

Differential microRNA profile underlies the divergent healing responses in skin and oral mucosal wounds

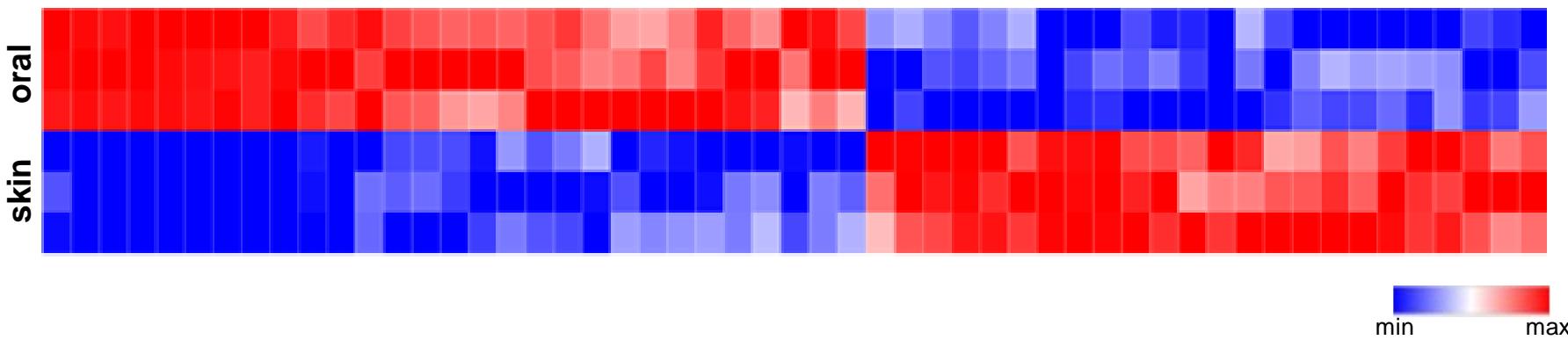
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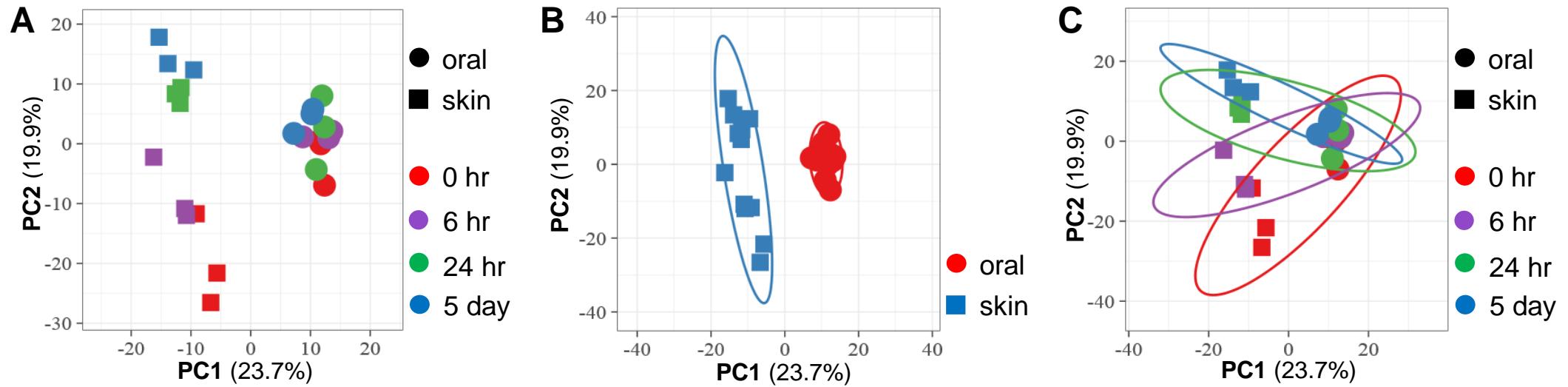
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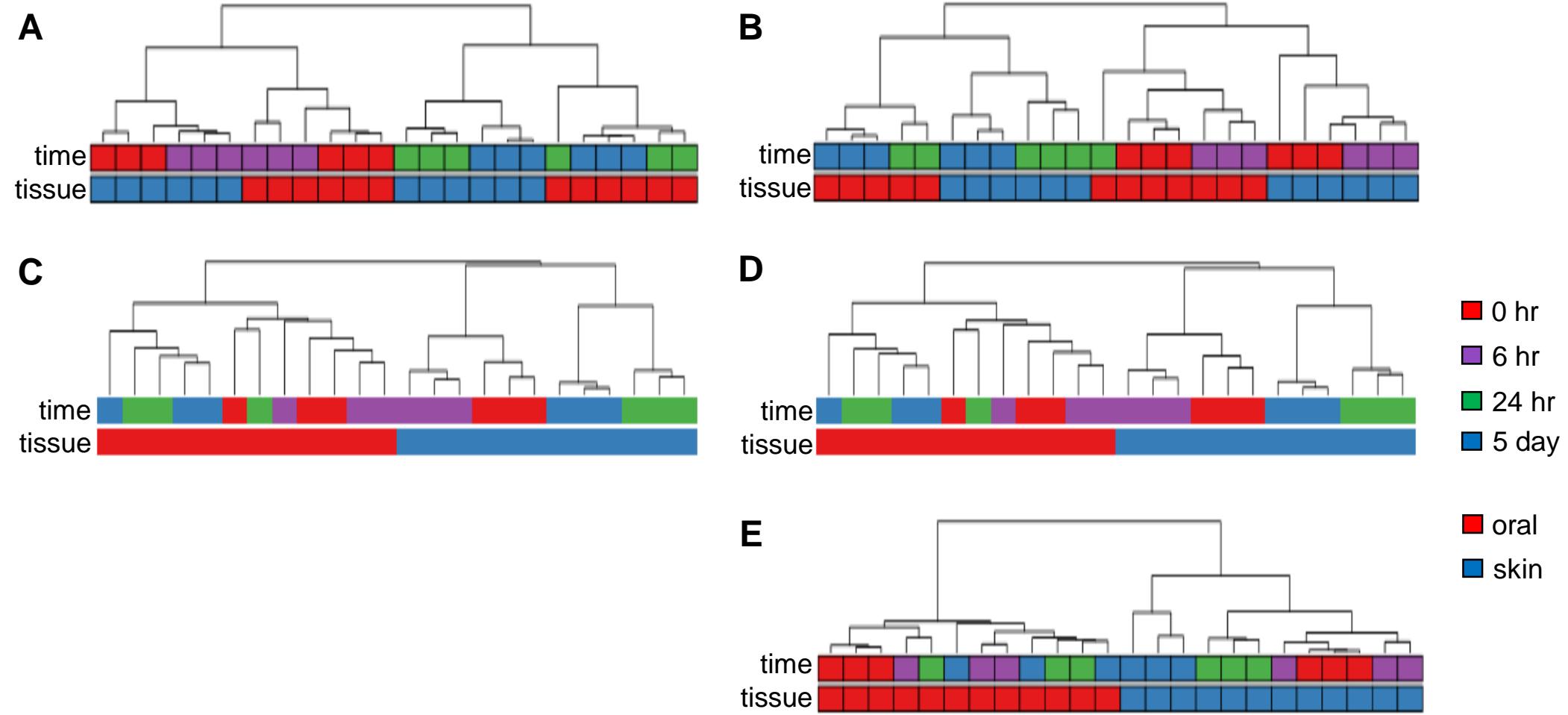
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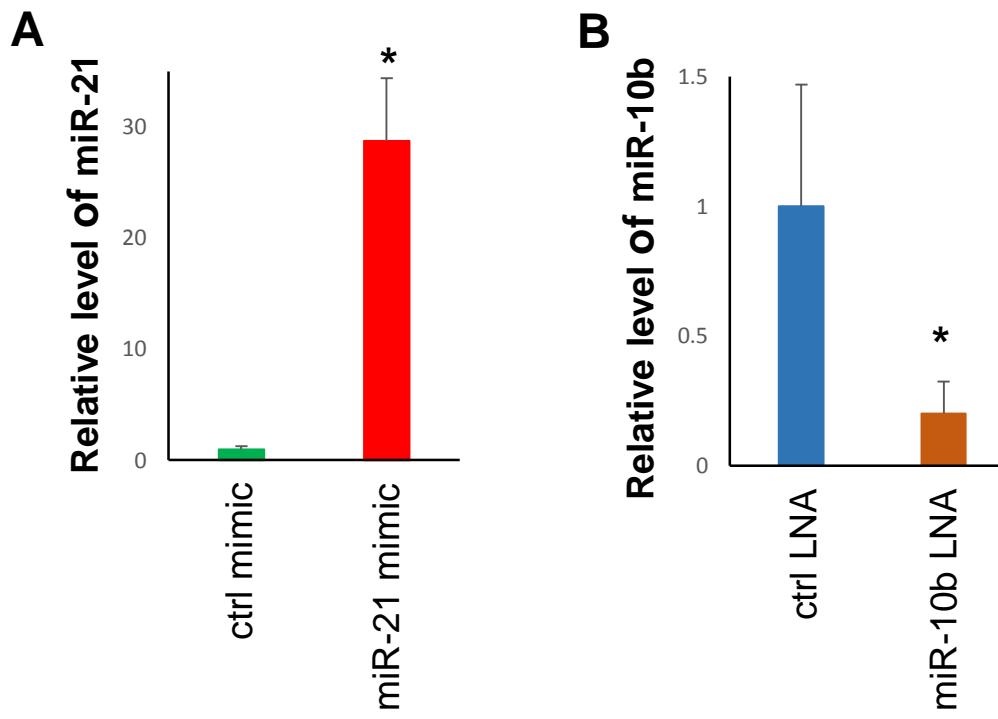
Supplementary Figure S1: Tissue specific microRNA signatures for skin and oral mucosa epithelium. MicroRNA profiling was performed on mouse skin and oral mucosal (palate) epithelium. Tissue specific microRNA signatures (presented as heatmap) for skin and oral mucosa epithelium were established by their differentially expression patterns (53 microRNAs, $p < 0.01$).



Supplementary Figure S2: Testing the classification potential of miRNome by unsupervised learning method to find grouping in samples set. Principal component analysis (PCA) was performed using the complete dataset ($n=24$), and PC1 and PC2 explain 23.7% and 19.9% of the variance, respectively (A). Prediction ellipses with probability of 0.95 were shown for tissue groups (B) and time group (C), indicate that PC1 can explain the variance associated with tissue specificity and PC2 appears to be associated with the variance related to wound healing time course.



Supplementary Figure S3: Testing the classification potential of microRNA signature sets by unsupervised learning method to find grouping in samples set. Hierarchical clustering analysis was performed on the whole sample set ($n=24$), using (A) 18 concurrently differentially expressed microRNAs associated with both skin and oral wound healing, (B) 33 differentially expressed microRNAs associated with oral wound healing, (C) 200 differentially expressed microRNAs associated with skin wound healing, (D) the combined 215 differentially expressed microRNAs in wound healing, (E) 53 baseline tissue specific differentially expressed microRNAs.



Supplementary Figure S4: Confirmation of the microRNA mimic mediated up-regulation of miR-21 and the LNA inhibitor mediated down-regulation of miR-10b in mouse skin wounds. (A) The miR-21 mimic or negative control mimic were delivered to the mouse skin wounds using an animal origin-free lipid nanoparticle (Invivofectamine 3.0 Reagent from Invitrogen) at the time of injury. The microRNA mimic mediated up-regulation of miR-21 was confirmed by TaqMan assay-based real time PCR quantification on day 1 wound samples. (B) The LNA inhibitor for miR-10b or negative control LNA were delivered to the mouse skin wounds using an animal origin-free lipid nanoparticle (Invivofectamine 3.0 Reagent from Invitrogen) at the time of injury. The LNA inhibitor mediated down-regulation of miR-10b was confirmed by TaqMan assay-based real time PCR quantification on wound tissue samples harvested at day 1 post-wounding (* p < 0.05).

