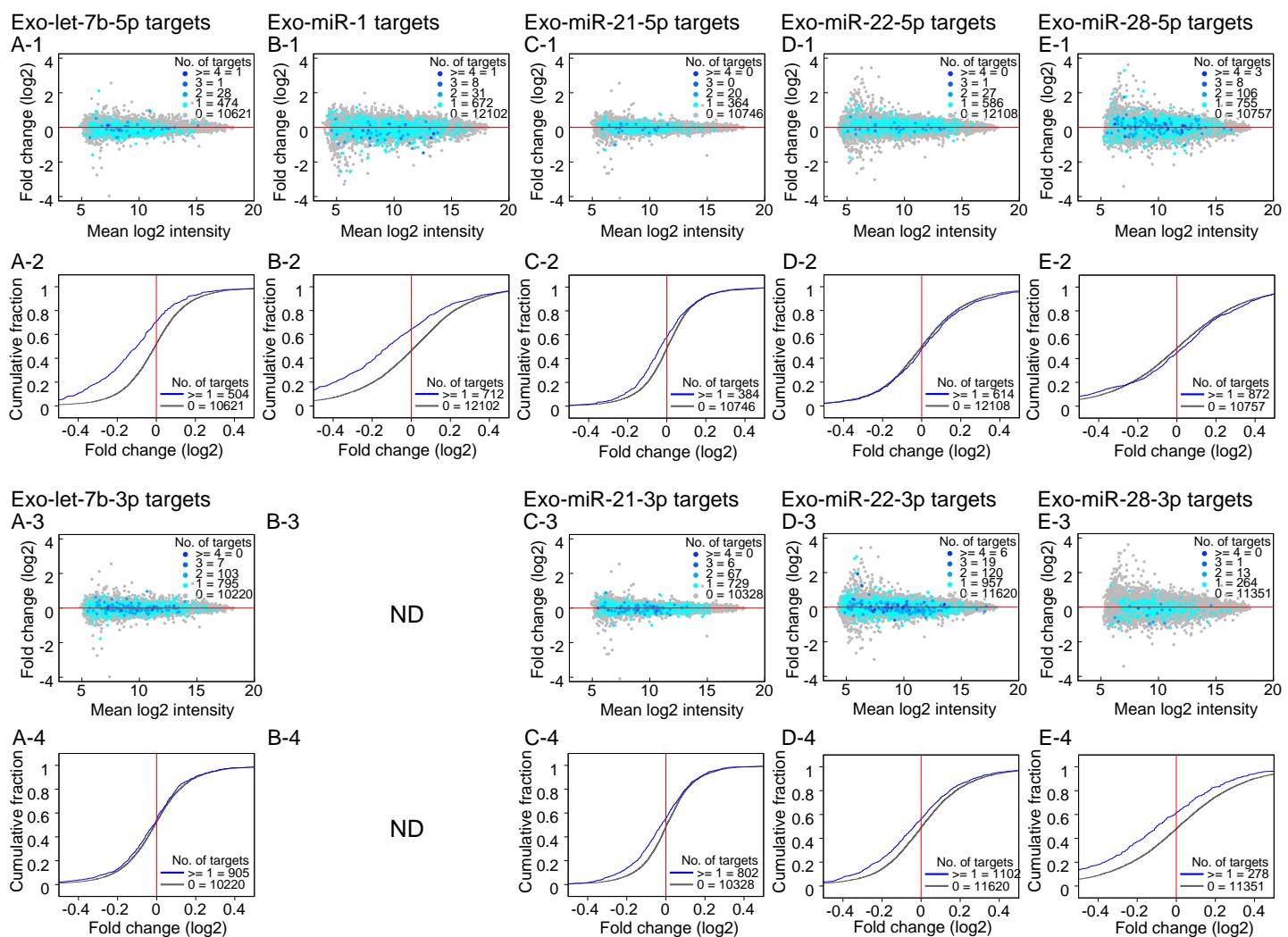
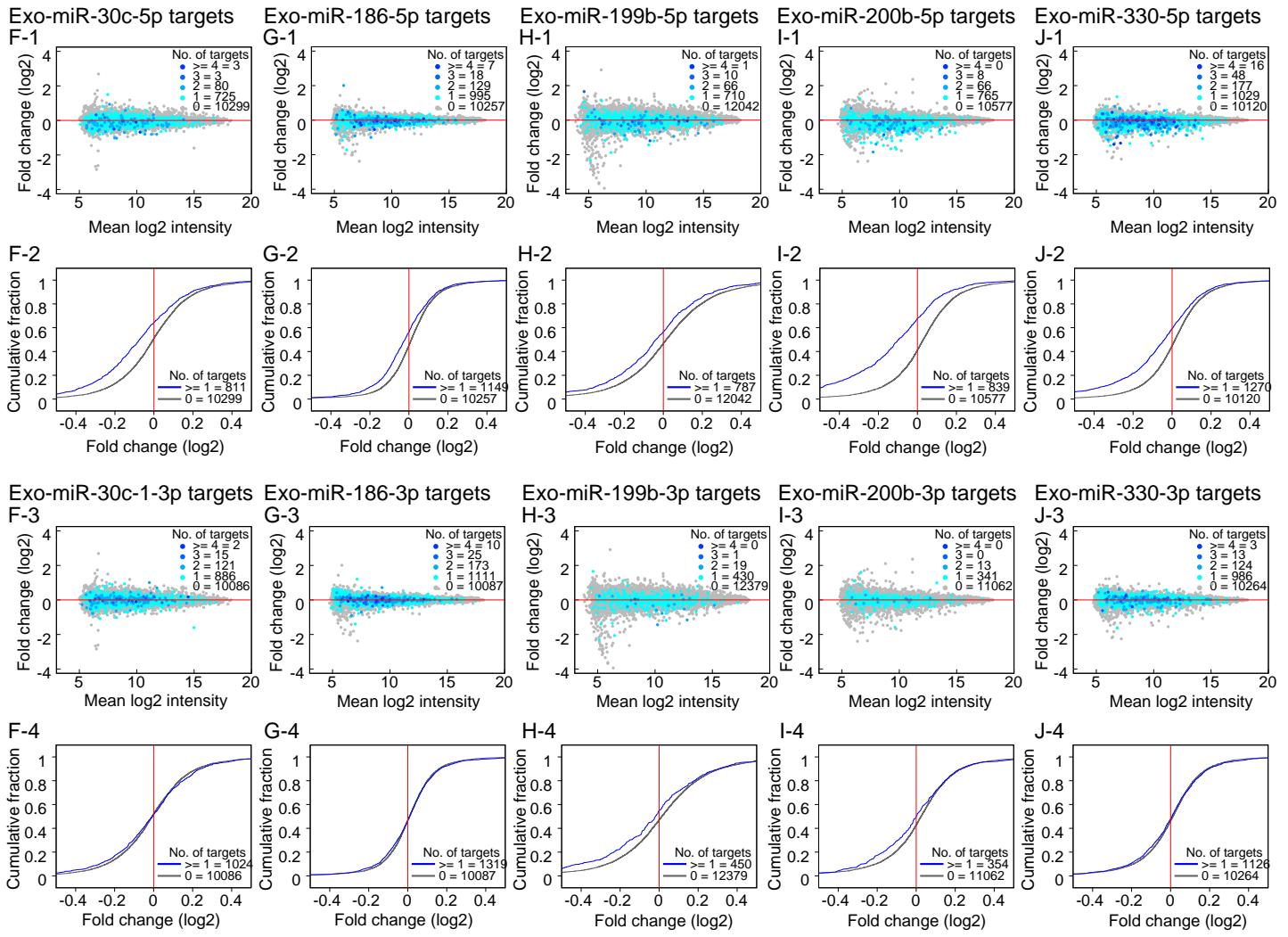


Supplementary Information

Figure S1. Microarray analysis of the expression of target genes of the transfected exo-miRNAs HeLa cells transfected with each of the fifteen exo-miRNA duplexes were subjected to microarray profiling. Data are presented as MA plots (A1-O1, A3-O3) and cumulative distributions (A2-O2, A4-O4) of seed-matched target genes of let-7b-5p (A-1 and A-2) and let-7b-3p (A-3 and A-4), miR-1 (B-1 and B-2), miR-21-5p (C-1 and C-2) and miR-21-3p (C-3 and C-4), miR-22-5p (D-1 and D-2) and miR-22-3p (D-3 and D-4), miR-28-5p (E-1 and E-2) and miR-28-3p (E-3 and E-4), miR-30c-5p (F-1 and F-2) and miR-30c-3p (F-3 and F-4), miR-186-5p (G-1 and G-2) and miR-186-3p (G-3 and G-4), miR-199b-5p (H-1 and H-2) and miR-199b-3p (H-3 and H-4), miR-200b-5p (I-1 and I-2) and miR-200b-3p (I-3 and I-4), miR-330-5p (J-1 and J-2) and miR-330-3p (J-3 and J-4), miR-335-5p (K-1 and K-2) and miR-335-3p (K-3 and K-4), miR-346 (L-1 and L-2), miR-446 (M-1 and M-2), miR-574-5p (N-1 and N-2) and miR-574-3p (N-3 and N-4), and miR-3126-5p (O-1 and O-2) and miR-3126-3p (O-3 and O-4). In MA plots, the vertical axis indicates the change in gene expression (\log_2 fold change relative to mock-transfected cells), and the horizontal axis indicates the mean signal intensity for the probes for each gene before and after miRNA transfection. The numbers of target sites in the 3' UTR are indicated by dots of different colors: navy blue, ≥ 4 seed-matched target sites in a single 3' UTR; dark blue, 3 target sites; blue, 2 sites; light blue, 1 site; gray, no seed-matched sites. (A-2~O-2, A-4~O-4). In cumulative distributions, blue and black lines indicate the cumulative fraction of seed-matched target transcripts (≥ 1 seed-matched target sites) and all transcripts, respectively. ND; Not determined.





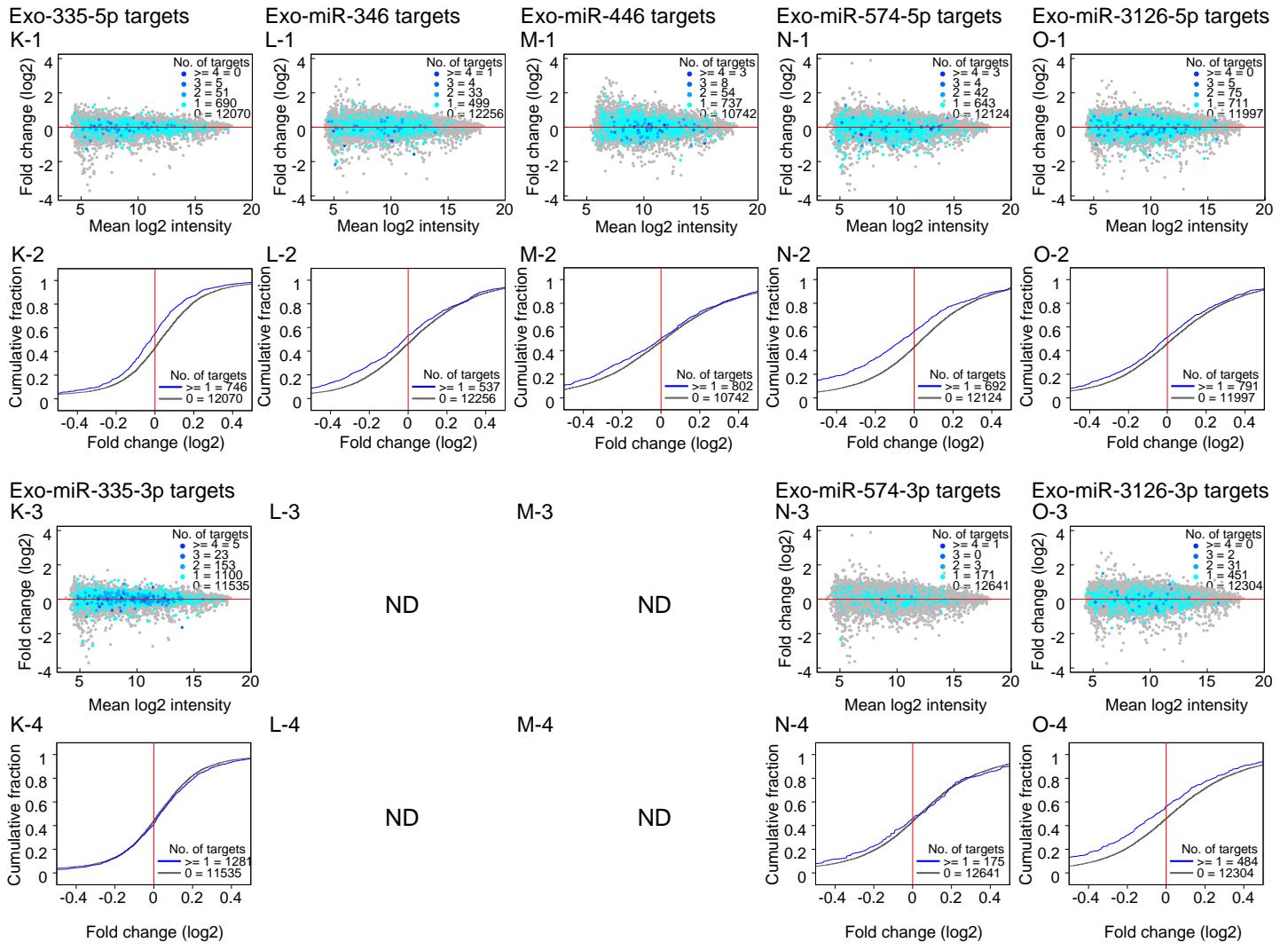


Figure S2. Microarray analysis of the expression of target genes of the endo-miRNA, let-7b-5p
 Microarray analysis of endo-let-7b-5p target gene expression in cells transfected with exo-let-7b (A), exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G), exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446 (M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative distributions (A-2~O-2) of seed-matched target genes of let-7b-5p.

