

Supplementary Information

Novel methods for the rapid and sensitive detection of Nipah virus based on CRISPR/Cas12a system

Xi Yang, Kexin Xu, Siying Li, Jiangnian Zhang, Yinli Xie*, Yongliang Lou*, Xingxing Xiao*.

*Corresponding author: Yinli Xie, xieyinli@wmu.edu.cn; Yongliang Lou, lyl@wmu.edu.cn; Xingxing Xiao, xiaoxingxing@wmu.edu.cn.

Supplementary Texts

1. Seven candidate primer pairs targeting the conserved regions of *N* gene.

F1/R1:

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F1: *****
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KY425655.1 ATGAGTGTATATCTTTGAAGAGGCGGCTAGT
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F6: ** ***** ***** ***** **
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MK673578.1 GAGAGTGAAC TAGAAGATGGGCTAAGTAC
MK673582.1 GAGAGTGAAC TAGAAGATGGGCTAAGTAC
MK673577.1 GAGAGTGAAC TAGAAGATGGGCTAAGTAC
KT163252.1 GAGAGTGAAC TAGAAGATGGGCTAAGTAC
KT163255.1 GAGAGTGAAC TAGAAGATGGGCTAAGTAC
KT163254.1 GAGAGTGAAC TAGAAGATGGGCTAAGTAC
KT163251.1 GAGAGTGAAC TAGAAGATGGGCTAAGTAC
KT163256.1 GAGAGTGAAC TAGAAGATGGGCTAAGTAC
FJ513078.1 GAGAGTGAAC TAGAAGATGGGCTAAGTAC
NC_002728.1 GAAAGTGAAC TAGAAGATGGGCTAAATAC
KT163250.1 GAAAGTGAAC TAGAAGATGGGCTAAATAC
KT163253.1 GAGAGTGAAC TAGAAGATGGGCTAAGTAC
JN808858.1 GAGAGTGAAC TAGAAGATGGGCTAAGTAC
JN808860.1 GAGAGTGAAC TAGAAGATGGGCTAAGTAC
AY029768.1 GAAAGTGAAC TAGAAGATGGGCTAAATAC
AY029767.1 GAAAGTGAAC TAGAAGATGGGCTAAATAC
AJ627196.1 GAAAGTGAAC TAGAAGATGGGCTAAATAC
AJ564623.1 GAAAGTGAAC TAGAAGATGGGCTAAATAC
FN869553.1 GAAAGTGAAC TAGAAGATGGGCTAAATAC
AJ564622.1 GAAAGTGAAC TAGAAGATGGGCTAAATAC

