

## Appendix

Table I. Primers and probes sequences used for respiratory viral targets.

Target Primers and Probe set	Sequence	Sequence Reference and supplier
<b><u>FLU A</u></b>		World Health Organisation, 2007. SA
11A forward	5'-AAGACCAATCCTGTACCTCTGA-3'	
11B reverse	5'-CAAAGCGTCTACGCTGCAGTCC-3'	
11P probe	5'-FAM-TTTGTGTTCCACGCTCACCGTGCC-BHQ-3'	
<b><u>FLU A H1</u></b>		Personal communication. Anon. HPA, Colindale. SOP V-5413. LT
16A forward	5'-GGAATAGCCCCCTACAATTG-3'	
16B reverse	5'-AATTTCGCATTCTGGGTTTCCTA-3'	
16P probe	5'-FAM-CGTTGCCGGATGGA-MGBNFQ-3'	
<b><u>FLU A H3</u></b>		Personal communication. Anon. HPA, Colindale. SOP V-5413. LT
17A forward	5'-CCTTTTTGTTGAACGCAGCAA-3'	
17B reverse	5'-CGGATGAGGCAACTAGTGACCTA-3'	
17P probe	5'-FAM-CCTACAGCAACTGTTACC-MGBNFQ-3	
<b><u>FLU A H5</u></b>		Health Protection Agency (2009). LT
9A forward	5'-GGATGGCAGGGAATGGTAGA-3'	
9B reverse	5'-TCTATTGCCTTTTGAGTGGATTCT T-3'	
9P probe	5'-FAM-TGGGTACCACCATAGCAAYGAGCAGG-MGBNFQ-3'	
<b><u>FLU B</u></b>		Van Elden, L J R et al, 2001. SA
4A forward	5'-AAATACGGTGGATTAATAAAAAGCAA-3'	
4B reverse	5'-CCAGCAATAGCTCCGAAGAAA-3'	
4P probe	FAM-5'-CACCCATATTGGGCAATTTCTATGGC-3'-TAMRA	
<b><u>RSV A</u></b>		Gunson, R N et al, 2005. SA
4A forward	5'-AGATCAACTTCTGTCCATCCAGCAA-3'	
4B reverse	5'-TTCTGCACATCATAATTAGGAG TATCAAT-3'	
4P probe	5'-FAM-CACCATCCAACGGAGCACAGGAGAT-BHQ-3'	
<b><u>RSV B</u></b>		Personal Communication, Anon. Specialist Virology Centre, Gartnavel. SA
5A forward	5'-AAGATGCAAATCATAAATTCACAGGA-3'	
5B reverse	5'-TGATATCCAGCATCTTTCAGTATCTTTATA-3'	
5P1 probe	5'-FAM-TTCCCTTCCTAACCTGGACATAGCATATAACATACCT-BHQ-3'	
<b><u>MPV 1A</u></b>		Kuypers, J et al. 2005. LT
3A forward	5'-GCCGTTAGCTTCAGTCAATTCAA-3'	
3B reverse	5'-TCCAGCATTGTCTGAAAATTGC-3'	
3P probe	5'-FAM-CAACATTTAGAAACCTTCT-MGBNFQ-3'	
<b><u>MPV 2B</u></b>		Kuypers, J et al. 2005. LT
3A forward	5'-GCCGTTAGCTTCAGTCAATTCAA-3'	
3B reverse	5'-TCCAGCATTGTCTGAAAATTGC-3'	
4P probe	5'-FAM-CGCACAACATTTAGGAATCTTCT-MGBNFQ-3'	
<b><u>PF1</u></b>		Personal Communication, Anon. Specialist Virology Centre, Gartnavel. LT
2A forward	5'-GTGATTTAAACCCGGTAATTTCTCA-3'	
2B reverse	5'-CCTTGTTCCCTGCAGCTATTACAAGA-3'	
2P probe	5'-FAM-ACCTATGACATCAACGAC-MGBNFQ-3'	
<b><u>PF2</u></b>		Personal Communication, Anon. Specialist Virology Centre, Gartnavel. LT
2A forward	5'-ATGAAAACCATTTACCTAAGTGATGGA-3'	
2B reverse	5'-CCTCCYGGTATRGAGTGAAC-3'	
2P probe	5'-FAM-TCAATCGCAAAGC-MGBNFQ-3'	

