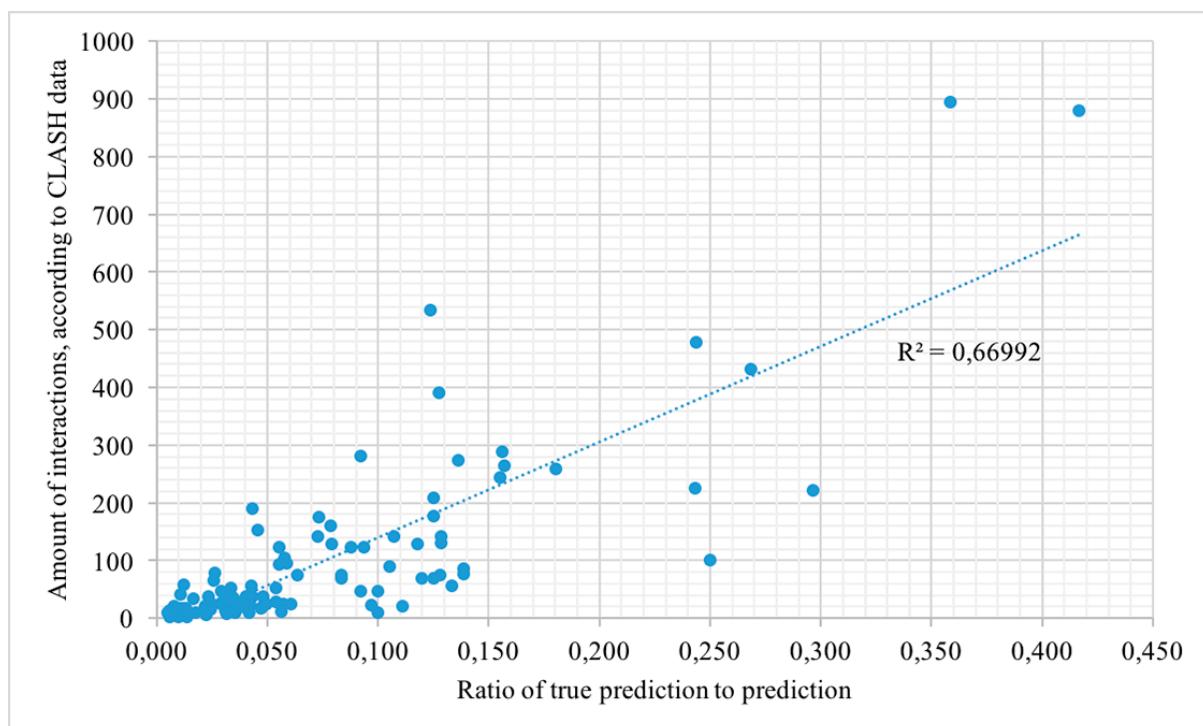


## Supplementary material

**Supplementary Figure S1. The correlation between the number of interactions (CLASH data) and between the ratios of true predictions to all predictions for each of the 102 miRNAs.** All five miRNA prediction programmes made 624 correctly predicted miRNA-mRNA interactions, which were formed by 102 miRNAs. For these 102 miRNAs, we defined the ratio of correctly predicted to all predicted pairs and marked it on the X-axis. On the Y-axis we marked the number of miRNA-mRNA interactions formed by the miRNA according to CLASH data.



**Supplementary Table S1. Characteristics of the 102 miRNAs that formed correctly predicted miRNA-mRNA interactions.** The number of their interactions with mRNAs according to CLASH data, predictions, true predictions and their prediction quality indexes.

