

This file contains five tables (I-V) listing the phase variation and Gene Scan data for all isolates for pilin modulation, pilin glycosylation, restriction-modification and Opa modules of phase-variable genes. The following information is listed for every gene for each isolate; size of PCR product as determined by GeneScan; derived repeat number (as determined from GeneScan relative to a product of known repeat number); derived expression state (as determined from a bioinformatic analysis of the open reading frame or relative to other isolates of known expression state). Two additional items of information are provided for the Opa genes; size of G tract and size of A tract (these two tracts are adjacent to the repetitive sequence and their length influences the expression state)

Supplementary Data Table I. Repeat numbers and expression states for pilin modulation and LOS phase-variable genes

Volunteer /time pt	Isolate	Phase Variable Gene											
		<i>lgtG</i>			<i>NMB1255</i>			<i>pilC1</i> ⁴			<i>pilC2</i>		
		Size ¹	Tract ²	Exp. State ³	Size ¹	Tract ²	Exp. State ³	Size ¹	Tract ²	Exp. State ³	Size ¹	Tract ²	Exp. State ³
cc174													
V43/1 st	N43.1.1	372.1	12C*	0	377.0	10G	1	256.5	14G	0	353.7	13G	1
	N43.1.2	372.0	12C	0	377.2	10G	1	256.4>255.5 {1.2}	14G	0	353.5	13G	1
	N43.1.3	372.2	12C	0	377.0	10G	1	256.4	14G	0	353.6	13G	1
	N43.1.4	372.0	12C	0	376.8	10G	1	256.3	14G	0	353.3	13G	1
	N43.1.5	371.9	12C	0	376.8	10G	1	256.3	14G	0	353.4	13G	1
	N43.1.6	372.1	12C	0	377.1	10G	1	256.3	14G	0	353.3	13G	1
V43/2 nd	N241.1	369.9	10C	0	376.0	9G	0	255.9 (380.4)	10G* (13G)	0	355.1	13G*	1
	N241.2	370.1	10C	0	375.9	9G	0	255.9 (381.4)	10G (14G)	1	355.2	13G	1
V43/3 rd	N349.1	369.9	10C	0	375.9	9G	0	255.9 (380.4)	10G* (13G)	0	355.2	13G*	1
	N349.2	370.2	10C	0	375.9	9G*	0	255.9 (380.5)	10G (13G)	0	355.1	13G	1
	N349.3	370.0	10C	0	375.7	9G	0	256.0 (380.4)	10G (13G)	0	355.1	13G	1
	N349.4	370.1	10C	0	376.0	9G	0	256.1 (379.5)	10G (12G)	0	355.1	13G	1
	N349.5	370.3	10C	0	376.0	9G	0	255.9 (380.4)	10G (13G)	0	355.1	13G	1
	N349.6	370.2	10C	0	376.0	9G	0	256.2 (380.4)	10G (13G)	0	355.1	13G	1
V43/4 th	Cleared												
V51/1 st	N51.1	370.0	10C	0	375.9	9G	0	256.1 (380.5)	10G* (13G)	0	356.0	14G*	0
	N51.2	370.1	10C	0	376.0	9G	0	256.3 (380.5)	10G (13G)	0	354.9	13G	1
	N51.3	369.8	10C	0	376.1	9G	0	256.3 (380.8)	10G (13G*)	0	355.9	14G	0

	N51.4	370.1	10C	0	376.2	9G	0	256.3 (380.6)	10G (13G)	0	355.9	14G	0
	N51.5	369.9	10C	0	376.0	9G	0	256.3 (380.5)	10G (13G*)	0	355.9	14G	0
	N51.6	370.2	10C	0	376.0	9G	0	256.3 (380.5)	10G (13G)	0	355.8	14G	0
V51/2 nd	N236.1	369.9	10C	0	375.9	9G	0	256.5 (380.6)	10G* (13G)	0	354.8	13G*	1
	N236.2	370.2	10C	0	376.0	9G	0	255.9 (380.6)	10G (13G)	0	355.0	13G	1
	N236.3	369.9	10C	0	375.8	9G	0	256.3 (380.5)	10G* (13G)	0	355.0	13G	1
	N236.4	370.2	10C	0	376.0	9G	0	256.3 (380.6)	10G (13G)	0	356.0	14G	0
	N236.5	370.2	10C	0	376.1	9G	0	256.4 (380.5)	10G (13G)	0	355.9	14G	0
	N236.6	370.2	10C	0	376.4	9G	0	256.4 (380.5)	10G (13G)	0	355.0	13G	1
V51/3 rd	N354.1	370.1	10C	0	376.1	9G	0	256.4 (380.5)	10G (13G)	0	355.8	14G	0
	N354.2	370.2	10C	0	375.6*	9G	0	256.0 (380.3)	10G (13G)	0	355.2	13G	1
	N354.3	370.2	10C	0	375.9	9G	0	255.9 (380.4)	10G (13G*)	0	354.9	13G	1
	N354.4	370.1	10C	0	376.3	9G	0	256.4 (380.6)	10G (13G)	0	354.9	13G	1
	N354.5	370.0	10C	0	376.1	9G	0	256.0 (380.5)	10G (13G)	0	355.2	13G	1
	N354.6	370.1	10C	0	376.3	9G	0	256.4 (380.5)	10G (13G)	0	355.0	13G	1
V51/4 th	N424.1	370.0	10C	0	375.9	9G	0	252.7 (379.6)	10G (12G)	0	353.4	13G	1
	N424.2	370.2	10C	0	376.1	9G	0	255.9 (379.6)	10G (12G)	0	355.2 (13)	13G	1
	N424.3	370.1	10C	0	375.9	9G	0	256.4 (379.6)	10G (12G)	0	354.9 (13)	13G	1
	N424.4	370.1	10C	0	375.9	9G	0	256.1 (379.6)	10G (12G)	0	355.1 (13)	13G	1

	N424.5	370.1	10C	0	375.8	9G	0	256.5 (379.6)	10G* (12G)	0	354.9 (13)	13G	1
	N424.6	370.1	10C	0	376.1	9G	0	256.1 (379.6)	10G (12G)	0	355.1 (13)	13G	1
V52/1 st	N52.1	370.6	10C*	0	376.0	9G	0	256.2 (380.6)	10G (13G)	0	356.0	14G*	0
	N52.2	370.1	10C	0	376.1	9G	0	256.0 (380.6)	10G (13G)	0	356.0	14G	0
	N52.3	370.4	10C	0	376.1	9G	0	255.9 (380.6)	10G (13G)	0	356.0	14G	0
	N52.4	370.1	10C	0	376.3	9G	0	256.0 (380.8)	10G (13G)	0	356.1	14G	0
	N52.5	370.2	10C	0	376.1	9G	0	255.9 (380.7)	10G (13G)	0	356.1	14G	0
	N52.6	370.1	10C	0	376.2	9G	0	256.0 (380.7)	10G (13G)	0	356.1	14G	0
V52/2 nd	N238.1	370.0	10C	0	375.9	9G	0	256.0 (380.4)	10G (13G)	0	355.2	13G*	1
	N238.2	370.6	10C	0	376.0	9G	0	255.9 (380.4)	10G* (13G)	0	355.2	13G	1
	N238.3	370.0	10C	0	375.8	9G	0	255.9 (380.4)	10G (13G)	0	356.2	14G	0
	N238.4	370.1	10C	0	376.2	9G	0	255.9 (380.5)	10G (13G)	0	355.2	13G	1
	N238.5	370.2	10C	0	376.0	9G	0	252.8 (380.4)	10G (13G)	0	353.3	13G	1
	N238.6	369.7	10C	0	376.2	9G	0	255.9 (380.6)	10G (13G)	0	356.1	14G	0
V52/3 rd	N342.1	370.2	10C	0	375.9	9G	0	256.1 (380.6)	10G (13G*)	0	355.1	13G	1
V52/4 th	Cleared												
V54/1 st	N54.1.1	371.1	11C	1	378.2	11G	0	257.8 (374.5)	12G (11G)	0	352.3	10G	1
	N54.1.2	370.1	10C	0	377.1	10G	1	257.8>256.9 {1.1} (375.0)	12G (12G)	0	352.3	10G	1
	N54.1.3	371.0	11C	1	377.2	10G	1	256.8>257.7	11G	0	352.3	10G	1

								{1.2} (375.5)	(12G)				
	N54.1.4	371.1	11C	1	377.0	10G	1	257.8>256.9 {1} (375.6)	12G (12G)	0	352.3	10G	1
	N54.1.5	371.0	11C	1	377.2	10G	1	257.8 (375.6)	12G (12G)	0	352.2	10G	1
	N54.1.6	370.9	11C	1	377.1	10G	1	254.4>253.5 {1} (375.5)	12G (12G)	0	350.4	10G*	1
V54/2 nd	N237.1	370.1	10C	0	378.2	11G*	0	254.3 (374.5)	12G (11G)	0	350.3	10G	1
	N237.2	371.0	11C*	1	378.0	11G	0	255.6 (374.6)	13G (11G)	1	350.4	10G	1
	N237.3	370.0	10C	0	378.1	11G	0	257.8 (374.5)	12G (11G)	0	352.3	10G	1
	N237.4	371.0	11C	1	378.0	11G	0	258.8 (374.6)	13G (11G)	1	352.3	10G	1
	N237.5	371.0	11C	1	377.1	10G	1	254.3 (374.5)	12G (11G)	0	350.3	10G	1
	N237.6	371.1	11C	1	377.0	10G*	1	254.3 (373.6)	12G (10G)	0	350.4	10G	1
V54/3 rd	N343.1	371.3	11C	1	379.1	12G	0	258.8 (375.5)	13G (12G)	1	352.2	10G	1
	N343.2	371.1	11C	1	378.1	11G	0	257.7 (375.6)	12G (12G)	0	352.3	10G	1
	N343.3	371.1	11C	1	378.3	11G	0	257.6 (375.5)	12G (12G)	0	352.4	10G	1
	N343.4	371.0	11C	1	379.2	12G*	0	255.5 (375.4)	13G (12G)	0	350.3	10G	1
	N343.5	370.1	10C	0	379.2	12G	0	255.4 (375.5)	13G (12G)	0	350.4	10G	1
	N343.6	371.0	11C	1	379.3	12G	0	255.4 (375.3)	13G (12G*)	0	350.1	10G	1
V54/4 th	Replaced												
V58/1 st	N58.1	370.1	10C	0	376.2	9G	0	256.0 (380.7)	10G* (13G)	0	355.9	14G*	0
	N58.2	368.0	8C*	1	376.0	9G	0	256.1 (380.7)	10G (13G)	0	356.1	14G	0
	N58.3	370.0	10C	0	376.0	9G	0	256.0	10G	0	356.0	14G	0

								(380.6)	(13G)				
	N58.4	370.1	10C	0	376.1	9G	0	255.9 (379.6)	10G (12G)	0	356.0	14G	0
	N58.5	370.3	10C	0	376.0	9G	0	256.0 (380.7)	10G (13G)	0	356.0	14G	0
	N58.6	370.2	10C	0	376.3	9G	0	255.9 (ND)	10G (13G)	ND	355.1>354.2 {1}	13G	1
V58/2 nd	N240.1	370.1	10C	0	375.9	9G	0	252.5 (380.4)	10G (13G)	0	354.4	14G	0
	N240.2	370.2	10C	0	375.8	9G	0	252.5 (380.4)	10G (13G)	0	353.1	13G	1
V58/3 rd	Cleared												
V58/4 th	N429.1	370.2	10C	0	376.0	9G	0	256.0 (379.6)	10G* (12G)	0	355.1	13G*	1
	N429.2	370.1	10C	0	376.1	9G	0	256.0 (379.6)	10G (12G)	0	355.1	13G	1
	N429.3	370.1	10C	0	375.6	9G	0	256.0 (380.5)	10G (13G)	0	355.1	13G	1
	N429.4	370.1	10C	0	375.7	9G	0	256.1 (380.3)	10G (13G)	0	355.0	13G	1
	N429.5	370.2	10C	0	376.1	9G	0	256.0 (379.6)	10G (12G)	0	353.4	13G*	1
	N429.6	370.1	10C	0	376.0	9G	0	256.1 (380.5)	10G (13G)	0	355.1	13G	1
V59/1 st	N59.1.1	370.1	10C	0	376.1	9G	0	256.4 (380.4)	10G* (13G)	0	355.0	13G*	1
	N59.1.2	370.1	10C	0	375.9	9G	0	255.8 (379.4)	10G (12G)	0	355.0	13G	1
	N59.1.3	370.2	10C	0	376.2	9G	0	256.3 (380.8)	10G (13G)	0	355.9	14G	0
	N59.1.4	370.1	10C	0	375.4	8G	0	256.3 (379.6)	10G (12G)	0	354.9	13G	1
	N59.1.5	370.0	10C	0	376.1	9G	0	256.0 (379.6)	10G (12G)	0	355.2	13G	1
	N59.1.6	370.3	10C	0	376.0	9G	0	256.6 (379.7)	10G* (12G)	0	354.9	13G	1
V59/2 nd	N253.1	370.0	10C	0	376.0	9G	0	256.2	10G*	0	355.0	13G*	1

								(379.4)	(12G)				
	N253.2	370.2	10C	0	376.1	9G	0	256.3 (379.4)	10G (12G)	0	355.0	13G	1
	N253.3	370.1	10C	0	375.9	9G	0	256.2 (379.7)	10G* (12G)	0	355.0	13G	1
	N253.4	370.2	10C	0	375.9	9G	0	256.3 (380.4)	10G (13G)	0	355.0	13G	1
	N253.5	370.0	10C	0	375.7	9G	0	256.3 (379.7)	10G (12G)	0	354.9	13G	1
	N253.6	370.1	10C	0	376.0	9G	0	256.5 (379.4)	10G (12G)	0	354.9	13G	1
V59/3 rd	N352.1	370.2	10C	0	375.6	9G	0	256.3 (380.5)	10G (13G)	0	355.0	13G	1
	N352.2	371.0	11C	1	376.0	9G	0	256.2 (380.6)	10G (13G)	0	355.0	13G	1
	N352.3	370.1	10C	0	375.7	9G	0	256.0 (380.6)	10G (13G)	0	355.1	13G	1
	N352.4	370.0	10C	0	376.0	9G	0	256.2 (379.6)	10G (12G)	0	355.0	13G	1
	N352.5	370.1	10C	0	376.6	10G	1	256.3 (380.5)	10G (13G)	0	354.9	13G	1
	N352.6	370.1	10C	0	375.8	9G	0	256.1 (379.4)	10G (12G)	0	355.0	13G	1
V59/4 th	N438.1	370.0	10C	0	375.9	9G	0	256.4 (380.4)	10G (13G*)	0	355.0	13G	1
	N438.2	370.1	10C	0	376.1	9G	0	256.3 (380.4)	10G (13G)	0	355.0	13G	1
	N438.3	370.2	10C	0	375.7	9G	0	256.3 (380.5)	10G* (13G)	0	354.0	12G	0
	N438.4	370.1	10C	0	376.0	9G	0	256.3 (380.5)	10G (13G)	0	354.9	13G	1
	N438.5	370.1	10C	0	375.7	9G	0	256.4 (380.6)	10G (13G)	0	355.1	13G	1
	N438.6	370.2	10C	0				256.4 (380.4)	10G (13G)	0	354.9	13G	1
V88/1 st	N88.1.1	372.1	12C	0	376.7	10G	1	259.5	13G	1	351.4	9G*	0
	N88.1.2	372.0	12C	0	377.0	10G	1	259.4	13G	1	351.4	9G	0

	N88.1.3	372.0	12C	0	376.7	10G	1	259.6	13G	1	351.4	9G	0
	N88.1.4	372.0	12C	0	377.0	10G	1	259.5	13G	1	351.4	9G	0
	N88.1.5	372.0	12C	0	376.7	10G	1	258.7	12G	0	351.3	9G	0
	N88.1.6	371.9	12C	0	376.8	10G	1	259.7	13G	1	351.3	9G	0
V88/2 nd	N272.1	371.9	12C	0	376.8	10G	1	259.6	13G	1	352.3	10G	1
	N272.2	372.0	12C	0	376.9	10G	1	259.5	13G	1	352.3	10G	1
	N272.3	372.1	12C	0	376.9	10G	1	259.5>258.6 {1}	13G	1	352.3	10G	1
	N272.4	372.5	12C	0	377.1	10G	1	259.5	13G	1	352.4	10G	1
	N272.5	372.0	12C	0	376.7	10G	1	259.5	13G	1	352.3	10G	1
	N272.6	372.0	12C	0	377.1	10G	1	256.1	14G	0	355.0	13G	1
V88/3 rd	N369.1	372.0	12C	0	376.9	10G	1	259.6>258.7 {1.2}	13G	1	352.3	10G	1
	N369.2	372.0	12C	0	377.2	10G	1	259.6	13G	1	352.3	10G	1
	N369.3	372.1	12C	0	377.7	11G	0	256.4	14G	0	350.2	10G	1
	N369.4	372.0	12C	0	378.1	11G	0	256.4	14G	0	352.4	10G	1
	N369.5	372.0	12C	0	376.8	10G	1	259.4>258.5 {1.2}	13G	1	352.3	10G	1
	N369.6	371.9	12C	0	377.1	10G	1	259.4	13G	1	352.4	10G	1
V88/4 th	N449.1	372.0	12C	0	377.9	11G	0	255.3	13G	1	350.3	10G	1
	N449.2	372.0	12C	0	378.1	11G	0	255.5	13G	1	350.4	10G	1
	N449.3	372.0	12C	0	377.9	11G	0	255.4	13G	1	350.4	10G	1
	N449.4	371.6	11C	1	378.1	11G	0	255.4	13G	1	350.4	10G	1
	N449.5	372.0	12C	0	377.9	11G	0	255.4	13G	1	350.4	10G	1
	N449.6	372.0	12C	0	378.1	11G	0	255.4	13G	1	350.2	10G	1
V138/1 st	N138.1	370.1	10C	0	375.7	9G	0	256.1 (379.4)	10G (12G)	0	353.3	13G	1
	N138.2	370.1	10C	0	376.0	9G	0	256.0 (380.5)	10G* (13G)	0	355.1	13G*	1
	N138.3	370.0	10C	0	375.9	9G	0	256.0 (380.5)	10G (13G)	0	353.2	13G	1
	N138.4	370.0	10C	0	376.2	9G	0	256.0 (380.5)	10G (13G)	0	355.1	13G	1
	N138.5	370.1	10C	0	375.7	9G	0	256.0 (380.5)	10G (13G)	0	355.2	13G	1
	N138.6	370.2	10C	0	376.1	9G	0	256.1 (381.6)	10G (14G)	1	355.2	13G	1

V138/2 nd	N288.1	368.9	10C	0	375.7	9G	0	256.2 (380.6)	10G* (13G)	0	355.0	13G*	1
	N288.2	370.0	10C	0	376.0	9G	0	256.2 (ND)	10G (ND)	ND	355.2	13G	1
	N288.3	370.1	10C	0	375.7	9G	0	256.2 (381.4)	10G (14G)	1	355.1	13G	1
	N288.4	369.9	10C	0	376.2	9G	0	256.0 (380.4)	10G (13G)	0	355.1	13G	1
	N288.5	370.1	10C	0	375.9	9G	0	256.0 (381.5)	10G (14G)	1	353.3	13G	1
	N288.6	370.1	10C	0	376.2	9G	0	256.1 (380.4)	10G (13G)	0	355.0	13G	1
V138/3 rd	N331.1	370.1	10C	0	375.8	9G	0	255.9 (379.6)	10G (12G)	0	355.2	13G	1
	N331.2	370.0	10C	0	376.1	9G	0	256.0 (380.4)	10G (13G)	0	355.1	13G	1
	N331.3	370.1	10C	0	375.6	9G	0	256.0 (380.3)	10G (13G)	0	355.2	13G	1
	N331.4	370.0	10C	0	375.8	9G	0	256.1 (380.5)	10G (13G)	0	355.0	13G	1
	N331.5	369.7	10C*	0	375.7	9G	0	256.1 (380.4)	10G (13G*)	0	355.1	13G	1
	N331.6	370.0	10C	0	376.0	9G	0	256.2 (380.4)	10G (13G)	0	355.0	13G	1
V138/4 th	Replaced												
cc167													
V124/1 st	N124.1	368.9	9C	0	377.9	11G	0	274.2	13G	1	386.0	11G	0
	N124.2	369.2	9C	0	378.2	11G	0	274.3	13G	1	386.1	11G	0
	N124.3	369.0	9C	0	378.0	11G	0	274.4	13G	1	386.2	11G	0
	N124.4	369.1	9C	0	378.2	11G	0	274.2	13G	1	385.2	10G	1
	N124.5	369.0	9C	0	378.0	11G	0	274.2	13G	1	385.1	10G	1
	N124.6	369.2	9C	0	378.3	11G	0	271.7	13G*	1	385.3	10G	1
V124/2 nd	N290.1	369.0	9C	0	377.9	11G	0	274.1	13G*	1	386.2	11G	0
	N290.2	369.1	9C*	0	378.3*	11G	0	274.2	13G	1	385.3	10G	1
	N290.3	368.7	9C	0	377.9	11G	0	274.2	13G	1	385.2	10G	1
	N290.4	369.1	9C	0	378.3	11G	0	274.3	13G	1	386.2	11G	0
	N290.5	368.9	9C	0	377.9	11G	0	273.4	12G	0	385.1	10G	1

