

# Supplementary Materials: Broad Impact of Exchange Protein Directly Activated by cAMP 2 (EPAC2) on Respiratory Viral Infections

**Table S1.** Primers for reverse transcription and qRT-PCR.

Primers for Reverse Transcription		
Name		Sequence (5' to 3')
HMPV genome		CGTCTCAGCCAATCCCTGGTGATTATGAGTAATTAAAAAAVTGGGACAAG
RSV genome		CTGCGATGAGTGGCAGGCACTACAGTGTATTAGACTTRACAGCAGAAG
HMPV-N		CGTCTCAGCCAATCCCTGGTTTTTTTTTTTTTAAATTACTC
RSV-N		CTGCGATGAGTGGCAGGCTTTTTTTTTTTTTAACTYAAAGCTC
Primers for qRT-PCR		
Name		Sequence (5' to 3')
HMPV genome	Forward	GCTTCATTACCCATGAAMAGAATATC
	Reverse	CGTCTCAGCCAATCCCTGG
RSV genome	Forward	GCATCTTCTCCATGRAATTCAGG
	Reverse	CTGCGATGAGTGGCAGGC
HMPV-N	Forward	CACAGARCTATTTTCWGCAGCAG
	Reverse	CGTCTCAGCCAATCCCTGG
RSV-N	Forward	ACTACAGTGTATTAGACTTRACAGCAGAAG
	Reverse	CTGCGATGAGTGGCAGGC
AdV-DBP	Forward	GCCATTGCGCCCAAGAAGAA
	Reverse	CTGTCCACGATTACCTCTGGTGAT
IP-10	Forward	GTGGCATTCAAGGAGTACCTC
	Reverse	GCCTTCGATTCTGGATTCAGACA
RANTES	Forward	CCTGCTGCTTTGCCTACATTGC
	Reverse	ACACACTTGGCGGTTCTTTCGG
18S	Forward	ACATCCAAGGAAGGCAGCAG
	Reverse	TCGTCACTACCTCCCCGG
GAPDH	Forward	CTCAAGATCATCAGCAATGCCT
	Reverse	AAGTTGTCATGGATGACCTTGG