Supplementary Table S1: Top 10 most abundant microRNAs in skin and oral mucosa during wound healing

wound healing time course	Skin		oral mucosa	
	miR name	% of miRNome	miR name	% of miRNome
0 hour (unwounded, baseline)	miR-10b	0.13453	miR-27b	0.096029
	miR-10a	0.076385	miR-26a	0.063469
	miR-26a	0.065607	miR-99a	0.059831
	miR-27b	0.060006	miR-143	0.04909
	miR-99a	0.040932	let-7c	0.045329
	miR-143	0.031033	miR-125b	0.045091
	miR-125b	0.030901	let-7f	0.034871
	let-7c	0.030461	miR-199a	0.032343
	let-7f	0.029307	miR-205	0.029648
	miR-1a	0.026512	miR-203	0.029631
6 hours post wounding	miR-21	0.010946 *	miR-21	0.013492 *
	miR-203	0.019338 *	miR-10b	0.028247 *
			miR-10a	0.000472 *
	miR-10b	0.116332	miR-27b	0.125659
	miR-1a	0.073676	miR-99a	0.066628
	miR-10a	0.065049	miR-26a	0.059918
	miR-27b	0.060364	miR-143	0.057865
	miR-99a	0.058471	let-7f	0.039192
	miR-26a	0.052577	let-7c	0.037748
	miR-143	0.039734	miR-125b	0.034433
24 hours post wounding	let-7f	0.037163	miR-203	0.033448
	miR-133a	0.027194	miR-199a	0.031841
	miR-99b	0.022871	miR-21	0.024844
	miR-21	0.020463 *		
	miR-21	0.064331	miR-27b	0.110127
	miR-10b	0.052373	miR-99a	0.053833
	miR-27b	0.050933	miR-26a	0.053736
	miR-10a	0.049371	miR-21	0.046823
	miR-99a	0.046014	let-7c	0.044605
	let-7f	0.04268	miR-143	0.042485
5 days post wounding	miR-26a	0.040699	let-7f	0.036187
	miR-146b	0.039564	miR-199a	0.028033
	miR-143	0.03701	miR-203	0.027665
	let-7c-5p	0.03682	miR-205	0.027606
	miR-21	0.129953	miR-27b	0.115615
	miR-27b	0.060603	let-7c	0.051672
	miR-10b	0.037804	miR-143	0.051497

let-7f	0.036286	miR-99a	0.051173
miR-26a	0.036118	miR-26a	0.047261
miR-143	0.033552	miR-21	0.040623
miR-199a	0.033522	let-7f	0.038901
let-7c	0.032773	miR-199a	0.031018
miR-99a	0.030027	miR-203	0.030965
miR-146b	0.030023	miR-125b	0.028131

note:

*: While not in top 10, these selected miRs were also presented for comparancing between tissues.

Supplementary Table S2: Baseline tissue-specific differentially expressed microRNAs between skin and oral mucosa epithelium

miR name	p-value	adj P	Skin		Oral		miR_seq
			Mean	StDev	Mean	StDev	
miR-378a-3p	1.29E-03	0.04304	15,167	183	6,013	400	ACTGGACTTGGAGTCAGAAGGC
miR-31-5p	1.85E-03	0.0523	437	140	9,980	739	AGGCAAGATGCTGGCATAGCT
miR-10a-5p	2.19E-03	0.05383	294,829	38,847	1,421	665	TACCCCTGTAGATCCGAATTGT
miR-10b-5p	2.52E-03	0.05931	519,259	82,583	85,095	24,872	TACCCCTGTAGAACCGAATTGT
miR-133a-3p	4.19E-03	0.0818	55,766	31,546	3,956	1,681	TTTGGTCCCCTCAACCAGCTGT
miR-146b-5p	4.92E-03	0.08283	66,712	6,568	10,796	3,023	TGAGAACTGAATTCCATAGGCTGT
miR-126a-5p	6.54E-03	0.096	3,437	491	6,408	985	CATTATTACTTTGGTAGCG
miR-34c-5p	7.34E-03	0.09894	122	43	10,229	8,453	AGGCAGTGTAGTTAGCTGATTGC
miR-30a-5p	8.14E-03	0.09918	47,463	4,893	29,008	1,047	TGTAAACATCCTCGACTGGAAGCT
miR-96-5p	8.19E-03	0.09918	11,975	1,738	6,919	917	TTTGGCACTAGCACATTTGCT
miR-1a-3p	8.30E-03	0.09918	102,330	27,364	9,517	5,590	TGGAATGTAAGAAGTATGTAT
miR-451a	9.75E-03	0.10566	823	249	16,627	11,565	AAACCGTTACCATTACTGAGT
miR-470-5p	2.90E-05	0.00801	0	0	344	40	TTCTTGGACTGGCACTGGTGAGT
miR-465b-3p	4.23E-05	0.00801	0	0	21	2	GATCAGGGCCTTCTAAGTAGA
miR-465c-5p	4.24E-05	0.00801	0	0	73	9	TATTTAGAATGGCGCTGATCTG
miR-10a-3p	7.10E-05	0.01005	51	8	0	0	CAAATTCTGTATCTAGGGGAAT
miR-881-3p	1.35E-04	0.01134	0	0	54	11	AACTGTGTCTTCTGAATAGA
miR-465a-5p	1.47E-04	0.01134	0	0	106	27	TATTTAGAATGGCACTGATGTGA
miR-107-3p	1.58E-04	0.01134	842	50	442	23	AGCAGCATTGTACAGGGCTATC
miR-1247-3p	1.60E-04	0.01134	0	0	24	6	CGGGAACGTCGAGACTGGAGC
miR-880-3p	2.50E-04	0.01574	0	0	55	15	TACTCCATCCTCTGTAGTAGA
miR-211-5p	3.99E-04	0.01965	423	87	42	10	TTCCCTTGTTCATCCTTGCCT
miR-1247-5p	4.36E-04	0.01965	12	5	601	286	ACCCGTCGGCTCGTCCCCGGA
miR-471-5p	4.46E-04	0.01965	0	0	22	9	TACGTAGTATAGTGCTTTAC
miR-351-5p	4.51E-04	0.01965	1,991	168	4,097	329	TCCCTGAGGAGCCCTTGAGCCT
miR-132-3p	8.69E-04	0.0335	34	1	122	13	TAACAGTCTACAGCCATGGTCG
miR-214-3p	8.88E-04	0.0335	2,337	229	4,735	453	ACAGCAGGCACAGACAGGCACT
miR-1983	1.20E-03	0.04229	30	6	111	17	CTCACCTGGAGCATGTTTCT
miR-34c	1.79E-03	0.0523	0	0	141	116	AGGCAGTGTAGTTAGCTGATTGT
miR-338-3p	1.84E-03	0.0523	470	59	1,124	77	TCCAGCATCAGTGATTTGTT
miR-872-3p	2.02E-03	0.05383	339	31	576	41	TGAACATTGCACTGCTCCT
miR-378b_R	2.10E-03	0.05383	3,066	241	1,296	186	ACTGGACTTGGAGTCAGAAGGCT
miR-150-5p	2.68E-03	0.06073	1,669	199	3,217	367	TCTCCAACCCTTGTACCAGT
miR-193b-3p	3.54E-03	0.07707	3,812	315	2,245	249	AACTGGCCCACAAAGTCCCGCT
miR-1a-1-5p	3.97E-03	0.0818	47	26	3	2	ACATACTCTTATATGCCATA
miR-467b-5p	4.21E-03	0.0818	5	3	86	50	TAAGTGCCTGCATGTATATGC
miR-218-5p	4.34E-03	0.0818	128	24	294	51	TTGTGCTTGATCTAACCATGT
miR-375-3p	4.67E-03	0.08283	277	43	1,377	450	TTTGTTCGTTGGCTCGCGTGA
miR-133a-5p	4.81E-03	0.08283	272	3	25	7	AGCTGGTAAAATGGAACCAAAT
miR-676-5p	4.98E-03	0.08283	76	10	38	2	ACTCTACACCTTAGGACTTGC
miR-672-5p	5.18E-03	0.0838	17	5	79	26	TGAGGTTGGGTACTGTGTGTGA
miR-511-5p	5.43E-03	0.08531	19	3	4	1	CATGCCTTTGCTCTGCACTCA
miR-128-3p	6.27E-03	0.09588	954	60	398	69	TCACAGTGAACCGGTCTCTT
miR-196a-5p	6.61E-03	0.096	186	216	0	0	TTAGGTAGTTCATGTTGTTGG
miR-223-3p	6.86E-03	0.09705	706	134	289	24	TGTCAGTTGTCAAATACCCCAT
miR-184-3p	7.25E-03	0.09894	41	5	22	3	TGGACGGAGAACTGATAAGGGT
miR-378d	7.63E-03	0.09918	1,332	134	368	96	ACTGGACTTGGAGTCAGAAGGT
miR-23a-5p	7.94E-03	0.09918	5	2	16	4	GGGGTTCTGGGGATGGGATT
miR-223-5p	8.56E-03	0.09918	105	15	19	6	CGTGTATTTGACAAGCTGAGTTGGACACT
miR-501-3p	8.59E-03	0.09918	3,835	392	1,609	327	AATGCACCCGGGCAAGGATTGG
miR-669a-5p	8.81E-03	0.09972	5	2	28	3	AGTTGTGTGCATGTTCATGTCT

miR-874-3p	9.87E-03	0.10566	146	30	308	48	CTGCCCTGGCCCCGAGGGACCGAC
miR-1198-5p	9.89E-03	0.10566	162	14	224	19	TATGTGTTCCCTGGCTGGCTTGG

notes:

Bold miR name: differentially expressed miRs that have the reads greater than the average (5458) of the dataset.

adj P: Benjamini-Hochberg adjusted P value.

Supplementary Table S3: Differentially expressed microRNAs in oral mucosal wound healing

