,GSE32678	839	0.668 32586	0.57 428 68	1.6	0.585	2.79	0.6599 35	9
hsa-miR-922	GSE29352,GSE71533,GSE60980,GSE32678,GSE31568,E-MTAB-753,GSE85589,GSE24279,GSE59856	839	0.787 25658	0.46 730 21	1.49	0.464	3.39	0.6598 02	9
hsa-miR-657	GSE71533,GSE32678,GSE60980,GSE29352,GSE85589,GSE31568,E-MTAB-753,GSE24279,GSE59856	839	0.474 21542	0.77 124 58	2.16	0.687	3.17	0.6593 01	9
hsa-miR-380-3p	GSE29352,GSE32678,GSE60980,E-MTAB-753,GSE85589,GSE31568,GSE59856,GSE71533,E-TABM-664,GSE24279	957	0.568 62505	0.69 052 58	1.89	0.628	3.04	0.6587 59	10
hsa-miR-323b-5p	GSE71533,GSE32678,E-TABM-664,E-MTAB-753,GSE60980,GSE31568,GSE85589,GSE24279,GSE59856,GSE29352	957	0.633 3466	0.61 395 79	1.64	0.604	2.79	0.6587 43	10
hsa-miR-490-5p	GSE60980,GSE71533,GSE32678,GSE31568,GSE24279,GSE29352,GSE59856,E-TABM-664,GSE85589,E-MTAB-753	957	0.508 1057	0.74 060 64	2	0.673	3.01	0.6584 42	10
hsa-miR-1323	GSE32678,GSE71533,GSE85589,GSE60980,GSE59856,GSE24279,GSE31568,GSE29352,E-MTAB-753	839	0.640 59478	0.60 404 93	1.64	0.601	2.77	0.6584 27	9
hsa-miR-495-3p	GSE71533,GSE60980,GSE29352,GSE24279,E-TABM-664,GSE31568,E-MTAB-753,GSE85589,GSE59856	925	0.481 4224	0.74 944 28	1.93	0.692	2.83	0.6580 67	9
hsa-miR-139-5p	E-MTAB-753,GSE60980,GSE29352,GSE71533,GSE31568,E-TABM-664,GSE85589,GSE59856,GSE24279	924	0.449 68011	0.76 500 04	1.93	0.719	2.72	0.6579 77	9
hsa-miR-193b-5p	GSE32678,GSE59856,GSE60980,E-MTAB-753,GSE31568,GSE24279,GSE29352,E-TABM-664,GSE85589,GSE71533	957	0.606 1282	0.67 563 53	1.92	0.588	3.32	0.6575 59	10
hsa-miR-491-5p	GSE71533,GSE31568,GSE29352,GSE59856,GSE24279,GSE60980,E-TABM-664,E-MTAB-753,GSE85589	925	0.627 00052	0.61 762 72	1.65	0.61	2.76	0.6575 38	9
hsa-miR-363-3p	GSE60980,GSE71533,E-MTAB-753,GSE29352,GSE31568,GSE85589,GSE24279,E-TABM-664,GSE59856	925	0.549 3157	0.68 302 61	1.73	0.658	2.68	0.6572 61	9

hsa-miR-9-3p	GSE71533,GSE60980,GSE32678,E-TABM-664,GSE29352,GSE31568,GSE85589,GSE24279,GSE59856,E-MTAB-753	957	0.604 93044	0.63 648 01	1.66	0.621	2.74	0.6570 83	10
hsa-miR-196b-5p	GSE29352,GSE31568,GSE71533,E-MTAB-753,GSE85589,E-TABM-664,GSE59856,GSE24279,GSE60980	925	0.470 11052	0.75 809 8	1.97	0.7	2.84	0.6566 56	9
hsa-miR-302b-3p	E-MTAB-753,GSE24279,GSE60980,GSE31568,GSE85589,E-TABM-664,GSE59856,GSE71533,GSE29352	925	0.440 97901	0.78 503 7	2.09	0.714	2.96	0.6565 19	9
hsa-miR-1251-5p	GSE24279,GSE32678,GSE71533,GSE31568,GSE85589,GSE59856,GSE60980,E-MTAB-753,GSE29352	839	0.526 51284	0.70 743 54	1.81	0.675	2.73	0.6565 14	9
hsa-miR-502-3p	E-MTAB-753,GSE24279,GSE60980,GSE71533,GSE29352,E-TABM-664,GSE85589,GSE32678,GSE31568,GSE59856	957	0.514 21089	0.72 444 03	1.88	0.678	2.81	0.6564 51	10
hsa-miR-219a-1-3p	GSE71533,GSE24279,E-MTAB-753,GSE32678,GSE85589,GSE60980,GSE31568,GSE59856,GSE29352	839	0.678 42287	0.54 190 53	1.58	0.62	2.61	0.6561	9
hsa-miR-26a-2-3p	GSE31568,GSE29352,E-TABM-664,GSE85589,GSE60980,GSE24279,GSE32678,E-MTAB-753,GSE59856,GSE71533	957	0.552 8793	0.68 654 17	1.8	0.659	2.76	0.6560 71	10