miRNA_name	Number_of_interactions_CLASH_data	Number_of_predictions	Number_of_true_predictions	ratio=true_predictions/predictions	Not_predicted_interactions	true_predictions/not_predicted_interactions	ratio=(true_predictions/predictions)/not_predicted_interactions	ratio=H/smallest ratio
miR-363-3p	2	72	1	1%	1	100%	0,0139	63,13
miR-340-3p	9	10	1	10%	8	13%	0,0125	56,82
miR-130a-3p	2	104	1	1%	1	100%	0,0096	43,71
miR-29c-3p	11	89	5	6%	6	83%	0,0094	42,56
miR-130b-3p	7	95	3	3%	4	75%	0,0079	35,89
miR-132-3p	9	48	2	4%	7	29%	0,0060	27,06
miR-330-3p	20	9	1	11%	19	5%	0,0058	26,58
miR-28-5p	23	31	3	10%	20	15%	0,0048	21,99
miR-210-3p	9	28	1	4%	8	13%	0,0045	20,29
miR-200c-3p	6	45	1	2%	5	20%	0,0044	20,20
miR-301a-3p	18	85	4	5%	14	29%	0,0034	15,28
miR-424-5p	25	181	9	5%	16	56%	0,0031	14,13
miR-218-5p	19	63	3	5%	16	19%	0,0030	13,53
miR-20b-5p	3	171	1	1%	2	50%	0,0029	13,29
miR-29a-3p	24	70	4	6%	20	20%	0,0029	12,99
miR-18b-5p	13	64	2	3%	11	18%	0,0028	12,91
miR-365a-3p	24	33	2	6%	22	9%	0,0028	12,52
miR-708-5p	13	32	1	3%	12	8%	0,0026	11,84
miR-361-5p	56	30	4	13%	52	8%	0,0026	11,66
miR-99a-5p	101	4	1	25%	100	1%	0,0025	11,36
miR-22-3p	19	50	2	4%	17	12%	0,0024	10,70
miR-454-3p	28	93	5	5%	23	22%	0,0023	10,63
miR-31-5p	46	65	6	9%	40	15%	0,0023	10,49
let-7g-5p	15	164	4	2%	11	36%	0,0022	10,08
miR-138-5p	9	57	1	2%	8	13%	0,0022	9,97
miR-17-3p	47	10	1	10%	46	2%	0,0022	9,88
miR-148b-3p	23	71	3	4%	20	15%	0,0021	9,60
miR-185-5p	69	92	11	12%	58	19%	0,0021	9,37
miR-19a-3p	7	83	1	1%	6	17%	0,0020	9,13
miR-96-5p	21	83	3	4%	18	17%	0,0020	9,13
miR-339-5p	70	56	7	13%	63	11%	0,0020	9,02
miR-19b-3p	13	92	2	2%	11	18%	0,0020	8,98
miR-505-3p	76	36	5	14%	71	7%	0,0020	8,89

miR-342-3p	74	39	5	13%	69	7%	0,0019	8,45
miR-29b-3p	25	104	4	4%	21	19%	0,0018	8,33
miR-18a-5p	86	72	10	14%	76	13%	0,0018	8,31
miR-27a-3p	23	116	4	3%	19	21%	0,0018	8,25
miR-15a-5p	38	188	9	5%	29	31%	0,0017	7,50
let-7i-5p	9	185	2	1%	7	29%	0,0015	7,02
miR-193a-3p	15	47	1	2%	14	7%	0,0015	6,91
miR-27b-3p	69	120	10	8%	59	17%	0,0014	6,42
miR-324-5p	222	27	8	30%	214	4%	0,0014	6,29
miR-182-5p	38	69	3	4%	35	9%	0,0012	5,65
miR-346	25	34	1	3%	24	4%	0,0012	5,57
miR-128-3p	37	99	4	4%	33	12%	0,0012	5,57
miR-378a-3p	90	38	4	11%	86	5%	0,0012	5,56
let-7f-5p	53	148	8	5%	45	18%	0,0012	5,46
miR-330-5p	26	34	1	3%	25	4%	0,0012	5,35
miR-125a-5p	75	48	4	8%	71	6%	0,0012	5,34
miR-10a-5p	226	37	9	24%	217	4%	0,0011	5,10
miR-196b-5p	41	71	3	4%	38	8%	0,0011	5,05
miR-15b-5p	141	187	24	13%	117	21%	0,0011	4,99
miR-7-5p	21	46	1	2%	20	5%	0,0011	4,94
miR-107	26	122	3	2%	23	13%	0,0011	4,86
miR-26a-5p	130	70	9	13%	121	7%	0,0011	4,83
miR-30e-5p	35	58	2	3%	33	6%	0,0010	4,75
miR-221-3p	128	68	8	12%	120	7%	0,0010	4,46
miR-181b-5p	74	110	7	6%	67	10%	0,0009	4,32
miR-106a-5p	40	158	5	3%	35	14%	0,0009	4,11
miR-20a-5p	141	177	19	11%	122	16%	0,0009	4,00
miR-34a-5p	56	117	5	4%	51	10%	0,0008	3,81
miR-183-5p	124	64	6	9%	118	5%	0,0008	3,61
let-7d-5p	18	159	2	1%	16	13%	0,0008	3,57
miR-17-5p	259	161	29	18%	230	13%	0,0008	3,56
miR-148a-3p	18	77	1	1%	17	6%	0,0008	3,47
miR-23b-3p	124	91	8	9%	116	7%	0,0008	3,44
miR-30c-5p	177	64	8	13%	169	5%	0,0007	3,36
miR-204-5p	17	88	1	1%	16	6%	0,0007	3,23
miR-24-3p	17	92	1	1%	16	6%	0,0007	3,09
miR-222-3p	244	71	11	15%	233	5%	0,0007	3,02
miR-181a-5p	37	86	2	2%	35	6%	0,0007	3,02
miR-16-5p	265	172	27	16%	238	11%	0,0007	3,00
miR-30b-5p	53	60	2	3%	51	4%	0,0007	2,97
miR-30d-5p	47	69	2	3%	45	4%	0,0006	2,93
miR-196a-5p	129	76	6	8%	123	5%	0,0006	2,92