	N290.6	369.2	9C	0	378.3	11G	0	271.0	10G	1	384.4	9G	0
V124/3 rd	N336.1	369.0	9C	0	376.7	10G	1	273.3	12G	0	385.0	10G	1
	N336.2	369.1	9C	0	378.2	11G	0	274.2	13G	1	385.2	10G	1
	N336.3	369.0	9C	0	376.9	10G	1	273.3	12G	0	385.1	10G	1
	N336.4	369.0	9C	0	378.3	11G	0	273.3	12G	0	385.2	10G	1
	N336.5	369.1	9C	0	377.1	10G	1	272.4	11G	0	385.3	10G	1
	N336.6	369.3	9C	0	377.3	10G	1	273.3	12G	0	385.3	10G	1
V64/1 st	N64.1	369.1	9C	0	377.9	11G	0	275.2	14G*	0	385.1	10G	1
	N64.2	369.1	9C	0	378.2	11G	0	274.3	13G*	1	385.1	10G*	1
	N64.3	368.9	9C	0	378.0	11G	0	274.2	13G	1	385.3	10G	1
	N64.4	369.1	9C	0	378.3	11G	0	274.1	13G	1	385.0	10G	1
	N64.5	368.9	9C	0	377.9	11G	0	274.2	13G	1	385.1	10G	1
	N64.6	369.1	9C	0	378.2	11G	0	274.2	13G	1	385.1	10G	1
V64/2 nd	N257.1	368.9	9C	0	377.8	11G	0	274.2	13G	1	385.1	10G*	1
	N257.2	369.2	9C	0	378.1	11G	0	274.3	13G	1	385.2	10G	1
	N257.3	369.1	9C	0	377.9	11G	0	274.3	13G	1	384.9	10G	1
	N257.4	369.2	9C	0	377.0	10G	1	275.2	14G	0	385.0	10G	1
	N257.5	369.0	9C	0	378.0	11G	0	274.3	13G	1	384.9	10G	1
	N257.6	369.1	9C	0	377.1	10G	1	275.3	14G	0	384.6	10G	1
V64/3 rd	N348.1	369.0	9C	0	377.9	11G	0	275.2	14G	0	385.2	10G	1
	N348.2	369.2	9C	0	378.2	11G	0	275.2	14G	0	385.2	10G	1
	N348.3	368.9	9C	0	378.3	11G	0	272.9	14G	0	384.5	10G	1
	N348.4	369.1	9C	0	380.4	13G	1	273.7	15G	0	384.3	10G	1
	N348.5	368.9	9C	0	377.4	11G	0	275.1	14G	0	385.0	10G	1
	N348.6	369.1	9C	0	378.3	11G	0	272.7	14G	0	384.2	10G	1
V117/1 st	N117.1	369.0	9C	0	378.4	11G	0	274.3	13G*	1	384.1	9G	0
	N117.2	369.1	9C	0	378.4	11G	0	274.3	13G*	1	384.8	10G*	1
V117/2 nd	N284.1	368.9	9C	0	378.2	11G	0	273.3	12G	0	385.3	10G	1
	N284.2	369.2	9C	0	378.3	11G	0	273.3	12G	0	385.3	10G	1
	N284.3	369.0	9C	0	378.3	11G	0	274.3	13G	1	384.2	9G	0
	N284.4	369.2	9C	0	377.4	11G	0	273.3	12G*	0	385.0	10G	1
	N284.5	369.0	9C	0	378.2	11G	0	274.3	13G	1	383.8	9G	0
	N284.6	369.1	9C	0	378.1	11G	0	273.2	12G	0	384.7	10G	1
V117/3 rd	N332.1	369.0	9C	0	377.0	10G	1	273.3	12G	0	385.1	10G	1
	N332.2	369.3	9C	0	377.4	10G	1	273.4	12G	0	385.1	10G	1

	N332.3	368.9	9C	0	377.2	10G	1	270.9	12G	0	384.2	10G	1
	N332.4	369.0	9C	0	378.3	11G	0	271.8	13G	1	384.2	10G*	1
	N332.5	369.0	9C	0	378.3	11G	0	274.2	13G	1	385.3	10G	1
	N332.6	369.1	9C	0	379.3	12G	0	274.2	13G	1	385.1	10G	1
V117/4 th	N417.1	369.0	9C	0	379.4	12G	0	274.2	13G	1	385.3	10G	1
	N417.2	369.2	9C	0	379.3	12G	0	274.2	13G	1	385.3	10G	1
	N417.3	369.0	9C	0	379.3	12G	0	272.9>271.9 {1.1}	14G	0	384.5	10G*	1
	N417.4	369.1	9C	0	379.2	12G	0	272.9>271.9 {1.1}	14G	0	384.6	10G	1
	N417.5	369.0	9C	0	379.3	12G	0	275.2	14G	0	385.0	10G	1
	N417.6	369.2	9C	0	379.3	12G	0	275.2	14G	0	384.6	10G	1
cc23													
V69/1 st	Non-carrier												
V69/2 nd	N258.1	370.9	11C	1	379.6	13G	1	277.6	12G*	1	424.6	8G*	0
	N258.2	370.6	11C	1	379.5	13G	1	279.6	12G	1	425.2	8G	0
	N258.3	370.9	11C	1	380.3	13G	1	279.7	12G	1	425.3	8G	0
	N258.4	370.8	11C	1	380.2	13G	1	279.7	12G	1	425.4	8G	0
	N258.5	371.0	11C	1	379.5	13G	1	279.6	12G	1	425.3	8G	0
	N258.6	370.6	11C	1	379.3??	13G	1	279.6	12G	1	425.4	8G	0
V69/3 rd	No sample												
V69/4 th	N431.1	370.7	11C	1	381.7	15G	0	276.6	11G	0	425.7	9G*	1
	N431.2	370.9	11C	1	380.5	14G	0	277.7	12G*	1	425.5	9G	1
	N431.3	370.8	11C	1	380.7	14G	0	279.6	12G	1	426.3	9G	1
	N431.4	370.9	11C	1	381.2	14G	0	279.6	12G	1	426.4	9G	1
	N431.5	370.7	11C	1	381.4>380.5 {1.2}	15G	0	278.8	11G	0	425.7	9G	1
	N431.6	370.9	11C	1	381.1	14G	0	278.7	11G	0	426.3	9G	1
V93/1 st	Non-carrier												
V93/2 nd	N264.1	370.8	11C	1	378.0	11G	0	281.5>282.4 {1.1}	14G	0	428.3	11G	0
	N264.2	370.9	11C	1	378.3	11G	0	281.4>282.4 {1.1}	14G*	0	428.3	11G*	0
	N264.3	371.7	12C	0	377.3	10G	1	281.4>282.3 {1.1}	14G	0	429.3	12G	1
	N264.4	370.9	11C	1	377.9	11G	0	281.3>282.3	14G	0	428.3	11G	0

								{1.0}					
	N264.5	370.7	11C	1	378.3	11G	0	281.4>282.3 {1.1}	14G	0	428.3	11G	0
	N264.6	370.8	11C	1	378.2	11G	0	281.4>282.3 {1.0}	14G	0	429.3	12G	1
V93/3 rd	N359.1	370.8	11C	1	377.1	10G	1	283.2>282.3 {1.0}	15G	1	429.3	12G	1
	N359.2	370.9	11C	1	377.3	10G	1	282.3>283.2 {1.2}	15G	1	429.4	12G	1
	N359.3	370.8	11C	1	377.2	10G	1	282.3>283.3 {1.1}	15G	1	429.3	12G	1
	N359.4	370.9	11C	1	376.9	10G	1	282.3>283.3 {1.1}	15G	1	429.3	12G	1
	N359.5	370.7	11C	1	376.9	10G	1	282.3>283.3 {1.1}	15G	1	429.3	12G	1
	N359.6	370.9	11C	1	376.9	10G	1	282.4>283.3 {1.2}	15G	1	429.2	12G	1
V96/1 st	Non-carrier												
V96/2 nd	N259.1	370.7	11C	1	380.2	13G	1	279.6	12G	1	426.3	9G	1
	N259.2	370.9	11C	1	379.5	13G	1	279.6	12G	1	426.5	9G	1
	N259.3	371.7	12C	0	378.1	11G	0	279.6	12G	1	427.2	10G	0
	N259.4	370.9	11C	1	380.2	13G	1	279.6	12G	1	426.3	9G	1
	N259.5	370.7	11C	1	379.3	12G	0	279.6	12G	1	426.4	9G	1
	N259.6	371.8	12C	0	377.1	11G	0	279.7	12G	1	427.3	10G	0
V96/3 rd	N370.1	372.2	12C	0	381.5>382.5 {1}	15G	0	277.9	10G*	0	426.3	9G	1
	N370.2	369.8	11C	1	380.4>379.4 {1.2}	14G	0	278.8	11G	0	426.3	9G	1
	N370.3	371.8>370.8 {1.1}	13C	0	378.3	11G	0	278.7	11G	0	431.3>432.4 {1.1}	14G	0
	N370.4	371.8>370.9 {1.1}	13C	0	381.6>382.6 {1}	15G	0	278.7	11G	0	426.4	9G	1
	N370.5	370.8	11C	1	379.3	13G	1	276.9	9G	1	425.4	8G	0
	N370.6	370.0	12C*	0	381.6>382.6 {1}	15G	0	277.7	10G	0	426.4	9G	1
V96/4 th	N445.1	372.3	12C	0	379.1	12G	0	278.8	11G	0	426.4	9G	1
	N445.2	374.8	16C	0	379.6	13G	1	276.6	11G	0	426.4	9G	1

	N445.3	375.7>3 76.7 {1}	17C	1	379.3	12G	0	278.8	11G	0	426.3	9G	1
	N445.4	374.8>3 75.7{1.1}	16C	0	379.1	12G	0	278.7	11G	0	426.3	9G	1
	N445.5	371.0>3 71.9 {1}	12C	0	378.3	11G	0	280.7>279.7 {1.1}	13G	0	426.3	9G	1
	N445.6	372.5	12C	0	377.2	10G	1	278.8	11G	0	426.2	9G	1
V222/1 st	N222.1	370.8	11C	1	379.1	13G	1	279.6	12G*	1	425.4 (8)	8G	0
	N222.2	370.7	11C*	1	380.2	13G	1	280.5 (13)	13G	0	426.4 (9)	9G*	1
	N222.3	371.8	12C*	0	379.3	12G	0	279.6 (12)	12G	1	425.4 (8)	8G	0
	N222.4	370.8	11C	1	379.4	13G	1	279.7 (12)	12G	1	425.4 (8)	8G	0
	N222.5	370.8	11C	1	379.5	13G	1	279.6 (12)	12G	1	425.2 (8)	8G	0
	N222.6	371.8	12C	0	379.1	12G	0	279.9 (12)	12G	1	425.3 (8)	8G	0
V222/2 nd	N309.1	370.8	11C	1	379.2	13G	1	279.7 (12)	12G	1	426.4 (9)	9G	1
	N309.2	370.8	11C	1	380.2	13G	1	279.7 (12)	12G	1	426.3 (9)	9G	1
	N309.3	370.9	11C	1	380.3	13G	1	279.6 (12)	12G	1	426.4 (9)	9G	1
	N309.4	370.7	11C	1	380.2	13G	1	279.6 (12)	12G	1	426.5 (9)	9G	1
	N309.5	370.8	11C	1	380.5	14G	0	279.5 (12)	12G	1	426.5 (9)	9G	1
	N309.6	370.7	11C	1	380.2	13G	1	279.6 (12)	12G	1	426.5 (9)	9G	1
V222/3 rd	N391.1	370.9	11C	1	379.3	13G	1	279.5 (12)	12G	1	426.5 (9)	9G	1
	N391.2	370.6	11C	1	379.6	13G	1	279.6 (12)	12G	1	426.5 (9)	9G	1
	N391.3	370.8	11C	1	380.2	13G	1	279.6 (12)	12G	1	426.3 (9)	9G	1
	N391.4	370.7	11C	1	380.1	13G	1	279.6 (12)	12G	1	426.3 (9)	9G	1
	N391.5	370.9	11C	1	380.1	13G	1	279.6 (12)	12G	1	426.4 (9)	9G	1
	N391.6	370.7	11C	1	380.1	13G	1	277.7 (12)	12G	1	425.5 (9)	9G	1
V222/4 th	N459.1	370.8	11C	1	379.1	12G	0	278.7	11G	0	426.5 (9)	9G	1
	N459.2	370.9	11C	1	379.1	12G	0	278.6 (11)	11G	0	426.6 (9)	9G	1
	N459.3	370.8	11C	1	379.1	12G	0	278.5 (11)	11G	0	426.3 (9)	9G	1
	N459.4	370.8	11C	1	379.1	12G	0	279.5 (12)	12G	1	426.2 (9)	9G	1
	N459.5	371.0	11C	1	379.2	12G	0	278.7 (11)	11G	0	426.5 (9)	9G	1
	N459.6	370.7	11C	1	379.1	12G	0	277.5 (12)	12G	1	425.4 (9)	8G	0
cc60													
V114/1 st	N114.1	370.6	11C	1	375.1	7G	0	274.8	10G*	1	401.9	9G*	0
	N114.2	370.8	11C	1	375.3	7G	0	274.7	10G	1	401.0	8G	0

	N114.3	370.7	11C	1	375.2	7G	0	274.8	10G	1	401.0	8G	0
	N114.4	370.9	11C	1	375.4	7G	0	274.7	10G	1	401.1	8G	0
	N114.5	370.7	11C	1	375.2	7G	0	274.8	10G	1	400.7	8G	0
	N114.6	370.9	11C	1	375.4	7G	0	274.7	10G	1	401.1	8G	0
V114/2 nd	N283.1	369.9	11C*	1	375.2	7G	0	274.9	10G	1	400.9	8G	0
	N283.2	369.9	11C	1	375.3	7G	0	274.7	10G	1	401.1	8G	0
	N283.3	369.7	11C	1	375.0	7G	0	272.2	10G	1	401.7	10G*	1
	N283.4	370.8	11C	1	375.3	7G	0	272.3	10G	1	401.6	10G	1
	N283.5	370.4	10C?	0	374.9	7G*	0	275.0	10G*	1	400.6	9G	0
	N283.6	370.6	11C	1	375.1	7G	0	274.7	10G	1	400.9	8G	0
V114/3 rd	N330.1	372.7	13C*	0	375.2	7G	0	273.1	8G	0	402.0	9G	0
	N330.2	370.0	10C	0	375.5	7G	0	272.9	8G	0	402.1	9G	0
	N330.3	370.0	10C*	0	375.3	7G	0	272.9	8G	0	401.9	9G	0
	N330.4	370.0	10C	0	375.4	7G	0	272.8	8G	0	402.2	9G	0
	N330.5	370.9	11C	1	375.4	7G	0	273.1	8G	0	402.0	9G	0
	N330.6	370.9	11C	1	375.5	7G	0	272.8	8G	0	402.0	9G	0
V134/1 st	N134.1	370.7	11C	1	375.4	7G	0	274.8	10G	1	400.9	8G	0
	N134.2	370.9	11C*	1	375.4	7G	0	274.6	10G	1	402.1	9G	0
	N134.3	370.6	11C	1	375.2	7G	0	274.9	10G*	1	402.0	9G	0
	N134.4	370.8	11C	1	375.3	7G	0	274.7	10G	1	402.1	9G	0
	N134.5	370.6	11C	1	375.1	7G	0	274.9	10G	1	401.9	9G	0
	N134.6	370.7	11C	1	375.3	7G	0	274.7	10G	1	401.9	9G	0
V134/2 nd	N295.1	370.5	11C?	1	375.1	7G	0	274.7	10G	1	401.0	8G	0
	N295.2	370.6	11C	1	375.4	7G	0	272.4	10G	1	400.5	9G*	0
	N295.3	370.6	11C	1	375.3	7G	0	274.8	10G	1	401.0	8G	0
	N295.4	370.5	11C	1	375.4	7G	0	274.6	10G	1	401.1	8G	0
	N295.5	370.4	11C?	1	375.3	7G	0	274.8	10G	1	400.9	8G	0
	N295.6	369.9	11C	1	375.4	7G	0	274.7	10G	1	401.1	8G	0
V134/3 rd	N333.1	370.9	11C	1	375.2	7G	0	274.8	10G	1	401.9	9G	0
	N333.2	370.8	11C	1	375.5	7G	0	274.6	10G	1	402.0	9G	0
	N333.3	370.9	11C	1	375.3	7G	0	274.9	10G	1	401.8	9G	0
	N333.4	371.0	11C	1	375.5	7G	0	274.6	10G	1	402.1	9G	0
	N333.5	370.9	11C	1	375.2	7G	0	275.0	10G	1	401.9	9G	0
	N333.6	371.0	11C	1	375.3	7G	0	274.7	10G	1	401.8	9G	0
V185/1 st	N185.1	371.0	11C*	1	376.2	8G	1	277.7	13G	1	402.9	10G	1

	N185.2	370.8	11C	1	376.4	8G	1	277.7	13G	1	402.6	10G	1
	N185.3	370.9	11C	1	376.3	8G	1	276.9	12G	0	403.8	11G*	0
	N185.4	370.9	11C	1	376.5	8G	1	277.5	13G	1	403.0	10G	1
	N185.5	370.9	11C	1	376.3	8G	1	277.5	13G	1	403.0	10G	1
	N185.6	370.9	11C	1	376.0	8G	1	275.4	13G	1	401.8	10G	1
V185/2 nd	N306.1	370.8	11C	1	376.2	8G	1	277.6	13G	1	402.8	10G	1
	N306.2	370.8	11C	1	376.4	8G	1	277.7	13G	1	402.9	10G	1
	N306.3	370.9	11C	1	376.3	8G	1	277.7	13G	1	402.8	10G	1
	N306.4	370.9	11C	1	376.4	8G	1	277.5	13G	1	403.1	10G	1
	N306.5	370.8	11C	1	376.2	8G	1	277.4	13G	1	403.0	10G	1
	N306.6	370.8	11C	1	376.4	8G	1	277.6	13G	1	402.9	10G	1
V185/4 th	N456.1	371.8	12C	0	376.2	8G	1	277.6	13G	1	401.9	9G	0
	N456.2	371.8	12C	0	376.3	8G	1	277.6	13G	1	401.8	9G	0
	N456.3	371.9	12C*	0	376.3	8G	1	277.7	13G	1	402.8	10G	1
	N456.4	371.8	12C	0	376.4	8G	1	277.5	13G	1	402.1	9G	0
	N456.5	371.7	12C	0	376.3	8G	1	274.4	13G*??	1	401.9	9G	0
	N456.6	371.7	12C	0	376.5	8G	1	277.5	13G	1	402.0	9G	0
V113/1 st	N113.1	369.8	11C*	1	376.2	8G	1	274.8	10G	1	401.9	9G	0
	N113.2	369.9	11C	1	376.4	8G	1	274.7	10G	1	401.0	8G	0
	N113.3	369.8	11C	1	376.4	8G	1	274.8	10G	1	401.0	8G	0
	N113.4	369.8	11C	1	376.4	8G	1	274.7	10G	1	401.1	8G	0
	N113.5	369.6	11C	1	376.2	8G	1	274.8	10G	1	400.7	8G	0
	N113.6	369.7	11C	1	376.4	8G	1	274.7	10G	1	401.1	8G	0
V113/2 nd	N281.1	369.7	11C	1	376.2	8G	1	274.7	10G	1	401.9	9G	0
	N281.2	369.8	11C	1	376.3	8G	1	274.6	10G	1	401.1	8G	0
	N281.3	369.9	11C	1	376.3	8G	1	274.9	10G	1	401.9	9G	0
	N281.4	369.9	11C	1	375.3	7G	0	273.8	9G	0	402.1	9G	0
	N281.5	369.8	11C	1	376.2	8G	1	274.9	10G	1	400.8	8G	0
	N281.6	369.8	11C	1	376.4	8G	1	274.7	10G	1	400.9	8G	0
V115/1 st	N115.1	369.9	11C	1	375.2	7G	0	274.9	10G	1	400.9	8G	0
	N115.2	369.8	11C	1	374.9	7G	0	274.7	10G	1	401.1	8G	0
	N115.3	369.8	11C	1	375.3	7G	0	274.9	10G	1	400.7	8G	0
	N115.4	369.8	11C	1	375.3	7G	0	274.8	10G	1	400.9	8G	0
	N115.5	369.9	11C	1	375.5	7G?	0	275.0	10G	1	401.9	9G	0
	N115.6	369.9	11C	1	375.6	7G?	0	274.7	10G	1	401.7	9G	0

V115/2 nd	N282.1	369.8	11C	1	374.8	7G	0	274.9	10G	1	401.8	9G	0
	N282.2	369.9	11C	1	374.8	7G	0	274.7	10G	1	401.0	8G	0
	N282.3	370.0	11C	1	375.0	7G	0	274.8	10G	1	402.0	9G	0
	N282.4	369.9	11C	1	374.8	7G	0	272.2	10G	1	401.9	10G	1
	N282.5	370.0	11C	1	375.0	7G	0	274.8	10G	1	401.9	9G	0
	N282.6	369.9	11C	1	374.9	7G	0	274.7	10G	1	402.1	9G	0
cc32													
V176/1 st	N176.1	369.8	10C	0	375.1	8G	0	278.1	13G*	1	429.4	12G*	1
	N176.2	369.6	10C	0	375.2	8G	0	278.0	13G	1	429.3	12G	1
	N176.3	369.8	10C	0	375.2	8G	0	278.0	13G	1	429.4	12G	1
	N176.4	369.6	10C	0	375.0	8G	0	278.0	13G	1	429.4	12G	1
	N176.5	369.7	10C	0	374.8	8G	0	278.1	13G	1	429.4	12G	1
	N176.6	369.6	10C	0	374.9	8G	0	278.1	13G	1	429.3	12G	1
V176/2 nd	N318.1	369.7	10C	0	374.8	8G	0	278.0	13G	1	429.4	12G	1
	N318.2	369.6	10C	0	374.9	8G	0	277.2	12G	0	429.3	12G	1
	N318.3	369.0	9C	0	375.0	8G	0	278.0	13G	1	429.3	12G	1
	N318.4	369.7	10C	0	374.8	8G	0	277.9	13G	1	429.4	12G	1
	N318.5	369.8	10C	0	374.9	8G	0	277.4	12G	0	429.4	12G	1
	N318.6	369.6	10C	0	374.8	8G	0	277.0	12G	0	429.5	12G	1
V176/3 rd	N399.1	369.8	10C	0	374.9	8G	0	277.0	12G*	0	429.4	12G*	1
	N399.2	369.5	10C	0	374.7	8G	0	277.1	12G	0	429.4	12G	1
	N399.3	369.8	10C	0	374.8	8G	0	277.1	12G	0	429.3	12G	1
	N399.4	369.5	10C	0	374.7	8G	0	277.2	12G	0	429.3	12G	1
	N399.5	369.7	10C	0	374.9	8G	0	277.1	12G	0	429.3	12G	1
	N399.6	369.7	10C	0	374.8	8G	0	277.2	12G	0	429.3	12G	1
V176/4 th	N408.1	369.9	10C	0	375.1	8G	0	277.3	12G*	0	429.2	12G*	1
	N408.2	369.6	10C	0	374.8	8G	0	277.2	12G	0	429.2	12G	1
	N408.3	369.8	10C	0	374.9	8G	0	277.2	12G	0	429.3	12G	1
	N408.4	369.7	10C	0	374.9	8G	0	277.2	12G	0	429.3	12G	1
	N408.5	369.9	10C	0	375.1	8G	0	277.2	12G	0	429.4	12G	1
	N408.6	369.6	10C	0	374.9	8G	0	277.2	12G	0	429.3	12G	1

¹Fragments sizes were determined using either a GS600LIZ (black type) or GS500LIZ (blue type) size standard; numbers in curly brackets indicate the ratio between the two major peaks where obvious multiple peaks were present (the largest peak was utilized if the ratio was >1.2; the larger repeat number equivalent peak was utilized if the ratio was <1.2); numbers in round brackets indicate the repeat number as determined from next generation whole genome sequence data; ²an asterisk indicates that the PCR fragment was sequenced by dideoxy sequencing and the repeat number determined by visual inspection of the trace file. ³Expression states, 1=1, 0=0 (all genes are subject to translational phase variation). ⁴A subset of cc174 isolates contained a second polyC tract withing pilC, the data for this tract is indicated in brackets whilst the expression state is a combination of data from both tracts.