miRNA	p-value	adj P	0hr		6hr		24h		5dy		miR_seq
			Fold	Chang	StDev	Fold	Chang	StDev	Fold	Chang	StDev
rno-miR-378a-3p	0.0008184	0.0867151	1	0.0664713	0.7328313	0.0347267	0.9548818	0.05151	1.0301608	0.09042	ACTGGACTTGGAGTCAGAAGGC
mmu-miR-148b-3p	0.0035625	0.1320058	1	0.0999046	1.3048705	0.4011814	2.4221855	0.9691142	3.2551242	0.6331512	TCAGTGCATCACAGAACCTTG
mmu-miR-181a-5p	0.0044725	0.1320058	1	0.1137351	0.8214249	0.0947066	0.6598383	0.0646125	0.7023587	0.0548996	AACATTCAACGCTCGGTGAGT
mmu-miR-31-5p	0.0066523	0.1516625	1	0.0740672	0.8015539	0.2209285	0.9452544	0.0772447	1.4866995	0.1154021	AGGCAAGATGCTGGCATAGCT
mmu-miR-23b-3p	0.0071932	0.1516625	1	0.111629	0.7357408	0.0466384	1.0309673	0.1224344	0.9944933	0.0684041	ATCACATTGCCAGGGATTACC
mmu-miR-200c-3p	0.0092364	0.1632779	1	0.1557818	0.8651189	0.2146033	1.7654788	0.5850174	1.8783663	0.1905794	TAATACTGCCGGTAATGATGGA
mmu-miR-195a-5p	0.0097908	0.1632779	1	0.1120782	0.9076189	0.0649711	0.8820817	0.1469965	0.6635547	0.0478765	TAGCAGCACAGAAATTGGC
mmu-miR-3068-5p	0.0002268	0.0752639	1	0.257308	1.0663981	0.3900415	3.5321486	0.8150827	3.6866457	0.1352283	TTGGAGTTCATGCAAGTTCTAAC
mmu-miR-101c	0.0002632	0.0752639	1	0.1031532	3.6350439	0.4766979	1.6863383	0.5710504	1.4593673	0.2365202	TACAGTACTGTGATAACTGAT
mmu-miR-30a-3p	0.0006925	0.0867151	1	0.0882855	1.0545809	0.0647274	0.7847491	0.0160182	0.8296518	0.0335745	CTTCAGTCGGATGTTGCAGC
mmu-miR-17-5p	0.0007147	0.0867151	1	0.1584791	0.8475679	0.0483152	1.5117758	0.1628031	0.9948067	0.0465453	CAAAGTGCTTACAGTGCAGGTAG
mmu-miR-219a-5p	0.0009096	0.0867151	1	0.2515456	0.8655548	0.1150959	1.9265423	0.2487612	0.9787669	0.078554	TGATTGTCCAACGCAATTCTCG
oan-miR-27b-3p	0.0013897	0.1075841	1	0.0529891	1.0604624	0.0815091	1.2212844	0.1148333	1.4813389	0.1309122	TTCACAGTGGCTAAGTCTGCGT
mmu-miR-676-5p	0.001521	0.1075841	1	0.0587988	1.2536092	0.1310737	1.5343393	0.1647377	1.5822301	0.1767254	ACTCTACAACCTTAGGACTTGC
mmu-miR-187-5p	0.0020207	0.1075841	1	0.1332469	0.9633736	0.1775188	1.0897682	0.1082475	1.7067577	0.0787297	AGGCTACAAACACAGGACCCGGGA
mmu-miR-136-5p	0.0020635	0.1075841	1	0.1548394	1.4044623	0.1419101	0.872338	0.052316	0.8988278	0.0852673	AGGACTCCATTGTTGTGATGAT
mmu-miR-362-5p	0.0020689	0.1075841	1	0.3961423	0.8407961	0.4010792	3.2770347	1.3747622	4.2970719	0.8784176	AATCCTTGGAACCTAGGTGTGAAT
mmu-miR-434-3p	0.0023305	0.1110887	1	0.0890199	1.317616	0.0777199	0.7998577	0.1350153	0.9871484	0.0587686	TTTGAACCACACTCGACTCCT
mmu-miR-20a-5p	0.002801	0.1232459	1	0.1382523	1.1142864	0.0608071	1.716838	0.3460544	1.4510558	0.0167519	AAAAGTGCTTATAGTGCAGGTAG
mmu-miR-425-3p	0.0035515	0.1320058	1	0.1509714	0.7956976	0.3416713	1.5800723	0.008423	1.9031004	0.134065	CATCGGGAAATGCGTGTCCGCC
mmu-miR-340-5p	0.0037138	0.1320058	1	0.0858626	2.052563	0.4916944	1.2777403	0.2398432	1.1007619	0.1596562	TTATAAAGCAATGAGACTGATT
mmu-miR-664-3p	0.0040783	0.1320058	1	0.0763262	0.9307747	0.0316889	0.7738323	0.146559	0.6382082	0.0549795	TATTCATTACTCCCCAGCCT
mdo-miR-181a-5p	0.0045652	0.1320058	1	0.4062099	0.4520095	0.1849618	1.5972577	0.5706788	1.6012956	0.1129886	AACATTCAACGCTCGGTGAGTTT
mmu-miR-700-5p	0.0046156	0.1320058	1	0.0989355	0.8064185	0.0942522	0.6528607	0.0928495	0.678155	0.0392187	TAAGGCTCCTCCTGTGCTTGC
mmu-miR-7a-5p	0.0049464	0.13473	1	0.1152172	1.4026239	0.3423426	2.0379229	0.3815397	1.533827	0.1159376	TGGAAGACTAGTGATTTGTTGTT
mmu-miR-193b-3p	0.0056089	0.1423618	1	0.1111418	0.5820916	0.0633109	0.6239657	0.1001622	0.5880681	0.1051606	AACTGGCCCACAAAGTCCCGCT
mmu-miR-141-5p	0.0057243	0.1423618	1	0.1374447	1.2939718	0.0981627	1.4636957	0.4518719	0.703996	0.0905252	CATCTCCAGTGCAGTGTGTTGGA
chi-let-7b-5p	0.0060559	0.1443321	1	0.2975176	0.5674505	0.268898	0.7429631	0.2223544	0.2416335	0.0560828	CGTGGTGAGGTAGTAGGTTGTGTTG
mmu-miR-369-3p	0.0072728	0.1516625	1	0.3373636	1.8783307	0.4679148	0.8015508	0.0371642	1.3602938	0.1336945	AATAATACATGGTTGATCTT
mmu-miR-223-3p	0.007424	0.1516625	1	0.0825289	2.6603169	0.8650528	3.7225306	2.4601821	1.0758219	0.174759	TGTCAAGTTGTCAAATACCCCAT
cgr-miR-2424	0.0086861	0.1632779	1	0.2404127	1.3771909	0.1189381	1.0305804	0.0909723	0.7900936	0.0090443	ACAGATCTTGGTAATCTGATGGCT
mmu-miR-185-5p	0.0089071	0.1632779	1	0.1768065	0.8605288	0.093547	1.1757479	0.1890558	1.4661178	0.1582404	TGGAGAGAAAGGCAGTCCCTGA
mmu-miR-1981-5p	0.0089076	0.1632779	1	0.1363764	0.7991036	0.0798437	1.5570912	0.6208528	2.1169091	0.5442181	GTAAAGGCTGGGCTTAGACGTGG

notes:

adj P: Benjamini-Hochberg adjusted P value.

Supplementary Table S4: Differentially expressed microRNAs in skin wound healing

miRNA	p-value	BH adj p-value	0hr		6hr		24h		5dy		miR_seq
			Fold	Chang	StDev	Fold	Chang	StDev	Fold	Chang	StDev
mmu-miR-362-5p	4.65E-08	1.37615E-05	1	0.0837141	0.6413249	0.0891709	4.2812828	0.4271781	5.5743529	0.866634	AATCCTTGGAACCTAGGTGTAAT
mmu-miR-378b	9.38E-07	0.000117099	1	0.5078893	101.84181	42.218462	68.367811	9.9917889	56.644035	8.8869892	CCTGGACTTGGAGTCAGAAGGC
mmu-miR-30a-5p	9.89E-07	0.000117099	1	0.1030876	1.2514485	0.111781	0.582879	0.0226006	0.453159	0.0342868	TGTAAACATCCCTGACTGGAAGCT
mmu-miR-21a-5p	1.90E-06	0.000187873	1	0.2539178	1.9940871	0.2800941	4.5685168	0.3442066	8.7049501	1.46660301	TAGCTTATCAGACTGATGTTGA
rno-miR-140-3p	2.82E-06	0.000238909	1	0.0324798	0.8162478	0.1297814	2.9362838	0.404611	1.6118772	0.037681	ACCACAGGGTAGAACACCACGGACT
mmu-miR-132-3p	4.93E-06	0.000340665	1	0.0352686	1.5800502	0.2569972	3.4745052	0.9094494	7.8466167	1.5319354	TAACAGTCTACAGCCATGGTCG
mmu-miR-142a-5p	5.18E-06	0.000340665	1	0.0255513	1.2848074	0.2651946	4.1437431	0.4405762	2.3621902	0.3453455	CCCATAAAAGTAGAAAGCACTAC
mmu-miR-574-5p	6.15E-06	0.000363805	1	0.1303411	1.0603561	0.0365244	1.7632596	0.1948124	2.3909489	0.1317313	TGAGTGTGTGTGTGAGTGT
mmu-miR-363-3p	8.53E-06	0.000459292	1	0.0952857	0.6652307	0.0891167	2.0053947	0.4497003	2.7139678	0.2054663	AATTGACGGTATCCATCTGTAT
mmu-miR-148b-3p	1.21E-05	0.000596292	1	0.1225519	1.4401436	0.1791595	3.6437393	0.7899089	2.8325129	0.2177709	TCAGTGATCACAGAACTTTG
mmu-miR-370-3p	1.40E-05	0.000638568	1	0.0605291	1.3253023	0.1824486	1.0811611	0.4500462	7.6443345	1.0795915	GCCTGCTGGGGTGGAACCTGGT
mmu-miR-503-5p	1.76E-05	0.00074296	1	0.0705028	2.5582538	1.259011	13.009299	3.9726796	24.044095	4.5673285	TAGCAGCGGGAACAGTACTGCAG
mmu-miR-130b-5p	1.89E-05	0.000747309	1	0.3830448	0.7955666	0.0347566	4.3723402	0.7893309	2.7859502	0.2058123	ACTCTTCCCTGTTGCACTACT
mmu-miR-223-5p	2.13E-05	0.000786602	1	0.1439168	1.0131108	0.2574665	4.9520961	0.3066759	1.9972864	0.5349822	CGTGTATTGACAAGCTGAGTTGGACACT
mmu-miR-193b-3p	2.63E-05	0.000843812	1	0.0826412	0.7382197	0.0453673	0.528423	0.1066424	0.2426527	0.0544516	AACTGGCCACAAAGTCCCGCT
mmu-miR-133a-5p	2.68E-05	0.000843812	1	0.0124961	1.6103408	0.6638912	0.6235684	0.0782252	0.2382276	0.0268378	AGCTGGAAAATGGAACCAAAT
mmu-miR-409-3p	2.82E-05	0.000843812	1	0.1951633	1.643613	0.2753653	1.6411885	0.5257005	7.3129923	1.1643407	GAATGTTGCTCGGTGAACCCCT
mmu-miR-30e-3p	2.85E-05	0.000843812	1	0.1336761	1.0853182	0.0504217	0.7121563	0.0388258	0.4821073	0.0605488	CTTCAGTCGGATGTTACAGC
mmu-miR-540-3p	3.20E-05	0.000902329	1	0.303299	1.4690526	0.4930735	1.3405418	0.3674875	8.9795826	1.373148	AGGTCAAGGGTCGATCCTGGC
mmu-miR-106b-3p	3.48E-05	0.00093549	1	0.1005209	1.0092353	0.1131692	3.1954698	0.7250982	2.2884583	0.2963492	CCGCACTGTTGGTACTTGCT
cgr-miR-139-5p	3.93E-05	0.00101248	1	0.1717709	1.5692417	0.2170089	6.2644141	1.1548753	1.8048574	0.5971037	TCTACAGTGACCGTCTCCAGT
mmu-miR-3068-5p	4.23E-05	0.001042265	1	0.16667673	0.6336216	0.051074	2.0222936	0.5268974	2.5359095	0.3824849	TTGGAGTTCATGCAAGTTCAACC
mmu-miR-1198-5p	5.01E-05	0.001096217	1	0.0848818	1.2767261	0.0491863	2.4768726	0.5033243	1.9572109	0.151669	TATGTGTTCTGGCTGGCTTGG
mmu-miR-7a-5p	5.29E-05	0.001096217	1	0.2993717	1.8734799	0.1347995	5.4050207	0.8134072	5.6690703	1.739785	TGGAAGACTAGTGATTTGTTGTT
mmu-miR-1983	5.38E-05	0.001096217	1	0.1999005	1.1325902	0.4877763	3.0117198	0.5608882	8.178836	2.1482012	CTCACCTGGAGCATTTTCT
mmu-miR-31-5p	5.45E-05	0.001096217	1	0.3202481	0.6691327	0.4225293	0.8408116	0.3884696	11.688945	1.7979082	AGGCAAGATGCTGGCATAGCT
mmu-miR-195a-5p	5.56E-05	0.001096217	1	0.1482205	1.0863882	0.1409315	0.6123434	0.0445061	0.4327755	0.0546023	TAGCAGCACAGAAATATTGGC
mmu-miR-485-3p	5.93E-05	0.001133174	1	0.2459614	1.9433503	0.0360904	1.2241166	0.5878268	6.8873522	0.8276695	GTCATACACGGCTCCTCTCT
mmu-miR-223-3p	7.01E-05	0.001296402	1	0.1896275	3.4725465	1.0452788	21.881903	7.0927223	5.3417263	2.7982239	TGTCAGTTGTCAAATACCCCAT
mmu-miR-667-3p	7.46E-05	0.001337691	1	0.1521384	0.8861784	0.191653	0.6646901	0.241756	3.5485176	0.349971	TGACACCTGCCACCCAGCCAAG
mmu-miR-425-3p	8.08E-05	0.001402056	1	0.2348775	0.6035792	0.1815773	2.1053006	0.0998882	2.2176167	0.2181035	CATGGGAATGTCGTGTCGCC
mmu-miR-335-3p	8.29E-05	0.001402056	1	0.3701342	1.8152725	0.652019	0.9940331	0.1049006	6.5290917	1.3450705	TTTTCATATTGCTCTGACC
mmu-miR-433-3p	8.93E-05	0.00146842	1	0.2100516	1.708403	0.338726	1.6380816	0.5173994	5.8043222	0.7944497	ATCATGATGGGCTCCTCGGTGT
mmu-miR-30a-3p	1.03E-04	0.001644349	1	0.183004	1.2628488	0.156489	0.6681405	0.1129742	0.3797533	0.064559	CTTCAGTCGGATGTTGCAGC
mmu-miR-34c-5p	1.23E-04	0.00191319	1	0.3544813	1.8903948	0.3133666	3.4416824	0.5934597	35.217775	29.988102	AGGCAGTGTAGTTAGCTGATTGC
mmu-miR-298-5p	1.35E-04	0.001986375	1	0.1104028	1.5461294	0.6010547	2.0791757	0.5955064	8.4553965	1.8720677	GGCAGAGGAGGGCTGTTCTCCC
mmu-miR-379-3p	1.38E-04	0.001986375	1	0.2696918	1.3416434	0.3230751	0.8762274	0.213002	4.2139488	0.7683176	TATGTAACATGGTCCACTAAC
mmu-let-7g-5p	1.44E-04	0.002025249	1	0.0380169	1.3297929	0.0565121	1.8645238	0.2929749	1.1722114	0.0564105	TGAGGTTAGTAGTTGTACAGTT
mmu-miR-3535	1.57E-04	0.002155566	1	0.2646319	1.1974208	0.3209042	0.4218387	0.082361	0.2401472	0.0560957	TGGATATGATGACTGATTACCTGAGA