R6: *** ***** ***** *****
AGTAGAAATAATCTCAGACATCGGAAACTA
KY425646.1 AGTAGAAATAATCTCAGACATCGGAAACTA
KY425655.1 AGTAGAAATAATCTCAGACATCGGAAACTA
MK673563.1 AGTAGAAATAATCTCAGACATCGGAAACTA
MK673561.1 AGTAGAAATAATCTCAGACATCGGAAACTA
MK673560.1 AGTAGAAATAATCTCAGACATCGGAAACTA
MK673559.1 AGTAGAAATAATCTCAGACATCGGAAACTA
MK673558.1 AGTAGAAATAATCTCAGACATCGGAAACTA
KT163249.1 AGTAGAAATAATCTCAGACATCGGAAACTA
KT163248.1 AGTAGAAATAATCTCAGACATCGGAAACTA
JN808861.1 AGTGGAAATAATCTCAGACATCGGAAACTA
AY858110.1 AGTAGAAATAATCTCAGACATCGGAAACTA
MK673565.1 AGTGGAAATAATCTCAGACATCGGAAACTA
MK673590.1 AGTGGAAATAATCTCAGACATCGGAAACTA
MK673589.1 AGTGGAAATAATCTCAGACATCGGAAACTA
JN808857.1 AGTGGAAATAATCTCAGACATCGGAAACTA
MK673570.1 AGTGGAAATAATCTCAGACATCGGAAACTA
MK673564.1 AGTGGAAATAATCTCAGACATCGGAAACTA
AY988601.1 AGTGGAAATAATCTCAGACATCGGAAACTA
JN808864.1 AGTGGAAATAATCTCAGACATCGGAAACTA
MK673583.1 AGTGGAAATAATCTCAGACATCGGAAACTA
MK673578.1 AGTGGAAATAATCTCAGACATCGGAAACTA
MK673582.1 AGTGGAAATAATCTCAGACATCGGAAACTA
MK673577.1 AGTGGAAATAATCTCAGACATCGGAAACTA
KT163252.1 AGTGGAAATAATCTCAGACATCGGAAACTA
KT163255.1 AGTGGAAATAATCTCAGACATCGGAAACTA
KT163254.1 AGTGGAAATAATCTCAGACATCGGAAACTA
KT163251.1 AGTGGAAATAATCTCAGACATCGGAAACTA
KT163256.1 AGTGGAAATAATCTCAGACATCGGAAACTA
FJ513078.1 AGTGGAAATAATCTCAGACATCGGAAACTA
NC_002728.1 AGTAGAAATAATCTCAGACATCGGAAACTA
KT163250.1 AGTAGAAATAATCTCAGACATCGGAAACTA
KT163253.1 AGTGGAAATAATCTCAGACATCGGAAACTA
JN808858.1 AGTGGAAATAATCTCAGACATCGGAAACTA
JN808860.1 AGTGGAAATAATCTCAGACATCGGAAACTA
AY029768.1 AGTAGAAATAATCTCAGACATCGGAAACTA
AY029767.1 AGTAGAAATAATCTCAGACATCGGAAACTA
AJ627196.1 AGTAGAAATAATCTCAGACATCGGAAACTA
AJ564623.1 AGTAGAAATAATCTCAGACATCGGAAACTA
FN869553.1 AGTAGAAATAATCTCAGACATCGGAAACTA
AJ564622.1 AGTAGAAATAATCTCAGACATCGGAAACTA

F7/R7:

F7: *** ***** ***** ***** **
GGAAAGTGAAACTAGAAGATGGGCTAAATAC
KY425646.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
KY425655.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
MK673563.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
MK673561.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
MK673560.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
MK673559.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
MK673558.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
KT163249.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
KT163248.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
JN808861.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
AY858110.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
MK673565.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
MK673590.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
MK673589.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
JN808857.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
MK673570.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
MK673564.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
AY988601.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
JN808864.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
MK673583.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
MK673578.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
MK673582.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
MK673577.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
KT163252.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
KT163255.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
KT163254.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
KT163251.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
KT163256.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
FJ513078.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
NC_002728.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
KT163250.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
KT163253.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
JN808858.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
JN808860.1 GGAGAGTGAAACTAGAAGATGGGCTAAGTAC
AY029768.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
AY029767.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
AJ627196.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
AJ564623.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
FN869553.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC
AJ564622.1 GGAAAGTGAAACTAGAAGATGGGCTAAATAC

R7: ***** ***** ***** *****
GCAGTAGAAATAATCTCAGACATCGGAAAC
KY425646.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
KY425655.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673563.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673561.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673560.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673559.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673558.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
KT163249.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
KT163248.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
JN808861.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
AY858110.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673565.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673590.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673589.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
JN808857.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673570.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673564.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
AY988601.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
JN808864.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673583.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673578.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673582.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
MK673577.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
KT163252.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
KT163255.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
KT163254.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
KT163251.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
KT163256.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
FJ513078.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
NC_002728.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
KT163250.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
KT163253.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
JN808858.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
JN808860.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
AY029768.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
AY029767.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
AJ627196.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
AJ564623.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
FN869553.1 GCAGTAGAAATAATCTCAGACATCGGAAAC
AJ564622.1 GCAGTAGAAATAATCTCAGACATCGGAAAC

2. Five candidate crRNAs targeting the conserved regions of *N* gene.

CR1:

