AX\_A8 Code

nohup IntaRNA -t AX\_A8\_1.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_1\_targets.txt &> nohuphsa-AX\_A8\_1.out &

nohup IntaRNA -t AX\_A8\_2.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_2\_targets.txt &> nohuphsa-AX\_A8\_2.out &

nohup IntaRNA -t AX\_A8\_3.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_3\_targets.txt &> nohuphsa-AX\_A8\_3.out &

nohup IntaRNA -t AX\_A8\_4.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_4\_targets.txt &> nohuphsa-AX\_A8\_4.out &

nohup IntaRNA -t AX\_A8\_5.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_5\_targets.txt &> nohuphsa-AX\_A8\_5.out &

nohup IntaRNA -t AX\_A8\_6.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_6\_targets.txt &> nohuphsa-AX\_A8\_6.out &

nohup IntaRNA -t AX\_A8\_7.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_7\_targets.txt &> nohuphsa-AX\_A8\_7.out &

nohup IntaRNA -t AX\_A8\_8.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_8\_targets.txt &> nohuphsa-AX\_A8\_8.out &

nohup IntaRNA -t AX\_A8\_9.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_9\_targets.txt &> nohuphsa-AX\_A8\_9.out &

nohup IntaRNA -t AX\_A8\_10.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_10\_targets.txt &> nohuphsa-AX\_A8\_10.out &

nohup IntaRNA -t AX\_A8\_11.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_11\_targets.txt &> nohuphsa-AX\_A8\_11.out &

nohup IntaRNA -t AX\_A8\_12.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_12\_targets.txt &> nohuphsa-AX\_A8\_12.out &

nohup IntaRNA -t AX\_A8\_13.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_13\_targets.txt &> nohuphsa-AX\_A8\_13.out &

nohup IntaRNA -t AX\_A8\_14.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_14\_targets.txt &> nohuphsa-AX\_A8\_14.out &

nohup IntaRNA -t AX\_A8\_15.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_15\_targets.txt &> nohuphsa-AX\_A8\_15.out &

nohup IntaRNA -t AX\_A8\_16.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_16\_targets.txt &> nohuphsa-AX\_A8\_16.out &

nohup IntaRNA -t AX\_A8\_17.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_17\_targets.txt &> nohuphsa-AX\_A8\_17.out &

nohup IntaRNA -t AX\_A8\_18.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_18\_targets.txt &> nohuphsa-AX\_A8\_18.out &

nohup IntaRNA -t AX\_A8\_19.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_19\_targets.txt &> nohuphsa-AX\_A8\_19.out &

nohup IntaRNA -t AX\_A8\_20.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_20\_targets.txt &> nohuphsa-AX\_A8\_20.out &

nohup IntaRNA -t AX\_A8\_21.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_21\_targets.txt &> nohuphsa-AX\_A8\_21.out &

nohup IntaRNA -t AX\_A8\_22.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_22\_targets.txt &> nohuphsa-AX\_A8\_22.out &

nohup IntaRNA -t AX\_A8\_23.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_23\_targets.txt &> nohuphsa-AX\_A8\_23.out &

nohup IntaRNA -t AX\_A8\_24.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_24\_targets.txt &> nohuphsa-AX\_A8\_24.out &

nohup IntaRNA -t AX\_A8\_25.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_25\_targets.txt &> nohuphsa-AX\_A8\_25.out &

nohup IntaRNA -t AX\_A8\_26.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_26\_targets.txt &> nohuphsa-AX\_A8\_26.out &

nohup IntaRNA -t AX\_A8\_27.txt -q human\_mature\_miRNA.fasta --outMode=C --threads=2 --out AX\_A8\_27\_targets.txt &> nohuphsa-AX\_A8\_27.out &