mmu-miR-96-5p	1.60E-04	0.002158314	1	0.1451501	1.059792	0.3840414	0.3436007	0.0457457	0.2286068	0.049049	TTTGGCACTAGCACATTTGCT
hsa-miR-411-3p	1.81E-04	0.002375894	1	0.3764971	1.3391691	0.4468603	0.8979431	0.2188992	6.635335	1.6070453	TATGTAACACGGCCACTAAC
mmu-miR-134-5p	1.93E-04	0.002477828	1	0.5045897	1.8306631	0.2552175	1.6669484	0.4641584	7.7421117	0.9986563	TGTGACTGGTTGACCAGAGGGG
mmu-miR-503-3p	1.97E-04	0.002477828	1	0.1669201	1.3625549	0.1656011	1.3606887	0.1713954	2.8326096	0.5163766	GAGTATTGTTCCACTGCCTGG
mmu-miR-351-3p	2.21E-04	0.002697113	1	0.5160102	3.1651568	0.5740431	4.2904132	0.3168941	12.748582	3.9546619	GGTCAAGAGGCGCCTGGGAAC
mmu-miR-30d-5p	2.31E-04	0.002697113	1	0.0801116	1.0830755	0.0613761	0.9045143	0.0719118	0.6733213	0.04992	TGTAAACATCCCCGACTGGAAGCT
mmu-miR-467b-5p	2.32E-04	0.002697113	1	0.5601718	2.8944241	0.3901845	8.115511	3.272596	5.5245781	0.405032	TAAGTGCCTGCATGTATATGC
mmu-miR-1a-3p	2.35E-04	0.002697113	1	0.2674081	2.9644116	1.6124386	0.7510039	0.0723657	0.3253727	0.0818349	TGGAATGTAAGAAGTATGTAT
mmu-miR-323-3p	2.39E-04	0.002697113	1	0.5585449	4.129499	0.7608464	3.4139647	1.6093895	15.024682	4.2041709	CACATTACACGGTCGACCTCT
mmu-miR-361-3p	2.41E-04	0.002697113	1	0.0792432	0.688706	0.0080252	1.0993344	0.162064	1.2191758	0.0198303	CCCCCAGGTGTGATTCTGATTGT
mmu-miR-541-5p	2.59E-04	0.002844113	1	0.1833458	1.3577702	0.1383864	1.0916668	0.415887	5.4283139	1.4219507	AAGGGATTCTGATGTTGGTCACACT
mmu-miR-26a-5p	2.85E-04	0.003068298	1	0.2251968	0.8548666	0.1153536	0.4822476	0.0365683	0.4036757	0.0567906	TTCAAGTAATCCAGGATAGGCT
mmu-miR-494-3p	3.16E-04	0.003343007	1	0.4294394	2.1115554	0.5560297	0.7715996	0.3377576	8.4825276	3.232242	TGAAACATACACGGAAACCTCT
mmu-miR-376b-3p	3.30E-04	0.003408051	1	0.3440436	1.6609537	0.6776486	0.7467926	0.2669873	5.7122662	0.8513435	ATCATAGAGGAACATCCACTT
mmu-miR-186-5p	3.34E-04	0.003408051	1	0.0690767	1.1457398	0.1549692	0.587867	0.0332535	0.6642662	0.106961	CAAAGAATTCTCCTTTGGGCT
mmu-miR-212-5p	3.59E-04	0.003562603	1	0.3830206	2.1547657	0.4243423	3.2183497	0.9333186	6.9008617	1.8257686	ACCTTGCTCTAGACTGCTTACT
mmu-miR-15b-3p	3.61E-04	0.003562603	1	0.0305556	0.8181714	0.1699805	2.3387035	0.6628641	2.2079041	0.2324212	CGAATCATTATTCGCTGCTCT
rno-miR-25-3p	3.67E-04	0.003562603	1	0.1632616	1.0748019	0.095391	2.2901618	0.4756695	1.6257507	0.0638482	CATTGCACTTGTCTCGGTCGA
mmu-miR-543-3p	3.80E-04	0.003566687	1	0.7012131	2.5617578	0.5445745	2.1204736	0.2965955	12.495648	2.3040777	AAACATTGCGGGTGCACTTCTT
mmu-miR-204-5p	3.86E-04	0.003566687	1	0.1804009	1.0925747	0.2258802	0.8357314	0.1992158	0.229482	0.09897	TTCCCTTGTCATCCTATGCC
mmu-miR-200a-3p	4.00E-04	0.003642225	1	0.1610251	1.1346017	0.0737134	1.0736307	0.0458806	0.4867077	0.1251561	TAACACTGTCTGGTAACGATGT
cfa-miR-194	4.25E-04	0.003812687	1	0.3220983	1.197963	0.4374698	5.4768064	1.0668365	3.079215	0.826779	TGTAACAGCAACTCCATGTTGGAT
mmu-miR-34b-3p	4.67E-04	0.004117098	1	0.0983948	2.4936436	0.4709371	3.3565731	1.2588176	31.724868	31.777169	AATCACTAACTCCACTGCCATC
mmu-miR-450b-5p	4.73E-04	0.004117098	1	0.4271449	1.822944	0.3300301	4.0121439	1.1600193	4.784724	1.1127844	TTTTGCAGTATGTTCTGAATAC
mmu-miR-3470a-p3	5.06E-04	0.004255833	1	0.3114472	5.1495352	2.9759897	8.2700526	1.2078015	2.1132654	0.5770552	TGCCCTGCGCTCTGAGTGC
mmu-miR-679-5p	5.06E-04	0.004255833	1	0.6020162	1.2697043	0.1811856	0.4040892	0.1382002	3.96496	1.2275843	TTGGACTGTGAGGTGACTCTGGT
cgr-miR-139-3p	5.10E-04	0.004255833	1	0.6151021	1.3568197	0.1397321	7.1855223	1.3224754	3.8051502	1.2796347	TGGAGACGCGGCCCTGTTGGAGT
mmu-miR-125a-3p	5.21E-04	0.004265844	1	0.2462619	1.5251663	0.1751887	2.4627273	0.4619914	2.3824232	0.061039	ACAGGTGAGGTTCTGGGAGC
hsa-mir-4450-p3	5.26E-04	0.004265844	1	0	423.0273	730.97273	30564.027	5904.1359	20063.524	6812.6168	CCCCCTGGCCCGTGG
mmu-miR-382-5p	5.34E-04	0.004274067	1	0.2798859	1.7164581	0.6450547	2.1983785	0.5719527	5.3355518	0.9304146	GAAGTTGTCGTGGTGGATTG
mmu-miR-152-5p	5.42E-04	0.004274247	1	0.0518277	0.7466772	0.0321612	0.7031543	0.1416423	1.5704891	0.2917459	TAGGTTCTGTGATACTCCGACT
mmu-miR-877-5p	6.10E-04	0.004692327	1	0.3079597	1.4961093	0.0473247	2.0203964	0.3501595	2.5746913	0.1153542	GTAGAGGAGATGGCGCAGGGGAC
mmu-miR-130b-3p	6.27E-04	0.004755625	1	0.5063375	0.3902219	0.1459471	3.0441247	0.9317452	2.8090063	0.99998	CAGTGCATGATGAAAGGGCAT
mmu-miR-125b-5p	6.75E-04	0.005008493	1	0.2615703	0.602216	0.1009059	0.3162235	0.0351576	0.4942301	0.0759235	TCCCTGAGACCCCTAACCTTGTA
mmu-miR-145a-3p	6.77E-04	0.005008493	1	0.0664913	1.3256934	0.1505205	0.6756536	0.0903052	0.6451689	0.1304375	GGATTCTGGAAATACTGTTCT
mmu-miR-1193-3p	7.06E-04	0.005131305	1	0.2535784	1.808854	0.6342052	0.6718077	0.2675109	3.7330045	0.8049799	TAGGTCAACCGTTTACTATC
mmu-miR-181c-5p	7.11E-04	0.005131305	1	0.171084	0.902911	0.1426848	0.5856032	0.1623561	0.3183347	0.0794569	AACATTCAACCTGTCGGTGAGT
mmu-miR-669c-5p	7.53E-04	0.005368925	1	0.3905979	0.8600733	0.229576	3.8504176	1.1115969	2.461776	0.3822116	ATAGTTGTTGTTGGATGTTG
mmu-miR-21a-3p	7.62E-04	0.005369281	1	0.9340874	5.2760077	2.7618646	15.400221	7.0625357	20.108678	4.3897202	CAACAGCAGTCGATGGGCTGT
hsa-miR-320b	7.95E-04	0.005511112	1	0.2189228	1.1937658	0.1912847	1.4762131	0.3631229	0.4490728	0.1131755	AAAAGCTGGGTTGAGAGGGT
mmu-miR-365-3p	8.01E-04	0.005511112	1	0.30881	0.5384289	0.0563338	0.3219968	0.0683889	0.309191	0.062356	TAATGCCCTAAACCTCTT
mmu-miR-361-5p	8.31E-04	0.005651865	1	0.2585363	1.214897	0.108786	2.5187528	0.3208922	1.7109224	0.2354898	TTATCAGAACCTCCAGGGTAC
mmu-miR-1a-1-5p	8.64E-04	0.005810023	1	0.5504198	1.6179643	0.6028675	0.7709123	0.4339074	0.1381317	0.0368482	ACATACTTCTTATGCCAT