<b><u>PF3</u></b>		Gunson, R N et al, 2005. SA
<b>3A forward</b>	5'-CCAGGGATATAYTAYAAAGGCAAAA-3'	
<b>3B reverse</b>	5'-CCGGGRCACCCAGTTGTG-3'	
<b>3P probe</b>	5'-FAM-TGGRTGTTCAAGACCTCCATAYCCGAGAAA-BHQ-3'	
<b><u>COV 229E</u></b>		Gunson, R N et al, 2005. SA
<b>6A forward</b>	5'-CAGTCAAATGGGCTGATGCA-3'	
<b>6B reverse</b>	5'-AAAGGGCTATAAAGAGAATAAGGTATTCT-3'	
<b>6P probe</b>	5'-FAM-CCCTGACGACCACGTTGTGGTTCA-BHQ-3'	
<b><u>COV OC43</u></b>		Gunson, R N et al, 2005. SA
<b>7A forward</b>	5'-CGATGAGGCTATTCCGACTAGGT-3'	
<b>7B reverse</b>	5'-CCTTCCTGAGCCTTCAATATAGTAACC-3'	
<b>7P probe</b>	5'-FAM-TCCGCCTGGCACGGTACTCCCT-BHQ-3'	
<b><u>COV NL63</u></b>		Gunson, R N et al, 2005. SA
<b>8A forward</b>	5'-ACGTACTIONCTATTATGAAGCATGATATTA-3'	
<b>8B reverse</b>	5'-AGCAGATCTAATGTTATACTTAAACTACG-3'	
<b>8P1 probe</b>	5'-FAM-ATTGCCAAGGCTCCTAACGTACAGGTGTT-BHQ-3'	
<b><u>COV HKU1</u></b>		Raymond, F et al, 2009. LT
<b>9A forward</b>	5'-TTGAAGGCTCAGGAAGGTCTGCT-3'	
<b>9B reverse</b>	5'-TGCTTAGTKACTTGCTGAGGTTTAG -3'	
<b>9P probe</b>	5'-6FAM-TAAAACAAGATTAGCGATCTC-MGBNFQ-3'	
<b><u>ADV</u></b>		Heim, A et al, 2003. SA
<b>3A forward</b>	5'-GCCACGGTGGGGTTTCTAAACTT-3'	
<b>3B reverse</b>	5'-GCCCCAGTGGTCTTACATGCACATC-3'	
<b>3P probe</b>	5'-FAM-TGCACCAGMCCSGGGCTCAGGTACTCCGA-BHQ-3'	
<b><u>HRV 1</u></b>		In-house development. Regional Virus Laboratory, Belfast. LT
<b>7A forward</b>	5'-AGCCTGCGTGGCTGCCTG-3'	
<b>7A2 forward</b>	5'-CCTGCGTGGCGGCCARC-3'	
<b>7B reverse</b>	5'-CCCAAAGTAGTYGGTCCCRCTCC-3'	
<b>7P probe</b>	5'-FAM-TCCTCCGGCYCCTGAATG-3'-MGBNFQ	
<b><u>HRV 2</u></b>		Scheltinga, S A et al, 2005. SA
<b>8A forward</b>	5'-GACARGGTGTGAAGAGCC-3'	
<b>8B forward</b>	5'-GACATGGTGTGAAGACYC-3'	
<b>8C reverse</b>	5'-CAAAGTAGTYGGTCCCATCC'-3	
<b>8P probe</b>	5'-FAM-TCCTCCGGCCCCCTGAATGYGGCTAA-3'-BHQ1	
<b><u>BOV</u></b>		Allander, T et al, 2007. SA
<b>2A forward</b>	5'-GGAAGAGACACTGGCAGACAA-3'	
<b>2B reverse</b>	5'-GGGTGTTCTGATGATATGAGC-3'	
<b>2P probe</b>	5'-FAM-CTGCGGCTCCTGCTCCTGTGAT-BHQ-3'	
<b><u>RNP</u></b>		World Health Organisation, 2009. SA
<b>1A forward</b>	5'-AGATTTGGACCTGCGAGCG-3'	
<b>1B reverse</b>	5'-GAGCGGCTGTCTCCACAAGT-3'	
<b>1P probe</b>	5'-6FAM-TTCTGACCTGAAGGCTCTGCGCG-BHQ-3'	

For each target in the test panel, the forward and reverse primers, probe sequences, reporters and quenchers are presented, along with their initial source references and supplier details.