Supplementary Data Table II. Repeat numbers and expression states for the phase-variable pilin glycosylation genes

Volunteer/ time pt	Isolate	Phase Variable Gene											
		<i>pgIA</i>			<i>pgIE</i>			<i>pgIH</i>			<i>pgII</i>		
		Size ¹	Tract ²	Exp. State ³	Size ¹	Tract ²	Exp. State ³	Size ¹	Tract ²	Exp. State ³	Size ¹	Tract ²	Exp. State ³
cc174													
V43/1 st	N43.1.1	425.4	12G	0	388.4	4CAAACAA*	1	187.1	14C	0	351.3	12G*	0
	N43.1.2	425.6	12G	0	388.5	4CAAACAA	1	187.1	14C	0	352.5	13G	1
	N43.1.3	425.3	12G	0	388.4	4CAAACAA	1	187.1	14C	0	352.4	13G	1
	N43.1.4	425.6	12G	0	388.7	4CAAACAA	1	187.2*	14C	0	352.5	13G	1
	N43.1.5	425.3	12G	0	388.6	4CAAACAA	1	187.1	14C	0	352.4	13G	1
	N43.1.6	425.6	12G	0	388.5	4CAAACAA	1	187.1	14C	0	352.5	13G	1
V43/2 nd	N241.1	426.1	12G	0	562.9	29CAAACAA*	0	187.9	15C	0	349.7 (10)	10G*	1
	N241.2	425.7>426.7 {1.2}	12G	0	562.4	29CAAACAA	0	188.0	15C	0	349.2	10G	1
V43/3 rd	N349.1	426.1>427.1 {1.1}	13G	0	562.3	29CAAACAA*	0	188.9>189.9 {1.1}	16C	1	349.7 (10)	10G*	1
	N349.2	426.1>427.1 {1.1}	13G	0	562.1	29CAAACAA	0	189.0	16C	1	349.4	10G	1
	N349.3	425.7>426.7 {1.1}	13G	0	562.1	29CAAACAA	0	189.0	16C	1	349.2	10G	1
	N349.4	426.0>427.0 {1.2}	13G	0	555.6	28CAAACAA	1	189.0	16C	1	349.3	10G	1
	N349.5	425.6	12G	0	570.1	30CAAACAA	0	189.0>189.9 {1.2}	16C	1	349.2	10G	1
	N349.6	426.0>427.0 {1.2}	12G	0	562.8	29CAAACAA	0	189.0>190.0 {1.2}	16C	1	349.3	10G	1
V43/4 th	Cleared												
V51/1 st	N51.1	424.6 (11)	11G*	1	563.1	29CAAACAA*	0	188.0*	15C	0	349.6 (10)	10G	1
	N51.2	424.3 (11)	11G	1	562.7	29CAAACAA	0	188.0>189.0 {1.2}	15C	0	349.3	10G	1
	N51.3	424.5 (11)	11G*	1	562.6	29CAAACAA	0	188.0	15C	0	349.3	10G	1
	N51.4	424.3 (11)	11G	1	561.8	29CAAACAA	0	188.0	15C	0	349.4	10G	1
	N51.5	424.5 (11)	11G	1	562.6	29CAAACAA	0	188.0>188.9 {1.2}	15C	0	349.4	10G	1
	N51.6	424.3 (11)	11G	1	562.1	29CAAACAA	0	188.1>189 {1.2}	15C	0	349.4	10G	1
V51/2 nd	N236.1	424.5	11G	1	563.0	29CAAACAA	0	187.9>188.8 {1.1}	15C	0	349.4	10G	1
	N236.2	424.3 (11)	11G	1	562.2	29CAAACAA	0	187.9	15C	0	349.3	10G	1
	N236.3	424.5 (11)	11G	1	562.7	29CAAACAA	0	187.9>188.9 {1.2}	15C	0	349.3	10G	1

	N236.4	424.2 (11)	11G	1	562.6	29CAAACAA	0	187.9	15C	0	349.7	10G	1
	N236.5	424.5 (11)	11G	1	561.9	29CAAACAA	0	187.9	15C	0	349.4	10G	1
	N236.6	424.2 (11)	11G	1	562.1	29CAAACAA	0	187.9	15C	0	349.1	10G*	1
V51/3 rd	N354.1	424.4	11G	1	554.8	28CAAACAA	1	190.9>191.8 {1.2}	18C	0	349.3	10G	1
	N354.2	424.4 (11)	11G	1	554.7	28CAAACAA	1	190.9>191.8 {1.2}* (10)	18C	0	349.5	10G	1
	N354.3	424.5 (11)	11G	1	555.2	28CAAACAA	1	190.9>191.9 {1.1}	18C	0	349.4	10G	1
	N354.4	424.4 (11)	11G	1	555.4	28CAAACAA	1	190.8*	18C	0	349.4	10G	1
	N354.5	424.5 (11)	11G	1	569.8	30CAAACAA	0	188.0	15C	0	349.5	10G	1
	N354.6	424.3 (11)	11G	1	554.9	28CAAACAA	1	190.9>190.0 {1.1}	17C	0	349.5	10G	1
V51/4 th	N424.1	423.5	10G	0	583.6 (13)	32CAAACAA	0	188.1	15C	0	349.7 (10)	10G	1
	N424.2	423.5	10G	0	582.9	32CAAACAA	0	187.9	15C	0	349.4	10G	1
	N424.3	423.2	10G	0	583.1	32CAAACAA	0	188.0	15C	0	349.5	10G	1
	N424.4	423.5	10G	0	583.6	32CAAACAA	0	187.9	15C	0	349.5	10G	1
	N424.5	423.2	10G	0	583.0	32CAAACAA	0	187.9>188.8 {1.1}	15C	0	349.3	10G	1
	N424.6	423.4	10G	0	576.5	31CAAACAA	1	188>188.9 {1.2}	15C	0	349.4	10G	1
V52/1 st	N52.1	426.2	12G	0	562.3	29CAAACAA	0	189.0>190 {1.2}	16C	1	349.6 (10)	10G	1
	N52.2	425.7	12G	0	562.1	29CAAACAA	0	189.1>190 {1.2}	16C	1	349.3	10G	1
	N52.3	426.0	12G	0	562.6	29CAAACAA	0	189>190 {1.2}	16C	1	349.3	10G	1
	N52.4	425.8	12G	0	562.4	29CAAACAA	0	189>189.9 {1.2}	16C	1	349.4	10G	1
	N52.5	426.1	12G	0	562.4	29CAAACAA	0	189>190 {1.2}	16C	1	349.5	10G	1
	N52.6	425.9	12G	0	562.5	29CAAACAA	0	189.0>189.9 {1.2}	16C	1	349.4	10G	1
V52/2 nd	N238.1	427.2	13G	0	533.7	25CAAACAA	1	189.0>189.8 {1.0}	16C	1	349.4	10G	1
	N238.2	427.9	14G	1	533.7	25CAAACAA	1	189>189.9 {1.1}	16C	1	349.4	10G	1
	N238.3	427.1	13G	0	576.3	31CAAACAA	1	188.9>189.9 {1.0}	16C	1	349.4	10G	1
	N238.4	427.1	13G	0	533.6	25CAAACAA	1	189.0	16C	1	349.5	10G	1
	N238.5	427.2	13G	0	533.7	25CAAACAA	1	188.9>189.9 {1.0}	16C	1	349.6	10G	1
	N238.6	427.9>426.8 {1.1}	14G	1	576.9	31CAAACAA	1	189.0	16C	1	349.4	10G	1
V52/3 rd	N342.1	427.1	13G	0	577.4	31CAAACAA*	1	188.0	15C	0	351.5	12G	0
V52/4 th	Cleared												
V54/1 st	N54.1.1	424.6 (11)	11G	1	388.8 (4)	4CAAACAA*	1	186.2	13C	1	349.6	10G	1
	N54.1.2	424.5	11G	1	388.8	4CAAACAA	1	186.3>185.2 {1.1}	12C	0	352.5>351.5 {1.0}	12G	0

	N54.1.3	424.3	11G	1	388.7	4CAAACAA	1	186.2	13C	1	352.5>351.5 {1.2}	13G	1
	N54.1.4	424.5	11G	1	388.7	4CAAACAA	1	186.2>185.2 {1.0}	12C	0	352.5>351.5 {1.1}	12G	0
	N54.1.5	424.4	11G	1	388.7	4CAAACAA	1	186.2>185.1 {1.1}	12C	0	352.5>351.5 {1.0}	12G	0
	N54.1.6	424.5	11G	1	388.8	4CAAACAA	1	186.1>185 {1.2}	13C	1	352.5>351.5 {1.0}	12G	0
V54/2 nd	N237.1	424.4	11G	1	388.8	4CAAACAA	1	185.1	12C	0	351.4	12G	0
	N237.2	425.6	12G	0	388.8	4CAAACAA	1	186.1	13C	1	351.5	12G	0
	N237.3	424.4	11G	1	388.7	4CAAACAA	1	184.2	11C	0	351.5	12G	0
	N237.4	424.6	11G	1	388.8	4CAAACAA	1	188>187 {1.1}	14C	0	351.5	12G	0
	N237.5	424.6	11G	1	388.7	4CAAACAA	1	184.1	11C	0	352.3	13G	1
	N237.6	424.5	11G	1	388.8	4CAAACAA	1	187>188 {1.1}	14C	0	352.5	13G	1
V54/3 rd	N343.1	424.5 (11)	11G	1	388.6 (4)	4CAAACAA	1	186.2	13C	1	351.5>352.6 {1.2}	12G	0
	N343.2	424.3	11G	1	388.7	4CAAACAA	1	185.1	12C	0	352.4	13G	1
	N343.3	424.5	11G	1	388.7	4CAAACAA	1	186.1	13C	1	352.5	13G	1
	N343.4	424.3	11G	1	388.6	4CAAACAA	1	185.2	12C	0	352.4	13G	1
	N343.5	424.5	11G	1	388.8	4CAAACAA	1	184.2	11C	0	352.5>351.4 {1.1}	12G	0
	N343.6	424.3	11G	1	388.7	4CAAACAA	1	186.1	13C	1	352.4	13G	1
V54/4 th	Replaced												
V58/1 st	N58.1	426.2	12G	0	562.9 (13)	29CAAACAA	0	188.1	15C	0	349.7 (10)	10G	1
	N58.2	426.1	12G	0	561.3	29CAAACAA	0	188.2	15C	0	349.5	10G	1
	N58.3	426.8>425.8 {1.1}	13G	0	562.2	29CAAACAA	0	188.0	15C	0	349.5	10G	1
	N58.4	426.1	12G	0	561.7	29CAAACAA	0	188.3	15C	0	349.6	10G	1
	N58.5	425.8	12G	0	562.1	29CAAACAA	0	188.2	15C	0	349.5	10G	1
	N58.6	425.7	12G	0	561.8	29CAAACAA	0	188.2	15C	0	349.5	10G	1
V58/2 nd	N240.1	427.9	14G	1	562.2	29CAAACAA	0	188.0	15C	0	349.5	10G	1
	N240.2	426.1	12G	0	562.4	29CAAACAA	0	187.9	15C	0	349.6	10G	1
V58/3 rd	Cleared												
V58/4 th	N429.1	425.9	12G	0	533.8 (13)	25CAAACAA	1	189.0	16C	1	349.7>348.6 {1.2}	10G	1
	N429.2	426.8>425.8 {1.1}	13G	0	533.7	25CAAACAA	1	189.0	16C	1	349.4	10G	1
	N429.3	427.1>426.1 {1.1}	13G	0	518.5	23CAAACAA	0	189.0	16C	1	349.6	10G	1
	N429.4	426.7>425.8 {1.2}	13G	0	518.8	23CAAACAA	0	189.0	16C	1	349.3	10G	1
	N429.5	426.1>427.1	13G	0	533.5	25CAAACAA	1	189.0	16C	1	349.6	10G	1

		{1.0}											
	N429.6	426.8>425.8 {1.1}	13G	0	518.7	23CAAACAA	0	189.1	16C	1	349.4	10G	1
V59/1 st	N59.1.1	426.1	12G	0	563.1	29CAAACAA	0	189.9	17C	0	349.6 (10)	10G	1
	N59.1.2	426.0>427.0 {1.1}	13G	0	562.7	29CAAACAA	0	189.9	17C	0	349.3	10G	1
	N59.1.3	425.8	12G	0	556.0	28CAAACAA	0	189.9	17C	0	349.5>348.6 {1.0}	9G	0
	N59.1.4	426.0>427.1 {1.2}	12G	0	562.6	29CAAACAA	0	190.0	17C	0	349.5	10G	1
	N59.1.5	425.8>426.8 {1.1}	13G	0	562.6	29CAAACAA	0	190.0	17C	0	349.4	10G	1
	N59.1.6	426.0>427.1 {1.2}	12G	0	562.3	29CAAACAA	0	190.0	17C	0	349.6	10G	1
	N59.1.7	426.2	12G	0	563.3	29CAAACAA	0	No data			349.6	10G	1
	N59.1.8	426.1	12G	0	562.9	29CAAACAA	0	No data			349.6	10G	1
	N59.1.9	426.1	12G	0	562.7	29CAAACAA	0	No data			349.6	10G	1
	N59.1.10	426.0	12G	0	562.8	29CAAACAA	0	No data			349.6	10G	1
V59/2 nd	N253.1	425.8	12G	0	562.3	29CAAACAA	0	190.0>189.0 {1.0}	16C	1	349.5	10G	1
	N253.2	426.0	12G	0	562.2	29CAAACAA	0	189.0>190.0 {1.1}	16C	1	349.4	10G	1
	N253.3	425.6>426.8 {1.0}	13G	0	562.6	29CAAACAA	0	190.0>189.0 {1.1}	16C	1	349.3	10G	1
	N253.4	426.0>427 {1.1}	13G	0	562.3	29CAAACAA	0	189.9>189.0 {1.0}	16C	1	349.3	10G	1
	N253.5	425.6>426.6 {1.1}	13G	0	562.9	29CAAACAA	0	189.9>189.0 {1.1}	16C	1	349.4	10G	1
	N253.6	426.0>427 {1.1}	13G	0	562.7	29CAAACAA	0	190.0>189.0 {1.0}	16C	1	349.3	10G	1
	N253.7	426.0>427.1 {1.1}	13G	0	562.5	29CAAACAA	0	No data			348.5>349.6 {1.1}	9G	0
	N253.8	426.1>427.1 {1.1}	13G	0	562.9	29CAAACAA	0	No data			348.5>349.5 {1.1}	9G	0
	N253.9	426.1>427.2 {1.1}	13G	0	562.7	29CAAACAA	0	No data			348.5>349.5 {1.1}	9G	0
	N253.10	426.1>427.2 {1.1}	13G	0	563.1	29CAAACAA	0	No data			348.4>349.5 {1.0}	9G	0
V59/3 rd	N352.1	425.7	12G	0	562.5	29CAAACAA	0	189.0>187.8 {1.1}	15C	0	349.3	10G	1
	N352.2	425.9>424.9 {1.2}	12G	0	562.5	29CAAACAA	0	189.0>188.0 {1.1}	15C	0	349.4	10G	1
	N352.3	425.7>426.7	13G	0	562.4	29CAAACAA	0	188.6>187.9 {1.1}	15C	0	349.3	10G	1

		{1.1}											
	N352.4	424.9	11G	1	576.6	31CAAACAA	1	189.0>187.9 {1.1}	15C	0	350.7	11G	0
	N352.5	425.7>426.8 {1.0}	13G	0	562.3	29CAAACAA	0	189.0>187.9 {1.1}	15C	0	349.4	10G	1
	N352.6	426.0>427.0 {1.0}	13G	0	562.9	29CAAACAA	0	189.0>187.9 {1.1}	15C	0	349.5	10G	1
	N352.7	424.9>426.0 {1.1}	12G	0	555.4	28CAAACAA	1	No data			348.4>349.4 {1.0}	9G	0
	N352.8	426.1	12G	0	562.8	29CAAACAA	0	No data			349.5	10G	1
	N352.9	425.1>426.1 {1.1}	12G	0	562.2	29CAAACAA	0	No data			349.5	10G	1
	N352.10	425.2>426.2 {1.1}	12G	0	563.5	29CAAACAA	0	No data			349.5	10G	1
V59/4 th	N438.1	426.0	12G	0	562.7 (13)	29CAAACAA	0	188.0>188.9 {1.0}	15C	0	349.7 (10)	10G	1
	N438.2	425.8>426.8 {1.0}	13G	0	562.6	29CAAACAA	0	188.0>188.9 {1.1}	15C	0	349.4	10G	1
	N438.3	427.0>426.0 {1.0}	13G	0	562.4	29CAAACAA	0	187.9>188.9 {1.1}	15C	0	349.4	10G	1
	N438.4	425.7	12G	0	562.5	29CAAACAA	0	188.0>188.9 {1.1}	15C	0	349.4	10G	1
	N438.5	426.0>427.0 {1.2}	12G	0	562.4	29CAAACAA	0	187.9>188.9 {1.1}	15C	0	349.4	10G	1
	N438.6	426.0>427.0 {1.0}	13G	0	562.0	29CAAACAA	0	187.9>188.9 {1.1}	15C	0	349.5	10G	1
	N438.7	426.1>427.2 {1.1}	13G	0	563.3	29CAAACAA	0	No data					
	N438.8	426.0>427.0 {1.1}	13G	0	563.3	29CAAACAA	0	No data			349.4	10G	1
	N438.9	426.0>427.0 {1.1}	13G	0	563.3	29CAAACAA	0	No data			349.6	10G	1
	N438.10	426.1>427.1 {1.1}	13G	0							349.3	10G	1
V88/1 st	N88.1.1	427.6	14G*	1	518.7	23CAAACAA*	0	188.9	16C	1	351.6>352.6 {1.0}	12G	0
	N88.1.2	427.6	14G	1	525.8	24CAAACAA	0	188.7	16C	1	352.5>351.4 {1.1}	12G	0
	N88.1.3	427.7	14G	1	518.5	23CAAACAA	0	188.9	16C	1	351.5>352.5 {1.1}	12G	0
	N88.1.4	427.6	14G	1	518.3	23CAAACAA	0	188.9	16C	1	352.5>351.5 {1.1}	12G	0
	N88.1.5	427.5	14G	1	518.5	23CAAACAA	0	189.1	16C	1	351.5>352.5 {1.0}	12G	0

	N88.1.6	427.7	14G	1	518.7	23CAAACAA	0	188.9	16C	1	351.4>352.5 {1.1}	12G	0
V88/2 nd	N272.1	426.6	13G*	0	518.7	23CAAACAA	0	188.9	16C	1	352.5>351.5 {1.1}	12G	0
	N272.2	426.6	13G	0	518.7	23CAAACAA	0	188.8	16C	1	351.5>352.5 {1.1}	12G	0
	N272.3	426.6	13G	0	518.4	23CAAACAA	0	188.9	16C	1	352.5>351.5 {1.0}	12G	0
	N272.4	426.6	13G	0	518.6	23CAAACAA	0	189.0	16C	1	351.5>352.5 {1.0}	12G	0
	N272.5	426.6	13G	0	518.6	23CAAACAA	0	188.9	16C	1	351.5>352.5 {1.1}	12G	0
	N272.6	426.7	13G	0	518.7	23CAAACAA	0	188.9	16C	1	351.5>352.6 {1.0}	12G	0
V88/3 rd	N369.1	427.8	14G	1	511.1	22CAAACAA	1	188.9	16C	1	351.5>352.5 {1.1}	12G	0
	N369.2	427.8	14G	1	511.3	22CAAACAA	1	189.0	16C	1	351.5>352.6 {1.0}	12G	0
	N369.3	427.7	14G	1	503.6	21CAAACAA	0	189.2	16C	1	351.5>350.6 {1.1}	12G	0
	N369.4	427.7	14G	1	503.6	21CAAACAA	0	189.1	16C	1	351.5>350.6 {1.1}	12G	0
	N369.5	427.7	14G	1	511.3	22CAAACAA	1	188.9	16C	1	352.4>351.5 {1.1}	13G	1
	N369.6	427.7	14G	1	511.2	22CAAACAA	1	189.0>188.0 {1.0}	15C	0	351.5>352.5 {1.0}	13G	1
V88/4 th	N449.1	427.7	14G	1	540.7 (13)	26CAAACAA	0	188.9	16C	1	351.6>352.3 (1.2)	13G	1
	N449.2	427.7	14G	1	533.5	25CAAACAA	1	188.9	16C	1	352.5>351.5 {1.1}	13G	1
	N449.3	427.6	14G	1	540.7	26CAAACAA	0	189.9	17C	0	352.5>351.5 {1.1}	13G	1
	N449.4	427.6	14G	1	540.7	26CAAACAA	0	189.0	16C	1	352.5>351.5 {1.1}	13G	1
	N449.5	427.7	14G	1	547.7	27CAAACAA	0	189.0	16C	1	351.5>352.5 {1.0}	13G	1
	N449.6	427.6	14G	1	533.2	25CAAACAA	1	189.0	16C	1	351.5>352.5 {1.1}	13G	1
V138/1 st	N138.1	427.1>426.1 {1.1}	13G	0	562.3	29CAAACAA	0	188.9	16C	1	349.5 (10)	10G	1
	N138.2	426.1>427.2 {1.0}	13G	0	562.3	29CAAACAA	0	188.9	16C	1	349.6	10G	1
	N138.3	426.1	12G	0	561.8	29CAAACAA	0	188.9>189.9 {1.1}	16C	1	349.7	10G	1
	N138.4	426.1>427.2 {1.0}	13G	0	562.5	29CAAACAA	0	188.9	16C	1	349.5	10G	1
	N138.5	427.1>426.1 {1.1}	13G	0	562.7	29CAAACAA	0	188.9>189.9 {1.2}	16C	1	349.6	10G	1
	N138.6	426.1	12G	0	561.8	29CAAACAA	0	189.0	16C	1	349.6	10G	1
V138/2 nd	N288.1	426.0	12G	0	562.4	29CAAACAA	0	188.9	16C	1	349.4	10G	1
	N288.2	427.1	13G	0	562.6	29CAAACAA	0	188.0>189.0 {1.1}	15C	0	349.5	10G	1
	N288.3	425.0	11G	1	562.2	29CAAACAA	0	188.0	15C	0	349.3	10G	1
	N288.4	427.1	13G	0	561.9	29CAAACAA	0	188.0>188.9 {1.1}	15C	0	349.6	10G	1
	N288.5	426.1	12G	0	561.9	29CAAACAA	0	188.0	15C	0	349.5	10G	1
	N288.6	427.1	13G	0	561.9	29CAAACAA	0	187.9>188.9 {1.2}	15C	0	350.6	11G	0
V138/3 rd	N331.1	427.9>426.9	14G*	1	561.9	29CAAACAA	0	188.9>187.9 {1.1}	15C	0	348.2 (9)	9G	0

