

<i>miRNA</i>	<i>N</i>	<i>Measure of pixel intensity</i>	<i>Time window (hours post-hatch)</i>	<i>Slope R²</i>	<i>Mean R²</i>	<i>Joint R²</i>	<i>Joint 95% CI</i>
<i>let-7</i>	283	99th percentile	72–233	0.002	0.007	0.013	[-0.016, 0.040]
<i>lin-4</i>	197	Mean over 99th percentile	72–212	0.162	0.014	0.165 (+)	[0.071, 0.26]
<i>mir-1</i>	72	Maximum (head)	72–192	0.086	0.002	0.091	[-0.035, 0.19]
<i>mir-47</i>	408	Maximum	72–201	0.257	0.104	0.287 (+)	[0.22, 0.35]
<i>mir-51</i>	83	Maximum	72–236	0.002	0.015	0.015	[-0.067, 0.11]
<i>mir-59</i>	104	95th percentile (vulva)	72–150	0.044	0.031	0.067	[-0.037, 0.14]
<i>mir-60</i>	218	99th percentile	72–210	0.117	0.101	0.174 (+)	[0.061, 0.27]
<i>mir-63</i>	138	99th percentile	72–231	0.121	0.025	0.138	[0.036, 0.24]
<i>mir-79</i>	86	Maximum	72–120	0.035	0.002	0.038	[-0.043, 0.11]
<i>mir-84</i>	142	99th percentile	72–231	0.006	0.059	0.059	[-0.04, 0.15]
<i>mir-85</i>	184	99th percentile	72–172	0.177	0.041	0.177 (+)	[0.10, 0.26]
<i>mir-90</i>	173	95th percentile	72–221	0.101	0.142	0.172 (+)	[0.076, 0.29]
<i>mir-228</i>	151	Mean over 99th percentile	72–170	0.122	0.019	0.125	[0.039, 0.22]
<i>mir-240-786</i>	160	95th percentile	72–228	0.178	0.187	0.232 (-)	[0.13, 0.33]
<i>mir-241</i>	97	99th percentile (head)	72–120	0.023	0.013	0.039	[-0.035, 0.10]
<i>mir-242</i>	98	Maximum	72–204	0.004	0.031	0.037	[-0.040, 0.11]
<i>mir-243</i>	343	Mean	72–176	0.305	0.091	0.326 (+)	[0.25, 0.40]
<i>mir-246</i>	193	95th percentile	72–200	0.120	0.024	0.128	[0.053, 0.20]
<i>mir-360</i>	87	Maximum	72–191	0.018	0.018	0.027	[-0.044, 0.093]
<i>mir-788</i>	187	Maximum	72–96	0.010	0.003	0.010	[-0.023, 0.039]
<i>mir-793</i>	177	Maximum	72–237	0.158	0.046	0.169 (+)	[0.079, 0.26]
<i>mir-794</i>	91	Maximum	72–96	0.001	0.055	0.058	[-0.045, 0.15]

Table S1: Fixed-time-window correlation of *PmiRNA::GFP* reporter expression with lifespan. A joint regression of slope and average expression against lifespan was performed using a window starting at 72 hours post-hatch and ending at the 90% survival timepoint. R² values exceeding 0.15 are indicated by **bolded** text. The 95% confidence interval (CI) for each joint correlation coefficient is also shown. The direction of correlation, derived from the individual regression on slope and mean expression (which we observed to always correlate in the same direction), is indicated by (+) or (-), respectively.