	CR1: PAM	Spacer sequence
	TTTN	ACACUCAUCUCUAUGUAUU
KY425646.1	TTT	CACACTCATCTCTATGTATT
KY425655.1	TTT	CACACTCATCTCTATGTATT
MK673563.1	TTT	CACACTCATCTCTATGTATT
MK673561.1	TTT	CACACTCATCTCTATGTATT
MK673560.1	TTT	CACACTCATCTCTATGTATT
MK673559.1	TTT	CACACTCATCTCTATGTATT
MK673558.1	TTT	CACACTCATCTCTATGTATT
KT163249.1	TTT	CACACTCATCTCTATGTATT
KT163248.1	TTT	CACACTCATCTCTATGTATT
JN808861.1	TTTT	ACACTCATCTCTATGTATT
AY858110.1	TTTT	ACACTCATCTCTATGTATT
MK673565.1	TTTT	ACACTCATCTCTATGTATT
MK673590.1	TTTT	ACACTCATCTCTATGTATT
MK673589.1	TTTT	ACACTCATCTCTATGTATT
JN808857.1	TTTT	ACACTCATCTCTATGTATT
MK673570.1	TTTT	ACACTCATCTCTATGTATT
MK673564.1	TTTT	ACACTCATCTCTATGTATT
AY988601.1	TTTT	ACACTCATCTCTATGTATT
JN808864.1	TTTT	ACACTCATCTCTATGTATT
MK673583.1	TTTT	ACACTCATCTCTATGTATT
MK673578.1	TTTT	ACACTCATCTCTATGTATT
MK673582.1	TTTT	ACACTCATCTCTATGTATT
MK673577.1	TTTT	ACACTCATCTCTATGTATT
KT163252.1	TTTT	ACACTCATCTCTATGTATT
KT163255.1	TTTT	ACACTCATCTCTATGTATT
KT163254.1	TTTT	ACACTCATCTCTATGTATT
KT163251.1	TTTT	ACACTCATCTCTATGTATT
KT163256.1	TTTT	ACACTCATCTCTATGTATT
FJ513078.1	TTTT	ACACTCATCTCTATGTATT
NC_002728.1	TTT	CACACTCATCTCTATGTATT
KT163250.1	TTT	CACACTCATCTCTATGTATT
KT163253.1	TTTT	ACACTCATCTCTATGTATT
JN808858.1	TTTT	ACACTCATCTCTATGTATT
JN808860.1	TTT	CACACTCATCTCTATGTATT
AY029768.1	TTT	CACACTCATCTCTATGTATT
AY029767.1	TTT	CACACTCATCTCTATGTATT
AJ627196.1	TTT	CACACTCATCTCTATGTATT
AJ564623.1	TTT	CACACTCATCTCTATGTATT
FN869553.1	TTT	CACACTCATCTCTATGTATT
AJ564622.1	TTT	CACACTCATCTCTATGTATT

CR2:

	CR2: PAM	Spacer sequence
	TTTN	GGAGUUAUCAAUCUAAGUU
KY425646.1	TTTAGGAGTTATCAATCTAAGTT	
KY425655.1	TTTAGGAGTTATCAATCTAAGTT	
MK673563.1	TTTAGGAGTTATCAATCTAAGTT	
MK673561.1	TTTAGGAGTTATCAATCTAAGTT	
MK673560.1	TTTAGGAGTTATCAATCTAAGTT	
MK673559.1	TTTAGGAGTTATCAATCTAAGTT	
MK673558.1	TTTAGGAGTTATCAATCTAAGTT	
KT163249.1	TTTAGGAGTTATCAATCTAAGTT	
KT163248.1	TTTAGGAGTTATCAATCTAAGTT	
JN808861.1	TTTAGGAGTTATCAATCTAAGTT	
AY858110.1	TTTAGGAGTTATCAATCTAAGTT	
MK673565.1	TTTAGGAGTTATCAATCTAAGTT	
MK673590.1	TTTAGGAGTTATCAATCTAAGTT	
MK673589.1	TTTAGGAGTTATCAATCTAAGTT	
JN808857.1	TTTAGGAGTTATCAATCTAAGTT	
MK673570.1	TTTAGGAGTTATCAATCTAAGTT	
MK673564.1	TTTAGGAGTTATCAATCTAAGTT	
AY988601.1	TTTAGGAGTTATCAATCTAAGTT	
JN808864.1	TTTAGGAGTTATCAATCTAAGTT	
MK673583.1	TTTAGGAGTTATCAATCTAAGTT	
MK673578.1	TTTAGGAGTTATCAATCTAAGTT	
MK673582.1	TTTAGGAGTTATCAATCTAAGTT	
MK673577.1	TTTAGGAGTTATCAATCTAAGTT	
KT163252.1	TTTAGGAGTTATCAATCTAAGTT	
KT163255.1	TTTAGGAGTTATCAATCTAAGTT	
KT163254.1	TTTAGGAGTTATCAATCTAAGTT	
KT163251.1	TTTAGGAGTTATCAATCTAAGTT	
KT163256.1	TTTAGGAGTTATCAATCTAAGTT	
FJ513078.1	TTTAGGAGTTATCAATCTAAGTT	
NC_002728.1	TTTAGGAGTTATCAATCTAAGTT	
KT163250.1	TTTAGGAGTTATCAATCTAAGTT	
KT163253.1	TTTAGGAGTTATCAATCTAAGTT	
JN808858.1	TTTAGGAGTTATCAATCTAAGTT	
JN808860.1	TTTAGGAGTTATCAATCTAAGTT	
AY029768.1	TTTAGGAGTTATCAATCTAAGTT	
AY029767.1	TTTAGGAGTTATCAATCTAAGTT	
AJ627196.1	TTTAGGAGTTATCAATCTAAGTT	
AJ564623.1	TTTAGGAGTTATCAATCTAAGTT	
FN869553.1	TTTAGGAGTTATCAATCTAAGTT	
AJ564622.1	TTTAGGAGTTATCAATCTAAGTT	