		{1.2}											
	N331.2	426.0>427.0 {1.1}	13G	0	576.1	31CAAACAA	1	188.9>189.8 {1.2}	16C	1	349.5	10G	1
	N331.3	426>427 {1.1}	13G	0	575.9	31CAAACAA	1	187.9>188.9 {1.1}	15C	0	349.5	10G	1
	N331.4	427.0>426.0 {1.2}	13G	0	575.9	31CAAACAA	1	188.9>187.9 {1.2}	16C	1	349.6	10G	1
	N331.5	426.0>427.1 (1.0)	13G	0	576.3	31CAAACAA	1	187.8>188.9 {1.1}	15C	0	349.4	10G	1
	N331.6	426.0>427.0 {1.1}	13G	0	576.3	31CAAACAA	1	188.9>189.8 {1.2}	16C	1	348.4	9G	0
V138/4 th	Replaced												
cc167													
V124/1 st	N124.1	420.7 (11)	11G	1	484.2	18CAAACAA*	0	187.1	11C	0	349.3 (10)	10G	1
	N124.2	420.7	11G	1	484.3	18CAAACAA	0	187.0	11C	0	349.3	10G	1
	N124.3	420.7	11G	1	484.2	18CAAACAA	0	187.1	11C	0	349.3	10G	1
	N124.4	420.7	11G	1	484.2	18CAAACAA	0	187.0	11C	0	349.3	10G	1
	N124.5	420.6	11G	1	484.1	18CAAACAA	0	187.1	11C	0	349.3	10G	1
	N124.6	420.6	11G	1	484.1	18CAAACAA*	0	186.9	11C	0	349.3	10G	1
V124/2 nd	N290.1	420.7	11G	1	484.3	18CAAACAA	0	187.1	11C	0	349.3	10G	1
	N290.2	420.8	11G	1	484.2	18CAAACAA	0	187.0	11C	0	349.4	10G	1
	N290.3	420.8	11G	1	484.3	18CAAACAA	0	187.1	11C	0	349.4	10G	1
	N290.4	420.8	11G	1	491.1	19CAAACAA	1	187.0	11C	0	349.3	10G	1
	N290.5	420.7	11G	1	484.2	18CAAACAA	0	187.1	11C	0	349.3	10G	1
	N290.6	420.7	11G	1	484.2	18CAAACAA	0	187.0	11C	0	349.3	10G	1
V124/3 rd	N336.1	420.6 (11)	11G	1	484.2	18CAAACAA	0	187.1	11C	0	349.3 (10)	10G	1
	N336.2	420.7	11G	1	506.3	21CAAACAA	0	187.0	11C	0	349.3	10G	1
	N336.3	420.7	11G	1	484.3	18CAAACAA	0	187.1	11C	0	349.3	10G	1
	N336.4	420.7	11G	1	484.2	18CAAACAA	0	186.9	11C	0	349.3	10G	1
	N336.5	420.7	11G	1	484.3	18CAAACAA	0	187.1	11C	0	349.3	10G	1
	N336.6	420.8	11G	1	484.2	18CAAACAA	0	186.9	11C	0	349.4	10G	1
V64/1 st	N64.1	420.6 (11)	11G	1	484.2	18CAAACAA	0	187.0	11C	0	349.3 (10)	10G	1
	N64.2	420.6	11G	1	484.1	18CAAACAA	0	186.9	11C	0	349.3	10G	1
	N64.3	420.7	11G	1	484.2	18CAAACAA	0	187.0	11C	0	349.3	10G	1
	N64.4	420.7	11G	1	484.2	18CAAACAA	0	186.9	11C	0	349.3	10G	1
	N64.5	420.7	11G	1	484.2	18CAAACAA	0	187.1	11C	0	349.3	10G	1
	N64.6	420.7	11G	1	484.3	18CAAACAA	0	186.9	11C	0	349.3	10G	1

V69/2 nd	N258.1	417.1 (8)	8G*	1	444.1 (12)	12CAAACAA*	0	187.4>188.4 {1.1}	16C*	1	347.1	10G*	1
	N258.2	417.0	8G	1	444.4	12CAAACAA	0	187.5>188.4 {1.1}	16C	1	347.2	10G	1
	N258.3	417.2	8G	1	444.2	12CAAACAA	0	187.4>188.4 {1.1}	16C	1	347.1	10G	1
	N258.4	417.1	8G	1	444.0	12CAAACAA	0	187.4>188.4 {1.2}	16C	1	347.1	10G	1
	N258.5	417.4	8G	1	444.1	12CAAACAA	0	187.3>188.2 {1.2}	16C	1	347.3	10G	1
	N258.6	417.1	8G	1	444.1	12CAAACAA	0	187.4>188.4 {1.0}	16C	1	347.2	10G	1
V69/3 rd	No sample												
V69/4 th	N431.1	417.4 (8)	8G	1	444.3 (12)	12CAAACAA	0	187.5>188.6 {1.2}	16C*	1	347.2 (10)	10G	1
	N431.2	417.3	8G	1	444.3	12CAAACAA	0	187.5>188.6 {1.1}	16C	1	347.2	10G	1
	N431.3	417.5	8G	1	444.2	12CAAACAA	0	187.6>188.5 {1.2}	16C	1	347.3	10G	1
	N431.4	417.3	8G	1	444.4	12CAAACAA	0	187.6>188.6 {1.1}	16C	1	347.2	10G	1
	N431.5	417.3	8G	1	444.4	12CAAACAA	0	187.5>188.5 {1.2}	16C	1	347.2	10G	1
	N431.6	417.2	8G	1	444.4	12CAAACAA	0	187.5>188.5 {1.1}	16C	1	347.3	10G	1
V93/1 st	Non-carrier												
V93/2 nd	N264.1	424.0	15G*??	1??	445.6 (11)	12CAAACAA	0	189.9	16C	1	349.4 (10)	10G	1
	N264.2	424.0	14G	1	445.5	12CAAACAA	0	189.8	16C	1	349.5	10G	1
	N264.3	425.0	15G	0	438.7	11CAAACAA	0	189.9	16C	1	349.4	10G	1
	N264.4	423.8	14G*	1	445.5	12CAAACAA	0	189.9	16C	1	349.5	10G	1
	N264.5	423.7	14G	1	445.4	12CAAACAA	0	189.9	16C	1	349.4	10G	1
	N264.6	424.3	15G	0	445.4	12CAAACAA	0	189.9	16C	1	349.4	10G	1
V93/3 rd	N359.1	425.4	16G*	0	445.5	12CAAACAA	0	190.9	17C	0	349.4	10G	1
	N359.2	425.4	16G	0	445.6	12CAAACAA	0	190.9	17C	0	349.4	10G	1
	N359.3	425.3	16G	0	445.4	12CAAACAA	0	190.9	17C	0	349.4	10G	1
	N359.4	425.2	16G	0	445.4	12CAAACAA	0	190.9	17C	0	349.4	10G	1
	N359.5	425.3	16G	0	452.1	13CAAACAA	1	189.9	16C	1	349.5	10G	1
	N359.6	425.3	16G	0	445.5	12CAAACAA	0	190.9	17C	0	349.4	10G	1
V96/1 st	Non-carrier												
V96/2 nd	N259.1	417.3 (8)	8G*	1	445.5 (12)	12CAAACAA*	0	185.1	11C	0	349.4 (10)	10G*	1
	N259.2	417.4	8G	1	445.5	12CAAACAA	0	185.0	11C	0	349.5	10G	1
	N259.3	417.5	8G	1	445.5	12CAAACAA	0	185.1	11C	0	349.5	10G	1
	N259.4	417.4	8G	1	445.5	12CAAACAA	0	184.9	11C	0	349.4	10G	1

								(13)					
	N259.5	417.4	8G	1	445.5	12CAAACAA	0	185.1	11C	0	349.4	10G	1
	N259.6	417.4	8G	1	445.5	12CAAACAA	0	184.9	11C	0	349.4	10G	1
V96/3 rd	N370.1	417.3	8G	1	445.5	12CAAACAA	0	185.1	11C	0	348.5	9G	0
	N370.2	417.3	8G	1	445.4	12CAAACAA	0	185.0 (10)	11C	0	348.4	9G	0
	N370.3	417.4	8G	1	445.4	12CAAACAA	0	185.1	11C	0	351.4	12G	0
	N370.4	417.5	8G	1	445.6	12CAAACAA	0	185.0	11C	0	349.4>350.4 {1.2}	10G	1
	N370.5	417.5	8G	1	445.5	12CAAACAA	0	185.1	11C	0	348.4	9G	0
	N370.6	417.5	8G	1	445.6	12CAAACAA	0	185.0	11C	0	348.4	9G	0
V96/4 th	N445.1	417.5 (8)	8G	1	445.5 (11)	12CAAACAA	0	185.2	11C	0	348.4 (9)	9G	0
	N445.2	417.4	8G	1	445.5	12CAAACAA	0	185.3	11C	0	348.5	9G	0
	N445.3	417.4	8G	1	445.5	12CAAACAA	0	185.3	11C	0	348.5	9G	0
	N445.4	417.4	8G	1	445.5	12CAAACAA	0	185.2	11C	0	348.5	9G	0
	N445.5	417.4	8G	1	445.5	12CAAACAA	0	185.1	11C	0	348.5	9G	0
	N445.6	417.4	8G	1	445.6	12CAAACAA	0	185.1	11C	0	349.4	10G	1
V222/1 st	N222.1	417.4 (8)	8G	1	445.6 (12)	12CAAACAA	0	190.0	16C	1	349.5	10G	1
	N222.2	417.4	8G	1	445.5	12CAAACAA	0	190.0	16C	1	349.5	10G	1
	N222.3	417.4	8G	1	445.4	12CAAACAA	0	190.0	16C	1	349.4	10G	1
	N222.4	417.4	8G	1	445.5	12CAAACAA	0	190.0	16C	1	349.4	10G	1
	N222.5	417.4	8G	1	445.5	12CAAACAA	0	190.0	16C	1	349.4	10G	1
	N222.6	417.4	8G	1	445.5	12CAAACAA	0	189.9	16C	1	349.5	10G	1
V222/2 nd	N309.1	417.4	8G	1	445.5	12CAAACAA	0	189.9	16C	1	349.4	10G	1
	N309.2	417.4	8G	1	445.6	12CAAACAA	0	189.9	16C	1	349.4	10G	1
	N309.3	417.2	8G	1	445.5	12CAAACAA	0	189.9	16C	1	349.4	10G	1
	N309.4	417.3	8G	1	445.5	12CAAACAA	0	189.9	16C	1	349.3	10G	1
	N309.5	417.1	8G	1	445.5	12CAAACAA	0	190.0	16C	1	349.4	10G	1
	N309.6	417.2	8G	1	445.4	12CAAACAA	0	190.0	16C	1	349.4	10G	1
V222/3 rd	N391.1	417.3	8G	1	445.5	12CAAACAA	0	189.1	15C	0	349.5	10G	1
	N391.2	417.4	8G	1	445.5	12CAAACAA	0	189.0	15C	0	349.5	10G	1
	N391.3	417.4	8G	1	445.6	12CAAACAA	0	189.0	15C	0	349.5	10G	1
	N391.4	417.4	8G	1	445.5	12CAAACAA	0	189.0	15C	0	349.4	10G	1
	N391.5	417.4	8G	1	445.5	12CAAACAA	0	189.0	15C	0	349.4	10G	1
	N391.6	417.4	8G	1	445.5	12CAAACAA	0	189.0	15C	0	349.4	10G	1

V222/4 th	N459.1	417.4 (8)	8G	1	445.5 (11)	12CAAACAA	0	189.0	15C	0	349.5 (10)	10G	1
	N459.2	417.3	8G	1	452.1	13CAAACAA	1	189.1	15C	0	349.4	10G	1
	N459.3	417.3	8G	1	445.4	12CAAACAA	0	189.0	15C	0	349.5	10G	1
	N459.4	417.3	8G	1	445.4	12CAAACAA	0	189.0	15C	0	349.5	10G	1
	N459.5	417.4	8G	1	458.3	14CAAACAA	0	189.0	15C	0	349.4	10G	1
	N459.6	417.4	8G	1	445.5	12CAAACAA	0	189.0	15C	0	349.4	10G	1
cc60													
V114/1 st	N114.1	424.1	14G*	1	464.6 (12)	15CAAACAA	0	Absent			430.7	13G	1
	N114.2	424.3	14G	1	464.7	15CAAACAA	0	Absent			430.7	13G	1
	N114.3	424.3	14G	1	464.7	15CAAACAA	0	Absent			430.7	13G	1
	N114.4	424.3	14G	1	464.7	15CAAACAA	0	Absent			430.7	13G	1
	N114.5	424.2>423.2 {1.2}	14G	1	464.7	15CAAACAA	0	Absent			430.7	13G	1
	N114.6	424.2	14G	1	464.7	15CAAACAA	0	Absent			430.7	13G	1
V114/2 nd	N283.1	424.4	14G	1	464.2	15CAAACAA	0	Absent			430.8	13G	1
	N283.2	424.4	14G	1	464.6	15CAAACAA	0	Absent			430.7	13G	1
	N283.3	425.2>424.3 [1.2]	15G	0	464.7	15CAAACAA	0	Absent			430.8	13G	1
	N283.4	424.4	14G	1	464.6	15CAAACAA	0	Absent			430.7	13G	1
	N283.5	425.5	15G	0	464.3	15CAAACAA	0	Absent			430.7	13G	1
	N283.6	424.4	14G	1	464.2	15CAAACAA	0	Absent			430.7	13G	1
V114/3 rd	N330.1	420.2 (10)	10G	0	464.7 (12)	15CAAACAA	0	Absent			427.5	11G	0
	N330.2	420.2	10G	0	464.5	15CAAACAA	0	Absent			427.8	11G	0
	N330.3	420.1	10G	0	464.6	15CAAACAA	0	Absent			429.7	12G	0
	N330.4	420.1	10G	0	464.6	15CAAACAA	0	Absent			428.6	11G	0
	N330.5	420.3	10G	0	464.7	15CAAACAA	0	Absent			433.3??	13G*	1
	N330.6	420.4	10G	0	464.7	15CAAACAA	0	Absent			428.5	12G	0
V134/1 st	N134.1	424.1	14G	1	464.7 (10)	15CAAACAA	0	Absent			430.7	13G	1
	N134.2	424.4	14G	1	464.7	15CAAACAA	0	Absent			430.7	13G	1
	N134.3	424.5	14G	1	464.7	15CAAACAA	0	Absent			430.7	13G	1
	N134.4	424.5	14G	1	464.6	15CAAACAA	0	Absent			430.8	13G	1
	N134.5	424.2	14G	1	464.7	15CAAACAA	0	Absent			430.8	13G	1

V113/1 st	N113.1	424.0	14G	1	464.2	15CAAACAA*	0	Absent			428.5	12G	0
	N113.2	424.1	14G	1	464.3	15CAAACAA	0	Absent			429.9	13G	1
	N113.3	424.1	14G	1	464.7	15CAAACAA	0	Absent			429.5	13G	1
	N113.4	424.1	14G	1	464.7	15CAAACAA	0	Absent			429.7	13G	1
	N113.5	424.1	14G	1	464.7	15CAAACAA	0	Absent			429.7	13G	1
	N113.6	424.2>423.2 {1.2}	14G	1	464.7	15CAAACAA	0	Absent			430.0	13G	1
V113/2 nd	N281.1	424.1	14G	1	464.6	15CAAACAA	0	Absent			429.4	13G*	1
	N281.2	424.2	14G	1	464.6	15CAAACAA	0	Absent			429.8>428.8 {1.2}	13G	1
	N281.3	424.1	14G	1	464.7	15CAAACAA	0	Absent			429.9	13G	1
	N281.4	423.1	13G	0	464.5	15CAAACAA	0	Absent			430.0	13G	1
	N281.5	423.9	14G	1	464.6	15CAAACAA	0	Absent			429.6	13G	1
	N281.6	423.9	14G	1	464.3	15CAAACAA	0	Absent			430.1	13G	1
V115/1 st	N115.1	424.1	14G	1	464.7	15CAAACAA	0	Absent			429.7	13G	1
	N115.2	424.2>423.2 {1.2}	14G	1	464.7	15CAAACAA	0	Absent			430.2	13G	1
	N115.3	424.2	14G	1	464.7	15CAAACAA	0	Absent			429.7	13G	1
	N115.4	424.3	14G	1	464.7	15CAAACAA	0	Absent			430.0	13G	1
	N115.5	424.0>423.0 {1.1}	14G	1	464.7	15CAAACAA	0	Absent			429.9>428.9 {1.2}	13G	1
	N115.6	424.1>423.1 {1.2}	14G	1	464.6	15CAAACAA	0	Absent			430.2	13G	1
V115/2 nd	N282.1	423.9	14G	1	464.3	15CAAACAA	0	Absent			429.6	13G	1
	N282.2	424.0>422.9 {1.2}	14G	1	464.2	15CAAACAA	0	Absent			430.1	13G	1
	N282.3	424.0	14G	1	464.3	15CAAACAA	0	Absent			430.0	13G	1
	N282.4	424.1	14G	1	464.4	15CAAACAA	0	Absent			429.6	13G	1
	N282.5	424.2	14G	1	464.5	15CAAACAA	0	Absent			429.7	13G	1
	N282.6	424.2	14G	1	464.4	15CAAACAA	0	Absent			430.0	13G	1
cc32													
V176/1 st	N176.1	429.8	16G	0	533.6 (14)	24CAAACAA	1	Absent			354.5	15G	0
	N176.2	429.9	16G	0	511.1	21CAAACAA	1	Absent			354.5	15G	0
	N176.3	429.9	16G	0	511.2	21CAAACAA	1	Absent			354.5	15G	0
	N176.4	429.9	16G	0	526.0	23CAAACAA	0	Absent			354.6	15G	0
	N176.5	429.9	16G	0	525.7	23CAAACAA	0	Absent			354.5	15G	0

	N176.6	429.9	16G	0	511.3	21CAAACAA	1	Absent			354.5	15G	0
V176/2 nd	N318.1	430.1	16G	0	533.7	24CAAACAA	1	Absent			355.4	16G	1
	N318.2	430.1	16G	0	525.9	23CAAACAA	0	Absent			355.5	16G	1
	N318.3	430.2	16G	0	533.3	24CAAACAA	1	Absent			355.5	16G	1
	N318.4	430.2	16G	0	533.6	24CAAACAA	1	Absent			355.4	16G	1
	N318.5	430.3	16G	0	526.0	23CAAACAA	0	Absent			355.6	16G	1
	N318.6	430.1	16G	0	525.9	23CAAACAA	0	Absent			355.4	16G	1
V176/3 rd	N399.1	429.9	16G	0	541.1	25CAAACAA	0	Absent			354.6	15G	0
	N399.2	428.9	15G	0	533.5	24CAAACAA	1	Absent			354.5	15G	0
	N399.3	428.9	15G	0	449.8	12CAAACAA	1	Absent			354.5	15G	0
	N399.4	428.9	15G	0	540.5	25CAAACAA	0	Absent			354.6	15G	0
	N399.5	429.9	16G	0	533.1	24CAAACAA	1	Absent			354.5	15G	0
	N399.6	429.9	16G	0	541.0	25CAAACAA	0	Absent			354.5	15G	0
V176/4 th	N408.1	429.0	15G	0	533.6 (12)	24CAAACAA	1	Absent			353.5	14G	0
	N408.2	429.2	15G	0	540.7	25CAAACAA	0	Absent			353.5	14G	0
	N408.3	429.1	15G	0	533.4	24CAAACAA	1	Absent			353.7	14G	0
	N408.4	429.2	15G	0	533.8	24CAAACAA	1	Absent			353.5	14G	0
	N408.5	429.2	15G	0	540.5	25CAAACAA	0	Absent			353.5	14G	0
	N408.6	429.1	15G	0	533.4	24CAAACAA	1	Absent			353.4	14G	0

¹Fragments sizes were determined using either a GS600LIZ (black type) or GS500LIZ (blue type) size standard; numbers in curly brackets indicate the ratio between the PCR fragment sizes where two obvious peaks were observed (where double peaks were observed, a ratio of >1.2 was utilised as a cut-off with the highest peak height being used to determine repeat number, if the ratio was <1.2 then the larger peak in bp was utilised to determine repeat number except in the cases of *pgII* and *pgIH* where sequencing indicated that the these double peaks followed the following rules 191/192, 17; 190/191, 16; 189/190, 15; 188/189, 14; 187/188, 14; 186/187, 13; 185/186, 12). ²an asterisk indicates that the PCR fragment was sequenced by dideoxy sequencing and the repeat number determined by visual inspection of the trace file. ³Expression states, 1=1, 0=0 (all genes are subject to translational phase variation).