mdo-miR-22-3p	9.34E-04	0.006215807	1	0.2597698	3.4874247	0.4413912	2.1055126	0.1832964	2.034302	0.6535229	CAGCTGCCAGTTGAAGAACCTGC
ptr-miR-203	9.53E-04	0.006216654	1	0.4441403	0.570919	0.0756663	0.3749706	0.0370697	0.2519709	0.0313367	TGAAATGTTAGGACCACTAGAAT
ptr-let-7i	9.56E-04	0.006216654	1	0.089807	1.0789105	0.0827996	1.5217702	0.2550633	1.624975	0.0551918	TGAGGTAGTAGTTGTGCTGTT
mmu-miR-30c-5p	9.80E-04	0.006306472	1	0.1339291	0.9809341	0.0244722	0.7775185	0.0205516	0.6348351	0.078099	TGTAAACATCCTACACTCTCAGCT
mmu-miR-99a-5p	9.96E-04	0.006316442	1	0.0842003	1.5237995	0.1828551	0.8738889	0.2086207	0.5379115	0.1481965	AACCCGTAGATCCGATCTTGT
mmu-miR-532-5p	1.00E-03	0.006316442	1	0.2361885	1.1308605	0.2618443	2.3098308	0.367316	2.316798	0.4282187	CATGCCTGAGTGTAGGACCGT
mmu-miR-203-3p	1.02E-03	0.006378673	1	0.0320972	0.7815985	0.0756983	0.6285213	0.0475543	0.5676929	0.1051031	TGAAATGTTAGGACCACTAGT
mmu-miR-369-5p	1.06E-03	0.006515823	1	0.1892327	1.906397	0.3355055	1.2184863	0.2815081	2.689006	0.4215147	AGATCGACCGTGTTATTCGC
mmu-mir-3470b-p5	1.11E-03	0.006661251	1	0.0282929	1.07393	0.0937758	1.9358043	0.4643978	0.7823614	0.1765447	TGTAGCCCTGGCTGTCC
mmu-miR-133a-3p	1.11E-03	0.006661251	1	0.5656919	2.0078005	1.0507723	0.6972099	0.3142804	0.1621058	0.0153855	TTTGGTCCCCCTCAACCAGCTGT
mmu-miR-872-3p	1.14E-03	0.006738455	1	0.0921454	0.7905882	0.0106498	0.7117445	0.09328	1.1642097	0.1323534	TGAACTATTGCAGTAGCCTCCT
hsa-miR-4448	1.17E-03	0.006738455	1	0.7641477	1.5148669	1.040295	13.416246	5.3000464	4.1536743	0.8637108	GGCTCGTTGGCTAGGGG
mmu-miR-10b-5p	1.17E-03	0.006738455	1	0.1590404	0.9224281	0.4031696	0.302635	0.0911346	0.2060538	0.0682445	TACCCGTAGAACCGAATTGT
mmu-miR-708-3p	1.17E-03	0.006738455	1	0.2572063	1.2745179	0.1359119	0.7035791	0.1237406	0.5078613	0.0593653	CAACTAGACTGTGAGCTTAGT
mmu-miR-125b-1-3p	1.19E-03	0.006757886	1	0.3020723	2.709657	0.0590845	1.7070669	0.2485054	1.6080386	0.1214718	ACGGGTTAGGCTCTGGGAGCT
mmu-miR-5099	1.20E-03	0.006757886	1	0.1587364	1.2776844	0.336601	3.3656113	0.9662659	1.6002579	0.1353409	TCTTAGATCGATGTGGTCTC
rno-miR-214-3p	1.23E-03	0.006857908	1	0.0981375	0.8395052	0.2117344	0.8169993	0.1111326	1.8510233	0.2991751	ACAGCAGGCACAGACAGGCAGT
mdo-miR-26-5pC	1.26E-03	0.006997458	1	0.316483	0.7956475	0.0435024	0.3702769	0.0857739	0.4072611	0.0795074	TTCAAGTAACCCAGGATAGGCT
mmu-miR-615-3p	1.30E-03	0.007111274	1	0.1234199	1.0080731	0.2256506	0.8355923	0.1584079	2.4102891	0.5818651	TCCGAGCCTGGGTCTCCCTCT
mmu-miR-340-3p	1.31E-03	0.007111274	1	0.223756	1.1499104	0.1054169	2.0666786	0.2216629	1.3222915	0.1005315	TCCGTCTCAGTTACTTTATAGC
mmu-miR-212-3p	1.34E-03	0.007211834	1	0.650503	1.7846136	0.7098364	5.9406308	2.5047141	7.6220823	1.8044612	TAACAGTCTCCAGTCACGCC
mmu-miR-192-5p	1.39E-03	0.007418295	1	0.1520014	1.2029768	0.1076329	1.3479674	0.0514551	0.7931981	0.0929728	CTGACCTATGAATTGACAGCC
mmu-miR-337-5p	1.40E-03	0.007418295	1	0.1743153	1.8870253	0.6741124	1.1635705	0.1782623	3.2104453	0.5461503	CGCGCGTATGCAGGAGTTGATT
mmu-miR-673-5p	1.45E-03	0.007570487	1	0.3511423	1.2538674	0.3364797	0.5105595	0.1806137	2.4970974	0.3003384	CTCACAGCTCTGGCCTTGGAGC
mmu-miR-146b-3p	1.48E-03	0.007673969	1	0.2384586	1.8645249	0.7113588	3.6254825	0.0704519	2.298739	0.3943435	GCCCTAGGGACTCAGTTCTGG
mmu-miR-299b-3p	1.58E-03	0.008119127	1	0.5459631	1.8728109	0.4820067	1.1168657	0.2777005	4.8652691	1.3935738	GTATGTGGACGGTAACCGCT
mmu-miR-329-5p	1.63E-03	0.008214548	1	0.2896125	1.6732116	0.198918	0.8809755	0.2324953	2.4263452	0.4331832	AGAGGTTTCTGGGTCTGTGTT
mmu-miR-423-3p	1.63E-03	0.008214548	1	0.1233289	0.8636987	0.0333471	1.3855005	0.264183	1.4785747	0.0413327	AGCTCGGTCTGAGGCCCTCAGT
mmu-miR-1195	1.64E-03	0.008214548	1	0.3114472	2.1630132	0.8272173	6.9081157	1.4070997	1.6468567	0.8384273	TGAGTTCGAGGCCAGCTGGTC
mdo-miR-26-5p	1.67E-03	0.008284584	1	0.4683106	0.5418045	0.1753635	0.2301134	0.0138412	0.229103	0.0290966	CGTGGTTCAAGTAATCCAGGATAGGCT
mmu-miR-199a-5p	1.72E-03	0.008486035	1	0.0562269	0.7437197	0.1400344	0.4944118	0.1129725	1.2556766	0.2676846	CCCAGTGTTCAGACTACCTGTT
mmu-miR-200c-3p	1.79E-03	0.008741844	1	0.2116411	0.5786912	0.1911222	1.2939722	0.2034201	1.6404653	0.1749315	TAATACTGCCGGTAATGATGGA
mmu-miR-329-3p	1.82E-03	0.008747475	1	0.4003871	1.3164219	0.4512617	0.7721013	0.0459532	2.7046698	0.2062814	AACACACCCAGCTAACCTTTT
mmu-miR-3068-3p	1.83E-03	0.008747475	1	0.0853513	1.3436764	0.1303298	0.8816311	0.1147326	0.7104505	0.1218952	GGTGAATTGCAGTACTCCAAC
mmu-miR-3473d	1.83E-03	0.008747475	1	0.0983948	0.6864788	0.4096885	1.5027072	0.221669	3.4271248	0.5691545	CACTGAGCCACTTCCAGCCCT
mmu-miR-340-5p	1.90E-03	0.008992678	1	0.4536075	2.6121444	1.098281	6.9295512	2.1219381	3.4206165	1.0923573	TTATAAAGCAATGAGACTGATT
mmu-miR-125a-5p	1.93E-03	0.00905272	1	0.3378152	0.5212082	0.1040373	0.2903637	0.0310685	0.4707845	0.0916847	TCCCTGAGACCCCTTAACCTGT
mmu-miR-328-3p	1.94E-03	0.00905272	1	0.1269239	1.2531525	0.108406	1.073184	0.156667	0.7256735	0.0490708	CTGGCCCTCTGCCCCCTCGTT
mmu-miR-181d-5p	1.96E-03	0.00905272	1	0.1154041	0.8370233	0.1188889	0.7635974	0.0925107	0.531843	0.0667381	AACATTCAATTGTTGTCGGTGGGT
mmu-miR-98-3p	1.99E-03	0.009087899	1	0.2002938	0.9050202	0.1561054	0.9469613	0.2892732	2.4031906	0.3844809	CTATACAACCTACTACCTTCT
mmu-miR-335-5p	2.00E-03	0.009087899	1	0.1500362	1.5836286	0.3175577	0.3961481	0.2564381	1.6974553	0.4220364	TCAAGAGCAATAACGAAAAATGT
mmu-miR-221-3p	2.05E-03	0.009198708	1	0.0451682	0.9441157	0.0433964	1.4548521	0.2312642	1.3346873	0.1339009	AGCTACATTGTTGCTGGTTTAC
mmu-miR-132-5p	2.07E-03	0.009198708	1	0.7180719	4.5208297	2.2352513	8.8163452	2.5033658	15.456815	2.1802049	ACCGTGGCTTCGATTGTTACT