CR3:

CR3: PAM Spacer sequence
TTTACAACCAAGAUAGGAU

KY425646.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
KY425655.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673563.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673561.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673560.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673559.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673558.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
KT163249.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
KT163248.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
JN808861.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
AY858110.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673565.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673590.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673589.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
JN808857.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673570.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673564.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
AY988601.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
JN808864.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673583.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673578.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673582.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
MK673577.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
KT163252.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
KT163255.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
KT163254.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
KT163251.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
KT163256.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
FJ513078.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
NC_002728.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
KT163250.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
KT163253.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
JN808858.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
JN808860.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
AY029768.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
AY029767.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
AJ627196.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
AJ564623.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
FN869553.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T
AJ564622.1	TTT	G	A	C	A	C	C	A	A	G	A	T	A	A	G	G	A	T	A	T

CR4:

CR4: PAM Spacer sequence
TTN CUGCAGGAGGUGUCUCAU

KY425646.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
KY425655.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673563.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673561.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673560.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673559.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673558.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
KT163249.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
KT163248.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
JN808861.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
AY858110.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673565.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673590.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673589.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
JN808857.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673570.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673564.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
AY988601.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
JN808864.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673583.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673578.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673582.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
MK673577.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
KT163252.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
KT163255.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
KT163254.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
KT163251.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
KT163256.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
FJ513078.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
NC_002728.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
KT163250.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
KT163253.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
JN808858.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
JN808860.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
AY029768.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
AY029767.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
AJ627196.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
AJ564623.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
FN869553.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT
AJ564622.1	TTT	GCTGC	AGG	AGG	TGT	GCTCAT

CR5:

	CR5:	Spacer sequence	PAM
		AGUUCUUUCCUCCUAGACG	NTTT
KY425646.1	T	CAAGAAAGGAGGATC	TGCTAAA
KY425655.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673563.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673561.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673560.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673559.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673558.1	T	CAAGAAAGGAGGATC	TGCTAAA
KT163249.1	T	CAAGAAAGGAGGATC	TGCTAAA
KT163248.1	T	CAAGAAAGGAGGATC	TGCTAAA
JN808861.1	T	CAAGAAAGGAGGATC	TGCTAAA
AY858110.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673565.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673590.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673589.1	T	CAAGAAAGGAGGATC	TGCTAAA
JN808857.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673570.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673564.1	T	CAAGAAAGGAGGATC	TGCTAAA
AY988601.1	T	CAAGAAAGGAGGATC	TGCTAAA
JN808864.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673583.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673578.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673582.1	T	CAAGAAAGGAGGATC	TGCTAAA
MK673577.1	T	CAAGAAAGGAGGATC	TGCTAAA
KT163252.1	T	CAAGAAAGGAGGATC	TGCTAAA
KT163255.1	T	CAAGAAAGGAGGATC	TGCTAAA
KT163254.1	T	CAAGAAAGGAGGATC	TGCTAAA
KT163251.1	T	CAAGAAAGGAGGATC	TGCTAAA
KT163256.1	T	CAAGAAAGGAGGATC	TGCTAAA
FJ513078.1	T	CAAGAAAGGAGGATC	TGCTAAA
NC_002728.1	T	CAAGAAAGGAGGATC	TGCTAAA
KT163250.1	T	CAAGAAAGGAGGATC	TGCTAAA
KT163253.1	T	CAAGAAAGGAGGATC	TGCTAAA
JN808858.1	T	CAAGAAAGGAGGATC	TGCTAAA
JN808860.1	T	CAAGAAAGGAGGATC	TGCTAAA
AY029768.1	T	CAAGAAAGGAGGATC	TGCTAAA
AY029767.1	T	CAAGAAAGGAGGATC	TGCTAAA
AJ627196.1	T	CAAGAAAGGAGGATC	TGCTAAA
AJ564623.1	T	CAAGAAAGGAGGATC	TGCTAAA
FN869553.1	T	CAAGAAAGGAGGATC	TGCTAAA
AJ564622.1	T	CAAGAAAGGAGGATC	TGCTAAA