hsa-miR-614	GSE32678,GSE60980,GSE71533,E-MTAB-753,GSE85589,GSE29352,GSE24279,GSE59856,GSE31568	839	0.588 76169	0.64 971 06	1.69	0.641	2.73	0.6559 22	9
hsa-miR-509-3p	GSE60980,GSE24279,GSE31568,GSE32678,E-TABM-664,GSE71533,E-MTAB-753,GSE29352,GSE85589,GSE59856	957	0.708 19756	0.53 519 63	1.54	0.561	2.83	0.6557 89	10
hsa-miR-432-5p	GSE60980,GSE71533,GSE31568,GSE24279,E-MTAB-753,E-TABM-664,GSE59856,GSE29352,GSE32678,GSE85589	957	0.543 75772	0.69 053 11	1.8	0.662	2.76	0.6556 8	10
hsa-miR-95-3p	E-TABM-664,GSE32678,GSE85589,GSE29352,GSE24279,GSE59856,GSE31568,E-MTAB-753,GSE60980,GSE71533	957	0.600 31232	0.64 077 98	1.68	0.625	2.73	0.6556 52	10
hsa-miR-34b-3p	GSE60980,E-MTAB-753,GSE31568,GSE29352,GSE24279,GSE85589,GSE71533,E-TABM-664,GSE59856	925	0.454 2345	0.75 317 66	1.86	0.724	2.61	0.6553 99	9
hsa-miR-519a-3p	GSE31568,GSE85589,GSE29352,GSE32678,E-MTAB-753,E-TABM-664,GSE71533,GSE24279,GSE59856	900	0.394 73624	0.81 226 99	2.12	0.748	2.86	0.6543 72	9
hsa-miR-33b-3p	GSE32678,GSE60980,GSE24279,GSE85589,E-MTAB-753,GSE71533,GSE29352,GSE59856,GSE31568	839	0.514 47729	0.72 866 28	1.92	0.669	2.91	0.6543 58	9
hsa-miR-1302	GSE32678,GSE71533,GSE85589,E-MTAB-753,GSE29352,GSE60980,GSE59856,GSE31568,GSE24279	839	0.536 15787	0.69 439 44	1.77	0.674	2.69	0.6542 41	9

hsa-miR-1270	GSE60980,GSE29352,GSE32678,GSE85589,E-MTAB-753,GSE71533,GSE31568,GSE59856,GSE24279	839	0.497 22383	0.71 760 45	1.77	0.702	2.56	0.6540 6	9
hsa-miR-381-3p	GSE71533,GSE24279,GSE85589,GSE29352,GSE60980,E-TABM-664,GSE59856,E-MTAB-753,GSE31568	925	0.675 54112	0.53 523 18	1.52	0.634	2.49	0.6539 81	9
hsa-miR-1226-3p	GSE71533,GSE29352,GSE31568,GSE60980,E-MTAB-753,GSE85589,GSE59856,GSE24279,GSE32678	839	0.633 5752	0.61 484 28	1.65	0.607	2.84	0.6538 22	9
hsa-miR-486-3p	GSE32678,GSE60980,GSE59856,E-MTAB-753,GSE29352,GSE85589,GSE31568,GSE24279,GSE71533	839	0.645 50671	0.58 886 97	1.58	0.609	2.62	0.6534 44	9
hsa-miR-548b-3p	GSE24279,GSE32678,GSE60980,E-MTAB-753,GSE31568,GSE29352,GSE85589,GSE71533,GSE59856	839	0.649 4543	0.58 458 97	1.63	0.616	2.69	0.6533 88	9
hsa-miR-105-5p	GSE32678,GSE60980,GSE29352,E-TABM-664,GSE24279,GSE31568,GSE85589,GSE71533,E-MTAB-753,GSE59856	957	0.640 59909	0.59 395 62	1.62	0.617	2.66	0.6533 78	10
hsa-miR-576-3p	GSE60980,GSE71533,GSE31568,GSE32678,E-MTAB-753,GSE85589,GSE59856,GSE24279,GSE29352	839	0.508 83206	0.69 942 27	1.68	0.703	2.48	0.6533 77	9
hsa-miR-548j-5p	GSE60980,GSE71533,E-MTAB-753,GSE31568,GSE29352,GSE59856,GSE32678,GSE24279,GSE85589	839	0.480 79905	0.75 617 67	2.02	0.686	2.97	0.6533 39	9
hsa-miR-642a-5p	GSE32678,GSE60980,GSE85589,GSE24279,GSE29352,GSE31568,GSE71533,GSE59856,E-MTAB-753	839	0.592 36406	0.68 711 28	1.93	0.599	3.27	0.6533 31	9
hsa-miR-518e-3p	GSE71533,GSE32678,GSE85589,GSE60980,GSE29352,GSE24279,GSE31568,E-TABM-664,E-MTAB-753,GSE59856	957	0.548 11672	0.67 764 95	1.7	0.666	2.6	0.6530 18	10
hsa-miR-525-3p	GSE29352,GSE60980,GSE71533,GSE85589,E-MTAB-753,E-TABM-664,GSE32678,GSE31568,GSE59856,GSE24279	957	0.515 95477	0.71 083 61	1.8	0.683	2.67	0.6526 88	10
