

TABLE S1

LIMITING DILUTION DATA FROM 5'UTR AND CODING REGION
NESTED PCRs USED FOR VIRUS TYPING

Virus Species	Region	RNA copies / reaction								Calculated copies/ μ l
		600000	60000	6000	600	60	6	0.6	0.06	
<i>Enterovirus</i>										
A	VP1	6/6	6/6	6/6	6/6	0/6	0/6	0/6	0/6	275.0
B	VP1	6/6	6/6	6/6	6/6	6/6	4/6	0/6	0/6	4.9
C	VP2	6/6	6/6	6/6	3/6	0/6	0/6	0/6	0/6	869.7
D	VP1	6/6	6/6	6/6	5/6	1/6	0/6	0/6	0/6	275.0
<i>Parechovirus</i>										
HPeV1	VP1	6/6	6/6	6/6	6/6	6/6	2/6	0/6	0/6	15.4
<i>Enterovirus</i>										
A	VP4	6/6	6/6	6/6	6/6	2/6	0/6	0/6	0/6	154.4
B	VP4	6/6	6/6	6/6	6/6	6/6	0/6	0/6	0/6	27.5
C	VP4	6/6	6/6	6/6	6/6	5/6	2/6	0/6	0/6	20.1
D	VP4	6/6	6/6	6/6	6/6	5/6	0/6	0/6	0/6	34.6
<i>Rhinovirus</i>										
HRVA	VP4	6/6	6/6	6/6	6/6	0/6	0/6	0/6	0/6	275.0
HRVB	VP4	6/6	6/6	6/6	6/6	6/6	6/6	0/6	0/6	2.7
HRVCc	VP4	6/6	6/6	6/6	4/6	0/6	0/6	0/6	0/6	489.7
HRVCa	VP4	6/6	6/6	6/6	4/6	3/6	0/6	0/6	0/6	178.1
<i>Enterovirus</i>										
A	5'UTR	6/6	6/6	6/6	6/6	6/6	6/6	2/6	0/6	1.5
B	5'UTR	6/6	6/6	6/6	6/6	6/6	6/6	4/6	0/6	0.5
C	5'UTR	6/6	6/6	6/6	6/6	6/6	0/6	0/6	0/6	27.5
D	5'UTR	6/6	6/6	6/6	6/6	6/6	5/6	1/6	0/6	2.7
<i>Rhinovirus</i>										
HRVA	5'UTR	6/6	6/6	6/6	1/6	0/6	0/6	0/6	0/6	2183.4
HRVB	5'UTR	6/6	6/6	6/6	6/6	6/6	3/6	1/6	0/6	6.1
HRVCa	5'UTR	6/6	6/6	6/6	6/6	2/6	0/6	0/6	0/6	154.4
HRVCc	5'UTR	6/6	6/6	6/6	6/6	6/6	5/6	4/6	0/6	0.5
<i>Parechovirus</i>										
HPeV-1	5'UTR	6/6	6/6	6/6	6/6	6/6	6/6	0/6	0/6	2.7