<i>miRNA</i>	<i>N</i>	<i>Measure of pixel intensity</i>	<i>Time window (days post-hatch)</i>	<i>Slope R²</i>	<i>Mean R²</i>	<i>Joint R²</i>	<i>Joint 95% CI</i>
<i>let-7</i>	307	99th percentile	6.0–7.5	0.014	0.024	0.036	[-0.007, 0.081]
<i>lin-4</i>	209	Mean over 99th percentile	4.5–8.0	0.200	0.021	0.203 (+)	[0.114, 0.293]
<i>mir-1</i>	79	Maximum (head)	6.0–7.5	0.102	0.025	0.117	[0.009, 0.237]
<i>mir-47</i>	433	Maximum	5.0–8.0	0.125	0.165	0.290 (+)	[0.219, 0.363]
<i>mir-51</i>	87	Maximum	8.0–8.5	0.025	0.048	0.098	[-0.016, 0.210]
<i>mir-59</i>	104	95th percentile (vulva)	4.5–6.5	0.002	0.092	0.092	[-0.017, 0.187]
<i>mir-60</i>	227	99th percentile	7.5–8.0	0.017	0.177	0.189 (+)	[0.073, 0.302]
<i>mir-63</i>	147	99th percentile	3.0–9.0	0.109	0.019	0.124	[0.033, 0.219]
<i>mir-79</i>	86	Maximum	3.5–4.0	0.005	0.010	0.025	[-0.033, 0.076]
<i>mir-84</i>	157	99th percentile	3.0–7.5	0.024	0.023	0.091	[-0.005, 0.172]
<i>mir-85</i>	194	99th percentile	4.0–6.5	0.115	0.048	0.146	[0.065, 0.228]
<i>mir-90</i>	181	95th percentile	8.0–8.5	0.014	0.215	0.229 (+)	[0.120, 0.347]
<i>mir-228</i>	164	Mean over 99th percentile	5.5–6.5	0.059	0.059	0.110	[0.036, 0.193]
<i>mir-240-786</i>	172	95th percentile	7.5–8.5	0.043	0.268	0.291 (-)	[0.167, 0.415]
<i>mir-241</i>	97	99th percentile (head)	3.0–3.5	0.013	0.037	0.051	[-0.040, 0.142]
<i>mir-242</i>	107	Maximum	4.5–5.5	0.096	0.012	0.096	[-0.008, 0.192]
<i>mir-243</i>	360	Mean	4.5–7.0	0.203	0.160	0.325 (+)	[0.255, 0.399]
<i>mir-246</i>	205	95th percentile	7.0–7.5	0.025	0.134	0.181 (+)	[0.087, 0.277]
<i>mir-360</i>	87	Maximum	4.0–5.0	0.061	0.009	0.071	[-0.025, 0.160]
<i>mir-788</i>	187	Maximum	3.0–3.5	0.000	0.005	0.006	[-0.029, 0.037]
<i>mir-793</i>	186	Maximum	5.5–9.5	0.120	0.070	0.161 (+)	[0.074, 0.243]
<i>mir-794</i>	91	Maximum	3.0–3.5	0.009	0.082	0.093	[-0.032, 0.209]

Table S2: Correlation of *PmiRNA::GFP* reporters with lifespan using a sliding window and 95% survival endpoint. A joint regression of slope and average expression against lifespan was performed using a sliding time window (minimum width of 12 hours) beginning at 3 days post-hatch and ending at the 95% survival timepoint. The optimal time window and highest correlation achieved is reported. R^2 values exceeding 0.15 are indicated by **bolded** text. The 95% confidence interval (CI) for each joint correlation coefficient is also shown. The direction of correlation, derived from the individual regression on slope and mean expression (which we observed to always correlate in the same direction), is indicated by (+) or (-), respectively.

<i>Reporter</i>	<i>N</i>	<i>Measure of pixel intensity</i>	<i>Time window (hours post-hatch)</i>	<i>Slope R²</i>	<i>Mean R²</i>	<i>Joint R²</i>	<i>Joint 95% CI</i>
<i>cpna-2</i>	65	99th percentile	168–192 (S)	0.112	0.006	0.123	[0.007, 0.242]
<i>cpna-2</i>	64	99th percentile	72–196 (F)	0.007	0.017	0.018	[-0.081, 0.093]
<i>myo-2</i>	171	Maximum	108–156 (S)	0.107	0.005	0.112	[0.024, 0.191]
<i>myo-2</i>	163	Maximum	72–168 (F)	0.069	0.009	0.071	[0.004, 0.144]

Table S3: Correlation with lifespan of promoter::GFP reporters for two non-miRNA promoters. A joint regression on slope and average expression against lifespan was performed using a sliding 12-hour window beginning at 3 days post-hatch (indicated by ‘S’). The optimal time window and highest correlation achieved is reported as the joint regression R². Results from using a fixed window from 3 days post-hatch to the 90% survival timepoint (indicated by ‘F’) are also shown.