hsa-miR-572	GSE32678,GSE85589,GSE31568,GSE29352,GSE24279,GSE59856,GSE60980,E-MTAB-753,GSE71533	839	0.530 6845	0.71 342 31	1.88	0.663	2.87	0.6521 16	9
hsa-miR-885-3p	GSE59856,GSE32678,GSE60980,GSE29352,GSE31568,GSE71533,GSE85589,E-MTAB-753,GSE24279	839	0.539 46818	0.67 469 41	1.65	0.684	2.47	0.6519 89	9
hsa-miR-424-3p	GSE24279,GSE60980,GSE32678,E-MTAB-753,GSE31568,GSE29352,E-TABM-664,GSE71533,GSE59856,GSE85589	957	0.541 69333	0.68 152 88	1.7	0.675	2.59	0.6518 42	10
hsa-miR-449a	GSE29352,E-MTAB-753,GSE71533,GSE85589,E-TABM-664,GSE24279,GSE60980,GSE31568,GSE59856	925	0.827 59988	0.40 813 79	1.4	0.437	3.4	0.6518 28	9

hsa-miR-137	GSE60980,GSE85589,GSE31568,GSE24279,GSE29352,GSE71533,GS E59856,E-TABM-664,E-MTAB-753	925	0.484 00875	0.72 439 65	1.77	0.714	2.52	0.6518 14	9
hsa-miR-361- 5p	GSE60980,GSE29352,E-MTAB-753,E-TABM- 664,GSE71533,GSE24279,GSE85589,GSE31568,GSE59856	925	0.624 80627	0.61 055 22	1.72	0.635	2.74	0.6517 26	9
hsa-miR-193b- 3p	GSE32678,GSE71533,GSE24279,GSE31568,E-TABM- 664,GSE59856,GSE29352,GSE85589,GSE60980,E-MTAB-753	957	0.540 31791	0.70 662 54	1.87	0.654	2.87	0.6513 39	10
hsa-miR-323a- 5p	E-TABM-664,GSE71533,GSE59856,E-MTAB- 753,GSE29352,GSE60980,GSE24279,GSE31568,GSE85589	925	0.608 58049	0.62 538 8	1.63	0.633	2.62	0.6512 13	9
hsa-miR-520f- 3p	GSE32678,GSE24279,E-MTAB- 753,GSE60980,GSE85589,GSE29352,GSE31568,E-TABM- 664,GSE59856,GSE71533	957	0.604 82624	0.62 926 28	1.64	0.633	2.65	0.6511 92	10
hsa-miR-661	GSE60980,GSE85589,GSE32678,GSE71533,GSE29352,GSE59856,E- MTAB-753,GSE31568,GSE24279	839	0.608 19056	0.62 528 25	1.63	0.632	2.68	0.6510 91	9
hsa-miR-367- 5p	GSE71533,GSE32678,GSE85589,GSE31568,GSE60980,E-TABM- 664,GSE24279,GSE29352,GSE59856,E-MTAB-753	957	0.507 64268	0.69 891 43	1.68	0.701	2.48	0.6510 28	10
hsa-miR-335- 3p	GSE71533,GSE31568,GSE60980,GSE29352,E-TABM- 664,GSE85589,E-MTAB-753,GSE59856,GSE24279	925	0.499 89477	0.71 732 68	1.78	0.699	2.59	0.6507 81	9
hsa-miR-450b- 3p	GSE60980,GSE31568,GSE29352,GSE32678,GSE24279,GSE85589,E- MTAB-753,GSE71533,GSE59856	839	0.444 75111	0.78 674 32	2.12	0.707	3.02	0.6502 5	9
hsa-miR-921	GSE32678,GSE71533,GSE29352,GSE60980,E-MTAB- 753,GSE31568,GSE85589,GSE24279,GSE59856	839	0.491 31727	0.72 397 3	1.79	0.707	2.57	0.6501 34	9
hsa-miR-412- 3p	GSE71533,GSE31568,E-MTAB-753,GSE24279,E-TABM- 664,GSE60980,GSE85589,GSE59856,GSE29352	925	0.551 77334	0.67 041 61	1.7	0.668	2.6	0.6496 39	9
hsa-miR-579- 3p	GSE29352,GSE32678,GSE31568,GSE60980,E-MTAB- 753,GSE24279,GSE71533,GSE59856,GSE85589	839	0.708 64362	0.51 659 98	1.5	0.582	2.65	0.6487 99	9
hsa-miR-330- 5p	GSE60980,GSE31568,E-TABM- 664,GSE24279,GSE85589,GSE59856,GSE71533,GSE29352,E-MTAB- 753	925	0.479 8677	0.75 083 68	1.98	0.698	2.86	0.6487 4	9
hsa-miR-511- 5p	GSE29352,GSE60980,GSE31568,GSE59856,GSE24279,GSE71533,E- TABM-664,GSE85589,GSE32678,E-MTAB-753	957	0.527 49186	0.72 363 85	1.95	0.661	2.97	0.6483 61	10
hsa-miR-432- 3p	GSE71533,GSE32678,E-MTAB- 753,GSE60980,GSE85589,GSE29352,E-TABM- 664,GSE24279,GSE31568,GSE59856	957	0.487 3123	0.71 782 71	1.73	0.72	2.45	0.6482 12	10

