MA\_ATCC Code

nohup IntaRNA -t MA\_ATCC\_1.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_1\_targets.txt &> nohuphsa-MA\_ATCC\_1.out &

nohup IntaRNA -t MA\_ATCC\_2.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_2\_targets.txt &> nohuphsa-MA\_ATCC\_2.out &

nohup IntaRNA -t MA\_ATCC\_3.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_3\_targets.txt &> nohuphsa-MA\_ATCC\_3.out &

nohup IntaRNA -t MA\_ATCC\_4.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_4\_targets.txt &> nohuphsa-MA\_ATCC\_4.out &

nohup IntaRNA -t MA\_ATCC\_5.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_5\_targets.txt &> nohuphsa-MA\_ATCC\_5.out &

nohup IntaRNA -t MA\_ATCC\_6.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_6\_targets.txt &> nohuphsa-MA\_ATCC\_6.out &

nohup IntaRNA -t MA\_ATCC\_7.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_7\_targets.txt &> nohuphsa-MA\_ATCC\_7.out &

nohup IntaRNA -t MA\_ATCC\_8.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_8\_targets.txt &> nohuphsa-MA\_ATCC\_8.out &

nohup IntaRNA -t MA\_ATCC\_9.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_9\_targets.txt &> nohuphsa-MA\_ATCC\_9.out &

nohup IntaRNA -t MA\_ATCC\_10.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_10\_targets.txt &> nohuphsa-MA\_ATCC\_10.out &

nohup IntaRNA -t MA\_ATCC\_11.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_11\_targets.txt &> nohuphsa-MA\_ATCC\_11.out &

nohup IntaRNA -t MA\_ATCC\_12.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_12\_targets.txt &> nohuphsa-MA\_ATCC\_12.out &

nohup IntaRNA -t MA\_ATCC\_13.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_13\_targets.txt &> nohuphsa-MA\_ATCC\_13.out &

nohup IntaRNA -t MA\_ATCC\_14.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_14\_targets.txt &> nohuphsa-MA\_ATCC\_14.out &

nohup IntaRNA -t MA\_ATCC\_15.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_15\_targets.txt &> nohuphsa-MA\_ATCC\_15.out &

nohup IntaRNA -t MA\_ATCC\_16.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_16\_targets.txt &> nohuphsa-MA\_ATCC\_16.out &

nohup IntaRNA -t MA\_ATCC\_17.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_17\_targets.txt &> nohuphsa-MA\_ATCC\_17.out &

nohup IntaRNA -t MA\_ATCC\_18.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_18\_targets.txt &> nohuphsa-MA\_ATCC\_18.out &

nohup IntaRNA -t MA\_ATCC\_19.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_19\_targets.txt &> nohuphsa-MA\_ATCC\_19.out &

nohup IntaRNA -t MA\_ATCC\_20.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out MA\_ATCC\_20\_targets.txt &> nohuphsa-MA\_ATCC\_20.out &