Endo-let-7b-5p targets

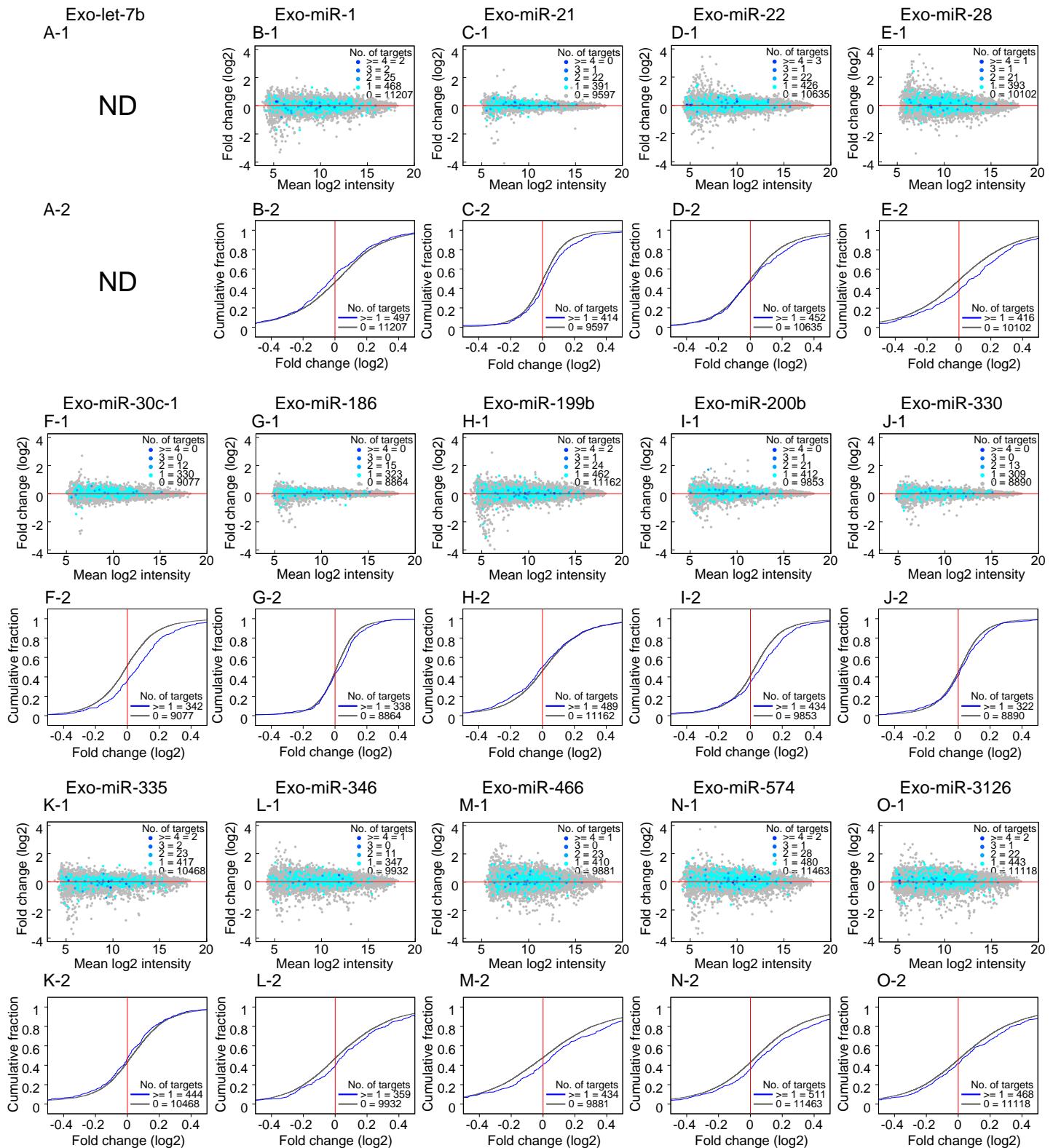


Figure S3. Microarray analysis of the expression of target genes of the endo-miRNA, miR-21-5p
 Microarray analysis of endo-miR-21-5p target gene expression in cells transfected with exo-let-7b (A),
 exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
 exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
 (M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
 distributions (A-2~O-2) of seed-matched target genes of miR-21-5p.

Endo-miR-21-5p targets

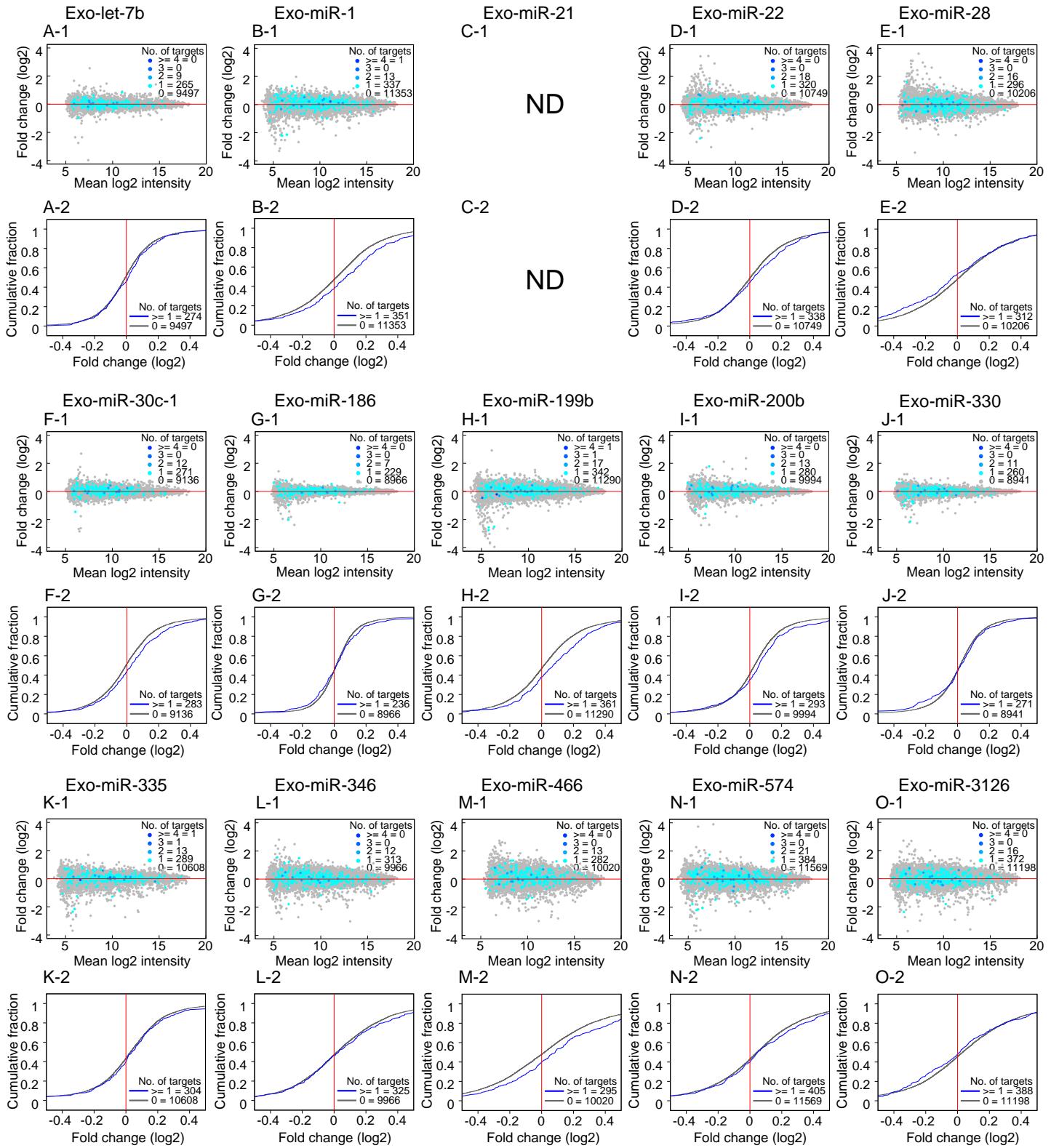


Figure S4. Microarray analysis of the expression of target genes of the endo-miRNA, miR-27a-3p
 Microarray analysis of endo-miR-27a-3p target gene expression in cells transfected with exo-let-7b (A),
 exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
 exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
 (M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
 distributions (A-2~O-2) of seed-matched target genes of miR-27a-3p.

Endo-miR-27a-3p targets

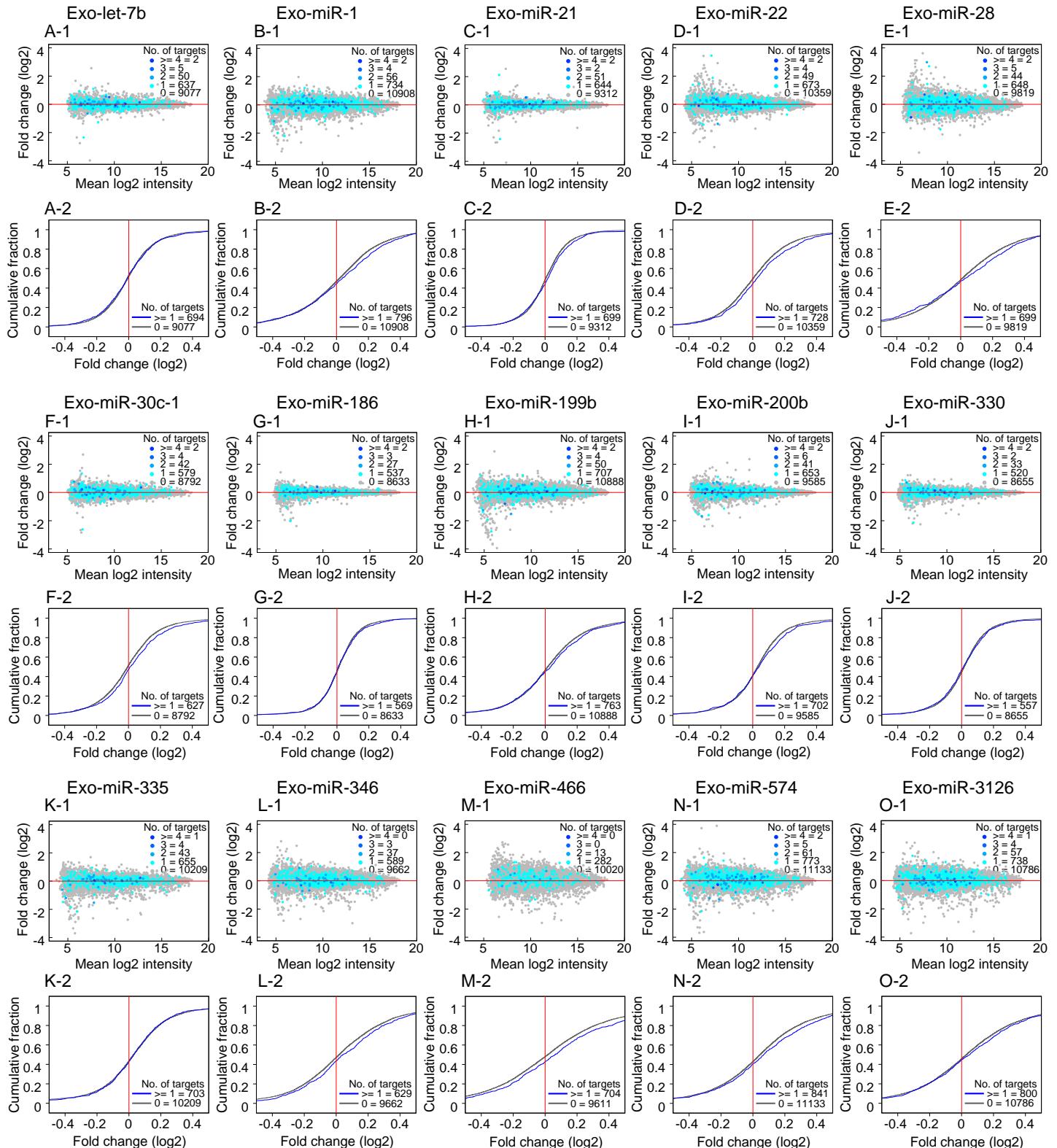


Figure S5. Microarray analysis of the expression of target genes of the endo-miRNA, miR-17-5p
Microarray analysis of endo-miR-17-5p target gene expression in cells transfected with exo-let-7b (A),
exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
(M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
distributions (A-2~O-2) of seed-matched target genes of miR-17-5p.

