

<b>A</b>	m1			m2			m4			m5			m6			p5			p6		
Clade 1	956	5	45	709	7	27	421	15	32	88	2	6	58	2	7	55	0	3	754	0	6
Clade 2	10	742	4	0	819	50	0	535	49	3	72	2	0	45	1	0	48	0	13	907	91
Other	189	8	2637	208	115	1548	135	59	3599	11	17	203	1	2	104	7	0	87	162	195	7977
	Major	Minor	Other	Major	Minor	Other	Major	Minor	Other	Major	Minor	Other	Major	Minor	Other	Major	Minor	Other	Major	Minor	Other

<b>B</b>	p1			p3		
Clade 1	27	2	1	17	429	354
Clade 2	0	14	18	434	0	182
Clade 3	3	0	2	70	0	160
Other	39	34	317	3	0	274
	Major	Minor	Other	Major	Minor	Other

C	p2				m3				p4		
Clade 1	0	0	22	Clade 1	0	0	155	Clade 1	0	0	1
Clade 2	0	0	19	Clade 2	0	0	280	Clade 2	0	0	2
Clade 3	0	0	5	Clade 3	0	0	206	Clade 3	0	0	8
Other	0	0	127	Clade 4	0	0	269	Other	0	0	187
				Clade 5	0	0	109				
				Other	0	0	2230				
	Major	Minor	Other		Major	Minor	Other		Major	Minor	Other

No clonal structure inferred previously