miR-193b-3p	432	41	11	27%	421	3%	0,0006	2,90
miR-497-5p	9	197	1	1%	8	13%	0,0006	2,88
miR-125b-5p	209	88	11	13%	198	6%	0,0006	2,87
miR-10b-5p	96	34	2	6%	94	2%	0,0006	2,84
miR-197-3p	93	36	2	6%	91	2%	0,0006	2,78
let-7a-5p	289	160	25	16%	264	9%	0,0006	2,69
miR-421	104	52	3	6%	101	3%	0,0006	2,60
let-7e-5p	274	176	24	14%	250	10%	0,0005	2,48
miR-320a	478	123	30	24%	448	7%	0,0005	2,47
miR-25-3p	142	96	7	7%	135	5%	0,0005	2,46
miR-93-5p	160	165	13	8%	147	9%	0,0005	2,44
miR-30a-5p	33	61	1	2%	32	3%	0,0005	2,33
miR-195-5p	13	170	1	1%	12	8%	0,0005	2,23
miR-106b-5p	123	181	10	6%	113	9%	0,0005	2,22
miR-615-3p	879	12	5	42%	874	1%	0,0005	2,17
miR-296-3p	176	41	3	7%	173	2%	0,0004	1,92
miR-92a-3p	895	92	33	36%	862	4%	0,0004	1,89
miR-320c	20	128	1	1%	19	5%	0,0004	1,87
miR-320b	66	116	3	3%	63	5%	0,0004	1,87
miR-328-3p	78	76	2	3%	76	3%	0,0003	1,57
miR-186-5p	391	110	14	13%	377	4%	0,0003	1,53
miR-149-5p	281	76	7	9%	274	3%	0,0003	1,53
let-7c-5p	152	175	8	5%	144	6%	0,0003	1,44
miR-326	42	94	1	1%	41	2%	0,0003	1,18
let-7b-5p	534	178	22	12%	512	4%	0,0002	1,10
miR-92b-3p	190	69	3	4%	187	2%	0,0002	1,06
miR-23a-3p	58	81	1	1%	57	2%	0,0002	0,98

**Supplementary Table S2. Types of interactions for 13'618 pairs from CLASH data formed by 73 miRNAs, which coupled 50 or more mRNAs.** For each miRNA-mRNA interaction from 13'618 pairs, we identify the number (percentage) of each type and number (percentage) summarized for canonical types and non-canonical types.

Type of a miRNA-mRNA interaction									
	6- mer	7- mer	8- mer	9- mer	Total canonical types	noncano- nical_se- ed	no- ne	nose- ed_9- nt_st- em	Total non canonical types
<b>Maximum</b>	17%	28%	22%	8%	62%	64%	71% %	48%	100%
<b>Average</b>	6%	9%	6%	2%	24%	38%	22% %	17%	76%
<b>Minimum</b>	0%	0%	0%	0%	0%	11%	2% %	0%	38%

**Supplementary Table S3. The number of correctly predicted pairs according to randomly generated mRNA-miRNA interaction datasets (average of 5 datasets) and CLASH data.**

Program	The average number of miRNA-mRNA interactions approved by five random patterns	The number of miRNA-mRNA interactions approved by CLASH data
<b>TargetScan</b>	20.8 (3.6)*	274
<b>PicTar</b>	33 (4.6)*	308
<b>RNA22</b>	10.8 (2)*	145
<b>PITA</b>	13.6 (2.4)*	199
<b>Miranda</b>	35.2 (6.4)*	420
<b>Total programmes</b>	65.8 (10)*	624

**Supplementary Table S4. Characteristics of the prediction programmes according to the CLASH data and the 5 randomly generated patterns (average).**

<b>Characteristics</b>	<b>Comparing with the CLASH data</b>	<b>Average of comparing with the 5 random patterns</b>
<b>Number of True positive interactions</b>	624	65.8
<b>Number of «experimental» interactions</b>	16'190	16'190
<b>Number of predicted interactions</b>	19'398	19'398
<b>Number of False positive interactions</b>	18'774	19'332.2
<b>Number of False Negative interactions</b>	15'566	16'124.2
<b>Sensitivity</b>	0.039	0.0041
<b>Positive predictive value</b>	0.032	0.0034