Supplementary Data Table III. Repeat numbers and expression states for the phase-variable restriction-modification genes

Volunteer /time pt	Isolate	Phase Variable Gene					
		<i>modA</i>			<i>modB</i>		
		Size ¹	Tract ²	Exp. State ³	Size ¹	Tract ²	Exp. State ³
cc174							
V43/1 st	N43.1.1	198.2	3AGCC	0	439.0	16GCCAA	1
	N43.1.2	198.2	3AGCC*	0	439.1	16GCCAA	1
	N43.1.3	198.3	3AGCC	0	439.1	16GCCAA	1
	N43.1.4	198.3	3AGCC	0	438.7	16GCCAA	1
	N43.1.5	198.3	3AGCC	0	438.7	16GCCAA	1
	N43.1.6	198.2	3AGCC	0	457.7	20GCCAA	0
V43/2 nd	N241.1	198.4	3AGCC	0	443.7	17GCCAA	0
	N241.2	198.3	3AGCC	0	443.6	17GCCAA*	0
V43/3 rd	N349.1	199.7	3AGCC*	0	438.7	16GCCAA	1
	N349.2	198.3	3AGCC	0	438.8	16GCCAA	1
	N349.3	198.3	3AGCC	0	438.7	16GCCAA	1
	N349.4	198.4	3AGCC	0	438.9	16GCCAA	1
	N349.5	198.3	3AGCC	0	439.4	16GCCAA	1
	N349.6	198.2	3AGCC	0	438.6	16GCCAA	1
V43/4 th	Cleared						
V51/1 st	N51.1	198.3	3AGCC	0	443.8	17GCCAA	0
	N51.2	198.3	3AGCC	0	443.6	17GCCAA	0
	N51.3	198.3	3AGCC	0	443.8	17GCCAA	0
	N51.4	198.5	3AGCC	0	444.0	17GCCAA	0
	N51.5	198.3	3AGCC	0	443.7	17GCCAA	0
	N51.6	198.3	3AGCC	0	444.0	17GCCAA	0
V51/2 nd	N236.1	198.3	3AGCC	0	443.8	17GCCAA	0
	N236.2	198.4	3AGCC	0	438.2	16GCCAA	1
	N236.3	198.4	3AGCC	0	438.8	16GCCAA	1
	N236.4	198.3	3AGCC	0	438.8	16GCCAA	1
	N236.5	198.3	3AGCC	0	444.0	17GCCAA	0
	N236.6	198.3	3AGCC	0	444.0	17GCCAA	0
V51/3 rd	N354.1	198.3	3AGCC	0	444.0	17GCCAA	0
	N354.2	198.2	3AGCC	0	444.0	17GCCAA	0
	N354.3	198.2	3AGCC	0	444.1	17GCCAA	0
	N354.4	198.2	3AGCC	0	444.1	17GCCAA	0

	N354.5	198.3	3AGCC	0	444.1	17GCCAA	0
	N354.6	198.2	3AGCC	0	444.0	17GCCAA	0
V51/4 th	N424.1	198.3	3AGCC	0	444.1	17GCCAA	0
	N424.2	198.4	3AGCC	0	444.0	17GCCAA	0
	N424.3	198.4	3AGCC	0	444.0	17GCCAA	0
	N424.4	198.4	3AGCC	0	458.0	20GCCAA	0
	N424.5	198.3	3AGCC	0	444.1	17GCCAA	0
	N424.6	198.2	3AGCC	0	448.9	18GCCAA	0
V52/1 st	N52.1	198.3	3AGCC	0	444.0	17GCCAA	0
	N52.2	198.5	3AGCC	0	444.0	17GCCAA	0
	N52.3	198.2	3AGCC	0	444.1	17GCCAA	0
	N52.4	198.2	3AGCC	0	444.1	17GCCAA	0
	N52.5	198.3	3AGCC	0	444.1	17GCCAA	0
	N52.6	198.3	3AGCC	0	444.1	17GCCAA	0
V52/2 nd	N238.1	198.3	3AGCC	0	444.1	17GCCAA	0
	N238.2	198.2	3AGCC	0	444.1	17GCCAA	0
	N238.3	198.4	3AGCC	0	448.9	18GCCAA	0
	N238.4	198.3	3AGCC	0	444.1	17GCCAA	0
	N238.5	198.3	3AGCC	0	444.0	17GCCAA	0
	N238.6	198.2	3AGCC	0	448.9	18GCCAA	0
V52/3 rd	N342.1	198.2	3AGCC	0	438.8	16GCCAA	1
V52/4 th	Cleared						
V54/1 st	N54.1.1	198.3	3AGCC	0	439.1 (16)	16GCCAA	1
	N54.1.2	198.3	3AGCC	0	439.1	16GCCAA	1
	N54.1.3	198.2	3AGCC	0	439.2	16GCCAA	1
	N54.1.4	198.3	3AGCC	0	439.1	16GCCAA	1
	N54.1.5	198.2	3AGCC	0	439.2	16GCCAA	1
	N54.1.6	198.3	3AGCC	0	439.1	16GCCAA	1
V54/2 nd	N237.1	198.3	3AGCC	0	439.2	16GCCAA	1
	N237.2	198.4	3AGCC	0	439.1	16GCCAA	1
	N237.3	198.5	3AGCC	0	439.1	16GCCAA	1
	N237.4	198.3	3AGCC	0	439.0	16GCCAA	1
	N237.5	198.3	3AGCC	0	439.0	16GCCAA	1
	N237.6	198.3	3AGCC	0	438.5*	16GCCAA	1
V54/3 rd	N343.1	198.2	3AGCC	0	439.2 (16)	16GCCAA	1

	N343.2	198.3	3AGCC	0	439.1	16GCCAA	1
	N343.3	198.4	3AGCC	0	439.2	16GCCAA	1
	N343.4	198.3	3AGCC	0	439.1	16GCCAA	1
	N343.5	198.2	3AGCC	0	439.1	16GCCAA	1
	N343.6	198.4	3AGCC	0	444.1	17GCCAA	0
V54/4 th	Replaced						
V58/1 st	N58.1	198.2	3AGCC	0	444.0 (17)	17GCCAA	0
	N58.2	198.4	3AGCC	0	444.1	17GCCAA	0
	N58.3	198.3	3AGCC	0	444.0	17GCCAA	0
	N58.4	198.4	3AGCC	0	444.0	17GCCAA	0
	N58.5	198.3	3AGCC	0	443.9	17GCCAA	0
	N58.6	198.4	3AGCC	0	444.0	17GCCAA	0
V58/2 nd	N240.1	198.4	3AGCC	0	453.5	19GCCAA	1
	N240.2	198.3	3AGCC	0	471.7	23GCCAA	0
V58/3 rd	Cleared						
V58/4 th	N429.1	198.3	3AGCC	0	394.4 (7)	8GCCAA	0
	N429.2	198.3	3AGCC	0	394.8	8GCCAA	0
	N429.3	198.2	3AGCC	0	467.1	22GCCAA	1
	N429.4	198.4	3AGCC	0	471.8	23GCCAA	0
	N429.5	198.3	3AGCC	0	394.5	8GCCAA	0
	N429.6	198.4	3AGCC	0	467.1	22GCCAA	1
V59/1 st	N59.1.1	198.4	3AGCC	0	443.7 (17)	17GCCAA	0
	N59.1.2	198.4	3AGCC	0	443.1	17GCCAA	0
	N59.1.3	198.3	3AGCC	0	444.0	17GCCAA	0
	N59.1.4	198.4	3AGCC	0	444.0	17GCCAA	0
	N59.1.5	198.3	3AGCC	0	444.0	17GCCAA	0
	N59.1.6	198.3 (3)	3AGCC	0	444.0	17GCCAA	0
V59/2 nd	N253.1	198.3	3AGCC	0	443.9 (17)	17GCCAA	0
	N253.2	198.4	3AGCC	0	444.0 (17)	17GCCAA	0
	N253.3	198.2	3AGCC	0	443.9	17GCCAA	0
	N253.4	198.4	3AGCC	0	444.0 (16)	17GCCAA	0
	N253.5	198.3	3AGCC	0	433.9 (15)	15GCCAA	0
	N253.6	198.4	3AGCC	0	443.9 (17)	17GCCAA	0
V59/3 rd	N352.1	198.3	3AGCC	0	438.5	16GCCAA	1
	N352.2	198.5	3AGCC	0	438.1	16GCCAA	1

	N352.3	198.3	3AGCC	0	448.1 (18)	18GCCAA	0
	N352.4	198.5	3AGCC	0	447.7	18GCCAA	0
	N352.5	198.3	3AGCC	0	442.8	17GCCAA	0
	N352.6	198.4	3AGCC	0	443.0	17GCCAA	0
V59/4 th	N438.1	198.3	3AGCC	0	462.7	21GCCAA	0
	N438.2	198.5	3AGCC	0	462.4	21GCCAA	0
	N438.3	198.5	3AGCC	0	443.1	17GCCAA	0
	N438.4	198.5	3AGCC	0	447.8	18GCCAA	0
	N438.5	198.3	3AGCC	0	462.6 (16)	21GCCAA	0
	N438.6	198.5	3AGCC	0	457.7	20GCCAA	0
V88/1 st	N88.1.1	198.3	3AGCC	0	525.9	34GCCAA*	1
	N88.1.2	198.4	3AGCC	0	525.5	34GCCAA	1
	N88.1.3	198.3	3AGCC	0	525.8	34GCCAA	1
	N88.1.4	198.5	3AGCC	0	525.5	34GCCAA	1
	N88.1.5	198.3	3AGCC	0	525.7	34GCCAA	1
	N88.1.6	198.5	3AGCC	0	525.6	34GCCAA	1
V88/2 nd	N272.1	198.3	3AGCC	0	525.8	34GCCAA	1
	N272.2	198.5	3AGCC	0	530.4	35GCCAA*	0
	N272.3	198.4	3AGCC	0	525.8	34GCCAA	1
	N272.4	198.5	3AGCC	0	530.3	35GCCAA	0
	N272.5	198.3	3AGCC	0	525.8	34GCCAA	1
	N272.6	198.5	3AGCC	0	525.6	34GCCAA	1
V88/3 rd	N369.1	198.3	3AGCC	0	501.5	29GCCAA	0
	N369.2	198.4	3AGCC	0	501.2	29GCCAA	0
	N369.3	198.3	3AGCC	0	530.4	35GCCAA	0
	N369.4	198.3	3AGCC	0	530.3	35GCCAA	0
	N369.5	198.3	3AGCC	0	501.2	29GCCAA	0
	N369.6	198.5	3AGCC	0	501.2	29GCCAA	0
V88/4 th	N449.1	198.3	3AGCC	0	539.9	37GCCAA	1
	N449.2	198.5	3AGCC	0	539.9	37GCCAA	1
	N449.3	198.4	3AGCC	0	539.9	37GCCAA	1
	N449.4	198.4	3AGCC	0	540>544.9 {1.1}	37GCCAA	1
	N449.5	198.4	3AGCC	0	540.0	37GCCAA	1
	N449.6	198.5	3AGCC	0	540.0	37GCCAA	1
V138/1 st	N138.1	198.3	3AGCC	0	443.4	17GCCAA*	0

	N138.2	198.5	3AGCC	0	443.0	17GCCAA	0
	N138.3	198.3	3AGCC	0	443.1	17GCCAA	0
	N138.4	198.5	3AGCC	0	442.6	17GCCAA	0
	N138.5	198.3	3AGCC	0	442.7	17GCCAA	0
	N138.6	198.5	3AGCC	0	443.0	17GCCAA	0
V138/2 nd	N288.1	198.3	3AGCC	0	438.1	16GCCAA	1
	N288.2	198.5	3AGCC	0	442.9	17GCCAA	0
	N288.3	198.4	3AGCC	0	443.0	17GCCAA	0
	N288.4	198.4	3AGCC	0	443.0	17GCCAA	0
	N288.5	198.2	3AGCC	0	472.3	23GCCAA	0
	N288.6	198.5	3AGCC	0	447.8	18GCCAA	0
V138/3 rd	N331.1	198.4	3AGCC	0	452.8 (16)	19GCCAA	1
	N331.2	198.5	3AGCC	0	452.7	19GCCAA	1
	N331.3	198.3	3AGCC	0	452.9	19GCCAA	1
	N331.4	198.5	3AGCC	0	452.6	19GCCAA	1
	N331.5	198.3	3AGCC	0	452.5	19GCCAA	1
	N331.6	198.5	3AGCC	0	452.6	19GCCAA	1
V138/4 th	Replaced						
cc167							
V124/1 st	N124.1	240.7 (13)	13AGCC	1	Absent		
	N124.2	241.3	13AGCC	1	Absent		
	N124.3	240.8	13AGCC	1	Absent		
	N124.4	241.2	13AGCC	1	Absent		
	N124.5	236.7	12AGCC	0	Absent		
	N124.6	241.1	13AGCC	1	Absent		
V124/2 nd	N290.1	240.7	13AGCC	1	Absent		
	N290.2	241.2	13AGCC	1	Absent		
	N290.3	240.7	13AGCC	1	Absent		
	N290.4	241.1	13AGCC	1	Absent		
	N290.5	240.8	13AGCC	1	Absent		
	N290.6	236.8	13AGCC*	1	Absent		
V124/3 rd	N336.1	240.9 (13)	13AGCC	1	Absent		
	N336.2	241.0	14AGCC	0	Absent		
	N336.3	240.6	13AGCC	1	Absent		
	N336.4	240.6	13AGCC	1	Absent		
	N336.5	240.6	13AGCC	1	Absent		

	N336.6	240.6	13AGCC	1	Absent		
					Absent		
V64/1 st	N64.1	240.5 (13)	13AGCC	1	Absent		
	N64.2	240.7	13AGCC	1	Absent		
	N64.3	240.7	13AGCC	1	Absent		
	N64.4	240.7	13AGCC	1	Absent		
	N64.5	240.6	13AGCC	1	Absent		
	N64.6	240.8	13AGCC	1	Absent		
V64/2 nd	N257.1	240.8	13AGCC	1	Absent		
	N257.2	240.6	13AGCC	1	Absent		
	N257.3	240.6	13AGCC	1	Absent		
	N257.4	244.8	14AGCC	0	Absent		
	N257.5	240.6	13AGCC	1	Absent		
	N257.6	244.8	14AGCC	0	Absent		
V64/3 rd	N348.1	240.5 (13)	13AGCC	1	Absent		
	N348.2	240.6	13AGCC	1	Absent		
	N348.3	244.7	14AGCC	0	Absent		
	N348.4	244.9	14AGCC	0	Absent		
	N348.5	236.5	12AGCC	0	Absent		
	N348.6	240.8	13AGCC	1	Absent		
					Absent		
V117/1 st	N117.1	240.8 (13)	13AGCC	1	Absent		
	N117.2	244.8	14AGCC	0	Absent		
V117/2 nd	N284.1	240.6	13AGCC	1	Absent		
	N284.2	240.6	13AGCC	1	Absent		
	N284.3	244.7	14AGCC	0	Absent		
	N284.4	240.6	13AGCC	1	Absent		
	N284.5	240.6	13AGCC	1	Absent		
	N284.6	240.6	13AGCC	1	Absent		
V117/3 rd	N332.1	240.5	13AGCC	1	Absent		
	N332.2	240.6	13AGCC	1	Absent		
	N332.3	240.6	13AGCC	1	Absent		
	N332.4	240.7	13AGCC	1	Absent		
	N332.5	245.0>240.7 (1)	14AGCC	0	Absent		
	N332.6	244.9	14AGCC	0	Absent		
V117/4 th	N417.1	240.6>236.4 (1) (13)	13AGCC	1	Absent		
	N417.2	240.5>236.3 (1.1)	13AGCC	1	Absent		

	N417.3	244.7>240.5 (1)	14AGCC	0	Absent		
	N417.4	240.6	13AGCC	1	Absent		
	N417.5	244.8	14AGCC	0	Absent		
	N417.6	244.8	14AGCC	0	Absent		
					Absent		
cc23							
V69/1 st	N1-carrier				Absent		
V69/2 nd	N258.1	256.8	17AGCC	0	Absent		
	N258.2	253.1 (16)	16AGCC	1	Absent		
	N258.3	249.8 (16)	16AGCC*	1	Absent		
	N258.4	253.2 (16)	16AGCC	1	Absent		
	N258.5	257.1 (17)	17AGCC	0	Absent		
	N258.6	253.2 (16)	16AGCC	1	Absent		
V69/3 rd	No sample				Absent		
V69/4 th	N431.1	257.1 (17)	17AGCC	0	Absent		
	N431.2	261.0 (18)	18AGCC	0	Absent		
	N431.3	260.9 (18)	18AGCC	0	Absent		
	N431.4	261.9 (19)	19AGCC	1	Absent		
	N431.5	256.9 (17)	17AGCC	0	Absent		
	N431.6	260.8 (18)	18AGCC	1	Absent		
V93/1 st	N1-carrier				Absent		
V93/2 nd	N264.1	249.0 (15)	15AGCC	0	Absent		
	N264.2	249.1	15AGCC	0	Absent		
	N264.3	253.1	16AGCC	1	Absent		
	N264.4	249.2	15AGCC	0	Absent		
	N264.5	249.3	15AGCC	0	Absent		
	N264.6	249.2	15AGCC	0	Absent		
V93/3 rd	N359.1	257.1	17AGCC	0	Absent		
	N359.2	253.9	16AGCC	1	Absent		
	N359.3	257.0	17AGCC	0	Absent		
	N359.4	257.0	17AGCC	0	Absent		
	N359.5	256.9	17AGCC	0	Absent		
	N359.6	256.9	17AGCC	0	Absent		
V96/1 st	N1-carrier				Absent		
V96/2 nd	N259.1	260.9	18AGCC	0	Absent		

	N259.2	260.8	18AGCC	0	Absent		
	N259.3	260.9	18AGCC	0	Absent		
	N259.4	261.0	18AGCC	0	Absent		
	N259.5	261.0	18AGCC	0	Absent		
	N259.6	257.1	17AGCC	0	Absent		
V96/3 rd	N370.1	264.9	19AGCC	1	Absent		
	N370.2	264.9	19AGCC	1	Absent		
	N370.3	260.9	18AGCC	0	Absent		
	N370.4	260.9	18AGCC	0	Absent		
	N370.5	260.9	18AGCC	0	Absent		
	N370.6	264.8	19AGCC	0	Absent		
V96/4 th	N445.1	264.8	19AGCC	0	Absent		
	N445.2	264.8	19AGCC	0	Absent		
	N445.3	264.8	19AGCC	0	Absent		
	N445.4	264.9	19AGCC	0	Absent		
	N445.5	264.9	19AGCC	0	Absent		
	N445.6	265.0	19AGCC	0	Absent		
V222/1 st	N222.1	224.0 (9)	9AGCC*	0	Absent		
	N222.2	215.5	7AGCC	1	Absent		
	N222.3	215.5	7AGCC	1	Absent		
	N222.4	215.4	7AGCC	1	Absent		
	N222.5	219.6	8AGCC	0	Absent		
	N222.6	219.7	8AGCC	0	Absent		
V222/2 nd	N309.1	215.4	7AGCC	1	Absent		
	N309.2	215.4	7AGCC	1	Absent		
	N309.3	215.3	7AGCC	1	Absent		
	N309.4	215.5	7AGCC	1	Absent		
	N309.5	212.3	7AGCC	1	Absent		
	N309.6	212.4	7AGCC	1	Absent		
V222/3 rd	N391.1	216.5	8AGCC	0	Absent		
	N391.2	216.7	8AGCC	0	Absent		
	N391.3	216.4	8AGCC	0	Absent		
	N391.4	216.5	8AGCC	0	Absent		
	N391.5	216.4	8AGCC	0	Absent		
	N391.6	220.6	9AGCC	0	Absent		
V222/4 th	N459.1	216.6 (8)	8AGCC	0	Absent		

	N459.2	216.8	8AGCC	0	Absent		
	N459.3	216.6	8AGCC	0	Absent		
	N459.4	216.6	8AGCC	0	Absent		
	N459.5	216.5	8AGCC	0	Absent		
	N459.6	216.6	8AGCC	0	Absent		
cc60							
V114/1 st	N114.1	257.1 (17)	17AGCC	0	419.1 (12)	12GCCAA	0
	N114.2	257.0	17AGCC	0	419.4	12GCCAA	0
	N114.3	257.1	17AGCC	0	419.2	12GCCAA	0
	N114.4	257.0	17AGCC	0	419.3	12GCCAA	0
	N114.5	257.0	17AGCC	0	419.3	12GCCAA	0
	N114.6	256.9	17AGCC	0	414.4	11GCCAA	0
V114/2 nd	N283.1	249.1	15AGCC	0	428.9	14GCCAA	0
	N283.2	260.8	18AGCC	0	429.1	14GCCAA	0
	N283.3	257.1	17AGCC	0	428.9	14GCCAA	0
	N283.4	249.1	15AGCC	0	428.8	14GCCAA	0
	N283.5	261.0	18AGCC	0	23.8	13GCCAA	1
	N283.6	257.1	17AGCC	0	428.9	14GCCAA	0
V114/3 rd	N330.1	257.0 (17)	17AGCC	0	424.1 (13)	13GCCAA	1
	N330.2	256.9	17AGCC	0	424.2	13GCCAA	1
	N330.3	257.0	17AGCC	0	424.2	13GCCAA	1
	N330.4	256.9	17AGCC	0	424.3	13GCCAA	1
	N330.5	257.0	17AGCC	0	424.1	13GCCAA	1
	N330.6	256.9	17AGCC	0	424.1	13GCCAA	1
V134/1 st	N134.1	257.0 (17)	17AGCC	0	424.0 (13)	13GCCAA	1
	N134.2	256.9	17AGCC	0	419.3	12GCCAA	0
	N134.3	257.1	17AGCC	0	419.2	12GCCAA	0
	N134.4	257.0	17AGCC	0	419.3	12GCCAA	0
	N134.5	257.1	17AGCC	0	418.9	12GCCAA	0
	N134.6	253.1	16AGCC	1	419.1	12GCCAA	0
V134/2 nd	N295.1	257.0	17AGCC	0	419.3	12GCCAA	0
	N295.2	257.0	17AGCC	0	419.2	12GCCAA	0
	N295.3	249.2	15AGCC	0	419.2	12GCCAA	0
	N295.4	256.8	17AGCC	0	419.3	12GCCAA	0
	N295.5	249.2	15AGCC	0	419.3	12GCCAA	0