Abbreviations: "ADV: adenovirus; BHQ: Blackhole Quencher;BOV: bocavirus; FAM: 6-carboxyfluorescein (reporter molecule); COV: coronavirus; FLU A: influenza A; FLU B: influenza B; LT: Life Technologies (Paisley, UK); MGBNFQ: minor groove binding nonfluorescent quencher; MPV: metapneumovirus; PF:

parainfluenza; RHV: rhinovirus; RNP: Ribonuclease P; RSV: Respiratory Syncytial Virus; SA: Sigma-Aldrich Company Ltd(Dorset, UK); TAMRA: Carboxytetramethylrhodamine”.

**Table 1. List of reagents used in the preparation of respiratory viral-related amplification mixes.**

Each provides instruction for the preparation of 1000 tests when reaction volume used is 8µl. All volumes shown are µl.

All RNA target TaqMan master mixes were prepared using SuperScript® III Platinum® One-Step Quantitative RT-PCR System (Life Technologies, Paisley, UK).

**Influenza A Matrix**

Nuclease free water	<b>1580</b>
Reaction mix x2	<b>5000</b>
BSA @ 2µg/µl (final 0.2µg/µl)	<b>1000</b>
MgSO <sub>4</sub> @ 50mM (final 4mM)	<b>160</b>
FLA 11A @ 200µM (final 0.4µM)	<b>20</b>
FLA 11B @ 200µM (final 0.4µM)	<b>20</b>
FLA 11P @ 100µM (final 0.2µM) FAM-BHQ	<b>20</b>
Superscript III RT/Platinum® Taq Mix	<b>200</b>

**Influenza A H1(seasonal)**

NFW	<b>1580</b>
Reaction mix x2	<b>5000</b>
BSA @ 2µg/µl (final 0.2µg/µl)	<b>1000</b>
MgSO <sub>4</sub> @ 50mM (final 4mM)	<b>160</b>
FLA 16A @ 200µM (final 0.4µM)	<b>20</b>
FLA 16B @ 200µM (final 0.4µM)	<b>20</b>
FLA 16P @ 100µM (final 0.2µM) FAM-TAMRA	<b>20</b>
Superscript III RT/Platinum® Taq Mix	<b>200</b>

**Influenza A H3**

NFW	<b>1580</b>
Reaction Mix x2	<b>5000</b>
BSA @ 2µg/µl (final 0.2µg/µl)	<b>1000</b>
MgSO <sub>4</sub> @ 50mM (final 4mM)	<b>160</b>
FLA 17A @ 200µM (final 0.4µM)	<b>20</b>
FLA 17B @ 200µM (final 0.4µM)	<b>20</b>
FLA 17P @ 100µM (final 0.2µM) FAM-TAMRA	<b>20</b>
Superscript III RT/Platinum® Taq Mix	<b>200</b>

**Influenza A H5**

NFW	<b>1580</b>
Reaction Mix x2	<b>5000</b>
BSA @ 2µg/µl (final 0.2µg/µl)	<b>1000</b>
MgSO <sub>4</sub> @ 50m (final 4mM)	<b>160</b>
FLA 9A @ 200µM (final 0.4µM)	<b>20</b>
FLA 9B @ 200µM (final 0.4µM)	<b>20</b>
FLA 9P @ 100µM (final 0.2µM) FAM-MGBNFQ	<b>20</b>
Superscript III RT/Platinum® Taq Mix	<b>200</b>

**TaqMan PCR: Influenza B**

NFW	<b>1580</b>
Reaction Mix x2	<b>5000</b>
BSA @ 2µg/µl (final 0.2µg/µl)	<b>1000</b>
MgSO <sub>4</sub> @ 50mM (final 4mM)	<b>160</b>

FLB 4A @ 200µM (final 0.4µM)	20
FLB 4B @ 200µM (final 0.4µM)	20
FLB 4P @ 100µM (final 0.2µM) FAM-TAMRA	20
Superscript III RT/Platinum® Taq Mix	200

#### TaqMan PCR: RSV Types A&B

NFW	1680
Reaction Mix x2	5000
BSA @ 2µg/µl (final 0.2µg/µl)	1000
RSV 4A @ 200µM (final 0.4µM)	20
RSV 4B @ 200µM (final 0.4µM)	20
RSV 4P @ 100µM (final 0.2µM) FAM-BHQ	20
RSV 5A @ 200µM (final 0.4µM)	20
RSV 5B @ 200µM (final 0.4µM)	20
RSV 5P1 @ 100µM (final 0.2µM) FAM-BHQ	20
Superscript III RT/Platinum® Taq Mix	200