Table S1. RAA primer pairs used in this study.

Primer	Sequence (5'-3')	Product size
F1	ATGAGTGATATCTTTGAAGAGGGCGGCTAGT	325
R1	TGACCATTGATCCAACATCTATTATTACAGCTTC	
F2	TAGGAGTTATCAATCTAAGTTAGGGAGAGATG	291
R2	ACCATTGATCCAACATCTATTATTACAGCTTC	
F3	GGAGTTATCAATCTAAGTTAGGGAGAGATG	213
R3	TCTCTGAATACATAGAGATGAGTGTGAAAG	
F4	CAAATAGACTGGGACTAAGTTCAGATCAAG	225
R4	ATGAAATACTCATCTCCCTTTGAAGGTAA	
F5	GAGGAATTGATCAGAACATGGCAAATAGAC	245
R5	TGAAATACTCATCTCCCTTTGAAGGTAAC	
F6	GAAAGTGAAACTAGAAGATGGGCTAAATAC	212
R6	TAGTTTCCGATGTCTGAGATTATTTCTACT	
F7	GGAAAGTGAAACTAGAAGATGGGCTAAATAC	211
R7	GTTTCCGATGTCTGAGATTATTTCTACTGC	

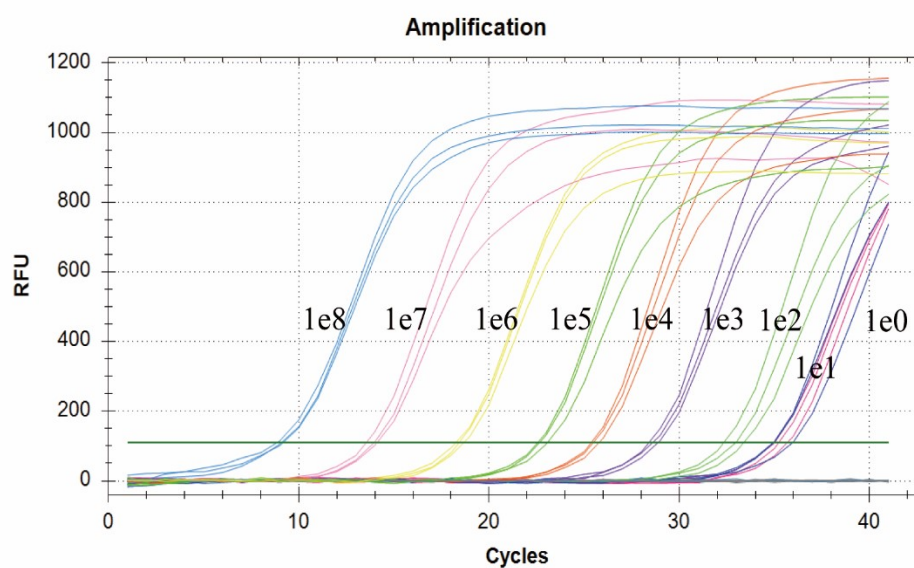


Figure S1. The sensitivity of the qPCR assay in NiV detection. qPCR was conducted with the CFX96 real-time PCR detection systems, and the amplification curves of each sample were shown in this figure.