Endo-miR-17-5p targets

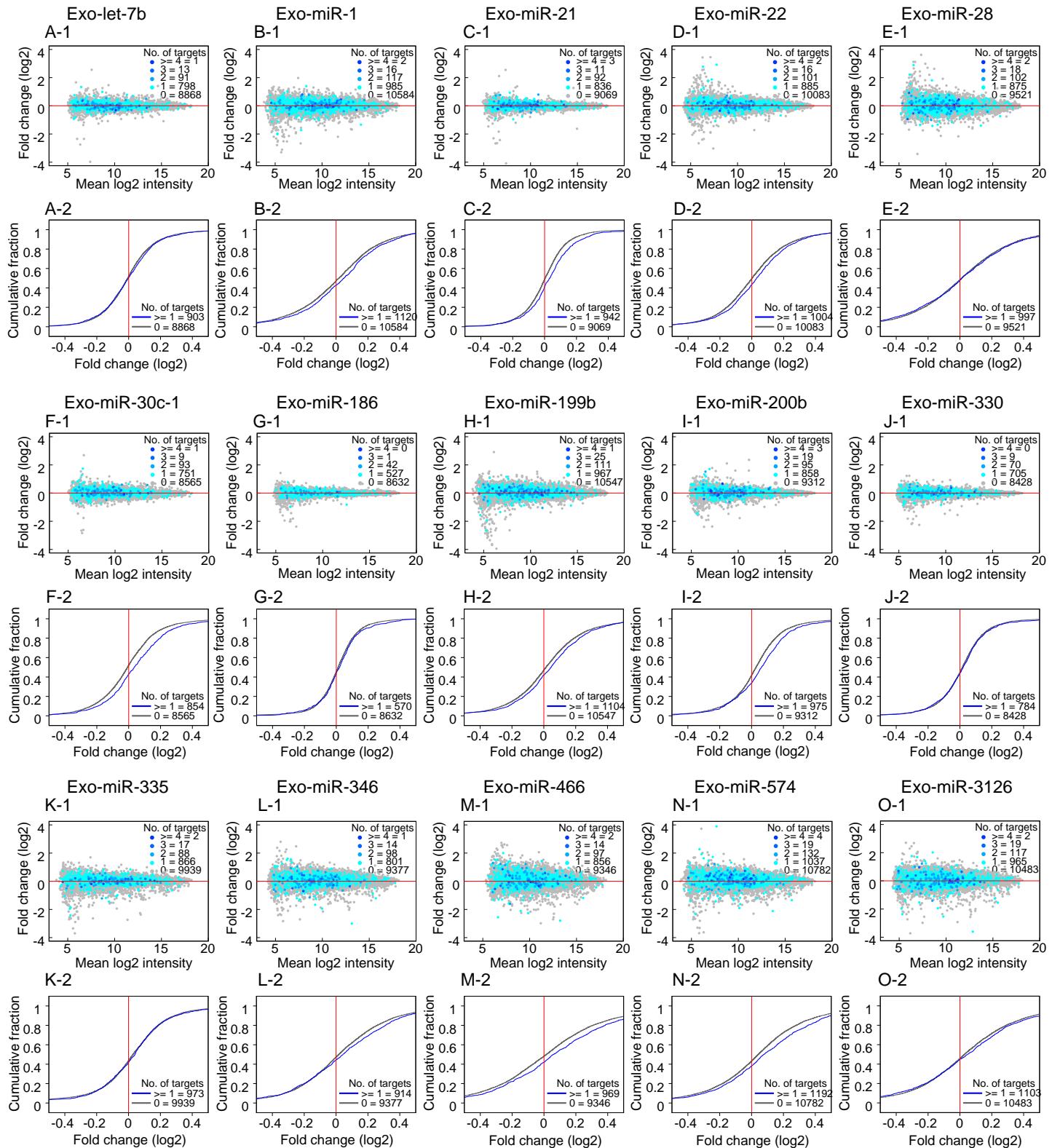


Figure S6. Microarray analysis of the expression of target genes of the endo-miRNA, miR-26a-5p
Microarray analysis of endo-miR-26a-5p target gene expression in cells transfected with exo-let-7b (A),
exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
(M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
distributions (A-2~O-2) of seed-matched target genes of miR-26a-5p.

Endo-miR-26a-5p targets

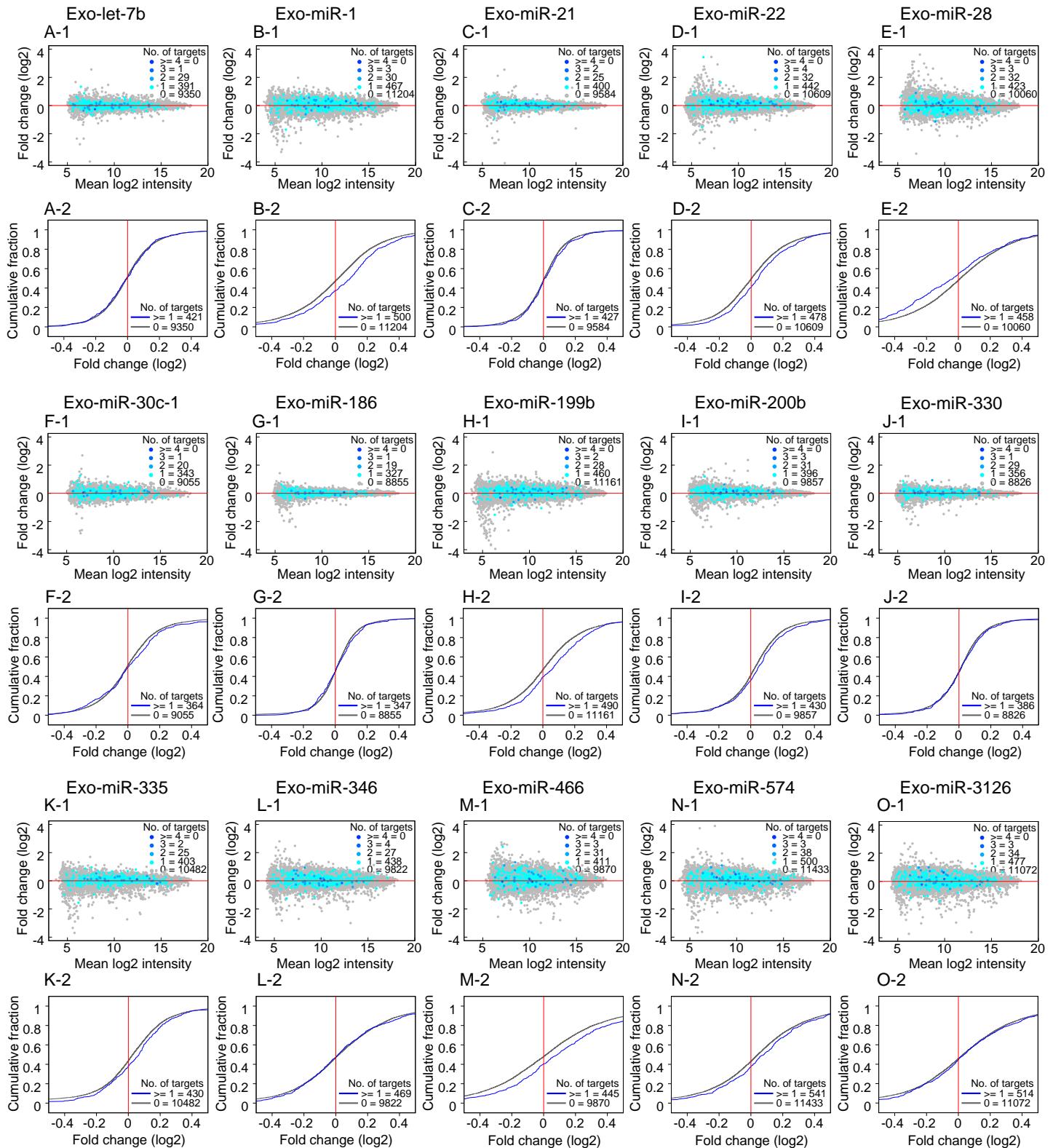


Figure S7. Microarray analysis of the expression of target genes of the endo-miRNA, miR-24-3p
Microarray analysis of endo-miR-24-3p target gene expression in cells transfected with exo-let-7b (A),
exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
(M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
distributions (A-2~O-2) of seed-matched target genes of miR-24-3p.

Endo-miR-24-3p targets

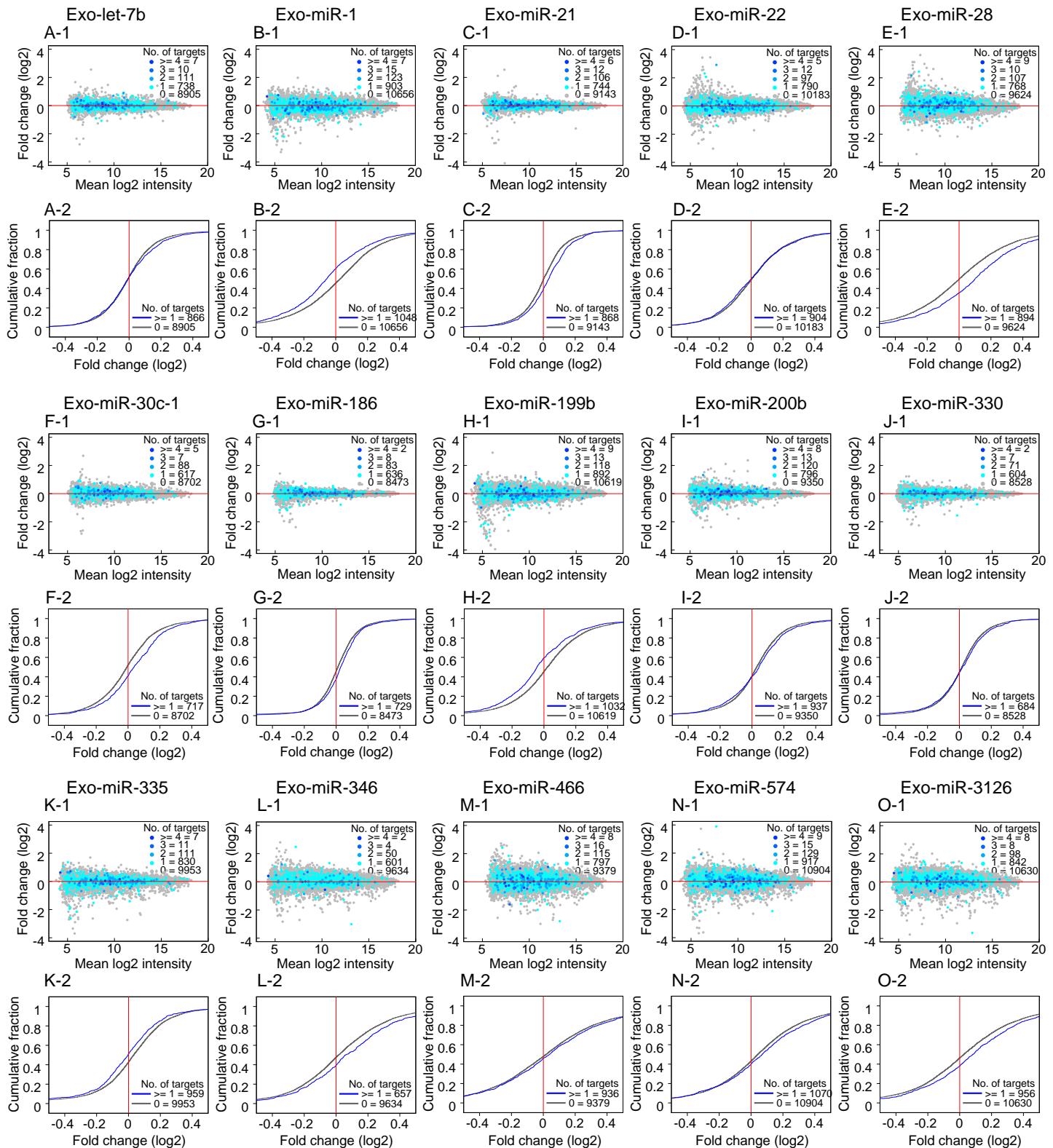


Figure S8. Microarray analysis of the expression of target genes of the endo-miRNA, miR-30a-5p
Microarray analysis of endo-miR-30a-5p target gene expression in cells transfected with exo-let-7b (A),
exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
(M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
distributions (A-2~O-2) of seed-matched target genes of miR-30a-5p.

Endo-miR-30a-5p targets

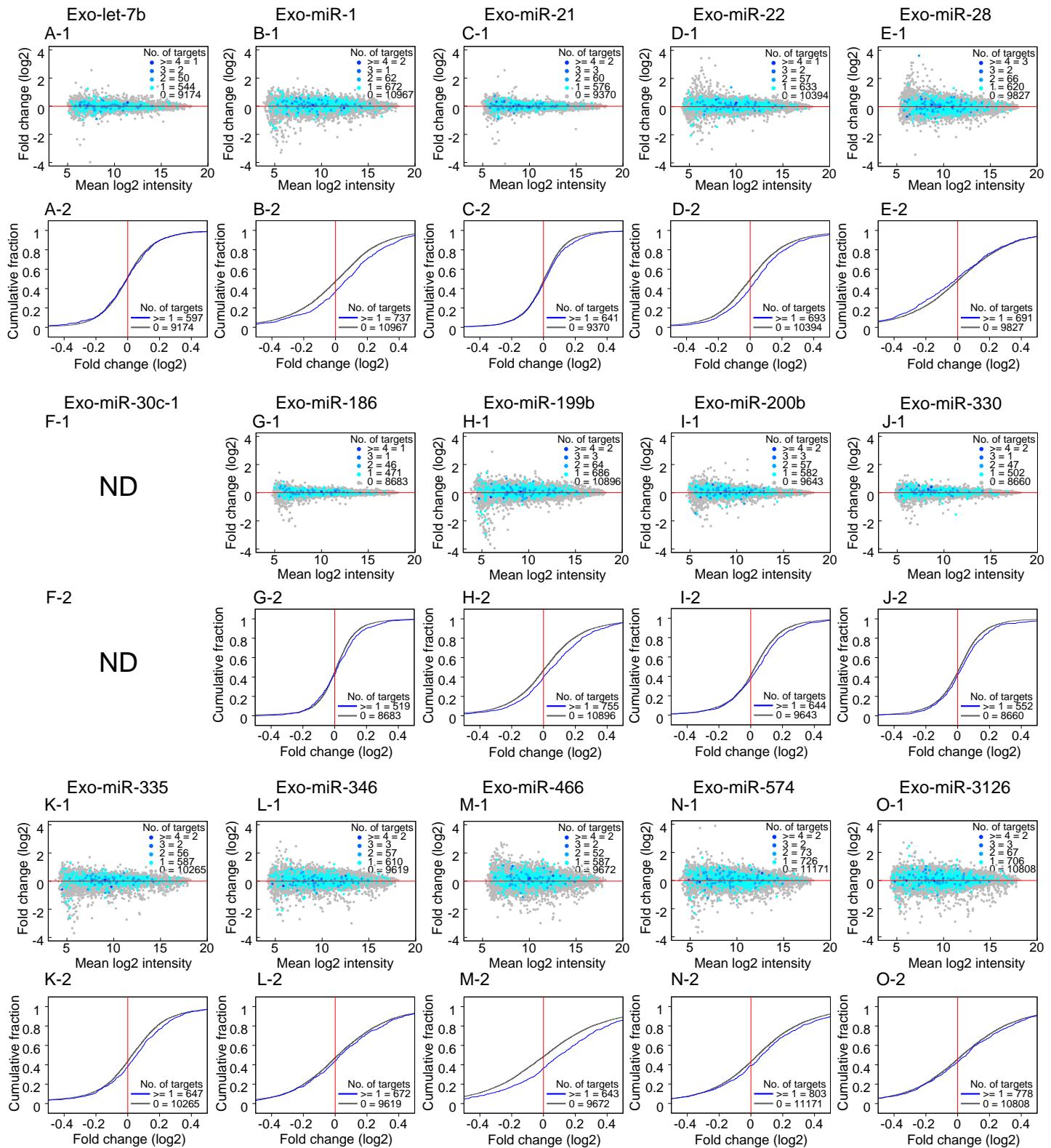


Figure S9. Microarray analysis of the expression of target genes of the endo-miRNA, miR-92a-3p
 Microarray analysis of endo-miR-92a-3p target gene expression in cells transfected with exo-let-7b (A),
 exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
 exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
 (M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
 distributions (A-2~O-2) of seed-matched target genes of miR-92a-3p.