#### TaqMan PCR: Metapneumovirus

NFW	1520
Reaction Mix x2	5000
BSA @ 2µg/µl (final 0.2µg/µl)	1000
MgSO <sub>4</sub> @ 50mM (final 4mM)	160
MPV 3A @ 200µM (final 0.4µM)	20
MPV 3B @ 200µM (final 0.4µM)	20
MPV 3P @ 100µM (final 0.2µM) FAM-MGB	20
MPV 3A @ 200µM (final 0.4µM)	20
MPV 4B @ 200µM (final 0.4µM)	20
MPV 4P @ 100µM (final 0.2µM) FAM-MGBNFQ	20
Superscript III RT/Platinum® Taq Mix	200

#### TaqMan PCR: Rhinovirus1

NFW	1560
Reaction mix x2	5000
BSA @ 2µg/µl (final 0.2µg/µl)	1000
MgSO <sub>4</sub> @ 50mM (final 4mM)	160
HRV 7A @ 200µM (final 0.4µM)	20
HRV 7A2 @ 200µM (final 0.4µM)	20
HRV 7B @ 200µM (final 0.4µM)	20
HRV 7P @ 100µM (final 0.2µM) FAM-MGB	20
Superscript III RT/Platinum® Taq Mix	200

#### TaqMan PCR: Rhinovirus2

NFW	1660
Reaction mix x2	5000
BSA @ 2µg/µl (final 0.2µg/µl)	1000
MgSO <sub>4</sub> @ 50mM (final 3.5mM)	80
HRV 8A @ 200µM (final 0.2µM)	10
HRV 8B @ 200µM (final 0.2µM)	10
HRV 8C @ 200µM (final 0.4µM)	20
HRV 8P @ 100µM (final 0.2µM) FAM-BHQ	20

Superscript III RT/Platinum® Taq Mix	<b>200</b>
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#### RNP

NFW	<b>1700</b>
Reaction mix x2	<b>5000</b>
BSA @ 2µg/µl (final 0.2µg/µl)	<b>1000</b>
RNP 1A @ 200µM (final 0.8µM)	<b>40</b>
RNP 1B @ 200µM (final 0.8µM)	<b>40</b>
RNP 1P @ 100µM (0.2µM) FAM-BHQ	<b>20</b>
Superscript III RT/Platinum® Taq Mix	<b>200</b>

#### TaqMan PCR: Single Mix: Parainfluenza 1

NFW	<b>1580</b>
Reaction mix x2	<b>5000</b>
BSA @ 2µg/µl (final 0.2µg/µl)	<b>1000</b>
MgSO <sub>4</sub> @ 50mM (final 4mM)	<b>160</b>
PF1 2A @ 200µM (final 0.4µM)	<b>20</b>
PF1 2B @ 200µM (final 0.4µM)	<b>20</b>
PF1 2P @ 100µM (final 0.2µM) FAM-MGBNFQ	<b>20</b>
Superscript III RT/Platinum® Taq Mix	<b>200</b>

#### TaqMan PCR: Single Mix: Parainfluenza 2

NFW	<b>1580</b>
Reaction mix x2	<b>5000</b>
BSA @ 2µg/µl (final 0.2µg/µl)	<b>1000</b>
MgSO <sub>4</sub> @ 50mM (final 4mM)	<b>160</b>
PF2 2A @ 200µM (final 0.4µM)	<b>20</b>
PF2 2B @ 200µM (final 0.4µM)	<b>20</b>
PF2 2P @ 100µM (final 0.2µM) FAM-MGBNFQ	<b>20</b>
Superscript III RT/Platinum® Taq Mix	<b>200</b>

#### TaqMan PCR: Single Mix: Parainfluenza 3

NFW	<b>1580</b>
Reaction Mix x2	<b>5000</b>
BSA @ 2µg/µl (final 0.2µg/µl)	<b>1000</b>
MgSO <sub>4</sub> @ 50mM (final 4mM)	<b>160</b>
PF3 3A @ 200µM (final 0.4µM)	<b>20</b>
PF3 3B @ 200µM (final 0.4µM)	<b>20</b>
PF3 3P @ 100µM (final 0.2µM) FAM-BHQ	<b>20</b>
Superscript III RT/Platinum® Taq Mix	<b>200</b>

