Table S2 – Prediction algorithms and databases available in miRSystem

Source	Version ^a	Original Records	Mappable Number ^b
DIANA-microT	10	1,450,653	1,413,583
miRanda	11	927,867	910,940
$mirBridge^{c}$	10	41,382	41,382
PicTar ^d	9.1	69,642	59,479
PITA	11	209,579	206,474
rna22	8	296,518	219,395
TargetScan ^e	16	110,284	110,284
TarBase and miRecords ^f	6-16	1,661	1,575

^a The version mapped to miRBase.

^b Number of mappable records in miRBase version 17.

^c The records of miRBridge were collected from the supplementary data.

^d The information about PicTar was obtained from UCSC Genome Browser.

^e Families IDs were converted to single miRNA IDs to get complete miRNA-gene matching data.

^f Merged together as experimentally validated datasets.