Endo-miR-92a-3p targets

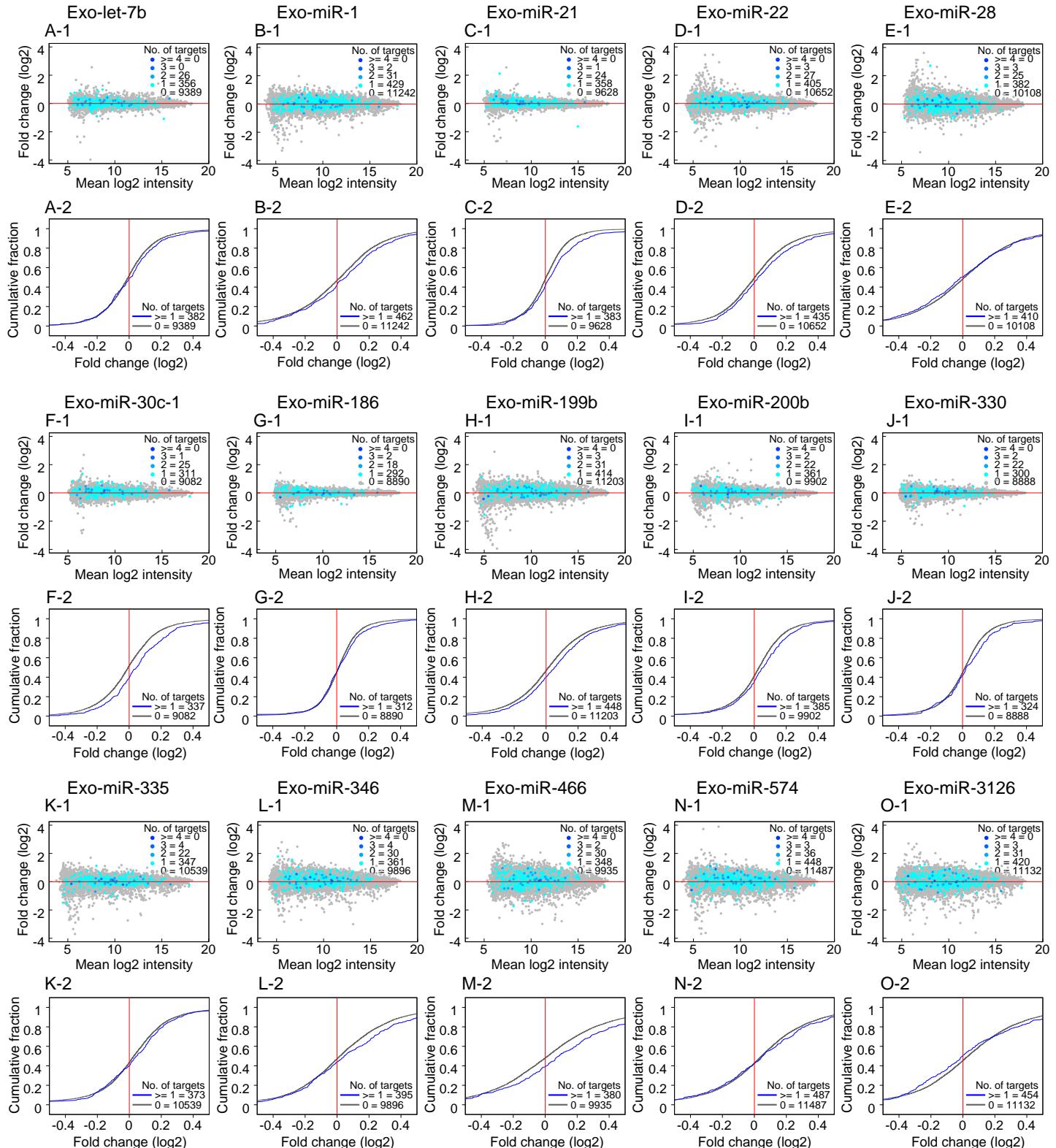


Figure S10. Microarray analysis of the expression of target genes of the endo-miRNA, miR-19a-3p
 Microarray analysis of endo-miR-19a-3p target gene expression in cells transfected with exo-let-7b (A),
 exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
 exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
 (M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
 distributions (A-2~O-2) of seed-matched target genes of miR-19a-3p.

Endo-miR-19a-3p targets

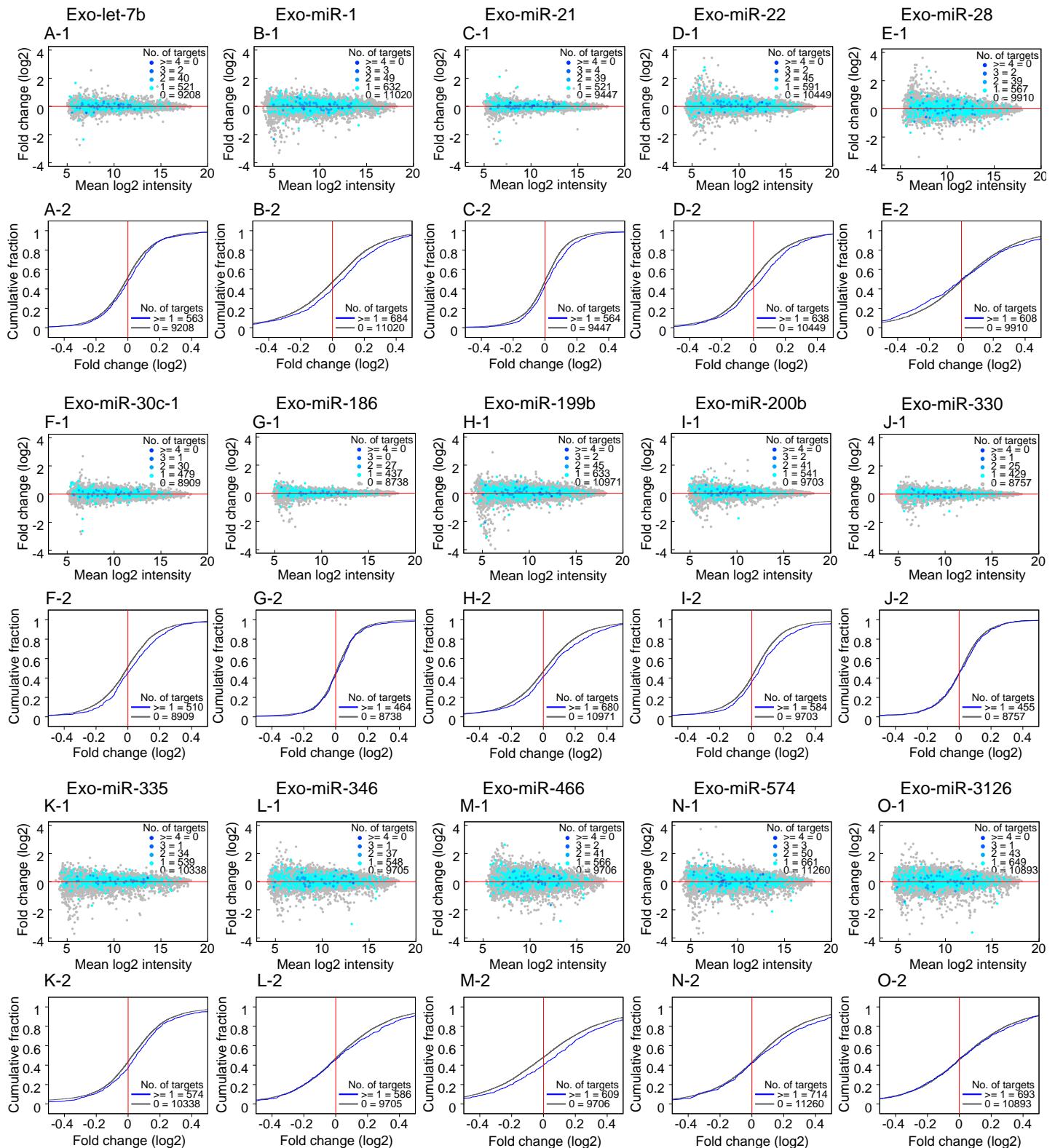


Figure S11. Microarray analysis of the expression of target genes of the endo-miRNA, miR-15a-5p
Microarray analysis of endo-miR-15a-5p target gene expression in cells transfected with exo-let-7b (A),
exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
(M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
distributions (A-2~O-2) of seed-matched target genes of miR-15a-5p.

Endo-miR-15a-5p targets

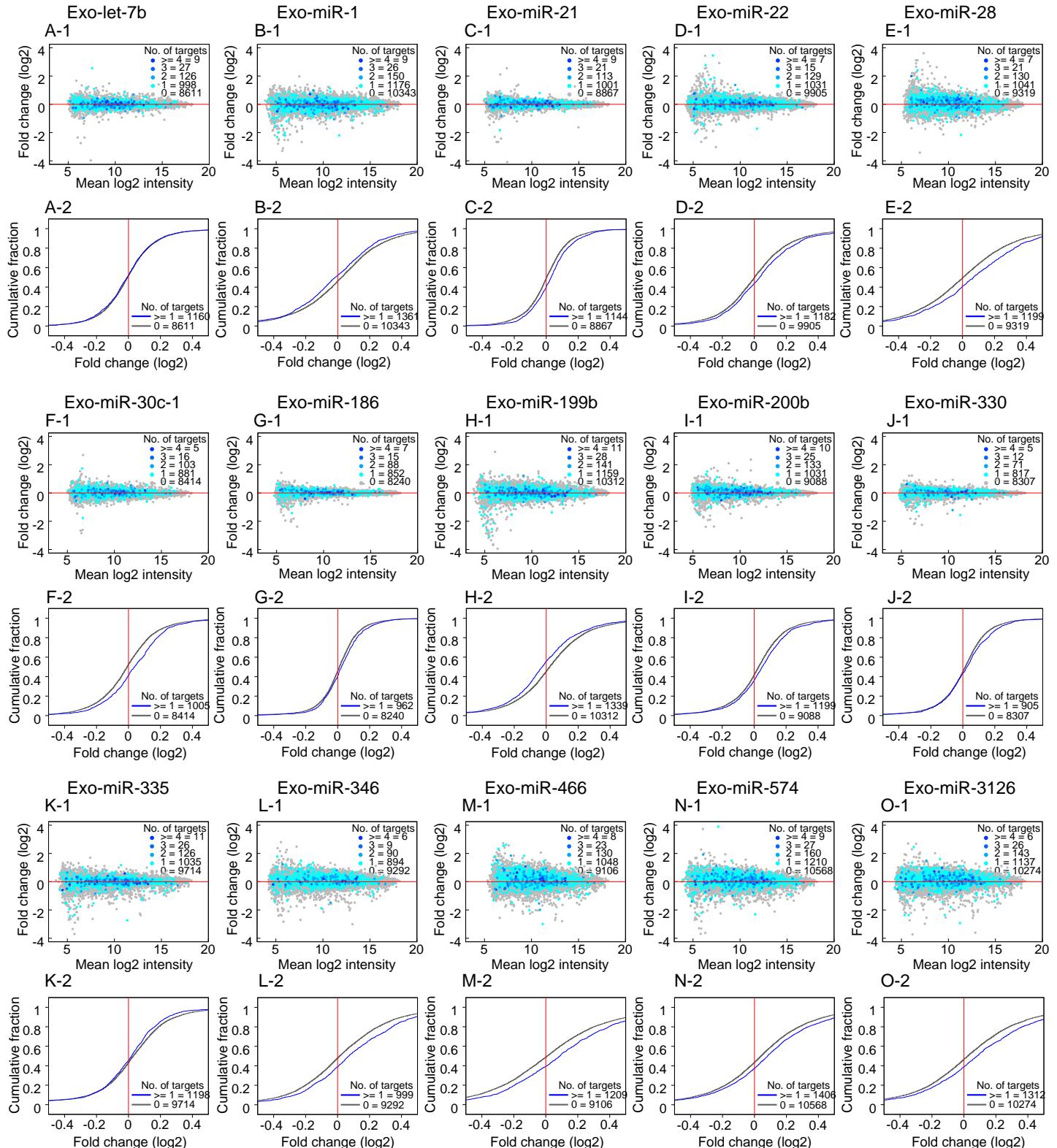


Figure S12. Microarray analysis of the expression of target genes of the endo-miRNA, miR-22-3p
Microarray analysis of endo-miR-22-3p target gene expression in cells transfected with exo-let-7b (A),
exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
(M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
distributions (A-2~O-2) of seed-matched target genes of miR-22-3p.