hsa-miR-15a-3p	GSE32678,GSE71533,GSE85589,GSE24279,GSE60980,GSE59856,E-MTAB-753,E-TABM-664,GSE31568,GSE29352	957	0.567 91972	0.66 922 7	1.76	0.655	2.72	0.6479 7	10
hsa-miR-24-1-5p	GSE60980,GSE71533,E-TABM-664,E-MTAB-753,GSE85589,GSE29352,GSE24279,GSE31568,GSE59856	925	0.730 4607	0.49 318 42	1.46	0.558	2.69	0.6477 34	9
hsa-miR-1299	GSE71533,GSE29352,GSE60980,GSE24279,E-MTAB-753,GSE31568,GSE59856,GSE85589,GSE32678	839	0.513 9909	0.70 335 72	1.75	0.694	2.56	0.6463 84	9
hsa-miR-296-5p	GSE71533,GSE60980,E-MTAB-753,GSE24279,E-TABM-664,GSE29352,GSE59856,GSE31568,GSE85589	925	0.646 00412	0.59 321 2	1.58	0.6	2.7	0.6462 5	9
hsa-miR-876-5p	GSE29352,GSE60980,GSE71533,GSE31568,GSE85589,E-MTAB-753,GSE32678,GSE59856,GSE24279	839	0.497 98573	0.70 979 07	1.72	0.706	2.48	0.6458	9
hsa-miR-17-3p	GSE60980,GSE24279,GSE29352,GSE71533,E-TABM-664,GSE85589,GSE59856,GSE31568,E-MTAB-753	925	0.655 90486	0.56 835 06	1.54	0.614	2.56	0.6457 9	9
hsa-miR-302c-5p	GSE32678,GSE60980,GSE71533,GSE85589,GSE29352,E-MTAB-753,GSE59856,E-TABM-664,GSE31568,GSE24279	957	0.540 28047	0.68 763 43	1.75	0.67	2.63	0.6457 29	10
hsa-miR-558	GSE32678,E-MTAB-753,GSE24279,GSE60980,GSE31568,GSE85589,GSE29352,GSE59856	839	0.734 84493	0.45 387 53	1.4	0.6	2.39	0.6456 51	9
hsa-miR-187-5p	GSE59856,GSE60980,GSE29352,GSE71533,GSE31568,E-MTAB-753,GSE32678,GSE85589,GSE24279	839	0.811 7692	0.53 592 36	1.74	0.36	5.36	0.6452 69	9
hsa-miR-626	GSE71533,GSE60980,GSE29352,GSE85589,GSE32678,E-MTAB-753,GSE24279,GSE31568,GSE59856	839	0.422 4637	0.82 210 04	2.46	0.71	3.48	0.6452 43	9
hsa-miR-875-3p	GSE71533,GSE24279,GSE31568,GSE32678,GSE60980,GSE85589,E-MTAB-753,GSE59856,GSE29352	839	0.566 26743	0.64 973 45	1.61	0.673	2.45	0.6452 38	9
hsa-miR-1226-5p	GSE31568,GSE85589,GSE29352,GSE71533,GSE32678,GSE59856,GSE24279,E-MTAB-753,GSE60980	839	0.663 38173	0.55 568 75	1.5	0.61	2.51	0.6450 85	9
hsa-miR-562	E-MTAB-753,GSE29352,GSE32678,GSE24279,GSE31568,GSE59856,GSE85589,GSE60980,GSE71533	839	0.692 9604	0.51 571 47	1.46	0.608	2.46	0.6447 8	9
hsa-miR-625-3p	GSE71533,E-MTAB-753,GSE32678,GSE59856,GSE24279,GSE60980,GSE31568,GSE29352,GSE85589	839	0.668 12426	0.55 877 72	1.52	0.595	2.6	0.6445 87	9
hsa-miR-621	GSE32678,GSE60980,GSE71533,GSE85589,GSE31568,GSE24279,GSE59856,E-MTAB-753,GSE29352	839	0.439 07556	0.77 591 59	2.02	0.726	2.81	0.6445 07	9

hsa-miR-616-3p	GSE32678,GSE71533,E-MTAB-753,GSE24279,GSE31568,GSE85589,GSE59856,GSE29352,GSE60980	839	0.467 80007	0.75 058 27	1.88	0.714	2.67	0.6438 61	9
hsa-miR-455-5p	GSE29352,E-MTAB-753,GSE85589,GSE31568,E-TABM-664,GSE71533,GSE60980,GSE59856,GSE24279	925	0.741 88988	0.49 105 64	1.46	0.538	2.82	0.6438 6	9
hsa-miR-526b-5p	GSE32678,GSE71533,GSE29352,GSE60980,E-MTAB-753,GSE31568,GSE85589,E-TABM-664,GSE24279,GSE59856	957	0.674 56584	0.55 405 76	1.51	0.587	2.65	0.6432 53	10
hsa-miR-1207-5p	E-MTAB-753,GSE32678,GSE71533,GSE31568,GSE60980,GSE29352,GSE59856,GSE24279,GSE85589	839	0.712 13141	0.50 566 61	1.46	0.578	2.57	0.6429 58	9