	N295.6	256.8	17AGCC	0	419.4	12GCCAA	0
V134/3 rd	N333.1	257.0 (17)	17AGCC	0	419.2 (12)	12GCCAA	0
	N333.2	256.9	17AGCC	0	419.3	12GCCAA	0
	N333.3	253.2	16AGCC	1	419.2	12GCCAA	0
	N333.4	256.9	17AGCC	0	414.4	11GCCAA	0
	N333.5	257.2	17AGCC	0	419.2	12GCCAA	0
	N333.6	257.0	17AGCC	0	419.3	12GCCAA	0
V185/1 st	N185.1	253.3 (16)	16AGCC	1	404.4 (9)	9GCCAA	0
	N185.2	253.2 (16)	16AGCC	1	409.4 (10)	10GCCAA	1
	N185.3	253.2	16AGCC	1	404.4	9GCCAA	0
	N185.4	257.0	17AGCC	0	404.5	9GCCAA	0
	N185.5	249.3	15AGCC	0	404.4	9GCCAA	0
	N185.6	253.1	16AGCC	1	404.5	9GCCAA	0
V185/2 nd	N306.1	249.2	15AGCC	0	404.4	9GCCAA	0
	N306.2	244.9	14AGCC	0	404.5	9GCCAA	0
	N306.3	249.3	15AGCC	0	404.4	9GCCAA	0
	N306.4	249.2	15AGCC	0	404.5	9GCCAA	0
	N306.5	249.4	15AGCC	0	404.3	9GCCAA	0
	N306.6	249.3	15AGCC	0	404.4	9GCCAA	0
V185/4 th	N456.1	245.2 (14)	14AGCC	0	404.3 (9)	9GCCAA	0
	N456.2	245.1 (14)	14AGCC	0	404.3 (9)	9GCCAA	0
	N456.3	240.8	13AGCC	1	409.3	10GCCAA	1
	N456.4	257.0	17AGCC	0	404.5	9GCCAA	0
	N456.5	253.2	16AGCC	1	404.5	9GCCAA	0
	N456.6	249.2	15AGCC	0	404.5	9GCCAA	0
V113/1 st	N113.1	257.1	17AGCC	0	419.3	12GCCAA	0
	N113.2	249.1	15AGCC	0	404.5	9GCCAA	0
	N113.3	257.0	17AGCC	0	419.3	12GCCAA	0
	N113.4	256.9	17AGCC	0	419.3	12GCCAA	0
	N113.5	257.2	17AGCC	0	419.2	12GCCAA	0
	N113.6	257.0	17AGCC	0	419.3	12GCCAA	0
V113/2 nd	N281.1	257.1	17AGCC	0	419.2	12GCCAA	0
	N281.2	257.1	17AGCC	0	419.3	12GCCAA	0
	N281.3	257.0	17AGCC	0	419.0	12GCCAA	0
	N281.4	256.9	17AGCC	0	419.2	12GCCAA	0

	N281.5	257.0	17AGCC	0	419.3	12GCCAA	0
	N281.6	256.9	17AGCC	0	419.2	12GCCAA	0
V115/1 st	N115.1	257.0	17AGCC	0	419.2	12GCCAA	0
	N115.2	253.6	17AGCC	0	418.6	12GCCAA*	0
	N115.3	257.1	17AGCC	0	419.2	12GCCAA	0
	N115.4	257.0	17AGCC	0	419.3	12GCCAA	0
	N115.5	257.2	17AGCC	0	419.3	12GCCAA	0
	N115.6	257.1	17AGCC	0	419.3	12GCCAA	0
V115/2 nd	N282.1	257.2	17AGCC	0	414.3	11GCCAA	0
	N282.2	228.1	10AGCC	1	419.2	12GCCAA	0
	N282.3	257.1	17AGCC	0	419.3	12GCCAA	0
	N282.4	257.0	17AGCC	0	414.4	11GCCAA	0
	N282.5	257.1	17AGCC	0	419.2	12GCCAA	0
	N282.6	257.0	17AGCC	0	414.3	11GCCAA	0
cc32							
V176/1 st	N176.1	266.1	20AGCC	0	458.0	19CCCAA	0
	N176.2	266.1	20AGCC	0	458.0	19CCCAA	0
	N176.3	266.2	20AGCC	0	458.0	19CCCAA	0
	N176.4	266.1	20AGCC	0	458.0	19CCCAA	0
	N176.5	202.6	4AGCC	1	457.9	19CCCAA	0
	N176.6	266.3	20AGCC	0	458.0	19CCCAA	0
V176/2 nd	N318.1	266.2	20AGCC	0	457.9	19CCCAA	0
	N318.2	266.2	20AGCC	0	458.0	19CCCAA	0
	N318.3	266.3	20AGCC	0	457.9	19CCCAA	0
	N318.4	266.1	20AGCC	0	458.0	19CCCAA	0
	N318.5	266.1	20AGCC	0	458.0	19CCCAA	0
	N318.6	266.4	20AGCC	0	458.0	19CCCAA	0
V176/3 rd	N399.1	266.4	20AGCC	0	458.1	19CCCAA	0
	N399.2	266.3	20AGCC	0	458.0	19CCCAA	0
	N399.3	266.4	20AGCC	0	457.9	19CCCAA	0
	N399.4	266.4	20AGCC	0	458.0	19CCCAA	0
	N399.5	266.3	20AGCC	0	457.9	19CCCAA	0
	N399.6	266.2	20AGCC	0	462.5	20CCCAA*	0
V176/4 th	N408.1	266.2	20AGCC	0	457.9	19CCCAA	0
	N408.2	266.3	20AGCC	0	457.7	19CCCAA	0

	N408.3	266.0	20AGCC	0	457.9	19CCCAA	0
	N408.4	266.2	20AGCC	0	452.8	18CCCAA	0
	N408.6	266.3	20AGCC	0	457.6	19CCCAA	0

¹Fragments sizes were determined using either a GS600LIZ (black type) or GS500LIZ (blue type) size standard; numbers in curly brackets indicate the ratio between the PCR fragment sizes where two obvious peaks were observed. ²an asterisk indicates that the PCR fragment was sequenced by dideoxy sequencing and the repeat number determined by visual inspection of the trace file. ³Expression states, 1=1, 0=0 (all genes are subject to translational phase variation).

V96/2nd	259.1	472.0	5	6	17	0	620.7	5	6	5	0
	259.2	477.1	5	6	18	1	620.7	5	6	5	0
	259.3	482.1	5	6	19	0	626.6	5	6	6	1
	259.4	477.1	5	6	18*	1	621.1	5	6	5	0
	259.5	477.2	5	6	18	1	621.1	5	6	5	0
	259.6	482.1	5	6	19	0	626.3	5	6	6	1
V96/3rd	370.1	ND	ND	ND	ND	ND	ND	ND	ND	ND	ND
	370.2	738.6	5	6	18	1	621.4	5	6	5	0
	370.3	487.8	5	6	20*	0	626.7	5	6	6	1
	370.4	748.4	5	6	20*	0	621.4	5	6	5	0
	370.5	738.6	5	6	18	1	621.5	5	6	5	0
	370.6	728.5	5	6	16	0	621.5	5	6	5	0
V96/4th	445.1	728.5	5	6	16	0	625.8	5	6	6	1
	445.2	728.6	5	6	16	0	625.6	5	6	6	1
	445.3	728.5	5	6	16*	0	626.0	5	6	6	1
	445.4	728.5	5	6	16	0	626.1	5	6	6	1
	445.5	728.5	5	6	16	0	626.2	5	6	6	1
	445.6	733.4	5	6	17	0	625.8	5	6	6	1
V69/2nd	258.1	482.0	5	6	19	0	641.7	5	6	9	1
	258.2	482.1	5	6	19	0	641.2	5	6	9	1
	258.3	482.1	5	6	19	0	641.4	5	6	9	1
	258.4	482.6	5	6	19	0	641.8	5	6	9	1
	258.5	482.6	5	6	19	0	641.5	5	6	9	1
	258.6	482.0	5	6	19	0	641.6	5	6	9	1
V69/3rd	431.1	472.2	5	6	17	0	652.1	5	6	11	0
	431.2	472.1	5	6	17	0	657.2	5	6	12	1
	431.3	472.4	5	6	17	0	657.1	5	6	12	1
	431.4	472.4	5	6	17	0	656.7	5	6	12	1
	431.5	472.4	5	6	17	0	672.1	5	6	15	1

	431.6	477.4	5	6	18	1	652.1	5	6	11	0
V93/2nd	264.1	472.5	5	6	17	0	641.6	5	6	9	1
	264.2	472.4	5	6	17	0	641.4	5	6	9	1
	264.3	461.9	5	6	15	1	626.8	5	6	6	1
	264.4	472.4	5	6	17	0	641.9	5	6	9	1
	264.5	472.3	5	6	17	0	641.0	5	6	9	1
	264.6	471.8	5	6	17	0	641.0	5	6	9	1
V93/3rd	359.1	482.7	5	6	19	0	642.0	5	6	9	1
	359.2	482.7	5	6	19	0	641.6	5	6	9	1
	359.3	482.4	5	6	19	0	642.1	5	6	9	1
	359.4	482.6	5	6	19	0	641.2	5	6	9	1
	359.5	487.7	5	6	20	0	641.6	5	6	9	1
	359.6	482.3	5	6	19	0	641.0	5	6	9	1
cc167											
V117/1st	117.1	616.5	5	6	4	0	652.8	5	6	12	1
	117.2	616.7	5	6	4*	0	652.8	5	6	12*	1
V117/2nd	284.1	617.0	5	6	4	0	647.9	5	6	11	0
	284.2	617.1	5	6	4	0	658.3	5	6	13	0
	284.3	617.1	5	6	4	0	653.1	5	6	12	1
	284.4	617.0	5	6	4	0	658.1	5	6	13	0
	284.5	616.9	5	6	4	0	652.9	5	6	12	1
	284.6	617.0	5	6	4	0	652.8	5	6	12	1
V117/3rd	332.1	616.4	5	6	4	0	683.2	5	6	18*	1
	332.2	616.6	5	6	4	0	673.2	5	6	16	0
	332.3	616.8	5	6	4	0	673.1	5	6	16	0
	332.4	616.8	5	6	4	0	668.0	5	6	15	1
	332.6	616.5	5	6	4	0	647.0	5	6	11	0
V117/4th	417.1	661.0	5	6	13	0	656.9	5	6	13	0
	417.2	661.8	5	6	13	0	657.6	5	6	13	0

	417.3	652.3	5	6	11	0	653.1	5	6	12	1
	417.4	645.7	5	6	10	0	651.3	5	6	12	1
	417.5	652.2	5	6	11	0	653.1	5	6	12	1
	417.6	652.2	5	6	11	0	653.1	5	6	12	1
V64/1st	64.1	616.6	5	6	4*	0	652.8	5	6	12	1
	64.2	616.7	5	6	4	0	652.7	5	6	12	1
	64.3	616.8	5	6	4	0	652.7	5	6	12	1
	64.4	616.8	5	6	4	0	652.8	5	6	12	1
	64.5	616.4	5	6	4	0	651.7	5	6	12	1
	64.6	616.6	5	6	4	0	652.7	5	6	12	1
V64/2nd	257.1	616.6	5	6	4	0	667.7	5	6	15	1
	257.2	617.0	5	6	4	0	668.3	5	6	15	1
	257.3	617.1	5	6	4	0	643.0	5	6	10	0
	257.4	617.1	5	6	4	0	668.3	5	6	15	1
	257.5	617.0	5	6	4	0	668.3	5	6	15	1
	257.6	616.9	5	6	4	0	668.2	5	6	15	1
V64/3rd	348.1	616.5	5	6	4	0	662.7	5	6	14	0
	348.2	616.7	5	6	4	0	662.8	5	6	14	0
	348.3	616.5	5	6	4	0	662.8	5	6	14	0
	348.4	616.7	5	6	4	0	662.7	5	6	14	0
	348.5	616.5	5	6	4	0	662.7	5	6	14	0
	348.6	616.6	5	6	4	0	662.0	5	6	14	0
V124/1st	124.1	617.0	5	6	4	0	657.1	5	6	12	1
	124.2	617.0	5	6	4*	0	657.3	5	6	12*	1
	124.3	616.3	5	6	4	0	656.0	5	6	12	1
	124.4	616.3	5	6	4	0	655.8	5	6	12	1
	124.5	616.3	5	6	4	0	656.0	5	6	12	1
V124/2nd	290.1	616.4	5	6	4	0	656.8	5	6	12	1

	290.3	616.9	5	6	4	0	657.3	5	6	12	1
	290.4	616.8	5	6	4	0	657.1	5	6	12	1
	290.5	616.5	5	6	4	0	656.8	5	6	12	1
	290.6	616.4	5	6	4	0	656.8	5	6	12	1
V124/3rd	336.1	616.6	5	6	4	0	657.0	5	6	12	1
	336.2	616.7	5	6	4	0	657.1	5	6	12	1
	336.3	617.0	5	6	4	0	657.2	5	6	12	1
	336.4	616.8	5	6	4	0	657.1	5	6	12	1
	336.5	616.7	5	6	4	0	656.9	5	6	12	1
	336.6	616.6	5	6	4	0	656.8	5	6	12	1
cc60											
V113/1st	113.1	458.6	5	6	13*	0	700.5	5	6	11*	0
	113.2	433.2	5	6	8	0	700.6	5	6	11	0
	113.3	458.7	5	6	13	0	700.7	5	6	11	0
	113.4	433.2	5	6	8	0	700.5	5	6	11	0
	113.5	443.3	5	6	10	0	700.7	5	6	11	0
	113.6	443.2	5	6	10	0	700.5	5	6	11	0
V113/2nd	281.1	443.6	5	6	10	0	700.9	5	6	11	0
	282.2	443.6	5	6	10	0	701.0	5	6	11	0
	282.3	443.7	5	6	10	0	701.0	5	6	11	0
	282.4	438.6	5	6	9	1	701.0	5	6	11	0
	282.5	443.6	5	6	10	0	700.9	5	6	11	0
	282.6	443.6	5	6	10	0	700.9	5	6	11	0
V114/1st	114.1	448.4	5	6	11	0	705.8	5	6	12	1
	114.2	438.3	5	6	9	1	690.7	5	6	9	1
	114.3	438.3	5	6	9	1	700.6	5	6	11	0
	114.4	438.3	5	6	9	1	700.8	5	6	11	0
	114.5	448.4	5	6	11	0	705.6	5	6	12	1
	114.6	448.4	5	6	11	0	705.6	5	6	12	1

	114.7	438.4	5	6	9	1	705.8	5	6	12	1
	114.8	443.4	5	6	10	0	706.0	5	6	12	1
	114.9	443.4	5	6	10	0	705.8	5	6	12	1
	114.10	443.4	5	6	10	0	705.8	5	6	12	1
	114.11	438.3	5	6	9	1	700.6	5	6	11	0
	114.12	438.4	5	6	9	1	700.8	5	6	11	0
	114.13	438.3	5	6	9	1	705.6	5	6	12	1
	114.14	438.3	5	6	9	1	700.6	5	6	11	0
	114.16	443.7	5	6	10	0	706.1	5	6	12	1
	114.18	443.8	5	6	10	0	706.1	5	6	12	1
V114/2nd	283.1	448.3	5	6	11	0	705.4	5	6	12	1
	283.2	448.8	5	6	11	0	706.2	5	6	12	1
	283.3	448.8	5	6	11	0	706.1	5	6	12	1
	283.4	448.4	5	6	11	0	705.6	5	6	12	1
	283.5	448.7	5	6	11	0	706.0	5	6	12	1
	283.6	448.8	5	6	11	0	706.0	5	6	12	1
V114/3rd	330.1	438.1	5	6	9	1	700.5	5	6	11	0
	330.2	438.2	5	6	9	1	700.4	5	6	11	0
	330.3	438.6	5	6	9	1	700.8	5	6	11	0
	330.4	438.5	5	6	9	1	700.7	5	6	11	0
	330.5	438.3	5	6	9	1	700.5	5	6	11	0
	330.6	438.1	5	6	9	1	700.5	5	6	11	0
	330.7	438.3	5	6	9	1	700.7	5	6	11	0
	330.8	438.4	5	6	9	1	700.8	5	6	11	0
	330.9	438.4	5	6	9	1	700.6	5	6	11	0
	330,10	438.4	5	6	9	1	695.7	5	6	10	0
	330.11	438.4	5	6	9	1	700.8	5	6	11	0
	330.12	438.3	5	6	9	1	700.7	5	6	11	0
	330.15	438.7	5	6	9	1	701.0	5	6	11	0
	330.16	438.7	5	6	9	1	701.1	5	6	11	0

	330.17	438.7	5	6	9	1	701.1	5	6	11	0
	330.18	438.8	5	6	9	1	701.0	5	6	11	0
	330.19	438.7	5	6	9	1	701.1	5	6	11	0
	330.20	438.7	5	6	9	1	701.1	5	6	11	0
V115/1st	115.1	433.2	5	6	8	0	705.6	5	6	12	1
	115.2	433.2	5	6	8	0	690.6	5	6	9	1
	115.3	443.3	5	6	10	0	705.7	5	6	12	1
	115.4	443.4	5	6	10	0	705.6	5	6	12	1
	115.5	443.3	5	6	10	0	700.6	5	6	11	0
	115.6	443.3	5	6	10	0	700.6	5	6	11	0
V115/2nd	282.1	433.3	5	6	8	0	700.5	5	6	11	0
	282.2	443.7	5	6	10	0	690.9	5	6	9	1
	282.3	433.6	5	6	8	0	701.0	5	6	11	0
	282.4	433.6	5	6	8	0	701.1	5	6	11	0
	282.5	433.6	5	6	8	0	701.0	5	6	11	0
	282.6	433.6	5	6	8	0	701.0	5	6	11	0
V134/1st	134.1	433.1	5	6	8	0	705.5	5	6	12	1
	134.2	433.2	5	6	8	0	700.6	5	6	11	0
	134.3	433.2	5	6	8	0	700.6	5	6	11	0
	134.4	433.2	5	6	8	0	700.6	5	6	11	0
	134.6	433.2	5	6	8	0	720.8	5	6	15	1
	134.7	433.2	5	6	8	0	700.7	5	6	11	0
	134.8	433.3	5	6	8	0	700.7	5	6	11	0
	134.9	433.3	5	6	8	0	700.7	5	6	11	0
	134.10	433.4	5	6	8	0	700.7	5	6	11	0
	134.11	433.3	5	6	8	0	700.7	5	6	11	0
	134.12	433.3	5	6	8	0	700.6	5	6	11	0
	134.13	433.3	5	6	8	0	700.7	5	6	11	0
	134.14	433.1	5	6	8	0	700.5	5	6	11	0

	134.16	433.6	5	6	8	0	701.2	5	6	11	0
	134.17	433.6	5	6	8	0	701.1	5	6	11	0
	134.18	443.8	5	6	10	0	706.1	5	6	12	1
V134/2nd	295.1	448.7	5	6	11	0	705.6	5	6	12	1
	295.2	448.8	5	6	11	0	706.0	5	6	12	1
	295.3	433.3	5	6	8	0	705.6	5	6	12	1
	295.4	448.4	5	6	11	0	705.5	5	6	12	1
	295.5	432.9	5	6	8	0	705.0	5	6	12	1
	295.6	433.6	5	6	8	0	720.6	5	6	15	1
V134/3rd	333.1	433.1	5	6	8	0	705.5	5	6	12	1
	333.2	433.1	5	6	8	0	705.5	5	6	12	1
	333.3	433.6	5	6	8	0	706.0	5	6	12	1
	333.4	433.5	5	6	8	0	706.0	5	6	12	1
	333.5	433.1	5	6	8	0	705.4	5	6	12	1
	333.6	433.0	5	6	8	0	705.4	5	6	12	1
	333.7	433.3	5	6	8	0	705.8	5	6	12	1
	333.8	433.4	5	6	8	0	705.9	5	6	12	1
	333.9	433.5	5	6	8	0	705.9	5	6	12	1
	333,10	433.5	5	6	8	0	705.9	5	6	12	1
	333.11	433.4	5	6	8	0	705.8	5	6	12	1
	333.12	433.4	5	6	8	0	705.8	5	6	12	1
	333.13	433.4	5	6	8	0	705.7	5	6	12	1
	333.14	433.2	5	6	8	0	705.6	5	6	12	1
	333.15	433.6	5	6	8	0	706.2	5	6	12	1
	333.16	433.7	5	6	8	0	706.2	5	6	12	1
	333.17	433.7	5	6	8	0	706.1	5	6	12	1
	333.18	433.7	5	6	8	0	706.1	5	6	12	1
	333.19	433.7	5	6	8	0	706.2	5	6	12	1
	333,20	433.7	5	6	8	0	706.2	5	6	12	1
V185/1st	185.1	463.6	5	6	13	0	438.2	5	6	8	0

	185.2	463.7	5	6	13	0	438.2	5	6	8	0
	185.3	463.6	5	6	13*	0	438.2	5	6	8	0
	185.4	463.7	5	6	13	0	438.2	5	6	8	0
	185.5	463.6	5	6	13	0	438.2	5	6	8*	0
	185.6	464.1	5	6	13	0	438.7	5	6	8	0
	185.8	463.7	5	6	13	0	438.2	5	6	8	0
	185.9	463.7	5	6	13	0	438.2	5	6	8	0
	185,10	463.8	5	6	13	0	438.4	5	6	8	0
	185.11	463.7	5	6	13	0	438.3	5	6	8	0
	185.12	463.8	5	6	13	0	438.3	5	6	8	0
	185.13	463.7	5	6	13	0	438.3	5	6	8	0
	185.14	463.7	5	6	13	0	438.2	5	6	8	0
	185.15	463.6	5	6	13	0	438.1	5	6	8	0
	185.16	464.0	5	6	13	0	438.6	5	6	8	0
	185.17	464.1	5	6	13	0	438.7	5	6	8	0
	185.18	464.1	5	6	13	0	438.7	5	6	8	0
	185.19	464.2	5	6	13	0	438.7	5	6	8	0
	185,20	464.1	5	6	13	0	438.7	5	6	8	0
V185/2nd	306.1	463.5	5	6	13	0	438.1	5	6	8	0
	306.2	473.8	5	6	15	1	438.2	5	6	8	0
	306.3	463.7	5	6	13	0	438.2	5	6	8	0
	306.4	463.7	5	6	13	0	438.2	5	6	8	0
	306.5	463.6	5	6	13	0	438.1	5	6	8	0
	306.6	463.6	5	6	13	0	438.2	5	6	8	0
V185/4th	456.1	453.6	5	6	11	0	438.4	5	6	8	0
	456.2	453.7	5	6	11	0	438.5	5	6	8	0
	456.3	458.8	5	6	12	1	438.5	5	6	8	0
	456.4	458.9	5	6	12	1	438.5	5	6	8	0
	456.5	453.7	5	6	11	0	438.5	5	6	8	0
	456.6	453.7	5	6	11	0	438.4	5	6	8	0
	456.7	453.9	5	6	11	0	438.3	5	6	8	0