Endo-miR-22-3p targets

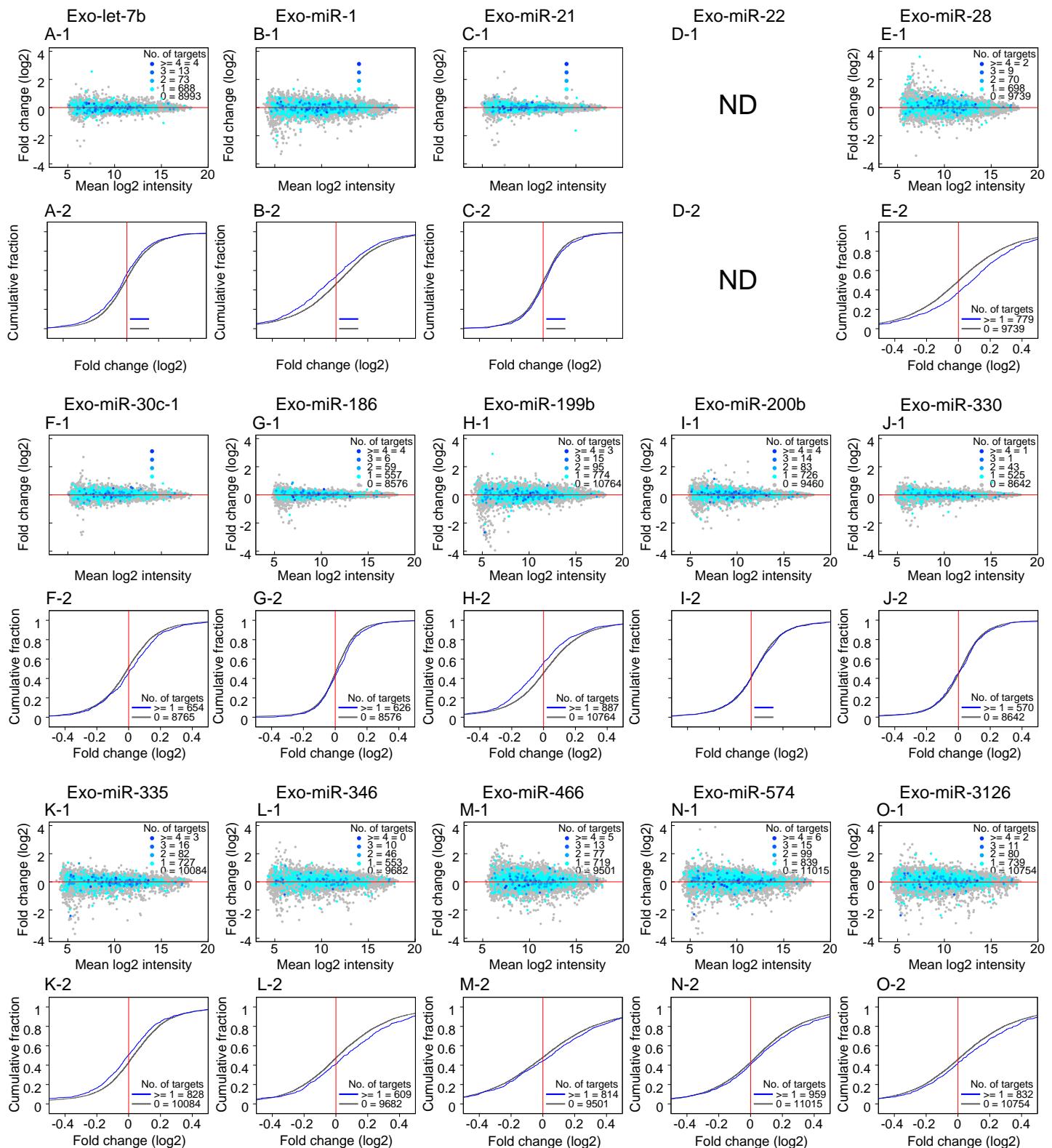


Figure S13. Microarray analysis of the expression of target genes of the endo-miRNA, miR-29a-3p
 Microarray analysis of endo-miR-29a-3p target gene expression in cells transfected with exo-let-7b (A),
 exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
 exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
 (M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
 distributions (A-2~O-2) of seed-matched target genes of miR-29a-3p.

Endo-miR-29a-3p targets

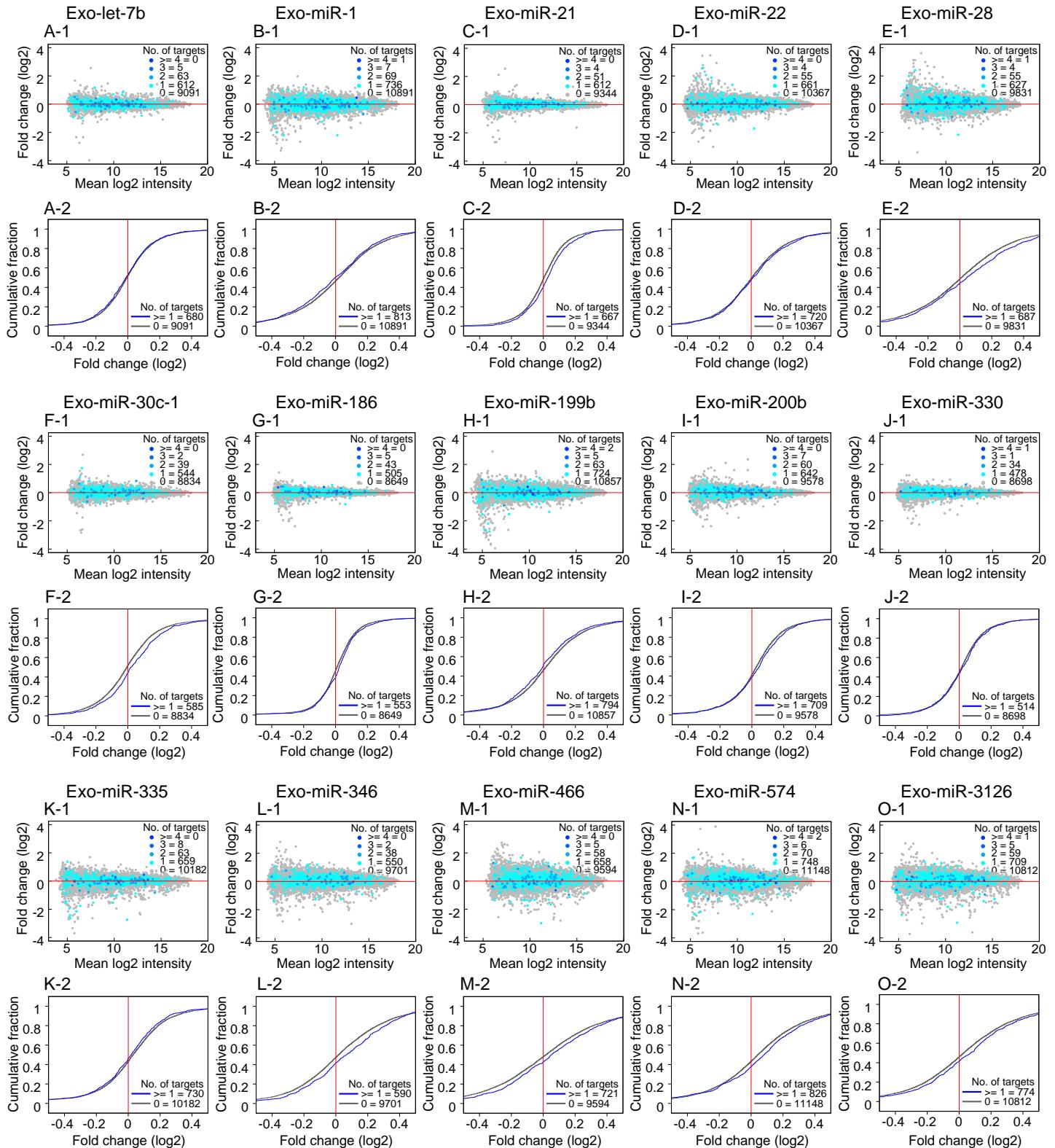


Figure S14. Microarray analysis of the expression of target genes of the endo-miRNA, miR-125a-5p
 Microarray analysis of endo-miR-125a-5p target gene expression in cells transfected with exo-let-7b (A),
 exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
 exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
 (M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
 distributions (A-2~O-2) of seed-matched target genes of miR-125a-5p.

Endo-miR-125a-5p targets

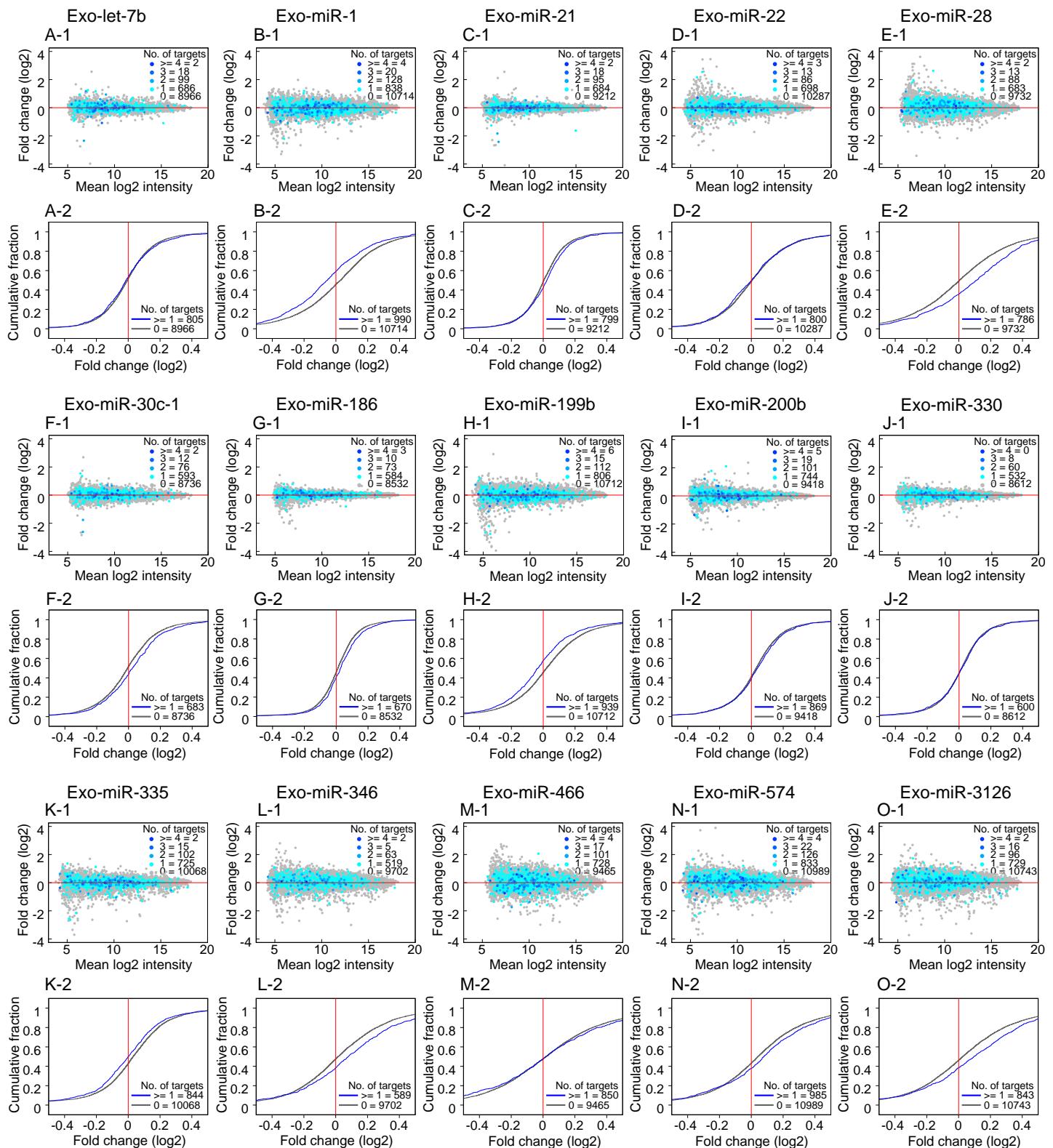


Figure S15. Microarray analysis of the expression of target genes of the endo-miRNA, miR-93-5p
Microarray analysis of endo-miR-93-5p target gene expression in cells transfected with exo-let-7b (A),
exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
(M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
distributions (A-2~O-2) of seed-matched target genes of miR-93-5p.

Endo-miR-93-5p targets

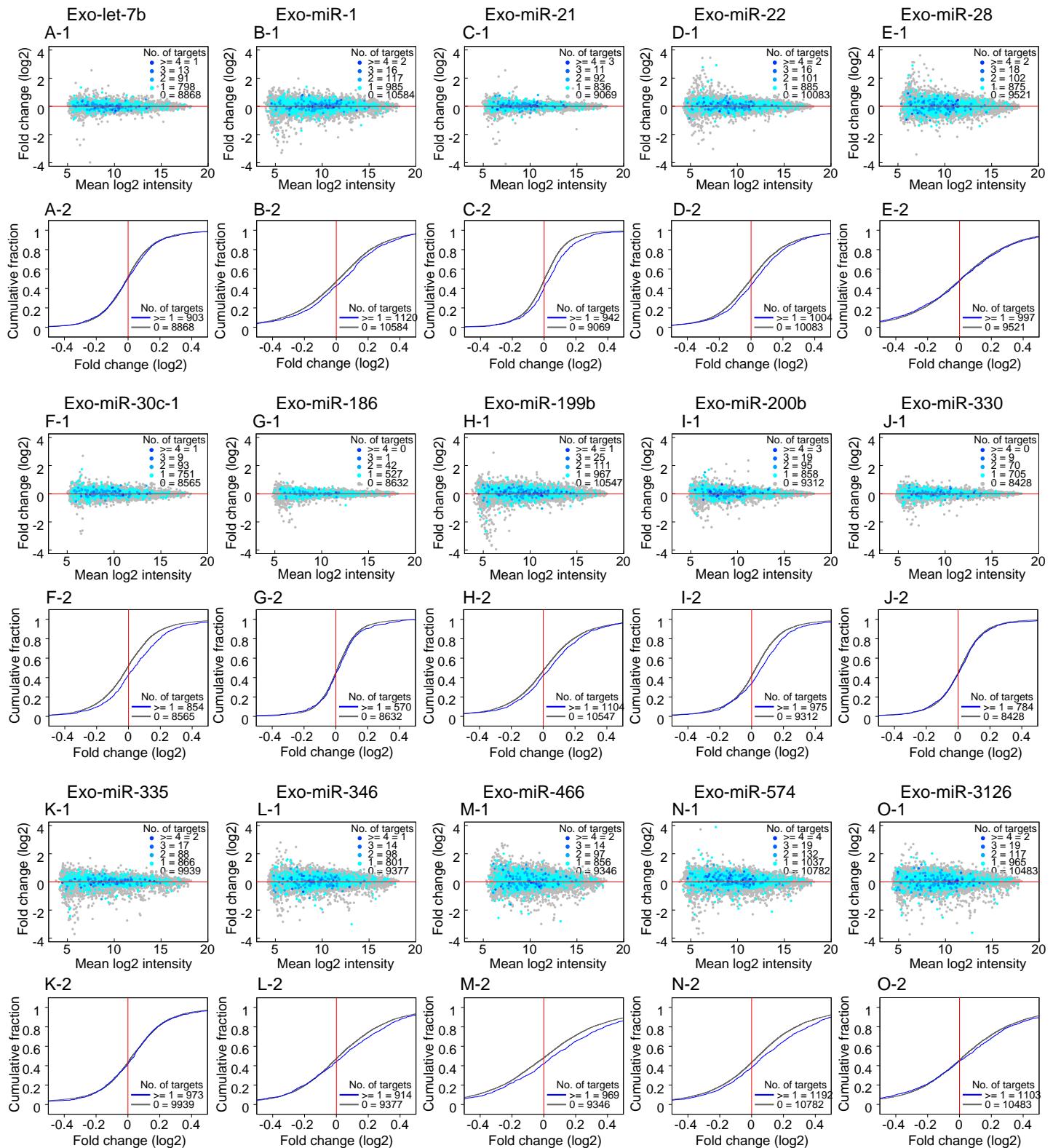


Figure S16. Microarray analysis of the expression of target genes of the endo-miRNA, miR-191-5p
 Microarray analysis of endo-miR-191-5p target gene expression in cells transfected with exo-let-7b (A),
 exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
 exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
 (M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
 distributions (A-2~O-2) of seed-matched target genes of miR-191-5p.

