

# Suggested pre-NGS wet-lab protocol

Flu A Multi-Segment RT-PCR			
RT-PCR			
ddH <sub>2</sub> O	10 µl		
2x RT-PCR Buffer	25 µl	42°	60'
Uni12/Inf1 10µM	0,8 µl	94°	2'
Uni12/Inf3 10µM	1,2 µl	94°	30"
Uni13/Inf1 10µM	2 µl	44°	30"
SSIII/Platinum Taq HiFi Enzyme Mix	1 µl	68°	3'
Mix	40 µl	94°	30"
RNA	10 µl	57°	30"
Total	50 µl	68°	3'
		68°	5'
		4°	∞

Flu B Multi-Segment RT-PCR			
RT-PCR			
ddH <sub>2</sub> O	7 µl	45°	60'
2x RT-PCR Buffer	12,5 µl	55°	30'
FluB universal primer cocktail*	2 µl	94°	2'
SSIII/Platinum Taq HiFi Enzyme Mix	0,5 µl	94°	20"
Mix	22 µl	40°	30"
RNA	3 µl	68°	3'30"
Total	25 µl	94°	20"
		58°	30"
		68°	3'30"
		68°	10'
		4°	∞

* Universal IBV-GA2 primer cocktail	
B-PBs-UniF (10µM)	100 µl
B-PBs-UniR (10µM)	100 µl
B-PA-UniF (10µM)	50 µl
B-PA-UniR (10µM)	50 µl
B-HANA-UniF (10µM)	100 µl
B-HANA-UniR (10µM)	100 µl
B-NP-UniF (10µM)	60 µl
B-NP-UniR (10µM)	60 µl
B-M-UniF (10µM)	30 µl
B-Mg-UniF (10µM)	30 µl
B-M-UniR (10µM)	60 µl
B-NS-UniF (10µM)	50 µl
B-NS-UniR (10µM)	50 µl
<b>FluB universal primer cocktail</b>	<b>840 µl</b>

Adapted\* from a RT-PCR assay described by Zhou and colleagues (Zhou et al, 2009, for Influenza A; Zhou et al, 2014, for Influenza B; Zhou and Wentworth DE, 2012).

## REFERENCES:

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- Zhou B, Lin X, Wang W, Halpin RA, Bera J, Stockwell TB, Barr IG, Wentworth DE. 2014. Universal influenza B virus genomic amplification facilitates sequencing, diagnostics, and reverse genetics. J Clin Microbiol, 52:1330-1337.
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