KP\_MGH Code

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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