mmu-miR-30c-2-3p	2.08E-03	0.009198708	1	0.1992231	0.9653194	0.0988994	0.6472745	0.1060463	0.453801	0.0974637	CTGGGAGAAGGCTGTTACTCT
mmu-miR-193b-5p	2.08E-03	0.009198708	1	0.0636888	0.9340729	0.098463	0.5763472	0.1918744	0.4791185	0.0592132	CGGGGTTTGAGGGCGAGATGA
cgr-miR-214-5p	2.11E-03	0.00923999	1	0.2179683	0.8060234	0.100771	0.7994889	0.1596177	1.6202021	0.0989944	TGCCCTGTCTACACTGCTGTGC
mmu-miR-152-3p	2.18E-03	0.009485439	1	0.2107579	1.331513	0.1110373	1.2382011	0.116605	1.7990415	0.0460345	TCAGTGCATGACAGAACTTGG
mmu-miR-10a-5p	2.23E-03	0.009632311	1	0.1317609	0.9084143	0.3592673	0.5024623	0.1548674	0.2383207	0.0880976	TACCCCTGAGATCCGAATTGT
mmu-miR-10a-3p	2.25E-03	0.009632311	1	0.1532046	0.9745423	0.1714622	1.9002149	0.2383496	1.2308395	0.1886834	CAAATTCTGATCTAGGGAAAT
mmu-miR-674-5p	2.27E-03	0.009649775	1	0.1573378	0.7600588	0.1507554	1.231239	0.0935557	1.5412692	0.2082108	GCACTGAGATGGGAGTGGTAGA
mmu-miR-434-5p	2.38E-03	0.010064861	1	0.3704331	1.8505936	0.1136602	0.622515	0.2264647	1.9643819	0.349352	GCTCGACTCATGGTTGAACCA
chi-miR-99b-3p	2.46E-03	0.010323635	1	0.1695248	0.9689348	0.1332949	1.5792321	0.1267033	2.1772204	0.6362701	CAAGCTCGTGTCTGGTCCGT
mmu-miR-351-5p	2.51E-03	0.010465544	1	0.0844893	1.4586638	0.3793304	0.9875856	0.2087979	2.4672966	0.6070181	TCCCTGAGGAGCCCTTGAGCCT
mmu-miR-379-5p	2.58E-03	0.010688997	1	0.6197519	2.6005561	0.6625438	2.7930513	0.5480395	4.5700396	0.6675896	TGGTAGACTATGGAACGTAGG
hsa-miR-4454	2.71E-03	0.011135146	1	0.311402	1.1938938	0.1763517	2.3062566	0.152184	1.8604314	0.3211017	TCGGATCCGAGTCACGGCACCA
mmu-miR-664-3p	2.86E-03	0.011691663	1	0.4108561	1.3226349	0.1018787	0.6945605	0.123068	0.3892615	0.0571229	TATTCATTTACTCCCCAGCCT
mmu-miR-665-3p	2.89E-03	0.011721564	1	0.6152815	1.8785085	1.0785012	1.2135596	0.7321664	7.6085242	0.9796893	ACCAGGAGGCTGAGGTCCCTTA
mmu-miR-148b-5p	2.94E-03	0.011779138	1	0.2371952	1.5091068	0.1941134	1.6384198	0.4292438	2.5139179	0.2235382	GAAGTTCTGTTACACTCAGGC
mmu-miR-107-3p	2.94E-03	0.011779138	1	0.0588604	0.7637262	0.043722	0.7815476	0.1566574	0.5728624	0.0601727	AGCAGCATTGTACAGGGCTATC
mmu-miR-574-3p	3.08E-03	0.012233153	1	0.159976	0.7226952	0.056914	0.6026445	0.0536147	0.7100194	0.0608214	CACGCTCATGCACACACCCACA
mmu-miR-20a-5p	3.18E-03	0.012458717	1	0.2842205	1.3300627	0.2187253	2.704159	0.4683716	2.3073174	0.6753191	TAAAGTGTTATAGTGCAGGTAG
rno-miR-181a-2-3p	3.20E-03	0.012458717	1	0.2653292	1.5079592	0.560857	1.0365957	0.1916379	0.4258602	0.1266605	ACCACCGACCGTTGACTGTACC
mmu-miR-24-2-5p	3.29E-03	0.012747155	1	0.0415376	0.7326629	0.0577254	0.5196467	0.1204124	0.7367385	0.0788055	GTGCCTACTGAGCTGAAACAGT
mmu-miR-210-3p	3.34E-03	0.012844095	1	0.3060559	0.7264773	0.0703653	1.4910294	0.5197572	2.3399945	0.5231143	CTGTGCGTGTGACAGCGGCTGA
mmu-miR-484	3.53E-03	0.013400543	1	0.1607864	0.6242116	0.0959694	0.5689294	0.1027721	0.9602006	0.0738081	TCAGGCTCAGTCCCCTCCCGAT
mmu-miR-221-5p	3.53E-03	0.013400543	1	0.123451	1.3830767	0.4084483	0.7507778	0.1707078	0.5391125	0.0653802	ACCTGGCATACAATGTAGATTCTGT
rno-miR-378a-5p	3.68E-03	0.013885205	1	0.1162085	1.2795807	0.4581227	0.7671961	0.1018661	0.4619925	0.1063637	CTCCTGACTCCAGTCCTGTGT
mmu-miR-15b-5p	3.92E-03	0.014544485	1	0.254236	0.9961632	0.1559138	1.9864855	0.2866396	1.6486097	0.2356136	TAGCAGCACATCATGGTTACA
mmu-mir-3102-p3	3.93E-03	0.014544485	1	0.1217697	1.1231961	0.0398356	1.2453359	0.1695152	1.7391794	0.2926564	GAGCACCCATTGGCTACCCACA
rno-miR-25-5p	4.02E-03	0.014777333	1	0.2642865	0.5820014	0.1125325	1.3843085	0.3165579	1.3872368	0.2148243	AGGCGGAGACTTGGCAATTGCT
rno-miR-874-3p	4.04E-03	0.014777333	1	0.203448	1.6711652	0.1747096	0.9866017	0.1144491	1.2519861	0.0609583	CTGCCCTGGCCCGAGGGACCGAC
mmu-miR-1839-5p	4.17E-03	0.015161566	1	0.0349554	1.2111516	0.0284152	1.3177189	0.1793922	0.9655972	0.063744	AAGGTAGATAGAACAGGTCTTGT
mmu-miR-3970	4.21E-03	0.015189612	1	0.0293925	1.6609021	0.6928908	2.5055393	0.1968514	1.2403072	0.1757495	TGAGGTAGTAGTTGTGCTT
mmu-miR-211-5p	4.24E-03	0.015211967	1	0.2066138	0.5482685	0.2593193	0.4384779	0.0998019	0.2386098	0.0661386	TTCCCTTTGTCATCCTTGGCCT
mmu-miR-495-3p	4.32E-03	0.015413002	1	0.408068	1.5209092	0.4491724	0.8112002	0.2079628	3.2814585	0.9815277	AAACAAACATGGTGCACCTCTT
mmu-miR-1247-5p	4.37E-03	0.015475064	1	0.3893578	0.8500563	0.6325552	1.5219218	0.494682	8.504674	2.8816599	ACCCGTCGGTTCGTCGGCGGA
mmu-miR-300-3p	4.56E-03	0.016053992	1	0.3589051	1.9686281	0.7736451	1.2421537	0.5566182	3.8856108	0.6304635	TATGCAAGGGCAAGCTCT
mmu-miR-28c	4.62E-03	0.016168779	1	0.038418	2.5820541	0.5936916	2.346044	0.6593746	2.1849821	0.7020198	AAGGAGCTCACAGTCATTGAA
mmu-miR-99a-3p	4.67E-03	0.016241055	1	0.3122041	0.5129538	0.0365709	0.34673	0.0542963	0.4371538	0.1392292	CAAGCTCGTTCTATGGGTCTGT
mmu-miR-23b-3p	4.69E-03	0.016241055	1	0.1794263	0.71159	0.0856286	0.6373715	0.1060278	0.4965126	0.0713242	ATCACATTGCCAGGGATTACC
mmu-miR-676-5p	4.75E-03	0.016241143	1	0.1366818	0.784872	0.1113213	0.6072587	0.0551504	0.6484576	0.0842404	ACTCTACACCTTAGGACTTGC
mmu-miR-224-5p	4.75E-03	0.016241143	1	0.1463129	1.2226207	0.2474027	1.0545799	0.3656132	2.3652663	0.1931441	TAAGTCACTAGTGGTCCGTT
mmu-miR-712-5p	4.87E-03	0.016575598	1	0.2597698	0.8459174	0.5202588	2.1886211	0.4080564	4.0927353	0.7872991	CTCCTTCACCCGGCGTACCCGC
mmu-miR-382-3p	4.95E-03	0.01673073	1	0.1506408	1.7382395	0.7143872	1.2238049	0.5702819	3.5159739	0.6977001	AATCATTCACGGACAAACACTT
mmu-miR-222-3p	4.99E-03	0.016771314	1	0.1176075	1.1978463	0.2587777	2.0301863	0.5529001	1.8339041	0.1542446	AGCTACATCTGGCTACTGGGTCT
hsa-miR-7977	5.20E-03	0.017391405	1	0.3892841	0.7436631	0.5078532	3.682947	0.9320637	6.1648424	4.1107003	CCCGGCCAACGACCCA

mmu-miR-582-3p	5.29E-03	0.017519612	1	0.290331	1.0502508	0.1731728	2.1214745	0.0970939	1.7353721	0.4044018	AACCTGTTGAACAACGTGAAACCC
mmu-miR-342-3p	5.30E-03	0.017519612	1	0.1179536	0.9414646	0.0667833	0.8569387	0.2121842	1.5077798	0.1103269	TCTCACAGAAAATCGCACCGT
mmu-miR-496a-3p	5.40E-03	0.017755774	1	0.4757403	2.0560373	0.5377885	1.0840162	0.4024521	3.6266225	0.7492749	AGTATTACATGCCAATCTCC
mmu-miR-409-5p	5.61E-03	0.018344963	1	0.3589175	2.4981159	0.2016965	0.7784754	0.2953417	1.6410004	0.3760255	AGGTTACCCGAGCAACTTGCAT
mmu-miR-194-5p	5.67E-03	0.018413039	1	0.2302868	1.386207	0.2040148	2.9052787	0.8708733	1.2588619	0.3962527	TGTAACAGCAAACCCATGTGGA
mmu-miR-6538	5.70E-03	0.018413039	1	0.2999702	1.9440028	1.1436939	7.041941	0.7296326	1.6147006	1.2666139	CGCGGGCTCCGGGGC
bta-miR-378	5.72E-03	0.018413039	1	0.1181157	2.224835	0.4944649	1.491496	0.2974698	1.4148635	0.2538082	CCTGGACTTGGAGTCAGAAGGCT
mmu-miR-17-5p	5.92E-03	0.018934664	1	0.3303307	1.225979	0.0878724	2.4548547	0.4177761	2.1884502	0.6963248	CAAAGTGCTTACAGTGCAGGTAG
mmu-miR-542-5p	6.09E-03	0.019328376	1	0.2671163	0.9438729	0.2503843	2.0347155	0.7322074	3.6774023	1.8229044	CTCGGGGATCATCATGTACGA
mmu-miR-144-3p	6.11E-03	0.019328376	1	1.015959	15.691404	5.7180296	14.157613	5.8400912	9.5186871	7.2860747	TACAGTATAAGATGATGTAC
mmu-miR-5126	6.17E-03	0.019437069	1	0.537093	1.3774477	1.2077884	5.6867813	2.6253156	11.344691	6.8201904	GCGCGGGCGGGGCGGGGG
mmu-miR-450a-5p	6.53E-03	0.020443501	1	0.1633147	1.3582107	0.3034625	1.7743298	0.5885427	2.4361341	0.2770026	TTTGGCGATGTGTTCTAATAT
chi-miR-326-5p	6.73E-03	0.020875338	1	0.135016	0.8610848	0.3711388	3.5847342	1.1754849	2.472339	1.33005	CCTCATCTGTCGTGTTGGCT
mmu-miR-145a-5p	6.74E-03	0.020875338	1	0.3201111	0.5908406	0.1324421	0.4066126	0.0381804	0.4020431	0.0867146	GTCCAGTTTCCCAGGAATCCCT
mmu-miR-26a-2-3p	7.08E-03	0.021675393	1	0.281269	0.8414142	0.1863195	0.4389714	0.0697446	0.6806672	0.0903391	CCTGTTCTTGATTACTGTTTC
mmu-miR-467c-5p	7.13E-03	0.021675393	1	1.7295242	14.536997	8.7645475	54.71416	17.898221	54.421037	16.03314	TAAGTGCCTGCATGTATGTG
mmu-miR-378d	7.14E-03	0.021675393	1	0.1008627	1.5725206	0.670377	1.0099429	0.1594194	0.5648632	0.0993916	ACTGGACTTGGAGTCAGAAGGT
mmu-miR-425-5p	7.23E-03	0.021826036	1	0.0602518	0.8879042	0.0848391	1.3755272	0.3227571	1.4868228	0.1988631	AATGACACGATCACTCCGTTG
oan-miR-1386	7.28E-03	0.021866895	1	0.3165177	0.6010565	0.1744502	1.1495189	0.1175332	1.4544761	0.1191957	TCGGCTCCTGGCTGGCTGCCA
mmu-miR-126a-3p	7.49E-03	0.022391448	1	0.0215421	0.9507635	0.149753	0.7171073	0.0887186	1.0394132	0.0551625	TCGTACCGTGAGTAATAATGC
mmu-miR-143-3p	7.68E-03	0.022860619	1	0.1535765	1.365799	0.0735043	0.9270947	0.122811	0.7927775	0.1444706	TGAGATGAAGCACTGTAGCT
mmu-miR-23a-5p	7.95E-03	0.023527499	1	0.3219745	2.5127578	0.5520956	4.1096851	1.8863033	3.4214251	1.0696368	GGGGTTCTGGGGATGGGATT
mmu-miR-652-3p	8.19E-03	0.024134283	1	0.2428944	0.5902822	0.0065492	1.1529782	0.2409497	0.7848517	0.0659197	AATGGGCCACTAGGGTTGTGA
mmu-miR-410-3p	8.43E-03	0.024518561	1	0.5015998	1.6942188	0.814943	0.6646572	0.1790523	2.7903132	0.6988702	AATATAACACAGATGGCCTGT
sha-miR-24	8.44E-03	0.024518561	1	0.4842972	0.4684974	0.145966	0.2489231	0.0897534	0.2578385	0.0529081	ATCTGGCTCAGTTCAGCAGGAAC
mmu-miR-182-5p	8.45E-03	0.024518561	1	0.2686847	0.7024421	0.2078117	0.4429616	0.1025267	0.3596214	0.0924438	TTTGGCAATGGTAGAACTCACACC
mmu-miR-191-3p	8.51E-03	0.024573195	1	0.4289188	1.4298961	0.2257558	2.2769702	0.5749977	2.576579	0.4763858	GCTGCACTGGATTCTGTT
mmu-miR-222-5p	8.73E-03	0.02508042	1	0.4709523	2.1596529	0.3088697	2.8272898	0.5811644	1.4341556	0.302107	GGCTCAGTAGCCAGTGTAGATC
mmu-miR-1948-3p	9.03E-03	0.02581485	1	0.3910427	0.4915103	0.2384145	0.6470618	0.2539182	2.4898556	0.8791734	TTTGGCAGAGCACTCGTACAG
mmu-miR-8112	9.22E-03	0.026240084	1	0.1599005	0.5739454	0.1055694	0.9834943	0.1051947	0.8473933	0.148724	TCTCCGCCACCTCCACCGCAGC
hsa-miR-1261	9.63E-03	0.027284151	1	0.3119386	2.0406684	1.2278387	4.9569668	0.3904987	2.7144925	1.080898	TGGATAAGGCATTGGC
mmu-miR-434-3p	9.89E-03	0.027871135	1	0.2642263	2.5101291	1.2272849	0.9134584	0.1174769	1.4230546	0.1173335	TTTGAACCACACTCGACTCCT