<i>miRNA</i>	<i>N</i>	<i>Measure of pixel intensity</i>	<i>Time window (hours post-hatch)</i>	<i>Maximal expression R²</i>	<i>p-value</i>	<i>Day of max expression R²</i>	<i>p-value</i>
<i>let-7</i>	283	99th percentile	72–233	0.000	0.989	0.001	0.681
<i>lin-4</i>	197	Mean over 99th percentile	72–212	0.000	0.853	0.001	0.638
<i>mir-1</i>	72	Maximum (head)	72–192	0.012	0.307	0.010	0.353
<i>mir-47</i>	408	Maximum	72–201	0.000	0.779	0.020	0.004
<i>mir-51</i>	83	Maximum	72–236	0.030	0.116	0.001	0.798
<i>mir-59</i>	104	95th percentile (vulva)	72–150	0.000	0.912	0.002	0.670
<i>mir-60</i>	218	99th percentile	72–210	0.003	0.425	0.025	0.019
<i>mir-63</i>	138	99th percentile	72–231	0.002	0.640	0.000	0.956
<i>mir-79</i>	86	Maximum	72–120	0.020	0.190	0.008	0.402
<i>mir-84</i>	142	99th percentile	72–231	0.005	0.397	0.005	0.388
<i>mir-85</i>	184	99th percentile	72–172	0.010	0.169	0.070	0.000
<i>mir-90</i>	173	95th percentile	72–221	0.013	0.138	0.028	0.028
<i>mir-228</i>	151	Mean over 99th percentile	72–170	0.022	0.071	0.027	0.044
<i>mir-240-786</i>	160	95th percentile	72–228	0.122	0.000	0.030	0.028
<i>mir-241</i>	97	99th percentile (head)	72–120	0.009	0.347	0.000	0.905
<i>mir-242</i>	98	Maximum	72–204	0.002	0.669	0.000	0.878
<i>mir-243</i>	343	Mean	72–176	0.001	0.534	0.028	0.002
<i>mir-246</i>	193	95th percentile	72–200	0.011	0.142	0.042	0.004
<i>mir-360</i>	87	Maximum	72–191	0.000	0.916	0.007	0.438
<i>mir-788</i>	187	Maximum	72–96	0.006	0.299	0.001	0.682
<i>mir-793</i>	177	Maximum	72–237	0.004	0.400	0.002	0.517
<i>mir-794</i>	91	Maximum	72–96	0.050	0.032	0.000	0.855

Table S4: Correlation coefficients and p-values for regression of lifespan against the whole-life maximum value of *PmiRNA::GFP* expression (measured as noted), or the day that that maximum value was achieved.

<i>miRNA</i>	<i>N</i>	<i>Time window (hours post-hatch)</i>	<i>GFP slope R²</i>	<i>GFP mean R²</i>	<i>AF slope R²</i>	<i>AF mean R²</i>	<i>Summed R²</i>	<i>Joint R²</i>
<i>lin-4</i>	197	72–212	0.162	0.014	0.140	0.037	0.353	0.302
<i>mir-47</i>	408	72–201	0.257	0.104	0.062	0.030	0.453	0.313
<i>mir-60</i>	218	72–210	0.117	0.101	0.145	0.059	0.421	0.241
<i>mir-90</i>	184	72–172	0.101	0.142	0.076	0.084	0.403	0.304
<i>mir-228</i>	173	72–221	0.122	0.019	0.090	0.100	0.331	0.204
<i>mir-240- 786</i>	160	72–228	0.178	0.187	0.009	0.031	0.405	0.237
<i>mir-243</i>	343	72–176	0.305	0.091	0.216	0.060	0.672	0.408
<i>mir-246</i>	193	72–200	0.120	0.024	0.101	0.058	0.303	0.191
<i>mir-793</i>	177	72–237	0.158	0.046	0.094	0.043	0.341	0.258

Table S5: Correlation coefficients for *PmiRNA::GFP* (slope and average), autofluorescence (slope and average of 95th percentile intensity), and joint *PmiRNA::GFP* and autofluorescence regression against lifespan.

