

Supplementary Table S3 Differential expression at the total miRNA level between SARS-CoV-2–infected patients requiring high-flow versus no oxygen support.

miRNA	Base mean^a	Fold Change^b	p adjusted^c
hsa-miR-671-5p	75.57	22.47	0.00519451
hsa-miR-6819-5p	11.13	14.22	0.00420542
hsa-miR-100-5p	277.51	3.05	0.01181046
hsa-miR-1306-3p	18.82	4.56	0.01227122
hsa-miR-3688-3p	20.97	-3.36	0.03054533
hsa-miR-4646-5p	12.31	3.71	0.03846312

^aThe average of the normalized count values according to DeSeq2.

^bThe effect size estimate according to DeSeq2.

^cP value adjusted for false discovery rate by the method of Benjamini and Hochberg.