	456.8	458.4	5	6	12	1	438.1	5	6	8	0
	456.9	458.5	5	6	12	1	438.2	5	6	8	0
	456.10	458.5	5	6	12	1	438.2	5	6	8	0
	456.11	453.5	5	6	11	0	438.2	5	6	8	0
	456.12	458.5	5	6	12	1	438.2	5	6	8	0
	456.13	453.5	5	6	11	0	438.2	5	6	8	0
	456.14	458.4	5	6	12	1	438.1	5	6	8	0
	456.16	458.9	5	6	12	1	438.5	5	6	8	0
	456.17	458.9	5	6	12	1	438.6	5	6	8	0
	456.18	453.9	5	6	11	0	438.6	5	6	8	0
	456.19	458.9	5	6	12	1	438.6	5	6	8	0
	456.20	458.9	5	6	12	1	438.6	5	6	8	0
cc174											
V51/1st	51.1	450.7	5	6	12*	1	655.7	5	6	10*	0
	51.2	450.7	5	6	12	1	655.7	5	6	10	0
	51.3	450.8	5	6	12	1	655.9	5	6	10	0
	51.4	450.8	5	6	12	1	655.7	5	6	10	0
	51.5	450.7	5	6	12	1	655.8	5	6	10	0
	51.6	450.8	5	6	12	1	655.9	5	6	10	0
V51/2nd	236.1	450.6	5	6	12	1	655.7	5	6	10	0
	236.2	460.9	5	6	14	0	650.5	5	6	9	1
	236.3	445.6	5	6	11	0	665.9	5	6	12	1
	236.4	455.8	5	6	13	0	655.8	5	6	10	0
	236.5	445.6	5	6	11	0	665.9	5	6	12	1
	236.6	445.7	5	6	11	0	665.9	5	6	12	1
V51/3rd	354.1	465.7	5	6	15	1	655.5	5	6	10	0
	354.2	465.8	5	6	15	1	655.4	5	6	10	0
	354.3	465.7	5	6	15	1	655.3	5	6	10	0
	254.4	465.7	5	6	15	1	655.6	5	6	10	0
	254.5	445.3	5	6	11	0	665.5	5	6	12	1

	254.6	465.7	5	6	15	1	655.5	5	6	10	0
V51/4th	424.1	445.6	5	6	11	0	665.8	5	6	12	1
	424.2	445.5	5	6	11	0	665.4	5	6	12	1
	424.3	445.6	5	6	11*	0	665.6	5	6	12*	1
	424.4	445.5	5	6	11	0	665.6	5	6	12	1
	424.5	445.5	5	6	11	0	665.7	5	6	12	1
	424.6	445.5	5	6	11	0	665.6	5	6	12	1
V52/1st	52.1	445.8	5	6	11	0	656.1	5	6	10	0
	52.2	445.8	5	6	11	0	651.0	5	6	9	1
	52.3	445.8	5	6	11	0	656.0	5	6	10	0
	52.4	445.8	5	6	11	0	656.0	5	6	10	0
	52.5	445.7	5	6	11	0	655.8	5	6	10	0
	52.6	445.7	5	6	11	0	655.9	5	6	10	0
V52/2nd	238.1	460.5	5	6	14	0	665.0	5	6	12	1
	238.2	445.8	5	6	11	0	666.1	5	6	12	1
	238.3	445.8	5	6	11	0	650.9	5	6	9	1
	238.4	445.8	5	6	11	0	651.0	5	6	9	1
	238.5	460.4	5	6	14	0	665.2	5	6	12	1
	238.6	445.8	5	6	11	0	650.9	5	6	9	1
V52/3rd	342.1	445.7	5	6	11	0	655.8	5	6	10	0
V58/1st	58.1	445.8	5	6	11	0	650.9	5	6	9	1
	58.2	445.8	5	6	11	0	650.8	5	6	9	1
	58.3	455.9	5	6	13	0	655.9	5	6	10	0
	58.4	445.8	5	6	11	0	650.8	5	6	9	1
	58.5	445.8	5	6	11	0	650.9	5	6	9	1
	58.6	456.0	5	6	13	0	656.1	5	6	10	0
V58/2nd	240.1	445.1	5	6	11	0	644.7	5	6	8	0
V58/4th	429.1	445.8	5	6	11	0	645.9	5	6	8	0
	492.2	445.8	5	6	11	0	645.9	5	6	8	0

	429.3	445.8	5	6	11	0	645.8	5	6	8	0
	429.4	445.8	5	6	11	0	645.6	5	6	8	0
	429.5	445.7	5	6	11	0	645.6	5	6	8	0
	429.6	445.8	5	6	11	0	645.7	5	6	8	0
V59/1st	59.1	450.5	5	6	12	1	655.4	5	6	10	0
	59.2	450.6	5	6	12	1	655.5	5	6	10	0
	59.3	450.6	5	6	12	1	655.5	5	6	10	0
	59.4	450.6	5	6	12	1	660.6	5	6	11	0
	59.5	450.0	5	6	12	1	655.5	5	6	10	0
	59.6	450.5	5	6	12	1	655.4	5	6	10	0
	59.11	451.1	5	6	12	1	655.8	5	6	10	0
	59.12	451.2	5	6	12	1	656.1	5	6	10	0
	59.13	451.1	5	6	12	1	656.1	5	6	10	0
	59.14	451.1	5	6	12	1	656.0	5	6	10	0
	59.15	451.1	5	6	12	1	656.0	5	6	10	0
	59.16	451.1	5	6	12	1	655.9	5	6	10	0
	59.17	451.1	5	6	12	1	655.9	5	6	10	0
	59.18	451.0	5	6	12	1	655.9	5	6	10	0
	59.20	451.4	5	6	12	1	656.5	5	6	10	0
V59/2nd	253.1	445.4	5	6	11	0	655.3	5	6	10	0
	253.2	445.4	5	6	11	0	655.4	5	6	10	0
	253.3	445.4	5	6	11	0	655.4	5	6	10	0
	253.4	445.4	5	6	11	0	655.4	5	6	10	0
	253.5	445.5	5	6	11	0	660.6	5	6	11	0
	253.6	445.4	5	6	11	0	655.4	5	6	10	0
V59/3rd	352.1	445.3	5	6	11	0	655.1	5	6	10	0
	352.2	445.3	5	6	11	0	655.1	5	6	10	0
	352.3	450.4	5	6	12	1	655.2	5	6	10	0
	352.4	445.3	5	6	11	0	655.1	5	6	10	0
	352.5	450.3	5	6	12	1	655.1	5	6	10	0

	352.6	450.4	5	6	12	1	655.1	5	6	10	0
	352.7	446.0	5	6	11	0	656.0	5	6	10	0
	352.8	451.1	5	6	12	1	656.1	5	6	10	0
	352.9	446.0	5	6	11	0	655.9	5	6	10	0
	352.10	446.0	5	6	11	0	661.2	5	6	11	0
	352.11	451.1	5	6	12	1	655.9	5	6	10	0
	352.12	446.0	5	6	11	0	656.0	5	6	10	0
	352.13	445.9	5	6	11	0	655.9	5	6	10	0
	352.14	440.8	5	6	10	0	655.8	5	6	10	0
	352.15	451.3	5	6	12	1	656.3	5	6	10	0
	352.16	456.5	5	6	13	0	656.3	5	6	10	0
	352.17	446.4	5	6	11	0	656.4	5	6	10	0
	352.18	446.4	5	6	11	0	656.5	5	6	10	0
	352.19	451.4	5	6	12	1	656.4	5	6	10	0
	352.20	451.4	5	6	12	1	656.4	5	6	10	0
V59/4th	438.1	445.3	5	6	11	0	655.3	5	6	10	0
	438.2	445.3	5	6	11	0	655.3	5	6	10	0
	438.3	450.4	5	6	12	1	655.2	5	6	10	0
	438.4	450.4	5	6	12	1	655.1	5	6	10	0
	438.5	445.2	5	6	11	0	655.1	5	6	10	0
	438.6	450.7	5	6	12	1	655.5	5	6	10	0
V138/1st	138.1	455.6	5	6	13	0	654.7	5	6	10	0
	138.2	460.6	5	6	14	0	653.7	5	6	10	0
	138.4	460.6	5	6	14	0	655.1	5	6	10	0
	138.5	445.3	5	6	11	0	654.1	5	6	10	0
	138.6	440.7	5	6	10	0	656.0	5	6	10	0
V138/2nd	288.1	445.7	5	6	11	0	666.1	5	6	12	1
	288.2	445.8	5	6	11	0	666.1	5	6	12	1
	288.3	445.7	5	6	11	0	666.0	5	6	12	1
	288.4	445.7	5	6	11	0	666.0	5	6	12	1

	288.5	445.6	5	6	11	0	665.9	5	6	12	1
	288.6	440.6	5	6	10	0	666.0	5	6	12	1
V88/1st	N88.1.1	672.6	5	6	9*	1	677.7	5	6	10	0
	N88.1.2	672.6	5	6	9	1	677.6	5	6	10	0
	N88.1.3	672.6	5	6	9	1	677.7	5	6	10	0
	N88.1.4	672.6	5	6	9	1	677.7	5	6	10	0
	N88.1.5	672.5	5	6	9	1	677.6	5	6	10	0
	N88.1.6	672.6	5	6	9	1	677.7	5	6	10*	0
	88.1.7	673.6	5	6	9	1	678.5	5	6	10	0
	88.1.8	673.6	5	6	9	1	678.6	5	6	10	0
	88.1.9	673.7	5	6	9	1	678.8	5	6	10	0
	88.1.10	673.5	5	6	9	1	678.5	5	6	10	0
	88.1	673.2	5	6	9	1	678.3	5	6	10	0
	88.2	673.2	5	6	9	1	678.4	5	6	10	0
	88.3	673.2	5	6	9	1	678.3	5	6	10	0
	88.4	673.2	5	6	9	1	678.3	5	6	10	0
	88.5	673.1	5	6	9	1	678.3	5	6	10	0
	88.6	673.1	5	6	9	1	678.2	5	6	10	0
	88.8	673.2	5	6	9	1	678.3	5	6	10	0
V88/2nd	N272.1	678.0	5	6	10	0	678.0	5	6	10	0
	N272.2	678.0	5	6	10	0	678.0	5	6	10	0
	N272.3	678.0	5	6	10	0	678.0	5	6	10	0
	N272.4	678.0	5	6	10	0	678.0	5	6	10	0
	N272.5	678.0	5	6	10	0	678.0	5	6	10	0
	N272.6	676.7	5	6	10	0	676.7	5	6	10	0
V88/3rd	N369.1	677.6	5	6	10	0	677.6	5	6	10	0
	N369.2	677.5	5	6	10	0	677.5	5	6	10	0
	N369.3	677.6	5	6	10	0	677.6	5	6	10	0
	N369.3	677.7	5	6	10	0	677.7	5	6	10	0
	N369.5	677.5	5	6	10	0	677.5	5	6	10	0

	N369.6	677.4	5	6	10	0	677.4	5	6	10	0
	369.7	678.4	5	6	10	0	678.4	5	6	10	0
	369.8	678.4	5	6	10	0	678.4	5	6	10	0
	369.9	678.6	5	6	10	0	678.6	5	6	10	0
	369.10	678.5	5	6	10	0	678.5	5	6	10	0
	369.11	678.3	5	6	10	0	678.3	5	6	10	0
	369.12	678.4	5	6	10	0	678.4	5	6	10	0
	369.14	678.3	5	6	10	0	678.3	5	6	10	0
	369.15	678.3	5	6	10	0	678.3	5	6	10	0
	369.16	678.7	5	6	10	0	678.7	5	6	10	0
	369.17	673.7	5	6	9	1	668.7	5	6	8	0
	369.18	678.7	5	6	10	0	678.7	5	6	10	0
	369.19	678.8	5	6	10	0	678.8	5	6	10	0
	369.20	678.6	5	6	10	0	678.6	5	6	10	0
V88/4th	N449.1	682.9	5	6	11*	0	687.7	5	6	12*	1
	449.2	677.8	5	6	10	0	682.9	5	6	11	0
	449.3	683.1	5	6	11	0	688.0	5	6	12	1
	449.4	683.1	5	6	11	0	688.0	5	6	12	1
	449.5	678.0	5	6	10	0	688.1	5	6	12	1
	449.6	677.9	5	6	10	0	682.9	5	6	11	0
V43/2nd	241.1	450.9	5	6	12	1	655.9	5	6	10	0
	241.2	456.0	5	6	13	0	655.9	5	6	10	0
V43/3rd	349.1	450.8	5	6	12	1	655.8	5	6	10	0
	349.2	450.9	5	6	12	1	655.9	5	6	10	0
	349.3	450.9	5	6	12	1	656.1	5	6	10	0
	349.4	450.9	5	6	12	1	656.0	5	6	10	0
	349.5	451.0	5	6	12	1	656.0	5	6	10	0
	349.6	450.9	5	6	12	1	656.1	5	6	10	0

¹Fragments sizes were determined using either a GS500LIZ size standard. ²Opa genes contain short C and A tracts adjacent to the 5'CTCTT repeat tract, these short tracts are invariant within a given strains as detected in all Sanger sequences of PCR products spanning the Opa repeats and in whole genome sequence data. ³Asterisks indicate repeat numbers

confirmed by dideoxy Sanger sequencing of PCR products. ⁴Expression states, 1=1, 0=0 (all genes are subject to translational phase variation).

Supplementary Data Table V. Repeat numbers and expression states for the phase-variable *opaD* and *opaJ* genes

Carrier/ Time pt	Isolate	<i>opaD</i>					<i>opaJ</i> (5'CTCTT)				
		Size ¹	C tract ²	A tract ²	Repeat No. ³	Exp. States ⁴	Size ¹	C tract ²	A tract ²	Repeat No. ³	Exp. States ⁴
cc23											
V222/1 st	222.1	426.7	5	6	8*	0	627.0	5	6	8*	0
	222.2	426.7	5	6	8	0	627.0	5	6	8	0
	222.3	426.1	5	6	8	0	626.4	5	6	8	0
	222.4	426.1	5	6	8	0	626.6	5	6	8	0
	222.5	426.1	5	6	8	0	625.5	5	6	8	0
	222.6	426.2	5	6	8	0	626.5	5	6	8	0
V222/2 nd	309.1	426.5	5	6	8	0	627.0	5	6	8	0
	309.2	426.7	5	6	8	0	627.1	5	6	8	0
	309.3	426.6	5	6	8	0	627.1	5	6	8	0
	309.4	426.7	5	6	8	0	632.2	5	6	9	1
	309.5	426.1	5	6	8	0	626.5	5	6	8	0
	309.6	426.6	5	6	8	0	627.1	5	6	8	0
V222/3 rd	391.1	426.5	5	6	8	0	639.5	3	6	8	1
	391.2	426.5	5	6	8	0	639.2	3	6	8	1
	391.3	426.8	5	6	8	0	639.5	3	6	8	1
	391.4	426.6	5	6	8	0	639.6	3	6	8	1
	391.5	426.5	5	6	8	0	639.4	3	6	8	1
	391.6	426.6	5	6	8	0	640.2	3	6	8	1
V222/4 th	459.1	426.6	5	6	8	0	639.4	3	6	8	1
	459.2	426.5	5	6	8	0	639.4	3	6	8	1
	459.4	426.1	5	6	8	0	640.0	3	6	8	1
	459.4	426.1	5	6	8	0	639.6	3	6	8	1
	459.5	426.1	5	6	8*	0	639.5	3	6	8*	1
	459.6	426.2	5	6	8	0	640.3	3	6	8	1
V96/2 nd	259.1	426.4	5	6	8	0	626.4	5	6	8	0
	259.2	426.3	5	6	8	0	626.4	5	6	8	0

	259.3	426.4	5	6	8	0	626.7	5	6	8	0
	259.4	426.4	5	6	8	0	626.8	5	6	8	0
	259.5	426.4	5	6	8	0	626.8	5	6	8	0
	259.6	426.3	5	6	8	0	627.1	5	6	8	0
V96/3rd	370.2	426.8	5	6	8	0	647.4	5	6	12	1
	370.3	421.9	5	6	7	0	627.5	5	6	8	0
	370.4	426.8	5	6	8	0	647.4	5	6	12	1
	370.5	426.8	5	6	8	0	622.4	5	6	7	0
	370.6	421.7	5	6	7	0	647.5	5	6	12	1
V96/4th	445.1	426.2	5	6	8	0	626.8	5	6	8	0
	445.2	426.2	5	6	8	0	626.3	5	6	8	0
	445.3	426.3	5	6	8	0	626.9	5	6	8	0
	445.4	426.3	5	6	8	0	626.8	5	6	8	0
	445.5	426.2	5	6	8	0	627.1	5	6	8	0
	445.6	426.3	5	6	8	0	626.7	5	6	8	0
V69/2nd	258.1	426.3	5	6	8	0	626.6	5	6	8	0
	258.2	426.4	5	6	8	0	626.9	5	6	8	0
	258.3	426.3	5	6	8	0	626.7	5	6	8	0
	258.4	426.8	5	6	8	0	627.3	5	6	8	0
	258.5	426.7	5	6	8	0	627.2	5	6	8	0
	258.6	426.2	5	6	8	0	626.6	5	6	8	0
V69/4th	431.1	431.6	5	6	9	1	626.8	5	6	8	0
	431.2	426.5	5	6	8	0	626.9	5	6	8	0
	431.3	426.6	5	6	8	0	626.8	5	6	8	0
	431.4	426.8	5	6	8	0	627.1	5	6	8	0
	431.5	426.8	5	6	8	0	627.0	5	6	8	0
	431.6	426.8	5	6	8	0	627.1	5	6	8	0
V93/2nd	264.1	421.7	5	6	7	0	622.0	5	6	7	0
	264.2	421.7	5	6	7	0	622.1	5	6	7	0

	264.3	421.3	5	6	7	0	621.7	5	6	7	0
	264.4	421.6	5	6	7	0	622.0	5	6	7	0
	264.5	421.6	5	6	7	0	621.7	5	6	7	0
	264.6	421.1	5	6	7	0	621.5	5	6	7	0
V93/3rd	359.1	421.9	5	6	7	0	622.2	5	6	7	0
	359.2	421.8	5	6	7	0	622.3	5	6	7	0
	359.3	421.5	5	6	7	0	621.9	5	6	7	0
	359.4	421.7	5	6	7	0	622.3	5	6	7	0
	359.5	421.8	5	6	7	0	622.2	5	6	7	0
	359.6	421.4	5	6	7	0	621.7	5	6	7	0
cc167											
V117/1st	117.1	420.6	5	6	4	0	616.5	5	6	4	0
	117.2	420.5	5	6	4*	0	616.7	5	6	4*	0
V117/2 nd	284.1	421.0	5	6	4	0	617.0	5	6	4	0
	284.2	421.1	5	6	4	0	612.0	5	6	3	1
	284.3	421.1	5	6	4	0	617.1	5	6	4	0
	284.4	421.0	5	6	4	0	611.8	5	6	3	1
	284.5	421.0	5	6	4	0	616.9	5	6	4	0
	284.6	421.0	5	6	4	0	611.8	5	6	3	1
V117/3rd	332.1	420.6	5	6	4*	0	611.5	5	6	3	1
	332.2	420.7	5	6	4	0	611.7	5	6	3	1
	332.3	420.7	5	6	4	0	611.7	5	6	3	1
	332.4	420.7	5	6	4	0	611.6	5	6	3	1
	332.6	420.6	5	6	4	0	611.6	5	6	3	1
V117/4th	417.1	420.4	5	6	4	0	616.4	5	6	4	0
	417.2	421.1	5	6	4	0	617.1	5	6	4	0
	417.3	421.1	5	6	4	0	617.2	5	6	4	0
	417.4	420.4	5	6	4	0	616.2	5	6	4	0
	417.5	421.2	5	6	4	0	617.0	5	6	4	0
	417.6	421.1	5	6	4	0	617.1	5	6	4	0

V64/1st	64.1	420.6	5	6	4	0	657.1	5	6	12*	1
	64.2	420.6	5	6	4	0	657.1	5	6	12	1
	64.3	420.7	5	6	4	0	657.0	5	6	12	1
	64.4	420.5	5	6	4	0	657.2	5	6	12	1
	64.5	420.5	5	6	4	0	656.9	5	6	12	1
	64.6	420.6	5	6	4	0	657.0	5	6	12	1
V64/2nd	257.1	420.5	5	6	4	0	651.8	5	6	11	0
	257.2	421.2	5	6	4	0	652.3	5	6	11	0
	257.3	421.2	5	6	4	0	652.2	5	6	11	0
	257.4	421.1	5	6	4	0	652.4	5	6	11	0
	257.5	421.1	5	6	4	0	652.2	5	6	11	0
	257.6	421.2	5	6	4	0	652.2	5	6	11	0
V64/3rd	348.1	420.5	5	6	4	0	641.7	5	6	9	1
	348.2	420.6	5	6	4	0	636.8	5	6	8	0
	348.3	420.5	5	6	4	0	636.8	5	6	8	0
	348.4	420.6	5	6	4	0	636.7	5	6	8	0
	348.5	420.5	5	6	4	0	641.6	5	6	9	1
	348.6	420.6	5	6	4	0	641.4	5	6	9	1
V124/1st	124.1	421.0	5	6	4	0	617.0	5	6	4	0
	124.2	421.1	5	6	4	0	617.0	5	6	4*	0
	124.3	420.5	5	6	4	0	616.3	5	6	4	0
	124.4	420.5	5	6	4	0	616.3	5	6	4	0
	124.5	420.5	5	6	4	0	616.3	5	6	4	0
V124/2 nd	290.1	420.5	5	6	4	0	616.4	5	6	4	0
	290.3	421.0	5	6	4	0	616.9	5	6	4	0
	290.4	420.9	5	6	4	0	616.8	5	6	4	0
	290.5	420.5	5	6	4	0	616.5	5	6	4	0
	290.6	420.4	5	6	4	0	616.4	5	6	4	0
V124/3rd	336.1	420.5	5	6	4	0	616.6	5	6	4	0

	336.2	420.7	5	6	4	0	616.7	5	6	4	0
	336.3	421.0	5	6	4	0	617.0	5	6	4	0
	336.4	420.8	5	6	4	0	616.8	5	6	4	0
	336.5	420.6	5	6	4	0	616.7	5	6	4	0
	336.6	420.5	5	6	4	0	616.6	5	6	4	0
cc60											
V113/1st	113.1	464.3	4	6	14*	1	413.4	4	6	4*	0
	113.2	464.4	4	6	14	1	413.5	4	6	4	0
	113.3	464.5	4	6	14	1	413.5	4	6	4	0
	113.4	464.3	4	6	14	1	413.4	4	6	4	0
	113.5	464.4	4	6	14	1	413.4	4	6	4	0
	113.6	464.3	4	6	14	1	413.5	4	6	4	0
V113/2 nd	281.1	464.7	4	6	14	1	413.8	4	6	4	0
	281.2	464.7	4	6	14	1	413.9	4	6	4	0
	281.3	464.8	4	6	14	1	413.9	4	6	4	0
	281.4	469.8	4	6	15	0	413.8	4	6	4	0
	281.5	434.3	4	6	8	1	413.8	4	6	4	0
	281.6	434.3	4	6	8	1	413.9	4	6	4	0
V114/1st	114.1	464.4	4	6	14	1	413.5	4	6	4	0
	114.2	469.5	4	6	15	0	413.5	4	6	4	0
	114.3	464.5	4	6	14	1	413.6	4	6	4	0
	114.4	459.4	4	6	13	0	413.5	4	6	4	0
	114.5	464.4	4	6	14	1	413.6	4	6	4	0
	114.6	464.5	4	6	14	1	413.5	4	6	4	0
	114.7	464.4	4	6	14	1	413.7	4	6	4	0
	114.8	464.5	4	6	14	1	413.5	4	6	4	0
	114.9	464.5	4	6	14	1	413.5	4	6	4	0
	114,10	464.5	4	6	14	1	413.6	4	6	4	0
	114.11	464.5	4	6	14	1	413.6	4	6	4	0