hsa-miR-198	GSE32678,GSE31568,GSE60980,GSE71533,E-TABM-664,E-MTAB-753,GSE29352,GSE59856,GSE24279,GSE85589	957	0.602 07161	0.62 191 17	1.61	0.644	2.53	0.6427 9	10
hsa-miR-129-2-3p	GSE29352,GSE71533,E-TABM-664,GSE24279,GSE32678,E-MTAB-753,GSE59856,GSE85589,GSE60980,GSE31568	957	0.399 45066	0.80 290 77	2.08	0.75	2.83	0.6425 84	10
hsa-miR-25-5p	GSE71533,GSE24279,GSE29352,GSE85589,GSE59856,GSE32678,GSE60980,GSE31568,E-MTAB-753	839	0.569 99333	0.65 663 07	1.67	0.662	2.55	0.6419 86	9
hsa-miR-186-3p	GSE71533,GSE60980,GSE32678,GSE85589,GSE24279,GSE31568,GSE29352,GSE59856,E-MTAB-753	839	0.566 44461	0.64 551 34	1.6	0.675	2.45	0.6416 23	9
hsa-miR-664a-5p	GSE71533,GSE24279,GSE85589,GSE32678,GSE31568,E-MTAB-753,GSE60980,GSE59856,GSE29352	839	0.624 61064	0.59 770 28	1.54	0.634	2.53	0.6414 92	9
hsa-miR-544a	GSE71533,GSE24279,E-MTAB-753,GSE29352,GSE60980,GSE59856,GSE31568,GSE32678,GSE85589	839	0.525 03766	0.68 946 88	1.73	0.695	2.53	0.6404 97	9
hsa-miR-152-3p	GSE60980,GSE71533,GSE29352,E-MTAB-753,GSE31568,E-TABM-664,GSE85589,GSE24279,GSE59856	924	0.492 13596	0.70 478 5	1.66	0.722	2.34	0.6403 83	9
hsa-miR-549a	GSE32678,GSE71533,GSE24279,GSE60980,GSE85589,GSE31568,GSE59856,GSE29352,E-MTAB-753	839	0.423 58319	0.77 940 49	1.94	0.744	2.64	0.6402 98	9
hsa-miR-588	GSE60980,GSE32678,E-MTAB-753,GSE31568,GSE24279,GSE29352,GSE71533,GSE59856,GSE85589	839	0.519 48463	0.67 751 95	1.6	0.712	2.3	0.6401 82	9
hsa-miR-325	GSE32678,E-MTAB-753,GSE24279,E-TABM-664,GSE85589,GSE31568,GSE71533,GSE59856,GSE29352,GSE60980	957	0.479 33737	0.73 125 18	1.8	0.716	2.54	0.6393 6	10
hsa-miR-1179	E-MTAB-753,GSE29352,GSE71533,GSE85589,GSE32678,GSE24279,GSE59856,GSE31568,GSE60980	839	0.380 92202	0.84 247 07	2.5	0.74	3.41	0.6392 82	9

hsa-miR-30d-3p	E-MTAB-753,GSE60980,GSE71533,GSE85589,GSE31568,E-TABM-664,GSE24279,GSE59856,GSE29352	925	0.766 45545	0.42 342 09	1.36	0.57	2.44	0.6390 16	9
hsa-miR-499a-3p	GSE71533,GSE31568,GSE60980,GSE29352,GSE32678,GSE85589,E-MTAB-753,GSE59856,E-TABM-664,GSE24279	957	0.507 27001	0.65 987 49	1.47	0.745	2.08	0.6384 37	10
hsa-miR-99a-3p	GSE71533,GSE24279,GSE60980,GSE29352,E-MTAB-753,GSE85589,GSE59856,GSE31568,GSE32678	839	0.509 70467	0.70 047 93	1.7	0.706	2.45	0.6382 99	9
hsa-miR-376b-3p	GSE60980,GSE29352,GSE32678,GSE31568,GSE85589,E-TABM-664,GSE24279,GSE59856,GSE71533,E-MTAB-753	957	0.762 63157	0.42 438 79	1.36	0.571	2.44	0.6380 88	10
hsa-miR-548k	GSE71533,GSE32678,GSE31568,GSE24279,GSE29352,E-MTAB-753,GSE59856,GSE85589,GSE60980	839	0.495 93376	0.70 461 22	1.7	0.715	2.43	0.6380 64	9
hsa-miR-622	GSE32678,E-MTAB-753,GSE29352,GSE60980,GSE31568,GSE85589,GSE59856,GSE71533,GSE24279	839	0.468 22266	0.74 137 78	1.86	0.723	2.61	0.6363 16	9
hsa-miR-483-5p	GSE60980,GSE29352,GSE71533,E-TABM-664,GSE24279,GSE32678,GSE59856,GSE85589,GSE31568,E-MTAB-753	957	0.556 44348	0.65 483	1.62	0.682	2.39	0.6360 94	10
hsa-miR-433-3p	GSE60980,GSE24279,GSE71533,E-TABM-664,GSE85589,GSE31568,GSE59856,GSE29352,E-MTAB-753	925	0.575 78476	0.63 172 74	1.58	0.672	2.38	0.6359 53	9