#### TaqMan PCR: Parainfluenza MULTIPLX

NFW	<b>1460</b>
Buffer @ x2	<b>5000</b>
BSA @ 2µg/µl (final 0.2µg/µl)	<b>1000</b>
MgSO <sub>4</sub> @ 50m (final 4mM)	<b>160</b>
PF1 2A @ 200µM (final 0.4µM)	<b>20</b>
PF1 2B @ 200µM (final 0.4µM)	<b>20</b>
PF1 2P @ 100µM (final 0.2µM) FAM-MGBNFQ	<b>20</b>
PF2 2A @ 200µM (final 0.4µM)	<b>20</b>

PF2 2B @ 200µM (final 0.4µM)	20
PF2 2P @ 100µM (final 0.2µM) FAM-MGBNFQ	20
PF3 3A @ 200µM (final 0.4µM)	20
PF3 3B @ 200µM (final 0.4µM)	20
PF3 3P @ 100µM (final 0.2µM) FAM-BHQ	20
Superscript III RT/Platinum® Taq Mix	200

**TaqMan PCR: Coronavirus 229E, OC43, NL63and HKU1 MULTIPLEX**

NFW	1400
Buffer @ x2	5000
BSA @ 2µg/µl (final 0.2µg/µl)	1000
MgSO <sub>4</sub> @ 50m (final 4mM)	160
COV 6A @ 200µM (final 0.4µM)	20
COV 6B @ 200µM (final 0.4µM)	20
COV 6P @ 100µM (final 0.2µM) FAM-MGBNFQ	20
COV 7A @ 200µM (final 0.4µM)	20
COV 7B @ 200µM (final 0.4µM)	20
COV 7P @ 100µM (final 0.2µM) FAM-MGBNFQ	20
COV 8A @ 200µM (final 0.4µM)	20
COV 8B @ 200µM (final 0.4µM)	20
COV 8P1@ 100µM (final 0.2µM) FAM-BHQ	20
COV 9A @ 200µM (final 0.4µM)	20
COV 9B @ 200µM (final 0.4µM)	20
COV 9P@ 100µM (final 0.2µM) FAM-MGB	20
Superscript III RT/Platinum® Taq Mix	200

**TaqMan PCR: Single Mix: Coronavirus 229E**

NFW	1580
Buffer @ x2	5000
BSA @ 2µg/µl (final 0.2µg/µl)	1000
MgSO <sub>4</sub> @ 50m (final 4mM)	160
COV 6A @ 200µM (final 0.4µM)	20
COV 6B @ 200µM (final 0.4µM)	20
COV 6P @ 100µM (final 0.2µM) FAM-MGBNFQ	20
Superscript III RT/Platinum® Taq Mix	200

**TaqMan PCR: Single Mix: Coronavirus OC43**

NFW	1580
Buffer @ x2	5000
BSA @ 2µg/µl (final 0.2µg/µl)	1000
MgSO <sub>4</sub> @ 50m (final 4mM)	160
COV 7A @ 200µM (final 0.4µM)	20
COV 7B @ 200µM (final 0.4µM)	20
COV 7P @ 100µM (final 0.2µM) FAM-MGBNFQ	20
Superscript III RT/Platinum® Taq Mix	200

**TaqMan PCR: Single Mix: Coronavirus NL63**

NFW	1580
Buffer @ x2	5000
BSA @ 2µg/µl (final 0.2µg/µl)	1000
MgSO <sub>4</sub> @ 50m (final 4mM)	160
COV 8A @ 200µM (final 0.4µM)	20
COV 8B @ 200µM (final 0.4µM)	20

COV 8P1 @ 100µM (final 0.2µM) FAM-BHQ1	<b>20</b>
Superscript III RT/Platinum® Taq Mix	<b>200</b>

**Taqman PCR: Single Mix: Coronavirus HKU1**

NFW	<b>1580</b>
Buffer @ x2	<b>5000</b>
BSA @ 2µg/µl (final 0.2µg/µl)	<b>1000</b>
MgSO <sub>4</sub> @ 50m (final 4mM)	<b>160</b>
COV 9A @ 200µM (final 0.4µM)	<b>20</b>
COV 9B @ 200µM (final 0.4µM)	<b>20</b>
COV 9P @ 100µM (final 0.2µM) FAM-BHQ1	<b>20</b>
Superscript III RT/Platinum® Taq Mix	<b>200</b>

All DNA target TaqMan amplification mixes were prepared with Platinum® Quantitative PCR SuperMix-UDG (Life Technologies, Paisley, UK).

