SP\_ATCC Code

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

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

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

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

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

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

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

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