hsa-miR-508-5p	GSE32678,GSE60980,GSE24279,E-MTAB-753,GSE71533,GSE31568,GSE29352,E-TABM-664,GSE85589,GSE59856	957	0.531 54556	0.68 386 14	1.7	0.69	2.49	0.6357 98	10
hsa-miR-302b-5p	GSE29352,GSE85589,GSE32678,E-TABM-664,GSE24279,E-MTAB-753,GSE31568,GSE59856,GSE71533,GSE60980	957	0.449 60972	0.76 962 09	1.99	0.716	2.8	0.6355 14	10
hsa-miR-499a-5p	GSE60980,GSE71533,E-MTAB-753,GSE29352,GSE31568,GSE85589,GSE59856,E-TABM-664,GSE24279	925	0.443 82498	0.75 860 82	1.88	0.741	2.56	0.6344 43	9
hsa-miR-514a-3p	GSE71533,GSE60980,GSE32678,GSE85589,E-TABM-664,E-MTAB-753,GSE29352,GSE24279,GSE31568,GSE59856	957	0.425 78873	0.80 621 94	2.28	0.718	3.19	0.6341 85	10
hsa-miR-609	GSE32678,GSE29352,GSE24279,E-MTAB-753,GSE31568,GSE59856,GSE71533,GSE85589,GSE60980	839	0.371 03241	0.83 134 12	2.31	0.758	3.07	0.6331 74	9
hsa-miR-369-5p	GSE71533,GSE60980,E-TABM-664,GSE85589,GSE31568,GSE59856,GSE24279,GSE29352,E-MTAB-753	925	0.757 67268	0.41 598 16	1.34	0.602	2.32	0.6331 46	9
hsa-miR-655-3p	GSE71533,GSE60980,GSE31568,GSE59856,GSE32678,GSE85589,E-MTAB-753,GSE24279,GSE29352	839	0.442 13762	0.81 004 75	2.39	0.688	3.51	0.6325 64	9

hsa-miR-518b	GSE71533,GSE60980,E-TABM-664,GSE59856,E-MTAB-753,GSE85589,GSE31568,GSE24279,GSE29352,GSE32678	957	0.534 24586	0.67 461 26	1.66	0.693	2.42	0.6325 64	10
hsa-miR-606	GSE71533,GSE85589,E-MTAB-753,GSE32678,GSE60980,GSE29352,GSE24279,GSE59856,GSE31568	839	0.778 56956	0.48 870 72	1.51	0.47	3.49	0.6319 52	9
hsa-miR-105-3p	GSE60980,GSE32678,GSE24279,GSE71533,GSE31568,E-TABM-664,GSE29352,GSE85589,GSE59856,E-MTAB-753	957	0.620 76026	0.58 036 39	1.55	0.674	2.34	0.6318 98	10
hsa-miR-933	GSE32678,GSE24279,GSE71533,GSE29352,GSE31568,GSE60980,GSE59856,E-MTAB-753,GSE85589	839	0.593 47815	0.61 064 23	1.53	0.674	2.3	0.6318 16	9
hsa-miR-448	GSE71533,GSE85589,GSE31568,E-MTAB-753,GSE29352,E-TABM-664,GSE60980,GSE24279,GSE59856	925	0.573 45494	0.62 562 63	1.53	0.676	2.33	0.6317 04	9
hsa-miR-219a-2-3p	GSE24279,E-MTAB-753,GSE31568,GSE60980,GSE85589,GSE71533,GSE59856,E-TABM-664,GSE29352	925	0.455 30787	0.75 863 64	1.92	0.721	2.7	0.6310 52	9
hsa-miR-422a	E-MTAB-753,GSE71533,GSE24279,GSE32678,GSE31568,E-TABM-664,GSE29352,GSE59856,GSE85589,GSE60980	957	0.648 07746	0.56 770 67	1.5	0.626	2.44	0.6306 89	10
hsa-miR-649	GSE29352,GSE60980,GSE85589,GSE31568,GSE24279,E-MTAB-753,GSE71533,GSE59856,GSE32678	839	0.817 45665	0.36 599 03	1.31	0.518	2.67	0.6304 54	9
hsa-miR-629-3p	GSE71533,GSE32678,E-MTAB-753,GSE85589,GSE59856,GSE29352,GSE24279,GSE31568,GSE60980	839	0.536 90457	0.68 290 75	1.7	0.684	2.53	0.6301 21	9
hsa-miR-548o-3p	GSE32678,GSE85589,GSE31568,GSE60980,GSE24279,GSE59856,E-MTAB-753,GSE71533,GSE29352	839	0.533 84334	0.67 060 15	1.66	0.701	2.4	0.6299 18	9
hsa-miR-29a-5p	GSE71533,GSE29352,E-MTAB-753,E-TABM-664,GSE31568,GSE60980,GSE59856,GSE24279,GSE85589	925	0.557 06843	0.63 432 27	1.52	0.697	2.23	0.6285 7	9
hsa-miR-15b-3p	E-MTAB-753,GSE85589,GSE60980,GSE71533,E-TABM-664,GSE31568,GSE24279,GSE59856,GSE29352	925	0.497 54198	0.67 686 69	1.53	0.743	2.11	0.6275 09	9
