PM Code

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

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

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

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

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

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

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

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

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

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