notes:

adj P: Benjamini-Hochberg adjusted P value.

Supplementary Table S5: 18-concurrent wound healing associated microRNA signature ^a

miRNA	tissue	p-value	adj P ^b	0 hr		6 hr		24 hr		5 day	
				fold	stdev	fold	stdev	fold	stdev	fold	stdev
mmu-miR-148b-3p	skin	1.21E-05	0.000596	1.00	0.12	1.44	0.18	3.64	0.79	2.83	0.22
	oral	3.56E-03	0.132005	1.00	0.1	1.3	0.4	2.42	0.97	3.25	0.63
mmu-miR-31-5p	skin	5.45E-05	0.001096	1.00	0.32	0.67	0.42	0.84	0.39	11.7	1.8
	oral	6.65E-03	0.151662	1.00	0.07	0.8	0.22	0.95	0.08	1.49	0.12
mmu-miR-23b-3p	skin	4.69E-03	0.016241	1.00	0.18	0.71	0.09	0.64	0.11	0.5	0.07
	oral	7.19E-03	0.151662	1.00	0.11	0.74	0.05	1.03	0.12	0.99	0.07
mmu-miR-200c-3p	skin	1.79E-03	0.008742	1.00	0.21	0.58	0.19	1.29	0.2	1.64	0.17
	oral	9.24E-03	0.163277	1.00	0.16	0.87	0.21	1.77	0.58	1.88	0.19
mmu-miR-195a-5p	skin	5.56E-05	0.001096	1.00	0.15	1.09	0.14	0.61	0.04	0.43	0.05
	oral	9.79E-03	0.163277	1.00	0.11	0.91	0.06	0.88	0.15	0.66	0.05
mmu-miR-3068-5p	skin	4.23E-05	0.001042	1.00	0.17	0.63	0.05	2.02	0.53	2.54	0.38
	oral	2.27E-04	0.075263	1.00	0.26	1.07	0.39	3.53	0.81	3.69	0.14
mmu-miR-30a-3p	skin	1.03E-04	0.001644	1.00	0.18	1.26	0.16	0.67	0.11	0.38	0.06
	oral	6.92E-04	0.086715	1.00	0.09	1.05	0.06	0.78	0.02	0.83	0.03
mmu-miR-17-5p	skin	5.92E-03	0.018935	1.00	0.33	1.23	0.09	2.46	0.42	2.19	0.7
	oral	7.15E-04	0.086715	1.00	0.16	0.85	0.05	1.51	0.16	0.99	0.05
mmu-miR-676-5p	skin	4.75E-03	0.016241	1.00	0.14	0.78	0.11	0.6	0.05	0.65	0.08
	oral	1.52E-03	0.107584	1.00	0.06	1.25	0.13	1.53	0.16	1.57	0.18
mmu-miR-362-5p	skin	4.65E-08	0.000014	1.00	0.08	0.64	0.09	4.28	0.43	5.58	0.87
	oral	2.07E-03	0.107584	1.00	0.4	0.84	0.4	3.27	1.37	4.29	0.88
mmu-miR-434-3p	skin	9.89E-03	0.027871	1.00	0.26	2.51	1.23	0.91	0.12	1.42	0.12
	oral	2.33E-03	0.111088	1.00	0.09	1.32	0.08	0.8	0.14	0.99	0.06
mmu-miR-20a-5p	skin	3.18E-03	0.012459	1.00	0.28	1.33	0.22	2.7	0.47	2.31	0.68
	oral	2.80E-03	0.123245	1.00	0.14	1.11	0.06	1.72	0.35	1.45	0.02
mmu-miR-425-3p	skin	8.08E-05	0.001402	1.00	0.24	0.61	0.18	2.12	0.1	2.23	0.22
	oral	3.55E-03	0.132006	1.00	0.15	0.8	0.34	1.59	0.01	1.91	0.13
mmu-miR-340-5p	skin	1.90E-03	0.008993	1.00	0.45	2.61	1.1	6.93	2.12	3.42	1.09
	oral	3.71E-03	0.132006	1.00	0.09	2.05	0.49	1.28	0.24	1.1	0.16
mmu-miR-664-3p	skin	2.86E-03	0.011692	1.00	0.41	1.32	0.1	0.69	0.12	0.39	0.06
	oral	4.08E-03	0.132006	1.00	0.08	0.93	0.03	0.77	0.15	0.64	0.05
mmu-miR-7a-5p	skin	5.29E-05	0.001096	1.00	0.3	1.87	0.13	5.41	0.81	5.67	1.74
	oral	4.95E-03	0.134730	1.00	0.12	1.4	0.34	2.04	0.38	1.53	0.12
mmu-miR-193b-3p	skin	2.63E-05	0.000844	1.00	0.08	0.74	0.05	0.53	0.11	0.24	0.05
	oral	5.61E-03	0.142362	1.00	0.11	0.58	0.06	0.62	0.1	0.59	0.11
mmu-miR-223-3p	skin	7.01E-05	0.001296	1.00	0.19	3.47	1.05	21.9	7.09	5.34	2.8
	oral	7.42E-03	0.151663	1.00	0.08	2.66	0.86	3.72	2.46	1.07	0.17

^aCut-off p-value of 0.01 was used. 18 microRNAs that were concurrently differentially expressed in both skin and oral mucosa wound healing were listed.

^bThe Benjamini-Hochberg adjusted P-values were computed for multiple hypothesis testing.

Supplementary Table S6: Expression changes of the top 10 baseline tissue specific microRNAs in skin and oral mucosa during wound healing

miR	miR-378a-3p			
time course	0hr	6hr	24hr	5day
skin (fold)	1	1.469336	0.939103	0.8457584
skin (stdev)	0.012098	0.518862	0.138471	0.087231
p value				5.26E-02
oral (fold)	0.396452	0.290532	0.378565	0.4084093
oral (stdev)	0.026353	0.013767	0.020421	0.0358472
p value				8.18E-04

miR	miR-31-5p			
time course	0hr	6hr	24hr	5day
skin (fold)	1	0.669133	0.840812	11.688945
skin (stdev)	0.320248	0.422529	0.38847	1.7979082
p value				5.45E-05
oral (fold)	22.85125	18.31651	21.60025	33.972947
oral (stdev)	1.692528	5.048494	1.765137	2.6370834
p value				6.65E-03

miR	miR-10a-5p			
time course	0hr	6hr	24hr	5day
skin (fold)	1	0.908414	0.502462	0.2383207
skin (stdev)	0.131761	0.359267	0.154867	0.0880976
p value				2.23E-03
oral (fold)	0.00482	0.006927	0.00965	0.0074661
oral (stdev)	0.002255	0.004842	0.005031	0.0046497
p value				5.64E-01

miR	miR-10b-5p			
time course	0hr	6hr	24hr	5day
skin (fold)	1	0.922428	0.302635	0.2060538
skin (stdev)	0.15904	0.40317	0.091135	0.0682445
p value				1.17E-03
oral (fold)	0.163878	0.150159	0.147039	0.1517026
oral (stdev)	0.0479	0.048789	0.039186	0.027659
p value				9.67E-01

miR	miR-133a-3p			
time course	0hr	6hr	24hr	5day
skin (fold)	1	2.0078	0.69721	0.1621058
skin (stdev)	0.565692	1.050772	0.31428	0.0153855

p value	1 . 11E-03			
oral (fold)	0.07094 0.009774 0.014111 0.014214			
oral (stdev)	0.030147 0.012295 0.012636 0.0068159			
p value	6 . 78E-02			

miR	miR-146b-5p			
time course	0hr	6hr	24hr	5day
skin (fold)	1	1.35334	1.779497	1.273712
skin (stdev)	0.098454	0.216424	0.375257	0.2865286
p value	2 . 65E-02			
oral (fold)	0.161831	0.219098	0.364548	0.2856465
oral (stdev)	0.04531	0.07541	0.148361	0.1047957
p value	1 . 06E-01			

miR	miR-126a-5p			
time course	0hr	6hr	24hr	5day
skin (fold)	1	1.53879	0.936007	1.5421504
skin (stdev)	0.142881	0.277857	0.32267	0.2875119
p value	4 . 75E-02			
oral (fold)	1.864485	2.169465	1.86865	1.3459816
oral (stdev)	0.286606	0.512187	0.720687	0.2903343
p value	2 . 11E-01			

miR	miR-34c-5p			
time course	0hr	6hr	24hr	5day
skin (fold)	1	1.890395	3.441682	35.217775
skin (stdev)	0.354481	0.313367	0.59346	29.988102
p value	1 . 23E-04			
oral (fold)	83.6016	156.4728	64.88194	46.964317
oral (stdev)	69.09045	154.963	77.05825	36.915668
p value	6 . 81E-01			

miR	miR-30a-5p			
time course	0hr	6hr	24hr	5day
skin (fold)	1	1.251449	0.582879	0.453159
skin (stdev)	0.103088	0.111781	0.022601	0.0342868
p value	9 . 89E-07			
oral (fold)	0.611179	0.775242	0.580714	0.5359318
oral (stdev)	0.022063	0.210435	0.080197	0.032217
p value	9 . 47E-02			

miR	miR-96-5p			
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time course	0hr	6hr	24hr	5day
skin (fold)	1	1.059792	0.343601	0.2286068
skin (stdev)	0.14515	0.384041	0.045746	0.049049
p value				1.60E-04
oral (fold)	0.577801	0.720975	0.692673	0.3836004
oral (stdev)	0.076565	0.080358	0.360337	0.0445133
p value				1.40E-01

Supplementary Table S7: Statistical analysis on TaqMan assay-based differential expression of miR-21 and miR-10l