Endo-miR-191-5p targets

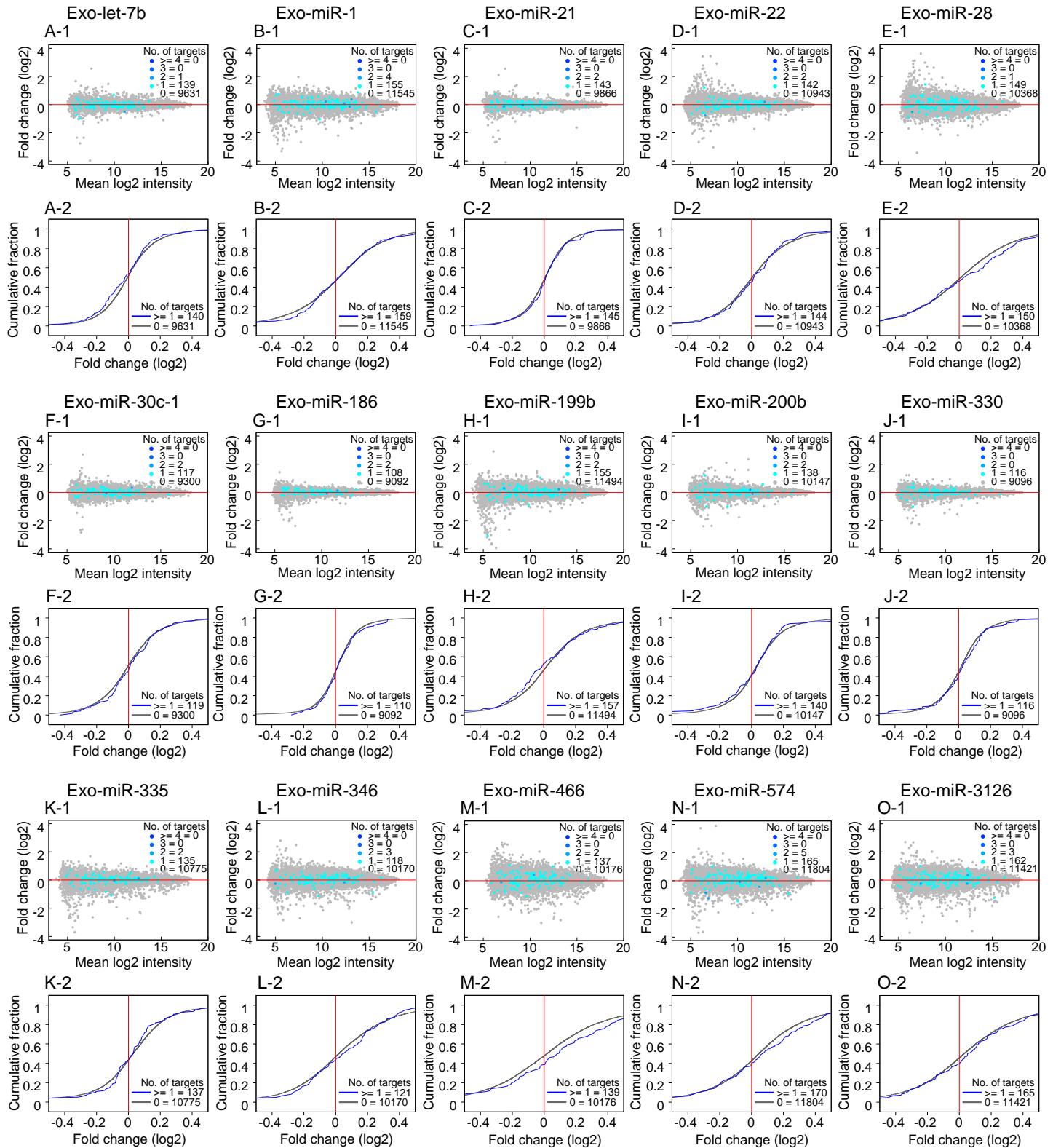


Figure S17. Microarray analysis of the expression of target genes of the endo-miRNA, miR-103a-3p. Microarray analysis of endo-miR-103a-3p target gene expression in cells transfected with exo-let-7b (A), exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G), exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446 (M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative distributions (A-2~O-2) of seed-matched target genes of miR-103a-3p.

Endo-miR-103a-3p targets

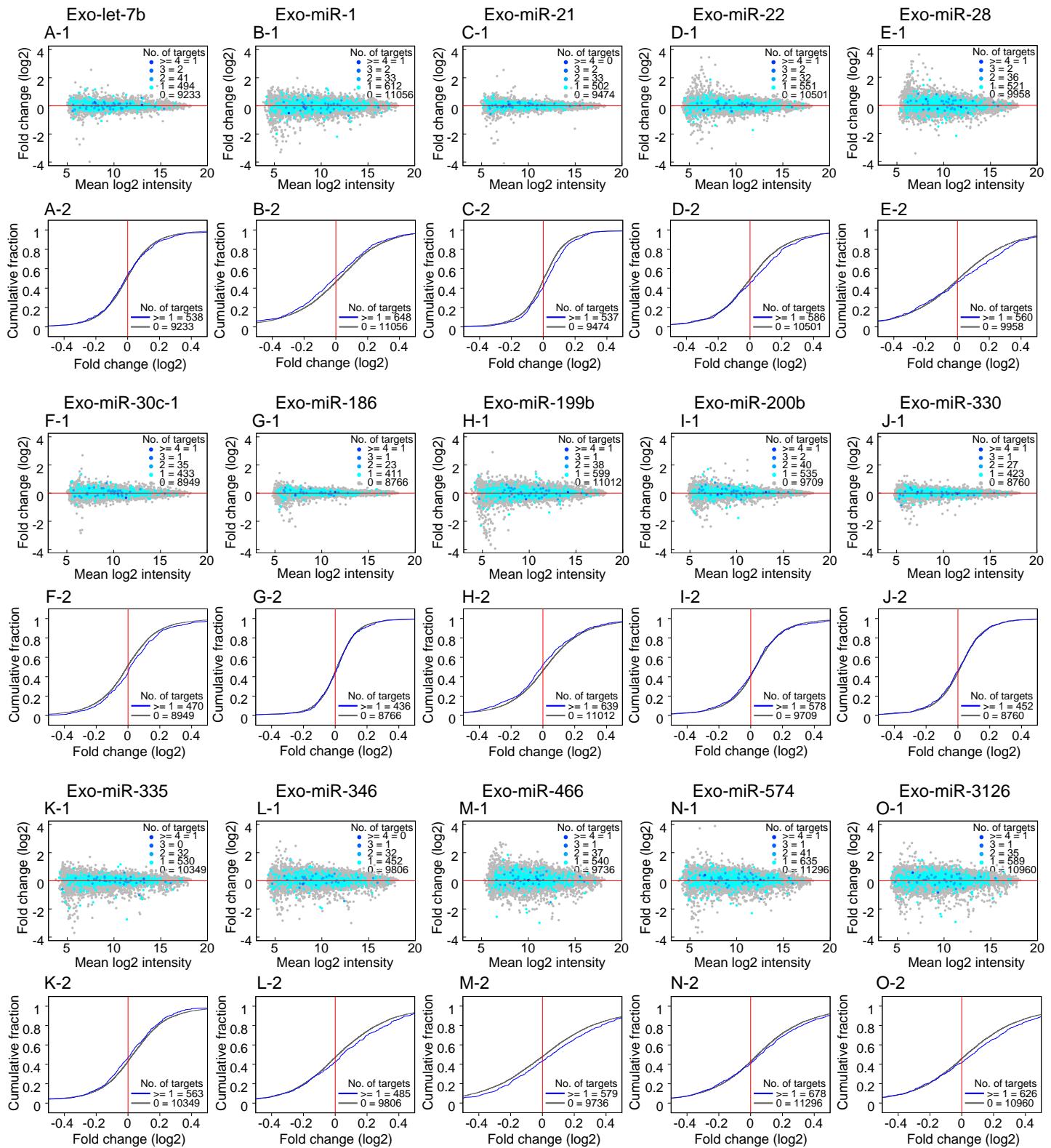


Figure S18. Microarray analysis of the expression of target genes of the endo-miRNA, miR-143-3p
Microarray analysis of endo-miR-143-3p target gene expression in cells transfected with exo-let-7b (A),
exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
(M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
distributions (A-2~O-2) of seed-matched target genes of miR-143-3p.

Endo-miR-143-3p targets

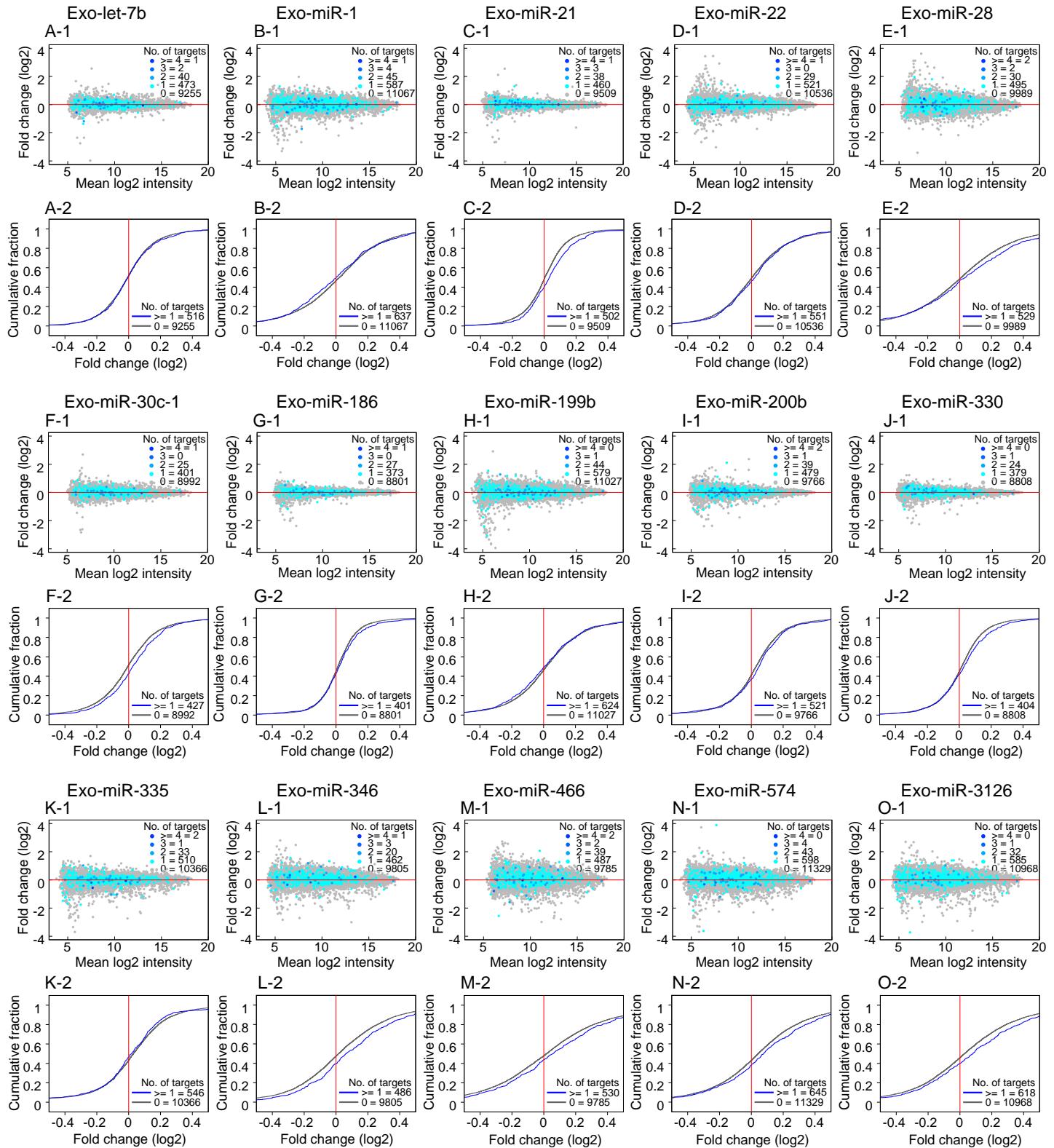


Figure S19. Microarray analysis of the expression of target genes of the endo-miRNA, miR-100-5p
 Microarray analysis of endo-miR-100-5p target gene expression in cells transfected with exo-let-7b (A),
 exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
 exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
 (M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
 distributions (A-2~O-2) of seed-matched target genes of miR-100-5p.