hsa-miR-1229-3p	GSE29352,GSE60980,GSE32678,E-MTAB-753,GSE71533,GSE59856,GSE85589,GSE31568,GSE24279	839	0.662 99539	0.54 807 2	1.46	0.622	2.43	0.6265 57	9
hsa-miR-149-5p	E-MTAB-753,GSE71533,GSE29352,GSE85589,GSE59856,GSE60980,E-TABM-664,GSE24279,GSE31568	925	0.546 15318	0.64 799 23	1.56	0.702	2.25	0.6264 35	9

hsa-miR-548m	GSE29352,E-MTAB-753,GSE60980,GSE32678,GSE24279,GSE85589,GSE71533,GSE59856,GSE31568	839	0.693 16139	0.48 218 52	1.39	0.648	2.19	0.6259 72	9
hsa-miR-1292-5p	GSE29352,GSE71533,GSE85589,GSE31568,GSE59856,GSE60980,E-MTAB-753,GSE24279,GSE32678	839	0.548 4081	0.64 683 15	1.57	0.701	2.26	0.6245 2	9
hsa-miR-130a-5p	E-MTAB-753,GSE29352,GSE31568,GSE32678,GSE24279,GSE85589,GSE59856,GSE71533,E-TABM-664,GSE60980	957	0.439 78405	0.78 653 51	2.1	0.714	2.96	0.6240 7	10
hsa-miR-18b-3p	GSE60980,GSE24279,GSE71533,E-MTAB-753,GSE85589,GSE59856,E-TABM-664,GSE29352,GSE31568,GSE32678	957	0.460 184	0.74 484 4	1.83	0.727	2.55	0.6237 65	10
hsa-miR-887-3p	GSE60980,GSE71533,E-MTAB-753,GSE32678,GSE24279,GSE31568,GSE29352,GSE59856,GSE85589	839	0.648 63355	0.56 445 49	1.49	0.627	2.43	0.6230 4	9
hsa-miR-617	GSE32678,GSE29352,E-MTAB-753,GSE24279,GSE60980,GSE71533,GSE85589,GSE31568,GSE59856	839	0.514 1172	0.68 084 27	1.64	0.715	2.32	0.6221 22	9
hsa-miR-373-3p	E-MTAB-753,E-TABM-664,GSE31568,GSE60980,GSE29352,GSE32678,GSE71533,GSE59856,GSE24279,GSE85589	957	0.489 02864	0.70 362 94	1.67	0.733	2.3	0.6219 89	10
hsa-miR-373-5p	GSE71533,GSE32678,GSE60980,GSE85589,GSE31568,E-TABM-664,GSE59856,GSE29352,E-MTAB-753,GSE24279	957	0.670 44587	0.56 176 4	1.52	0.587	2.68	0.6206 43	10
hsa-miR-520d-3p	E-MTAB-753,GSE32678,GSE31568,GSE29352,E-TABM-664,GSE85589,GSE59856,GSE24279,GSE71533,GSE60980	957	0.406 25236	0.77 310 74	1.8	0.772	2.36	0.6204 83	10
hsa-miR-185-3p	GSE71533,GSE29352,GSE85589,GSE60980,E-TABM-664,GSE31568,GSE32678,GSE59856,GSE24279,E-MTAB-753	957	0.558 59265	0.62 344 64	1.49	0.71	2.13	0.6194 59	10
hsa-miR-496	GSE71533,GSE29352,GSE60980,E-MTAB-753,GSE24279,E-TABM-664,GSE85589,GSE59856,GSE31568	921	0.509 72871	0.67 735 64	1.61	0.734	2.21	0.6170 72	9
hsa-miR-573	GSE71533,E-MTAB-753,GSE31568,GSE32678,GSE60980,GSE85589,GSE29352,GSE59856,GSE24279	839	0.715 82805	0.44 759 09	1.32	0.647	2.1	0.6169 48	9
hsa-miR-518c-3p	GSE29352,GSE31568,E-MTAB-753,GSE24279,GSE60980,E-TABM-664,GSE85589,GSE71533,GSE59856,GSE32678	957	0.694 68364	0.47 475 09	1.34	0.653	2.08	0.6156 61	10
hsa-miR-1289	GSE32678,GSE29352,GSE24279,GSE60980,E-MTAB-753,GSE85589,GSE31568,GSE59856,GSE71533	839	0.369 98164	0.81 602 52	2.08	0.779	2.68	0.6120 51	9

hsa-miR-106a-3p	GSE60980,GSE32678,E-MTAB-753,GSE71533,GSE85589,GSE31568,GSE24279,E-TABM-664,GSE59856,GSE29352	957	0.539 16665	0.64 041 48	1.53	0.728	2.12	0.6098 96	10
hsa-miR-1182	GSE32678,GSE60980,GSE71533,GSE59856,GSE85589,GSE29352,GSE31568,GSE24279,E-MTAB-753	839	0.424 05032	0.76 007 57	1.82	0.764	2.41	0.6086 14	9
