BC\_J2315 Code

nohup IntaRNA -t BC\_J2315\_1.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_1\_targets.txt &> nohuphsa-BC\_J2315\_1.out &

nohup IntaRNA -t BC\_J2315\_2.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_2\_targets.txt &> nohuphsa-BC\_J2315\_2.out &

nohup IntaRNA -t BC\_J2315\_3.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_3\_targets.txt &> nohuphsa-BC\_J2315\_3.out &

nohup IntaRNA -t BC\_J2315\_4.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_4\_targets.txt &> nohuphsa-BC\_J2315\_4.out &

nohup IntaRNA -t BC\_J2315\_5.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_5\_targets.txt &> nohuphsa-BC\_J2315\_5.out &

nohup IntaRNA -t BC\_J2315\_6.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_6\_targets.txt &> nohuphsa-BC\_J2315\_6.out &

nohup IntaRNA -t BC\_J2315\_7.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_7\_targets.txt &> nohuphsa-BC\_J2315\_7.out &

nohup IntaRNA -t BC\_J2315\_8.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_8\_targets.txt &> nohuphsa-BC\_J2315\_8.out &

nohup IntaRNA -t BC\_J2315\_9.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_9\_targets.txt &> nohuphsa-BC\_J2315\_9.out &

nohup IntaRNA -t BC\_J2315\_10.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_10\_targets.txt &> nohuphsa-BC\_J2315\_10.out &

nohup IntaRNA -t BC\_J2315\_11.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_11\_targets.txt &> nohuphsa-BC\_J2315\_11.out &

nohup IntaRNA -t BC\_J2315\_12.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_12\_targets.txt &> nohuphsa-BC\_J2315\_12.out &

nohup IntaRNA -t BC\_J2315\_13.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_13\_targets.txt &> nohuphsa-BC\_J2315\_13.out &

nohup IntaRNA -t BC\_J2315\_14.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_14\_targets.txt &> nohuphsa-BC\_J2315\_14.out &

nohup IntaRNA -t BC\_J2315\_15.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_15\_targets.txt &> nohuphsa-BC\_J2315\_15.out &

nohup IntaRNA -t BC\_J2315\_16.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_16\_targets.txt &> nohuphsa-BC\_J2315\_16.out &

nohup IntaRNA -t BC\_J2315\_17.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_17\_targets.txt &> nohuphsa-BC\_J2315\_17.out &

nohup IntaRNA -t BC\_J2315\_18.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_18\_targets.txt &> nohuphsa-BC\_J2315\_18.out &

nohup IntaRNA -t BC\_J2315\_19.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_19\_targets.txt &> nohuphsa-BC\_J2315\_19.out &

nohup IntaRNA -t BC\_J2315\_20.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_20\_targets.txt &> nohuphsa-BC\_J2315\_20.out &

nohup IntaRNA -t BC\_J2315\_21.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_21\_targets.txt &> nohuphsa-BC\_J2315\_21.out &

nohup IntaRNA -t BC\_J2315\_22.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_22\_targets.txt &> nohuphsa-BC\_J2315\_22.out &

nohup IntaRNA -t BC\_J2315\_23.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_23\_targets.txt &> nohuphsa-BC\_J2315\_23.out &

nohup IntaRNA -t BC\_J2315\_24.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_24\_targets.txt &> nohuphsa-BC\_J2315\_24.out &

nohup IntaRNA -t BC\_J2315\_25.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_25\_targets.txt &> nohuphsa-BC\_J2315\_25.out &

nohup IntaRNA -t BC\_J2315\_26.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_26\_targets.txt &> nohuphsa-BC\_J2315\_26.out &

nohup IntaRNA -t BC\_J2315\_27.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_27\_targets.txt &> nohuphsa-BC\_J2315\_27.out &

nohup IntaRNA -t BC\_J2315\_28.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_28\_targets.txt &> nohuphsa-BC\_J2315\_28.out &

nohup IntaRNA -t BC\_J2315\_29.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_29\_targets.txt &> nohuphsa-BC\_J2315\_29.out &

nohup IntaRNA -t BC\_J2315\_30.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out BC\_J2315\_30\_targets.txt &> nohuphsa-BC\_J2315\_30.out &