SM\_K279A Code

nohup IntaRNA -t SM\_K279A\_1.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_1\_targets.txt &> nohuphsa-SM\_K279A\_1.out &

nohup IntaRNA -t SM\_K279A\_2.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_2\_targets.txt &> nohuphsa-SM\_K279A\_2.out &

nohup IntaRNA -t SM\_K279A\_3.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_3\_targets.txt &> nohuphsa-SM\_K279A\_3.out &

nohup IntaRNA -t SM\_K279A\_4.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_4\_targets.txt &> nohuphsa-SM\_K279A\_4.out &

nohup IntaRNA -t SM\_K279A\_5.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_5\_targets.txt &> nohuphsa-SM\_K279A\_5.out &

nohup IntaRNA -t SM\_K279A\_6.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_6\_targets.txt &> nohuphsa-SM\_K279A\_6.out &

nohup IntaRNA -t SM\_K279A\_7.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_7\_targets.txt &> nohuphsa-SM\_K279A\_7.out &

nohup IntaRNA -t SM\_K279A\_8.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_8\_targets.txt &> nohuphsa-SM\_K279A\_8.out &

nohup IntaRNA -t SM\_K279A\_9.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_9\_targets.txt &> nohuphsa-SM\_K279A\_9.out &

nohup IntaRNA -t SM\_K279A\_10.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_10\_targets.txt &> nohuphsa-SM\_K279A\_10.out &

nohup IntaRNA -t SM\_K279A\_11.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_11\_targets.txt &> nohuphsa-SM\_K279A\_11.out &

nohup IntaRNA -t SM\_K279A\_12.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_12\_targets.txt &> nohuphsa-SM\_K279A\_12.out &

nohup IntaRNA -t SM\_K279A\_13.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_13\_targets.txt &> nohuphsa-SM\_K279A\_13.out &

nohup IntaRNA -t SM\_K279A\_14.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_14\_targets.txt &> nohuphsa-SM\_K279A\_14.out &

nohup IntaRNA -t SM\_K279A\_15.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_15\_targets.txt &> nohuphsa-SM\_K279A\_15.out &

nohup IntaRNA -t SM\_K279A\_16.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_16\_targets.txt &> nohuphsa-SM\_K279A\_16.out &

nohup IntaRNA -t SM\_K279A\_17.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_17\_targets.txt &> nohuphsa-SM\_K279A\_17.out &

nohup IntaRNA -t SM\_K279A\_18.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out SM\_K279A\_18\_targets.txt &> nohuphsa-SM\_K279A\_18.out &