**TaqMan PCR: Bocavirus DNA**

NFW	<b>1740</b>
Platinum® qPCR Supermix-UDG	<b>5000</b>
BSA @ 2µg/µl (final 0.2µg/µl)	<b>1000</b>
MgCl <sub>2</sub> 50mM (final 4mM)	<b>200</b>
BOV 2A @ 200µM (final 0.4µM)	<b>20</b>
BOV 2B @ 200µM (final 0.4µM)	<b>20</b>
BOV 2P @ 100µM (final 0.2µM) FAM-TAMRA	<b>20</b>

**TaqMan PCR: Adenovirus DNA**

NFW	<b>1740</b>
Platinum® qPCR Supermix-UDG	<b>5000</b>
BSA @ 2µg/µl (final 0.2µg/µl)	<b>1000</b>
MgCl <sub>2</sub> @ 50mM (final 4mM)	<b>200</b>
ADV 3A @ 200µM (final 0.4µM)	<b>20</b>
ADV 3B @ 200µM (final 0.4µM)	<b>20</b>
ADV 3P @ 100µM (final 0.2µM) FAM-BHQ	<b>20</b>

Abbreviations: "ADV: adenovirus; BHQ: Blackhole Quencher; BOV: bocavirus; BSA: Bovine serum albumin; COV: coronavirus; FAM: 6-carboxyfluorescein; FLU A: influenza A; FLU B: influenza B; MGBNFQ: minor groove binding nonfluorescent quencher; MgCl<sub>2</sub>: Magnesium chloride; MgSO<sub>4</sub>: Magnesium sulphate; MPV: metapneumovirus; NFW: nuclease free water; PF: parainfluenza; RHV: rhinovirus; RSV: Respiratory Syncytial Virus; TAMRA: Carboxytetramethylrhodamine.



**Table 3. Primers used in reverse-tagging step (PCR 2).**

<b>Name</b>	<b>Sequence</b>	<b>Length</b>
806R_f1	GTGAaGGAGHcAGAcGTGTGCTCCGATCTN NNNNACGGACTACHV GGG TW TCTAAT	61
806R_f2	GTGACTGGAGUCAGACGTGTGCTCUCCGATCTNNTNNNACGGACTACHV GGGTWTCTAAT	62
806R_f3	GTGAaGGAGUcAGACGTGTGAmCCGATCT NNCTNN NAC GGACTACHV GCGTWTCTAAT	63
806R_f4	GTGACTGGAGUCAGACGTGTGCTmCCGATCTNNACTNNNACGGACTACHV GGGTWTCTAA	64
806R_f5	GTGACTGGAGTrCAGACGTGTGCTCTrCCGATCTNNGAaNNNACGGAaACHW GGGTWTCTAAT	65
806R_f6	GTGACTGGAGTTCAGACGTGTGCTCTCCGATCTNNGACTNNNACGGACTACHV GGGTWTCTAAT	66

**Table 4. Primers for forward-tagging step (PCR 3).**

<b>Name</b>	<b>Sequence</b>	<b>Length</b>
515F_f1	GCCTCCCTCGCGCCATCAGAGATGTG TATAAGAGACAG NNNN NNNNGA GTGCCA GCMGCCGCGGTAA	67
515F_f2	GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAG NNNNNNNN GAGTGCCA GCM GCCGCGGTAA	68
515F_f3	GCCTCCCTCGCGCCATCAGAGATGTG TATAAGAGACAG NNNN T NNNN GA GTGCCAGCMGCCGCGGTAA GCCTCCCTCGCGCCATCAGAGATGTG	69
515F_f4	TATAAGAGACAG NNNN CT NNNN GAGTGCCAGCMGCCGC GGTAAGCCTCCCTC GCGC C ATCAGAGATGTG TATAAGAGACAG NNNN ACT NNNN GA	70
515F_f5	GTGCCAGCMGCCGCGGTAA GCCTCCCTCGCGCCATC AGAGATGTG TATAAGA GACAG NNNN GACT NNNN GA	71
515F_f6	GTGCCAGCMGCCGCGGTAAGCCTCCCTCGCGCCATCAGAGATGTG TATAAGAG ACAG NNNNTGACT NNNN GA GTGCCAGCMGCCGCGGTAA	72