<i>Experiment</i>	<i>N</i>	<i>Measure of pixel intensity</i>	<i>Time window (hours post-hatch)</i>	<i>Slope R²</i>	<i>Mean R²</i>	<i>Joint R²</i>	<i>Joint 95% CI</i>
<i>let-7_20170526</i>	101	99th percentile	72–213	0.004	0.001	0.004	[-0.029, 0.047]
<i>let-7_20170517</i>	100	99th percentile	72–208	0.042	0.001	0.045	[-0.048, 0.127]
<i>let-7_20170216</i>	48	99th percentile	72–217	0.005	0.047	0.320	[0.148, 0.520]
<i>let-7_20170208</i>	31	99th percentile	72–182	0.078	0.401	0.401	[0.180, 0.691]
<i>lin-4_20171222</i>	99	Mean over 99th percentile	72–209	0.075	0.013	0.124	[-0.024, 0.275]
<i>lin-4_20171219</i>	97	Mean over 99th percentile	72–214	0.152	0.004	0.152	[-0.034, 0.133]
<i>mir-1_20170518</i>	72	Maximum (head)	72–192	0.084	0.002	0.086	[-0.026, 0.182]
<i>mir-47_20190213</i>	74	Maximum	72–185	0.041	0.205	0.261	[0.137, 0.416]
<i>mir-47_20181101</i>	79	Maximum	72–196	0.246	0.203	0.404	[0.293, 0.547]
<i>mir-47_20180511</i>	89	Maximum	72–196	0.208	0.082	0.234	[0.072, 0.416]
<i>mir-47_20170728</i>	96	Maximum	72–201	0.196	0.127	0.419	[0.294, 0.562]
<i>mir-47_20170504</i>	71	Maximum	72–213	0.219	0.000	0.226	[0.066, 0.394]
<i>mir-51_20180530</i>	83	Maximum	72–236	0.004	0.016	0.016	[-0.068, 0.110]
<i>mir-59_20170811</i>	104	95th percentile (vulva)	72–144	0.026	0.026	0.064	[-0.031, 0.137]
<i>mir-60_20180917</i>	109	99th percentile	72–204	0.064	0.101	0.145	[-0.007, 0.277]
<i>mir-60_20180518</i>	111	99th percentile	72–208	0.184	0.114	0.249	[0.135, 0.386]
<i>mir-63_20161215</i>	62	99th percentile	72–245	0.061	0.004	0.068	[-0.077, 0.199]
<i>mir-63_20161202</i>	75	99th percentile	72–229	0.019	0.029	0.249	[0.088, 0.429]
<i>mir-79_20171027</i>	86	Maximum	72–120	0.034	0.001	0.034	[-0.045, 0.097]
<i>mir-84_20181018</i>	77	99th percentile	72–214	0.015	0.012	0.015	[-0.077, 0.096]
<i>mir-84_20180928</i>	65	99th percentile	72–236	0.045	0.073	0.115	[-0.028, 0.307]
<i>mir-85_20180914</i>	87	99th percentile	72–207	0.138	0.040	0.145	[0.025, 0.260]
<i>mir-85_20180525</i>	97	99th percentile	72–181	0.214	0.066	0.220	[0.083, 0.362]
<i>mir-90_20190430</i>	89	95th percentile	72–212	0.027	0.151	0.177	[0.045, 0.333]
<i>mir-90_20180928</i>	80	95th percentile	72–218	0.087	0.070	0.114	[-0.031, 0.248]
<i>mir-228_20180914</i>	73	Mean over 99th percentile	72–218	0.125	0.008	0.139	[-0.001, 0.288]

<i>mir-228_20180504</i>	77	Mean over 99th percentile	72–176	0.223	0.048	0.223	[0.046, 0.393]
<i>mir-240-786_20181004</i>	100	95th percentile	72–228	0.219	0.242	0.258	[0.127, 0.399]
<i>mir-240-786_20181128</i>	60	95th percentile	72–212	0.234	0.160	0.237	[0.065, 0.435]
<i>mir-241_20170809</i>	97	99th percentile (head)	72–120	0.027	0.013	0.039	[-0.048, 0.103]
<i>mir-242_20181105</i>	98	Maximum	72–204	0.012	0.034	0.036	[-0.047, 0.103]
<i>mir-243_20190122</i>	90	Mean	72–184	0.453	0.084	0.479	[0.341, 0.631]
<i>mir-243_20181107</i>	109	Mean	72–200	0.193	0.068	0.255	[0.116, 0.388]
<i>mir-243_20180613</i>	60	Mean	72–166	0.342	0.001	0.363	[0.210, 0.524]
<i>mir-243_20171128</i>	82	Mean	72–181	0.244	0.286	0.331	[0.160, 0.554]
<i>mir-246_20180612</i>	100	95th percentile	72–184	0.214	0.101	0.223	[0.100, 0.366]
<i>mir-246_20171116</i>	93	95th percentile	72–209	0.091	0.039	0.104	[0.007, 0.201]
<i>mir-360_20180418</i>	87	Maximum	72–191	0.018	0.018	0.027	[-0.051, 0.103]
<i>mir-788_20180427</i>	100	Maximum	72–96	0.018	0.005	0.018	[-0.035, 0.072]
<i>mir-788_20171027</i>	87	Maximum	72–96	0.030	0.011	0.041	[-0.048, 0.123]
<i>mir-793_20190606</i>	79	Maximum	72–235	0.084	0.026	0.119	[0.005, 0.235]
<i>mir-793_20181121</i>	97	Maximum	72–238	0.205	0.063	0.219	[0.058, 0.375]
<i>mir-794_20190320</i>	91	Maximum	72–96	0.005	0.057	0.059	[-0.044, 0.153]

Table S6: Regression results from individual biological replicates using raw data. A joint regression on slope and average expression against lifespan was performed using a window starting at 72 hours post-hatch and ending at the 90% survival timepoint.