	114.12	464.6	4	6	14	1	413.6	4	6	4	0
	114.13	474.6	4	6	16	0	413.5	4	6	4	0
	114.14	464.4	4	6	14	1	413.5	4	6	4	0
	114.16	464.8	4	6	14	1	414.4	4	6	4	0
	114.18	464.9	4	6	14	1	414.3	4	6	4	0
V114/2 nd	283.1	459.3	4	6	13	0	413.4	4	6	4	0
	283.2	469.9	4	6	15	0	414.1	4	6	4	0
	283.3	459.7	4	6	13	0	414.0	4	6	4	0
	283.4	459.3	4	6	13	0	413.1	4	6	4	0
	283.5	459.6	4	6	13	0	413.9	4	6	4	0
	283.6	459.7	4	6	13	0	414.0	4	6	4	0
V114/3 rd	330.1	469.3	4	6	15	0	413.5	4	6	4	0
	330.2	469.3	4	6	15	0	413.3	4	6	4	0
	330.3	469.7	4	6	15	0	413.8	4	6	4	0
	330.4	469.6	4	6	15	0	413.6	4	6	4	0
	330.5	469.4	4	6	15	0	413.5	4	6	4	0
	330.6	469.4	4	6	15	0	413.3	4	6	4	0
	330.7	469.6	4	6	15	0	413.5	4	6	4	0
	330.8	469.7	4	6	15	0	413.6	4	6	4	0
	330.9	469.6	4	6	15	0	413.5	4	6	4	0
	330,10	469.6	4	6	15	0	413.7	4	6	4	0
	330.11	469.5	4	6	15	0	413.6	4	6	4	0
	330.12	469.5	4	6	15	0	413.6	4	6	4	0
	330.15	469.9	4	6	15	0	414.1	4	6	4	0
	330.16	470.0	4	6	15	0	414.2	4	6	4	0
	330.17	470.0	4	6	15	0	414.2	4	6	4	0
	330.18	469.9	4	6	15	0	414.2	4	6	4	0
	330.19	470.0	4	6	15	0	414.1	4	6	4	0
	330.20	470.0	4	6	15	0	414.1	4	6	4	0
V115/1 st	115.1	464.4	4	6	14	1	413.5	4	6	4	0

	115.2	433.9	4	6	8	1	413.5	4	6	4	0
	115.3	464.5	4	6	14	1	413.5	4	6	4	0
	115.4	464.4	4	6	14	1	413.6	4	6	4	0
	115.5	464.3	4	6	14	1	413.6	4	6	4	0
	115.6	464.3	4	6	14	1	413.5	4	6	4	0
V115/2 nd	282.1	464.4	4	6	14	1	413.4	4	6	4	0
	282.2	469.9	4	6	15	0	414.1	4	6	4	0
	282.3	464.8	4	6	14	1	414.0	4	6	4	0
	282.4	464.8	4	6	14	1	414.1	4	6	4	0
	282.5	464.8	4	6	14	1	414.0	4	6	4	0
	282.6	464.9	4	6	14	1	414.0	4	6	4	0
V134/1 st	134.1	464.3	4	6	14	1	413.4	4	6	4	0
	134.2	459.3	4	6	13	0	413.5	4	6	4	0
	134.3	459.2	4	6	13	0	413.4	4	6	4	0
	134.4	459.3	4	6	13	0	413.5	4	6	4	0
	134.6	459.2	4	6	13	0	413.5	4	6	4	0
	134.7	459.4	4	6	13	0	413.6	4	6	4	0
	134.8	459.4	4	6	13	0	413.5	4	6	4	0
	134.9	459.4	4	6	13	0	413.6	4	6	4	0
	134,10	459.4	4	6	13	0	413.6	4	6	4	0
	134.11	459.4	4	6	13	0	413.5	4	6	4	0
	134.12	459.3	4	6	13	0	413.5	4	6	4	0
	134.13	459.3	4	6	13	0	413.3	4	6	4	0
	134.14	459.3	4	6	13	0	413.5	4	6	4	0
	134.16	459.7	4	6	13	0	414.2	4	6	4	0
	134.17	459.8	4	6	13	0	414.2	4	6	4	0
	134.18	464.9	4	6	14	1	414.3	4	6	4	0
V134/2 nd	295.1	464.3	4	6	14	1	413.5	4	6	4	0
	295.2	464.8	4	6	14	1	414.0	4	6	4	0
	295.3	459.3	4	6	13	0	413.5	4	6	4	0

	295.4	464.4	4	6	14	1	413.5	4	6	4	0
	295.5	459.2	4	6	13	0	413.6	4	6	4	0
	295.6	464.7	4	6	14	1	413.8	4	6	4	0
V134/3rd	333.1	459.3	4	6	13	0	413.4	4	6	4	0
	333.2	459.3	4	6	13	0	413.3	4	6	4	0
	333.3	459.7	4	6	13	0	413.8	4	6	4	0
	333.4	459.5	4	6	13	0	413.7	4	6	4	0
	333.5	459.3	4	6	13	0	413.4	4	6	4	0
	333.6	459.2	4	6	13	0	413.4	4	6	4	0
	333.7	459.4	4	6	13	0	413.7	4	6	4	0
	333.8	459.4	4	6	13	0	413.6	4	6	4	0
	333.9	459.5	4	6	13	0	413.7	4	6	4	0
	333.10	459.6	4	6	13	0	413.7	4	6	4	0
	333.11	459.6	4	6	13	0	413.6	4	6	4	0
	333.12	459.4	4	6	13	0	413.7	4	6	4	0
	333.13	464.5	4	6	14	0	413.6	4	6	4	0
	333.14	459.4	4	6	13	0	413.5	4	6	4	0
	333.15	459.7	4	6	13	0	414.1	4	6	4	0
	333.16	459.8	4	6	13	0	414.2	4	6	4	0
	333.17	459.9	4	6	13	0	414.2	4	6	4	0
	333.18	459.9	4	6	13	0	414.2	4	6	4	0
	333.19	459.8	4	6	13	0	414.1	4	6	4	0
	333.20	459.7	4	6	13	0	414.1	4	6	4	0
V185/1st	185.1	438.8	4	6	9	0	428.8	4	6	7	0
	185.2	438.9	4	6	9	0	429.0	4	6	7	0
	185.3	438.9	4	6	9*	0	428.9	4	6	7	0
	185.4	439.0	4	6	9	0	429.0	4	6	7	0
	185.5	439.0	4	6	9	0	428.9	4	6	7	0
	185.6	438.9	4	6	9	0	428.9	4	6	7	0
	185.8	438.9	4	6	9	0	428.9	4	6	7	0

	185.9	439.2	4	6	9	0	429.1	4	6	7	0
	185.10	439.1	4	6	9	0	428.9	4	6	7	0
	185.11	439.0	4	6	9	0	428.8	4	6	7	0
	185.12	439.0	4	6	9	0	429.0	4	6	7	0
	185.13	438.9	4	6	9	0	428.9	4	6	7	0
	185.14	438.9	4	6	9	0	429.0	4	6	7	0
	185.15	438.8	4	6	9	0	428.6	4	6	7	0
	185.16	439.3	4	6	9	0	429.3	4	6	7	0
	185.17	439.3	4	6	9	0	429.3	4	6	7	0
	185.18	439.4	4	6	9	0	429.3	4	6	7	0
	185.19	439.4	4	6	9	0	429.3	4	6	7	0
	185.20	439.4	4	6	9	0	429.4	4	6	7	0
V185/2 nd	306.1	443.9	4	6	10	0	428.9	4	6	7	0
	306.2	443.9	4	6	10	0	428.8	4	6	7	0
	306.3	449.1	4	6	11	1	428.8	4	6	7	0
	306.4	444.0	4	6	10	0	428.9	4	6	7	0
	306.5	443.9	4	6	10	0	428.8	4	6	7	0
	306.6	449.0	4	6	11	1	428.8	4	6	7	0
V185/4 th	456.1	448.9	4	6	11	1	428.7	4	6	7	0
	456.2	448.9	4	6	11	1	428.7	4	6	7	0
	456.3	459.2	4	6	13	0	429.0	4	6	7	0
	456.4	459.2	4	6	13	0	428.7	4	6	7	0
	456.5	459.2	4	6	13	0	428.8	4	6	7	0
	456.6	449.0	4	6	11	1	428.7	4	6	7	0
	456.7	449.2	4	6	11	1	429.5	4	6	7	0
	456.8	459.2	4	6	13	0	428.7	4	6	7	0
	456.9	459.3	4	6	13	0	428.6	4	6	7	0
	456.10	459.2	4	6	13	0	428.8	4	6	7	0
	456.11	449.1	4	6	11	1	428.9	4	6	7	0
	456.12	459.2	4	6	13	0	428.9	4	6	7	0
	456.13	449.1	4	6	11	1	428.7	4	6	7	0

	456.14	459.2	4	6	13	0	428.9	4	6	7	0
	456.16	459.5	4	6	13	0	428.9	4	6	7	0
	456.17	459.6	4	6	13	0	429.0	4	6	7	0
	456.18	449.5	4	6	11	1	429.1	4	6	7	0
	456.19	459.6	4	6	13	0	429.3	4	6	7	0
	456.20	459.7	4	6	13	0	429.3	4	6	7	0
cc174											
V51/1ST	51.1	640.8	5	6	8*	0	433.7	5	5	7*	0
	51.2	641.1	5	6	8	0	438.7	5	5	8	1
	51.3	646.0	5	6	9	1	438.9	5	5	8	1
	51.4	646.1	5	6	9	1	438.7	5	5	8	1
	51.5	640.9	5	6	8	0	438.7	5	5	8	1
	51.6	646.0	5	6	9	1	438.6	5	5	8	1
V51/2nd	236.1	641.0	5	6	8	0	438.7	5	5	8	1
	236.2	641.0	5	6	8	0	443.8	5	5	9	0
	236.3	641.1	5	6	8	0	433.8	5	5	7	0
	236.4	641.1	5	6	8	0	438.7	5	5	8	1
	236.5	641.1	5	6	8	0	433.7	5	5	7	0
	236.6	641.1	5	6	8	0	433.7	5	5	7	0
V51/3rd	354.1	640.8	5	6	8	0	443.6	5	5	9	0
	354.2	641.0	5	6	8	0	443.6	5	5	9	0
	354.3	640.9	5	6	8	0	443.6	5	5	9	0
	254.4	641.0	5	6	8	0	443.7	5	5	9	0
	254.5	641.0	5	6	8	0	433.6	5	5	7	0
	254.6	640.9	5	6	8	0	443.6	5	5	9	0
V51/4th	424.1	640.9	5	6	8	0	433.6	5	5	7	0
	424.2	640.9	5	6	8	0	433.5	5	5	7	0
	424.3	640.9	5	6	8*	0	433.6	5	5	7*	0
	424.4	640.9	5	6	8	0	433.4	5	5	7	0
	424.5	640.9	5	6	8	0	433.6	5	5	7	0

	424.6	640.8	5	6	8	0	438.5	5	5	8	1
V52/1st	52.1	641.3	5	6	8	0	439.0	5	5	8	1
	52.2	641.1	5	6	8	0	438.9	5	5	8	1
	52.3	641.2	5	6	8	0	439.0	5	5	8	1
	52.4	641.2	5	6	8	0	439.0	5	5	8	1
	52.5	641.1	5	6	8	0	438.8	5	5	8	1
	52.6	641.1	5	6	8	0	438.9	5	5	8	1
V52/2nd	238.1	640.2	5	6	8	0	432.9	5	5	7	0
	238.2	641.2	5	6	8	0	433.9	5	5	7	0
	238.3	641.2	5	6	8	0	449.2	5	5	10	0
	238.4	641.3	5	6	8	0	434.0	5	5	7	0
	238.5	640.0	5	6	8	0	443.2	5	5	9	0
	238.6	641.3	5	6	8	0	449.1	5	5	10	0
V52/3rd	342.1	641.1	5	6	8	0	438.8	5	5	8	1
V58/1st	58.1	646.3	5	6	9	1	444.0	5	5	9	0
	58.2	646.2	5	6	9	1	444.0	5	5	9	0
	58.3	641.2	5	6	8	0	438.8	5	5	8	1
	58.4	646.2	5	6	9	1	443.8	5	5	9	0
	58.5	645.6	5	6	9	1	443.0	5	5	9	0
	58.6	641.3	5	6	8	0	439.0	5	5	8	1
V58/2nd	240.1	640.3	5	6	8	0	437.9	5	5	8	1
V58/3rd	429.1	641.2	5	6	8	0	438.9	5	5	8	1
	492.2	641.1	5	6	8	0	438.9	5	5	8	1
	429.3	641.2	5	6	8	0	438.9	5	5	8	1
	429.4	641.1	5	6	8	0	438.8	5	5	8	1
	429.5	641.1	5	6	8	0	438.8	5	5	8	1
	429.6	641.1	5	6	8	0	438.8	5	5	8	1
V59/1st	59.1	640.7	5	6	8	0	433.5	5	5	7	0

	59.2	640.7	5	6	8	0	438.5	5	5	8	1
	59.3	640.8	5	6	8	0	443.5	5	5	9	0
	59.4	640.7	5	6	8	0	438.4	5	5	8	1
	59.5	640.5	5	6	8	0	433.2	5	5	7	0
	59.6	640.6	5	6	8	0	433.3	5	5	7	0
	59.11	640.6	5	6	8	0	433.2	5	5	7	0
	59.12	640.8	5	6	8	0	433.4	5	5	7	0
	59.13	640.8	5	6	8	0	433.2	5	5	7	0
	59.14	640.8	5	6	8	0	433.3	5	5	7	0
	59.15	640.7	5	6	8	0	433.3	5	5	7	0
	59.16	645.7	5	6	9	0	438.2	5	5	8	1
	59.17	640.6	5	6	8	0	433.1	5	5	7	0
	59.18	640.5	5	6	8	0	433.1	5	5	7	0
	59.20	641.2	5	6	8	0	438.7	5	5	8	1
V59/2nd	253.1	640.7	5	6	8	0	433.4	5	5	7	0
	253.2	640.7	5	6	8	0	433.4	5	5	7	0
	253.3	635.7	5	6	7	0	433.5	5	5	7	0
	253.4	640.8	5	6	8	0	433.4	5	5	7	0
	253.5	640.8	5	6	8	0	433.4	5	5	7	0
	253.6	640.7	5	6	8	0	433.3	5	5	7	0
V59/3rd	352.1	640.6	5	6	8	0	438.2	5	5	8	1
	352.2	640.6	5	6	8	0	438.3	5	5	8	1
	352.3	640.6	5	6	8	0	433.2	5	5	7	0
	352.4	640.6	5	6	8	0	438.2	5	5	8	1
	352.5	640.7	5	6	8	0	433.3	5	5	7	0
	352.6	640.6	5	6	8	0	433.1	5	5	7	0
	352.7	640.5	5	6	8	0	438.2	5	5	8	1
	352.8	640.6	5	6	8	0	433.3	5	5	7	0
	352.9	640.7	5	6	8	0	433.4	5	5	7	0
	352,10	640.7	5	6	8	0	438.2	5	5	8	1
	352.11	640.6	5	6	8	0	433.2	5	5	7	0

	352.12	640.5	5	6	8	0	438.2	5	5	8	1
	352.13	640.6	5	6	8	0	438.1	5	5	8	1
	352.14	640.6	5	6	8	0	438.0	5	5	8	1
	352.15	641.0	5	6	8	0	433.6	5	5	7	0
	352.16	641.1	5	6	8	0	433.6	5	5	7	0
	352.17	641.1	5	6	8	0	433.6	5	5	7	0
	352.18	641.2	5	6	8	0	438.7	5	5	8	1
	352.19	641.2	5	6	8	0	433.6	5	5	7	0
	352,20	641.1	5	6	8	0	433.6	5	5	7	0
V59/4th	438.1	645.8	5	6	9	1	438.4	5	5	8	1
	438.2	645.6	5	6	9	1	438.2	5	5	8	1
	438.3	645.6	5	6	9	1	438.3	5	5	8	1
	438.4	645.7	5	6	9	1	433.3	5	5	7	0
	438.5	645.6	5	6	9	1	438.3	5	5	8	1
	438.6	645.5	5	6	9	1	437.9	5	5	8	1
V138/1st	138.1	640.3	5	6	8	0	432.8	5	5	7	0
	138.2	640.5	5	6	8	0	433.0	5	5	7	0
	138.4	640.3	5	6	8	0	432.9	5	5	7	0
	138.5	640.4	5	6	8	0	437.8	5	5	8	1
	138.6	641.3	5	6	8	0	433.8	5	5	7	0
V138/2 nd	288.1	641.1	5	6	8	0	433.8	5	5	7	0
	288.2	641.1	5	6	8	0	433.7	5	5	7	0
	288.3	641.0	5	6	8	0	433.7	5	5	7	0
	288.4	645.3	5	6	9	1	433.7	5	5	7	0
	288.5	641.0	5	6	8	0	433.6	5	5	7	0
	288.6	640.9	5	6	8	0	433.6	5	5	7	0
V88/1st	N88.1. 1	671.1	5	6	14	0	441.3	5	5	7	0
	N88.1.	676.4	5	6	15	1	441.4	5	5	7	0

	2										
	N88.1.3	671.2	5	6	14	0	441.5	5	5	7	0
	N88.1.4	671.1	5	6	14	0	441.4	5	5	7	0
	N88.1.5	681.2	5	6	16	0	441.4	5	5	7	0
	N88.1.6	671.1	5	6	14	0	441.3	5	5	7	0
	88.1.7	671.6	5	6	14	0	441.4	5	5	7	0
	88.1.8	671.7	5	6	14	0	441.5	5	5	7	0
	88.1.9	671.8	5	6	14	0	441.4	5	5	7	0
	88.1.10	676.6	5	6	15	1	441.3	5	5	7	0
	88.1	671.2	5	6	14	0	441.0	5	5	7	0
	88.2	671.2	5	6	14	0	441.0	5	5	7	0
	88.3	671.3	5	6	14	0	441.0	5	5	7	0
	88.4	671.3	5	6	14	0	441.0	5	5	7	0
	88.5	671.1	5	6	14	0	440.9	5	5	7	0
	88.6	671.1	5	6	14	0	441.0	5	5	7	0
	88.8	671.1	5	6	14	0	440.7	5	5	7	0
V88/2nd	N272.1	671.5	5	6	14*	0	441.0	5	5	7*	0
	N272.2	671.6	5	6	14	0	441.0	5	5	7	0
	N272.3	671.6	5	6	14	0	441.0	5	5	7	0
	N272.4	671.5	5	6	14	0	440.9	5	5	7	0
	N272.5	671.5	5	6	14	0	440.9	5	5	7	0
	N272.6	671.5	5	6	14	0	441.0	5	5	7	0
V88/3rd	N369.1	665.9	5	6	13	0	446.3	5	5	8	1
	N369.2	665.9	5	6	13	0	446.4	5	5	8	1
	N369.3	661.1	5	6	12	1	441.4	5	5	7	0
	N369.3	660.8	5	6	12	1	441.3	5	5	7	0
	N369.5	665.9	5	6	13	0	446.4	5	5	8	1
	N369.6	665.9	5	6	13	0	446.3	5	5	8	1

	369.7	661.2	5	6	12	1	441.1	5	5	7	0
	369.8	666.2	5	6	13	0	446.2	5	5	8	1
	369.9	661.2	5	6	12	0	441.2	5	5	7	0
	369,10	666.4	5	6	13	0	446.2	5	5	8	1
	369.11	666.2	5	6	13	0	446.1	5	5	8	1
	369.12	671.3	5	6	14	0	446.1	5	5	8	1
	369.14	666.2	5	6	13	0	446.1	5	5	8	1
	369.15	661.1	5	6	12	1	440.9	5	5	7	0
	369.16	666.5	5	6	13	0	446.5	5	5	8	1
	369.17	681.9	5	6	16	0	446.6	5	5	8	1
	369.18	661.6	5	6	12	1	441.5	5	5	7	0
	369.19	661.5	5	6	12	1	441.6	5	5	7	0
	369,20	661.4	5	6	12	1	441.5	5	5	7	0
V88/4th	N449.1	681.2	5	6	16	0	440.4	5	5	7	0
	449.2	676.4	5	6	15	1	440.4	5	5	7	0
	449.3	676.6	5	6	15	1	440.5	5	5	7	0
	449.4	676.4	5	6	15	1	440.5	5	5	7	0
	449.5	671.3	5	6	14*	0	440.7	5	5	7*	0
	449.6	676.5	5	6	15	1	440.6	5	5	7	0
V43/2nd	241.1	651.3	5	6	10	0	438.9	5	5	8	1
	241.2	651.3	5	6	10	0	438.9	5	5	8	1
V43/3rd	349.1	646.1	5	6	9	1	443.9	5	5	9	0
	349.2	646.1	5	6	9	1	444.0	5	5	9	0
	349.3	646.4	5	6	9	1	444.0	5	5	9	0
	349.4	646.4	5	6	9	1	444.1	5	5	9	0
	349.5	646.3	5	6	9	1	444.0	5	5	9	0
	349.6	646.3	5	6	9	1	444.0	5	5	9	0

¹Fragments sizes were determined using a GS500LIZ size standard. ²Opa genes contain short C and A tracts adjacent to the 5'CTCTT repeat tract, these short tracts are invariant within a given strains as detected in all Sanger sequences of PCR products spanning the Opa repeats and in whole genome sequence data. ³Asterisks indicate repeat numbers

confirmed by dideoxy Sanger sequencing of PCR products. ⁴Expression states, 1=1, 0=0 (all genes are subject to translational phase variation).