A: Validation of differential expression in tissue samples

A.1: miR-21 in tissue samples

A.1.a: 2way ANOVA - miR-21 in tissue samples

Two-way ANOVA		Ordinary				
Alpha		SS	DF	MS	F (DFn, DFd)	P value
Interaction		51.13	3	17.04	F (3, 76) = 5.132	P = 0.0028
Row Factor (time)		156.7	3	52.22	F (3, 76) = 15.72	P < 0.0001
Column Factor (tissue type)		27.26	1	27.26	F (1, 76) = 8.209	P = 0.0054
Residual		252.4	76	3.321		

A.1.b: Multiple t tests - miR-21 in tissue samples

		Mean (skin)	Mean (oral)	(statistical significance determined by Holm-Sidak method)			
				P value	significance	Difference	SE of diff ratio df
0 hr	Row 1	1	1.03525	0.880149	-0.0352453	0.23117	0.1525 23
6 hr	Row 2	1.10854	1.03525	0.784571	0.0732943	0.26471	0.2769 21
24 hr	Row 3	1.79576	1.09139	0.0493723 *	0.704372	0.33124	2.1265 16
5 day	Row 4	6.47986	2.49812	0.0415756 *	3.98174	1.79721	2.2155 16

A.1.c: Shapiro-Wilk tests - miR-21 in tissue samples

		skin	oral		
		P value	Null	P value	Null
0 hr	Row 1	0.427686	Accept	0.436846	Accept
6 hr	Row 2	0.090062	Accept	0.000623	Reject
24 hr	Row 3	0.569615	Accept	0.371813	Accept
5 day	Row 4	0.015668	Reject	0.145458	Accept

A.2: miR-10b in tissue samples

A.2.a: 2way ANOVA - miR-10b in tissue samples

Two-way ANOVA		Ordinary				
Alpha		SS	DF	MS	F (DFn, DFd)	P value
Interaction		2.886	3	0.962	F (3, 72) = 13.15	P < 0.0001
Row Factor (time)		2.887	3	0.9625	F (3, 72) = 13.16	P < 0.0001
Column Factor (tissue type)		7.154	1	7.154	F (1, 72) = 97.81	P < 0.0001
Residual		5.266	72	0.07314		

A.2.b: Multiple t tests - miR-10b in tissue samples

		Mean (skin)	Mean (oral)	(statistical significance determined by Holm-Sidak method)			
				P value	significance	Difference	SE of diff ratio df
0 hr	Row 1	1	0.061636	0.000666664 *	0.938364	0.21134	4.4401 13
6 hr	Row 2	1.07812	0.047931	2.59E-07 *	1.03019	0.13262	7.7682 19
24 hr	Row 3	0.296832	0.0577083	0.03538784 *	0.239124	0.07249	3.2986 20
5 day	Row 4	0.268708	0.0485352	0.0162982 *	0.220173	0.08394	2.6229 20

A.2.c: Shapiro-Wilk tests - miR-10b in tissue samples

		skin	oral		
		P value	Null	P value	Null
0 hr	Row 1	0.23429	Accept	0.000416	Reject
6 hr	Row 2	0.988214	Accept	0.000035	Reject
24 hr	Row 3	0.008166	Reject	0.001878	Reject
5 day	Row 4	0.000341	Reject	0.004668	Reject

B: Validation of differential expression in LCM samples

B.1: miR-21 in LCM samples

B.1.a: 2way ANOVA - miR-21 in LCM samples

Two-way ANOVA		Ordinary				
Alpha		SS	DF	MS	F (DFn, DFd)	P value
Interaction		23.54	3	7.846	F (3, 24) = 7.114	P = 0.0014
Row Factor (time)		44.02	3	14.67	F (3, 24) = 13.30	P < 0.0001
Column Factor (tissue type)		23.56	1	23.56	F (1, 24) = 21.36	P = 0.0001
Residual		26.47	24	1.103		

B.1.b: Multiple t tests - miR-21 in LCM samples

		Mean (skin)	Mean (oral)	(statistical significance determined by Holm-Sidak method)			
				P value	significance	Difference	SE of diff ratio df
0 hr	Row 1	1.1444	0.235591	0.0153373 *	0.908811	0.27094	3.3545 6
6 hr	Row 2	0.887953	0.131738	0.0163974 *	0.756214	0.22912	3.3005 6
24 hr	Row 3	1.65043	1.12877	0.530282	0.521658	0.78349	0.6658 6
5 day	Row 4	5.80185	1.12458	0.00833249 *	4.67727	1.21076	3.8631 6

B.1.c: Shapiro-Wilk tests - miR-21 in LCM samples

		skin	oral		
		P value	Null	P value	Null
0 hr	Row 1	0.947874	Accept	0.025577	Reject
6 hr	Row 2	0.75682	Accept	0.083015	Accept
24 hr	Row 3	0.147838	Accept	0.018483	Reject
5 day	Row 4	0.897928	Accept	0.344031	Accept

B.2: miR-10b in LCM samples

B.2.a: 2way ANOVA - miR-10b in LCM samples

Two-way ANOVA		Ordinary				
Alpha		SS	DF	MS	F (DFn, DFd)	P value
Interaction		1.859	3	0.6195	F (3, 24) = 7.070	P = 0.0014
Row Factor (time)		1.836	3	0.612	F (3, 24) = 6.984	P = 0.0015
Column Factor (tissue type)		4.11	1	4.11	F (1, 24) = 46.90	P < 0.0001
Residual		2.103	24	0.08763		

B.2.b: Multiple t tests - miR-10b in LCM samples

		Mean (skin)	Mean (oral)	(statistical significance determined by Holm-Sidak method)			
				P value	significance	Difference	SE of diff ratio df
0 hr	Row 1	1.17113	0.0627785	0.00229105 *	1.10835	0.21867	5.0686 6
6 hr	Row 2	1.3093	0.028406	0.00742183 *	1.28089	0.32317	3.9635 6
24 hr	Row 3	0.310278	0.04828	0.0565111	0.261998	0.11116	2.3571 6
5 day	Row 4	0.25989	0.044054	0.0814784	0.215836	0.10322	2.0911 6

B.2.c: Shapiro-Wilk tests - miR-10b in LCM samples

		skin	oral		
		P value	Null	P value	Null
0 hr	Row 1	0.223427	Accept	0.469074	Accept
6 hr	Row 2	0.680263	Accept	0.412278	Accept
24 hr	Row 3	0.275825	Accept	0.22837	Accept
5 day	Row 4	0.048727	Reject	0.002357	Reject

Supplementary Table S8: Statistical analysis on closure of the wounds treated with miR-21 and miR-10b reagents

A: wounds treated with control and miR-21 mimic

A.1: 2way ANOVA - wounds treated with control and miR-21 mimic

Two-way ANOVA	Ordinary			
Alpha	0.05			
ANOVA table	SS	DF	MS	F (DFn, DFd)
Interaction	335.2	9	37.24	F (9, 100) = 1.581 P = 0.1312
Row Factor (time)	540038	9	6004	F (9, 100) = 254.8 P < 0.0001
Column Factor (treatment)	1216	1	1216	F (1, 100) = 51.59 P < 0.0001
Residual	2356	100	23.56	

A.2: Multiple t tests - wounds treated with control and miR-21 mimic

	Mean (ctrl)	Mean (miR-21)	P value	(statistical significance determined by Holm-Sidak method)			
				significance	Difference	SE of different ratio	df
day 1	Row 1	28.5867	32.735	0.311891	-4.14833	3.89494	1.06506 10
day 2	Row 2	47.0717	51.395	0.210477	-4.32333	3.23082	1.33815 10
day 3	Row 3	54.3733	59.775	0.261995	-5.40167	4.54393	1.18876 10
day 4	Row 4	61.3967	72.4817	0.00110855 *	-11.085	2.4525	4.51988 10
day 5	Row 5	68.465	79.685	0.00036866 *	-11.22	2.13354	5.25886 10
day 6	Row 6	75.3533	84.1083	0.0126509 *	-8.755	2.88828	3.03122 10
day 7	Row 7	82.2717	90.9317	0.00292527 *	-8.66	2.21627	3.90747 10
day 8	Row 8	88.1333	94.6467	0.0391902 *	-6.51333	2.74674	2.3713 10
day 9	Row 9	95.9417	98.8817	0.00687099 *	-2.94	0.866929	3.39128 10
day 10	Row 10	98.97	99.5783	0.163996	-0.608334	0.405013	1.50201 10

A.3: Shapiro-Wilk tests - wounds treated with control and miR-21 mimic

	ctrl	miR-21	
		P value	Null
day 1	Row 1	0.880938 Accept	0.015552 Reject
day 2	Row 2	0.048305 Reject	0.354856 Accept
day 3	Row 3	0.965486 Accept	0.774883 Accept
day 4	Row 4	0.676812 Accept	0.591661 Accept
day 5	Row 5	0.691616 Accept	0.832817 Accept
day 6	Row 6	0.340818 Accept	0.486321 Accept
day 7	Row 7	0.065263 Accept	0.810627 Accept
day 8	Row 8	0.482778 Accept	0.409187 Accept
day 9	Row 9	0.127901 Accept	0.279401 Accept
day 10	Row 10	0.129791 Accept	0.180812 Accept

B: wounds treated with control and miR-10b LNA

B.1: 2way ANOVA - wounds treated with control and miR-10b LNA

Two-way ANOVA	Ordinary			
Alpha	0.05			
ANOVA table	SS	DF	MS	F (DFn, DFd)
Interaction	529.1	9	58.79	F (9, 87) = 0.7714 P = 0.6430
Row Factor (time)	84927	9	9436	F (9, 87) = 123.8 P < 0.0001
Column Factor (treatment)	1251	1	1251	F (1, 87) = 16.42 P = 0.0001
Residual	6630	87	76.21	

B.2: Multiple t tests - wounds treated with control and miR-10b LNA

	Mean (ctrl)	Mean (miR-10b)	P value	(statistical significance determined by Holm-Sidak method)			
				significance	Difference	SE of different ratio	df
day 1	Row 1	11.2085	23.95	0.151825	-12.7414	8.13706	1.56585 9
day 2	Row 2	24.4597	33.2662	0.245699	-8.80655	7.09181	1.24179 9
day 3	Row 3	36.2235	48.8293	0.100135	-12.6058	6.87992	1.83227 9
day 4	Row 4	47.6599	52.779	0.434157	-5.11904	6.25346	0.818593 9
day 5	Row 5	54.4978	60.861	0.283508	-6.36319	5.57918	1.14052 9
day 6	Row 6	63.9945	70.4322	0.279583	-6.43774	5.41722	1.18838 6
day 7	Row 7	74.8959	87.2205	0.00296294 *	-12.3246	3.0565	4.03225 9
day 8	Row 8	93.4787	97.2499	0.296344	-3.77115	3.4017	1.10861 9
day 9	Row 9	97.7562	98.1437	0.77031	-0.387475	1.28763	0.300922 9
day 10	Row 10	99.302	100	0.00186593	-0.69804	0.160682	4.34422 9

B.3: Shapiro-Wilk tests - wounds treated with control and miR-10b LNA

	ctrl	miR-10b	
		P value	Null
day 1	Row 1	0.079956 Accept	0.829051 Accept
day 2	Row 2	0.230435 Accept	0.450255 Accept
day 3	Row 3	0.096018 Accept	0.758435 Accept
day 4	Row 4	0.081085 Accept	0.355892 Accept
day 5	Row 5	0.240936 Accept	0.131831 Accept
day 6	Row 6	0.495443 Accept	0.904362 Accept
day 7	Row 7	0.659012 Accept	0.248763 Accept
day 8	Row 8	0.016694 Reject	0.050619 Accept
day 9	Row 9	0.907885 Accept	0.017018 Reject
day 10	Row 10	0.565286 Accept	NaN Reject