Endo-miR-100-5p targets

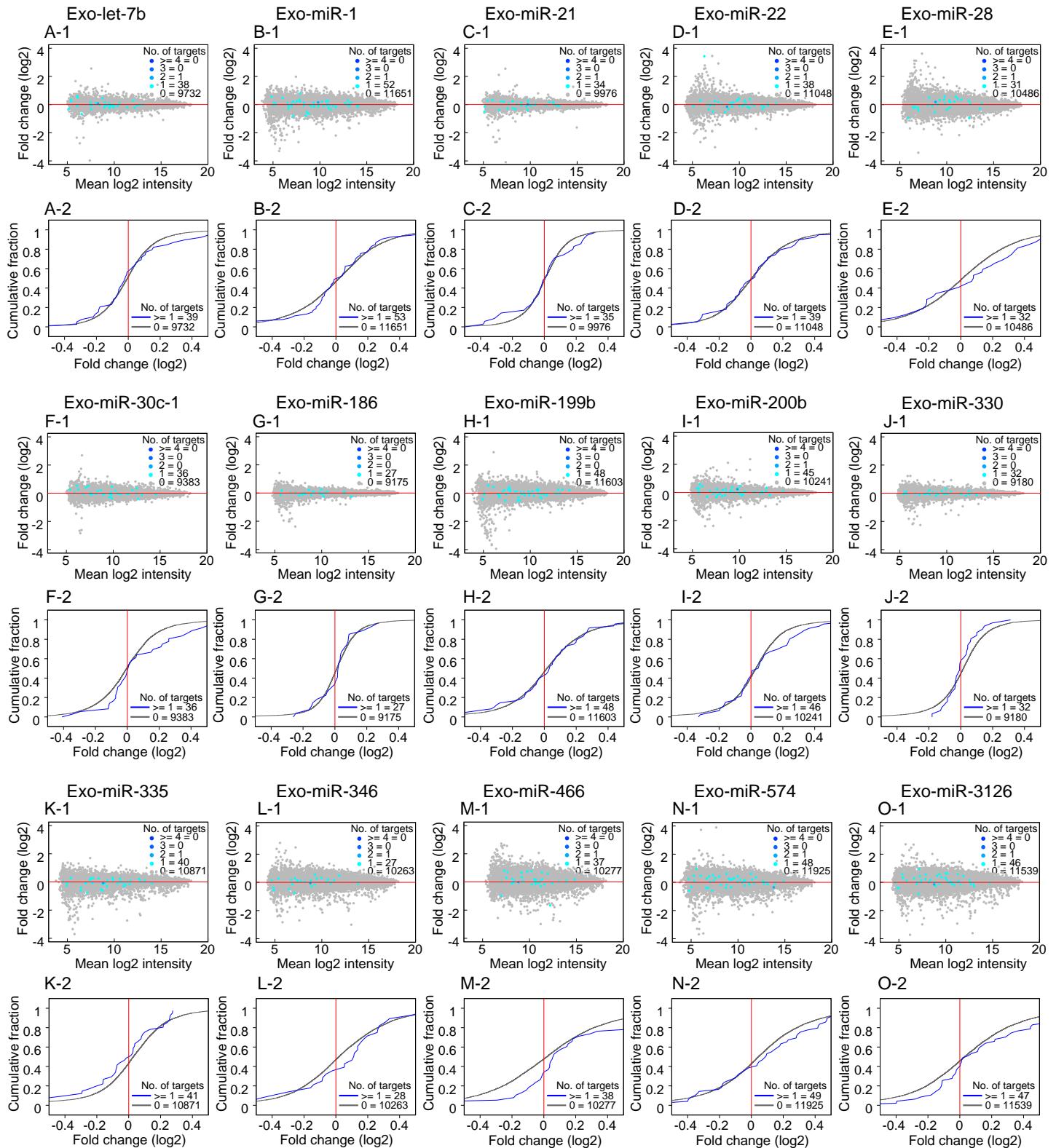


Figure S20. Microarray analysis of the expression of target genes of the endo-miRNA, miR-23a-3p
Microarray analysis of endo-miR-23a-3p target gene expression in cells transfected with exo-let-7b (A),
exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
(M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
distributions (A-2~O-2) of seed-matched target genes of miR-23a-3p.

Endo-miR-23a-3p targets

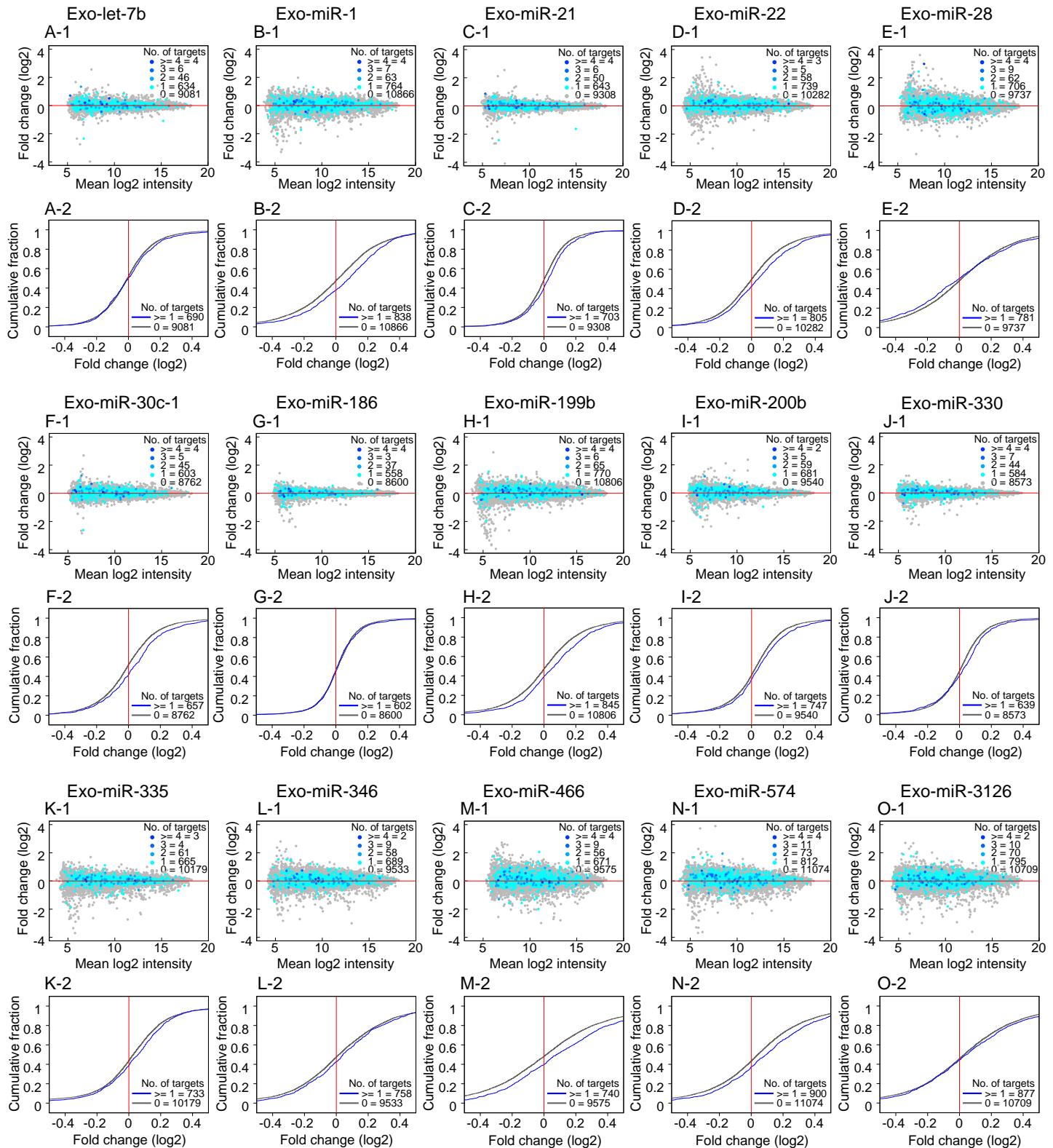


Figure S21. Microarray analysis of the expression of target genes of the endo-miRNA, miR-186-5p
 Microarray analysis of endo-miR-186-5p target gene expression in cells transfected with exo-let-7b (A),
 exo-miR-1 (B), exo-miR-21 (C), exo-miR-22 (D), exo-miR-28 (E), exo-miR-30c-1 (F), exo-miR-186 (G),
 exo-miR-199b (H), exo-miR-200b (I), exo-miR-330 (J), exo-miR-335 (K), exo-miR-346 (L), exo-miR-446
 (M), exo-miR-574 (N), exo-miR-3126 (O). Data are presented as MA plots (A-1~O-1) and cumulative
 distributions (A-2~O-2) of seed-matched target genes of miR-186-5p.