hsa-miR-1203	GSE29352,GSE85589,GSE59856,GSE31568,GSE32678,GSE71533,E-MTAB-753,GSE60980,GSE24279	839	0.463 3307	0.77 811 45	2.14	0.69	3.12	0.6078 74	9
hsa-miR-483-3p	GSE32678,GSE29352,GSE60980,GSE24279,GSE31568,E-TABM-664,GSE85589,GSE59856,GSE71533,E-MTAB-753	957	0.634 14246	0.55 695 39	1.43	0.662	2.2	0.6010 48	10
hsa-miR-567	GSE29352,GSE71533,E-MTAB-753,GSE60980,GSE31568,GSE59856,GSE85589,GSE24279,GSE32678	839	0.342 68128	0.85 505 87	2.46	0.773	3.2	0.5965 89	9
hsa-miR-133b	GSE71533,GSE60980,GSE24279,GSE85589,GSE29352,GSE31568,GSE59856,E-MTAB-753,GSE32678	839	0.563 29488	0.67 872 81	1.78	0.645	2.81	0.5878 99	9
hsa-miR-1227-3p	GSE32678,E-MTAB-753,GSE60980,GSE31568,GSE29352,GSE85589,GSE71533,GSE59856,GSE24279	839	0.464 4041	0.75 603 99	1.95	0.715	2.77	0.5870 35	9
hsa-miR-936	GSE32678,GSE24279,GSE85589,GSE29352,GSE59856,GSE31568,E-MTAB-753,GSE71533,GSE60980	839	0.492 6211	0.68 797 95	1.6	0.741	2.18	0.5867 83	9

Supplemental table 2. Univariate and multivariate survival analysis of the clinical/pathological and miR-30a of PDAC.

Factor		Univariate analysis		Univariate analysis	
		HR(95% CI)	<i>p</i> -value	HR(95% CI)	<i>p</i> -value
Age		1 (1.01-1.09)	0.018	1.04(1.00-1.09)	0.06860549
Sex	Male/Female	1.2 (0.59-2.63)	0.56		
Tumor dimension		1 (0.85-1.180)	0.97		
Histological grade	G1/2 VS. G3/4	2.7 (1.32-5.50)	0.0065	1.51(0.65-3.57)	0.33636418
Positive lymph node number		1.1 (0.98-1.15)	0.12		
Radiation	Yes/No	0.45 (0.18-1.09)	0.075		
Resection status	Rx/0 VS. R1/2	2.6 (1.26-5.32)	0.0095	1.47(0.62-3.48)	0.38500155
Pathological stage	I VS. II	5.1 (1.20-21.85)	0.027	0.88(0.085-9.12)	0.91670943
Metastasis	M0 VS. M1	0.84 (0.41-1.76)	0.65		
Regional lymph node	N0 VS. N1	2.5 (1.07-5.84)	0.034	1.77(0.54-5.79)	0.34572052
Tumor	T1/2 VS. T3/4	3.4 (1.18-9.91)	0.023	2.07(0.44-9.72)	0.35544273
Molecular target therapy	Yes VS. No	0.83 (0.41-1.68)	0.6		
miR-30a	High VS. low	2.3 (1.15-4.80)	0.019	2.43(1.05-5.59)	0.03722112

Supplemental table 3. Univariate and multivariate survival analysis of the clinical/pathological and miR-203 of PDAC.

Factor		Univariate analysis		Univariate analysis	
		HR(95% CI)	<i>p</i> -value	HR(95% CI)	<i>p</i> -value
Age		1 (1.00779-1.08537)	0.018	1.04(1.00-1.09)	0.060844
Sex	Male/Female	1.2 (0.590345-2.63435)	0.56		
Tumor dimension		1 (0.853533-1.17979)	0.97		
Histological grade	G1/2 VS. G3/4	2.7 (1.31917-5.49521)	0.0065	1.75(0.77-3.96)	0.179074
Positive lymph node number		1.1 (0.983305-1.15448)	0.12		
Radiation	Yes/No	0.45 (0.182766-1.08511)	0.075		
Resection status	Rx/0 VS. R1/2	2.6 (1.26232-5.31859)	0.0095	1.77(0.79-4.00)	0.16626
Pathological stage	I VS. II	5.1 (1.20031-21.8451)	0.027	1.31(0.13-12.75)	0.818848
Metastasis	M0 VS. M1	0.84 (0.4054-1.75739)	0.65		
Regional lymph node	N0 VS. N1	2.5 (1.07385-5.83838)	0.034	1.93(0.59-6.24)	0.270545
Tumor	T1/2 VS. T3/4	3.4 (1.18481-9.91132)	0.023	1.32(0.29-6.11)	0.721071
Molecular target therapy	Yes VS. No	0.83 (0.408158-1.67961)	0.6		
miR-203	High VS. low	2.3 (1.12404-4.71535)	0.023	3.14(1.28-7.71)	0.012479