EC\_K12 Code

nohup IntaRNA -t EC\_K12\_1.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_1\_targets.txt &> nohuphsa-EC\_K12\_1.out &

nohup IntaRNA -t EC\_K12\_2.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_2\_targets.txt &> nohuphsa-EC\_K12\_2.out &

nohup IntaRNA -t EC\_K12\_3.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_3\_targets.txt &> nohuphsa-EC\_K12\_3.out &

nohup IntaRNA -t EC\_K12\_4.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_4\_targets.txt &> nohuphsa-EC\_K12\_4.out &

nohup IntaRNA -t EC\_K12\_5.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_5\_targets.txt &> nohuphsa-EC\_K12\_5.out &

nohup IntaRNA -t EC\_K12\_6.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_6\_targets.txt &> nohuphsa-EC\_K12\_6.out &

nohup IntaRNA -t EC\_K12\_7.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_7\_targets.txt &> nohuphsa-EC\_K12\_7.out &

nohup IntaRNA -t EC\_K12\_8.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_8\_targets.txt &> nohuphsa-EC\_K12\_8.out &

nohup IntaRNA -t EC\_K12\_9.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_9\_targets.txt &> nohuphsa-EC\_K12\_9.out &

nohup IntaRNA -t EC\_K12\_10.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_10\_targets.txt &> nohuphsa-EC\_K12\_10.out &

nohup IntaRNA -t EC\_K12\_11.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_11\_targets.txt &> nohuphsa-EC\_K12\_11.out &

nohup IntaRNA -t EC\_K12\_12.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_12\_targets.txt &> nohuphsa-EC\_K12\_12.out &

nohup IntaRNA -t EC\_K12\_13.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_13\_targets.txt &> nohuphsa-EC\_K12\_13.out &

nohup IntaRNA -t EC\_K12\_14.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_14\_targets.txt &> nohuphsa-EC\_K12\_14.out &

nohup IntaRNA -t EC\_K12\_15.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_15\_targets.txt &> nohuphsa-EC\_K12\_15.out &

nohup IntaRNA -t EC\_K12\_16.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_16\_targets.txt &> nohuphsa-EC\_K12\_16.out &

nohup IntaRNA -t EC\_K12\_17.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out EC\_K12\_17\_targets.txt &> nohuphsa-EC\_K12\_17.out &