Endo-miR-186-5p targets

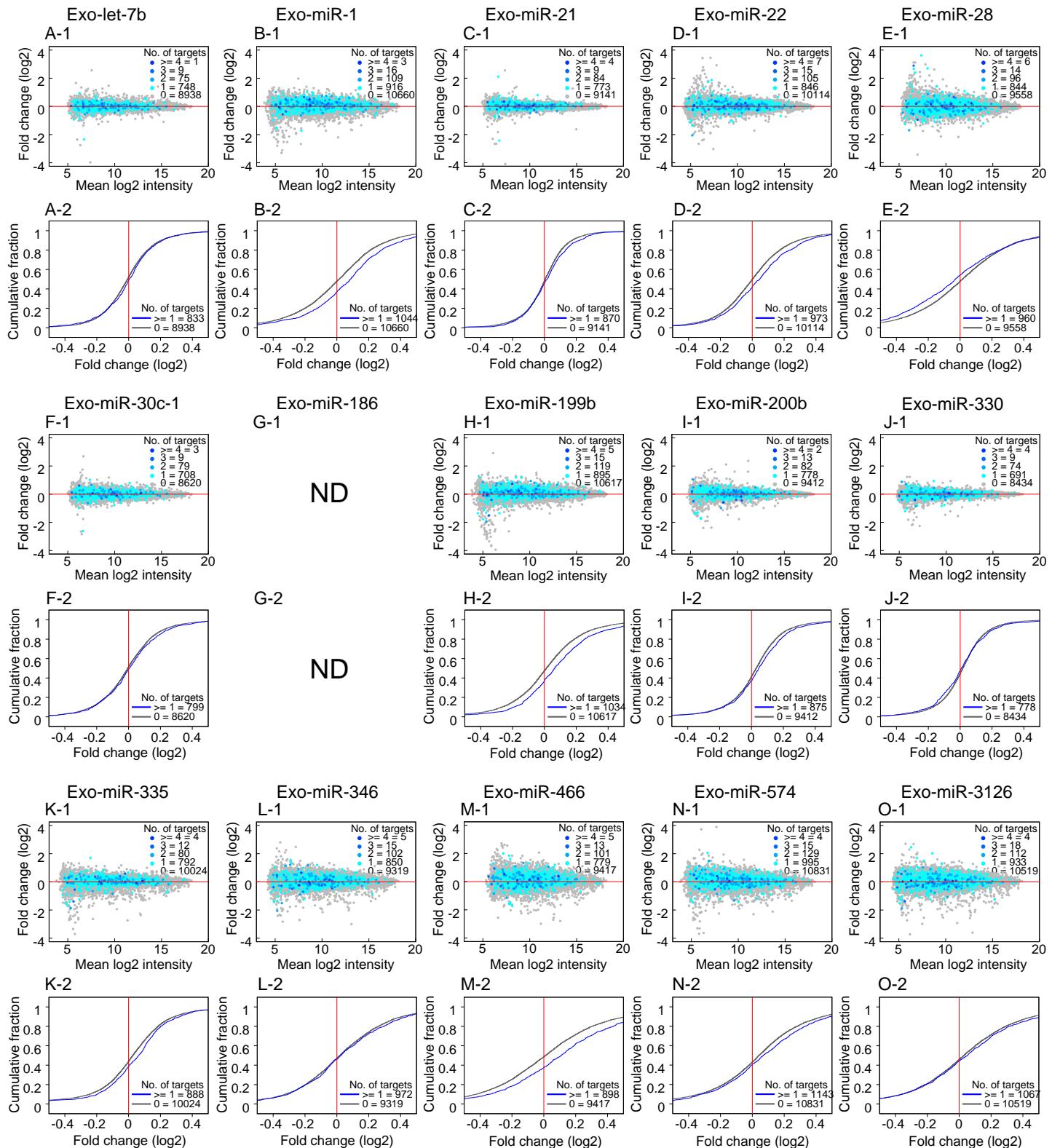
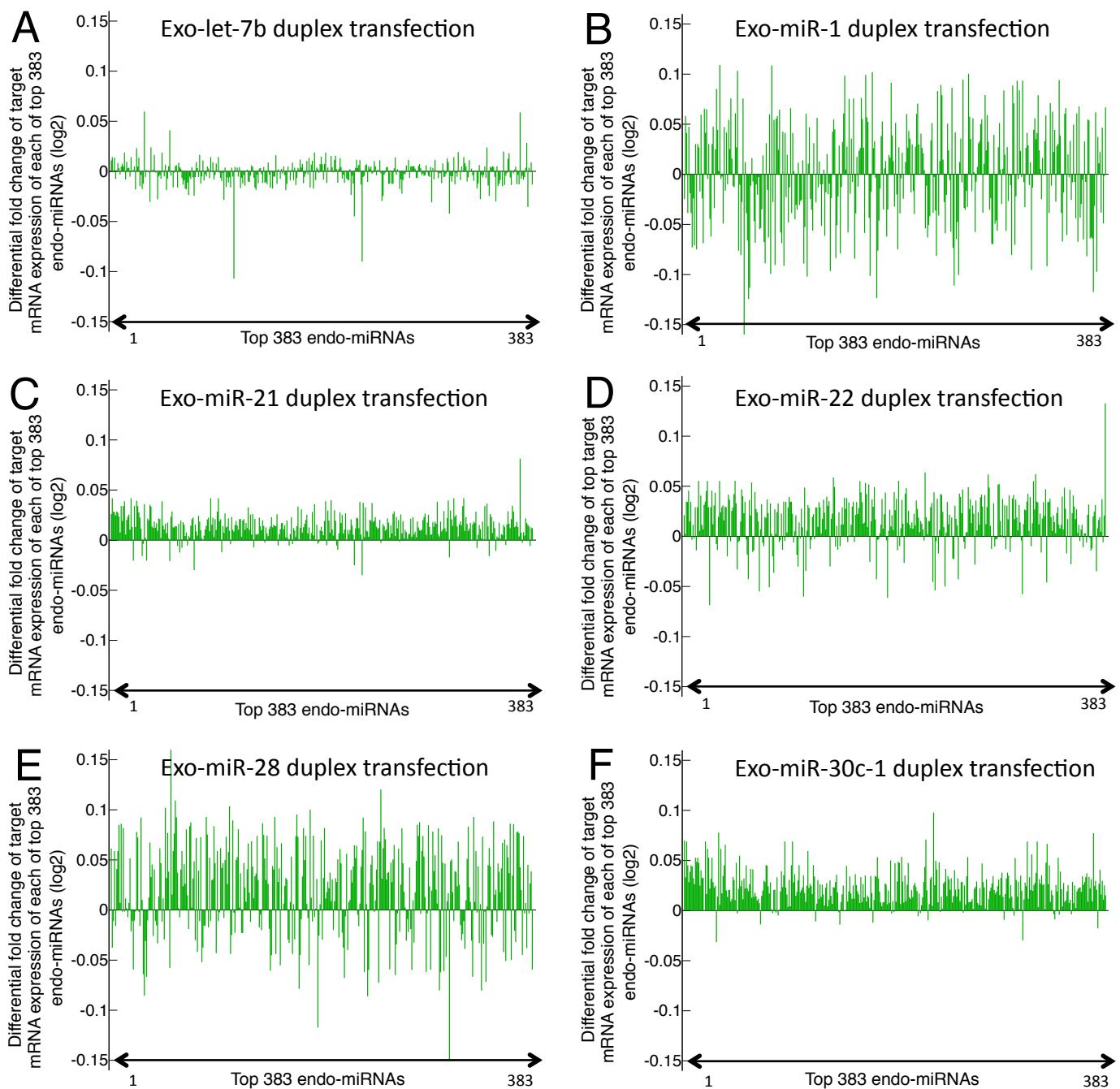
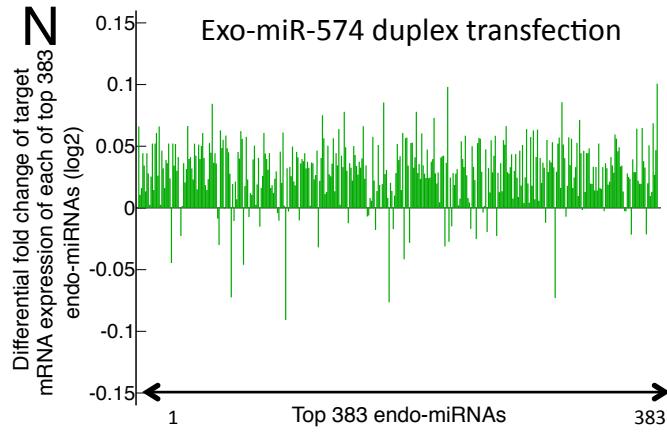
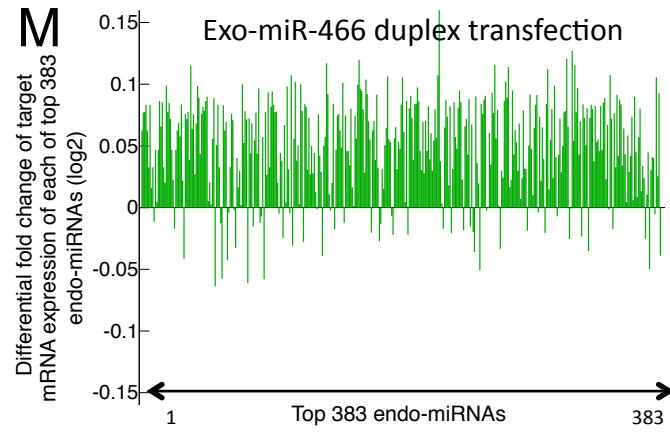
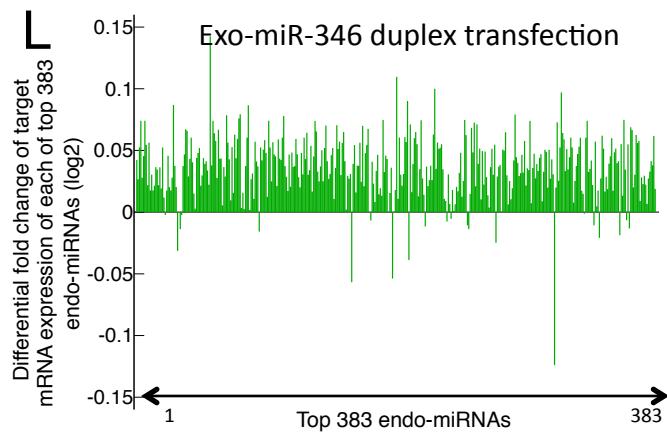
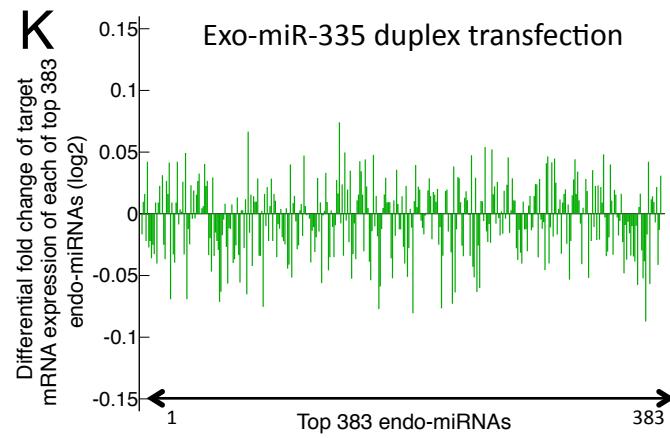
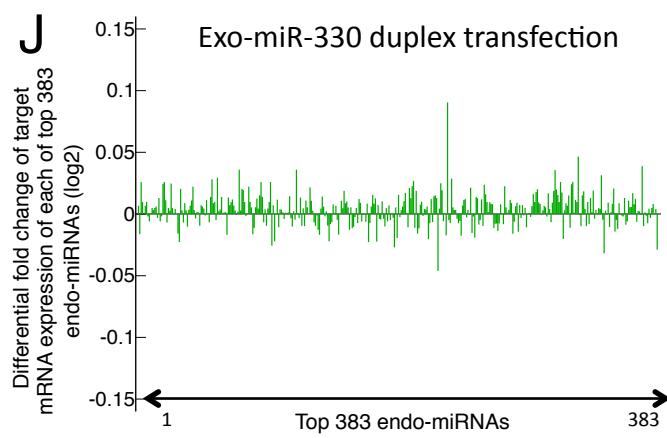
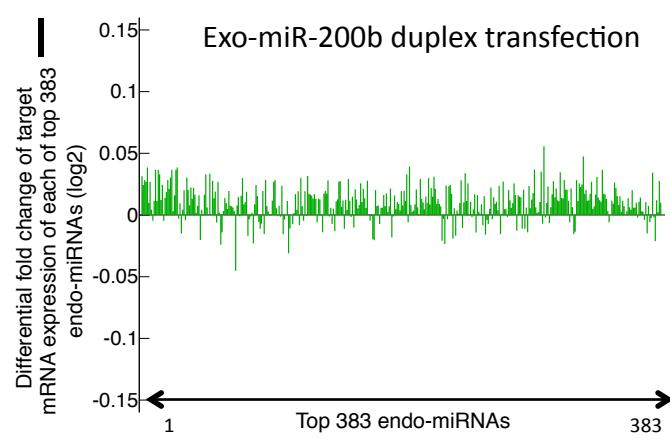
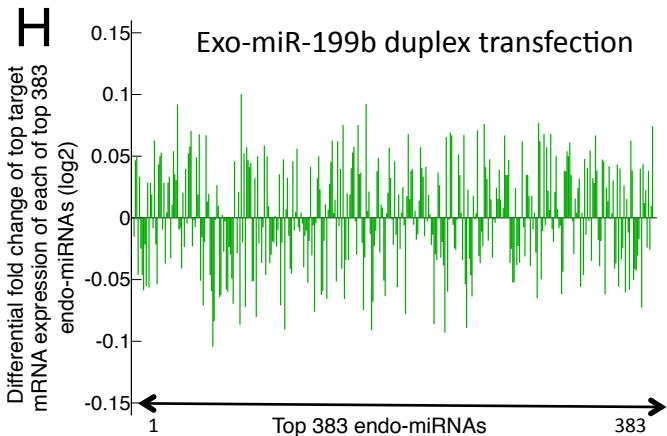
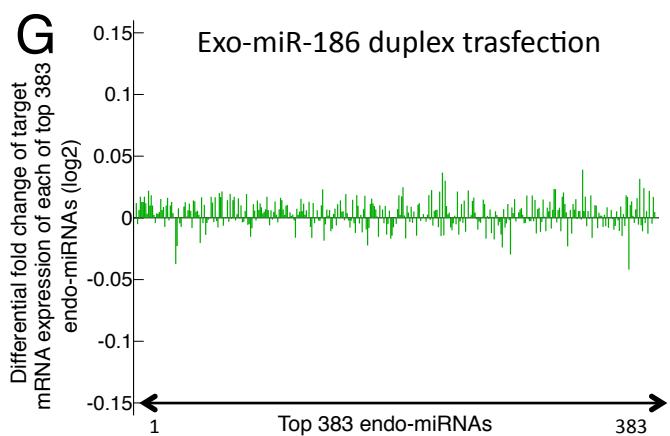


Figure S22. Mean fold changes of the expression levels of endo-miRNA target transcripts by the transfection of different exo-miRNAs. HeLa cells were transfected with each of fifteen exo-miRNA duplexes, \log_2 mean differential fold changes of the expression levels of seed-matched target genes of each of top 20 endo-miRNAs were calculated. Data were shown with respect to endo-miRNAs, let-7b-5p (**A**), miR-21 (**B**), mR-27a-3p (**C**), miR-17-5p (**D**), miR-26a-5p (**E**), miR-24-3p (**F**), miR-30a-5p (**G**), miR-92a-5p (**H**), miR-19a-5p (**I**), miR-15a-5p (**J**), miR-22-3p (**K**), miR-29a-3p (**L**), miR-125a-5p (**M**), miR-93-5p (**N**), miR-191-5p (**O**), miR-103a-3p (**P**), miR-143-3p (**Q**), miR-100-5p (**R**), miR-23a-3p (**S**), and mR-186-5p (**T**), and their averaged values were shown in (**U**). Note that each exo-miRNA increased the endo-miRNA target genes in different degrees. The individual data are shown in Figures S2-S21. Data represent the mean \pm S.E (*, P < 0.05).





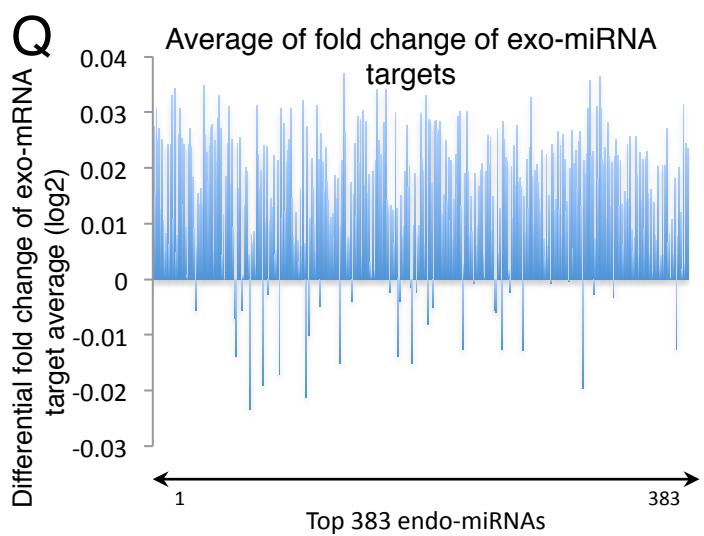
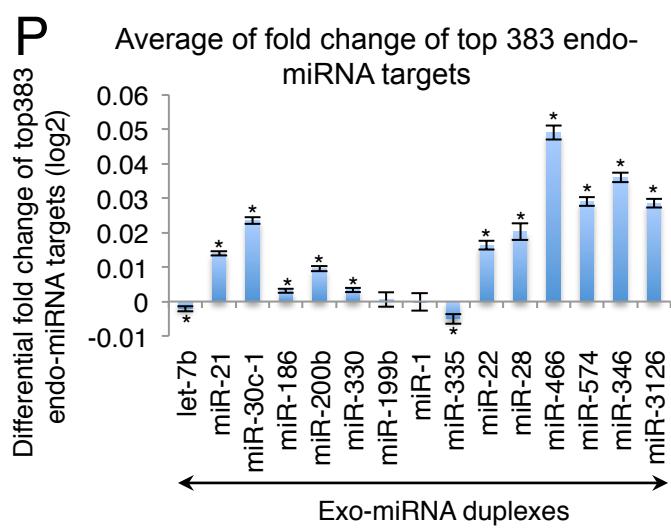
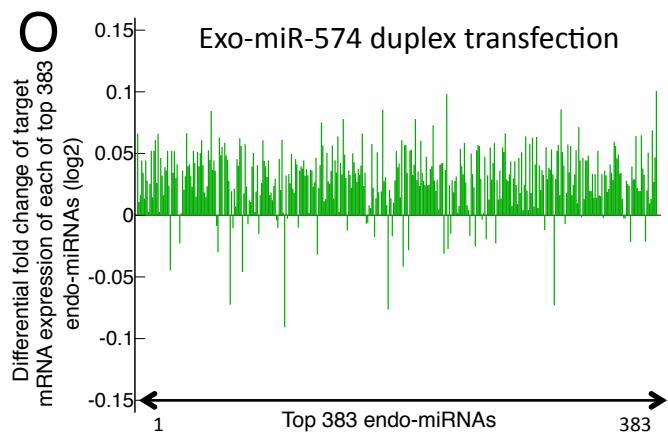


Figure S23. Confirmation of microarray data by quantitative RT-PCR. A total of 28 genes were subjected to quantitative analysis, and the results were compared with the corresponding microarray data. HeLa cells were transfected with miRNA duplexes, let-7b, miR-21, miR-30c-1, miR-186, miR-200b, and the expressions of the following were measured by quantitative RT-PCR: let-7b-5p targets, HDHD1A, UBQLN, PBX2, and ZC3H11A; miR-21-5p targets, SKA2, CASP7, SLC11A2, and UBR3; miR-30c-5p targets, RARG, SPAST, METTL7A, and SEC23A; miR-186-5p targets, ZNF652, SLC39A1, OTUD4, and USP37; miR-200b-5p targets, AP1S2, OSTM1, SHCBP1, and PGM2L1. The correlation coefficient between the quantitative RT-PCR and microarray expression data was 0.74.

