

**Table S1.** Allele sizes at eight loci for each multilocus genotype (MG) observed within clonal lineages EU1, NA1, and NA2.

MG	Microsatellite locus							
	PrMS6 (CGA) <sub>8</sub>	Pr9C3 (CA) <sub>15</sub>	PrMS39b (GA) <sub>4</sub> (GATA) <sub>33</sub>	PrMS43a (CAGA) <sub>71</sub>	PrMS43b (CAGA) <sub>75</sub> (...) (CAGA) <sub>17</sub>	PrMS45 (TCCG) <sub>11</sub>	18 (AG) <sub>39</sub>	64 (CT) <sub>16</sub>
<b><i>EU1 lineage</i></b>								
1	165/165	218/224	136/140	162/186	134/145	null	217/261	350/397
2	165/165	218/224	136/140	162/186	134/145	null	217/261	350/401
<b><i>NA1 lineage</i></b>								
1	165/168	216/226	246/246	372/372	489/489	166/186	219/275	342/379
2	165/168	216/226	250/250	368/368	485/485	166/186	219/275	342/379
3	165/168	216/226	246/246	372/372	485/485	166/186	219/275	342/379
4	165/168	216/226	250/250	368/368	489/489	166/186	219/275	342/379
5	165/168	216/226	250/250	364/364	485/485	166/186	219/275	342/379
6	165/168	216/226	246/246	368/368	489/489	166/186	219/275	342/379
7	165/168	216/226	246/246	376/376	489/489	166/186	219/275	342/379
8	165/168	216/226	250/250	364/364	485/485	166/186	219/277	342/379
9	165/168	216/226	246/246	372/372	null	166/186	219/275	342/379
10	165/168	216/226	246/246	376/376	485/485	166/186	219/275	342/379
11	165/168	216/226	246/246	372/372	493/493	166/186	219/275	342/379
12	165/168	216/226	254/254	368/368	485/485	166/186	219/275	342/379
13	165/168	216/226	250/250	360/360	485/485	166/186	219/275	342/379
14	165/168	216/226	250/250	368/368	481/481	166/186	219/275	342/379
15	165/168	216/226	250/250	372/372	489/489	166/186	219/275	342/379
16	165/168	216/226	250/250	372/372	485/485	166/186	219/279	342/379
17	165/168	216/226	246/246	372/372	489/489	166/186	219/275	342/383
18	165/168	216/226	250/250	364/364	489/489	166/186	219/275	342/379
19	165/168	216/226	242/242	372/372	485/485	166/186	219/275	342/379
20	165/168	216/226	246/246	368/368	485/485	166/186	219/275	342/379
21	165/168	216/226	246/246	376/376	493/493	166/186	219/275	342/379
22	165/168	216/226	254/254	364/364	485/485	166/186	219/275	342/379
23	165/168	216/226	250/250	349/349	485/485	166/186	219/275	342/379
24	165/168	216/226	250/250	364/364	485/485	166/186	219/279	342/379
25	165/168	216/226	250/250	364/364	489/489	166/186	219/277	342/379
26	165/168	216/226	246/246	376/376	null	166/186	219/275	342/379
27	165/168	216/226	246/246	368/368	493/493	166/186	219/279	342/379
28	165/168	216/226	250/250	360/360	481/481	166/186	219/275	342/379
29	165/168	216/226	246/246	372/372	477/477	166/186	219/275	342/379
30	165/168	216/226	246/246	372/372	481/481	166/186	219/275	342/379
31	165/168	216/226	246/246	372/372	489/489	166/186	219/271	342/379
32	165/168	216/226	250/250	376/376	489/489	166/186	219/279	342/379
33	165/168	216/226	250/250	345/345	481/481	166/186	219/275	342/379

34	165/168	216/226	250/250	364/364	477/477	166/186	219/275	342/379
35	165/168	216/226	246/246	368/368	489/489	166/186	219/275	342/342
36	165/168	216/226	246/246	372/372	418/418	166/186	219/275	342/379
37	165/168	216/226	246/246	372/372	489/489	166/186	219/279	342/379
38	165/168	216/226	250/250	376/376	489/489	166/186	219/275	342/379
39	165/168	216/226	246/246	380/380	485/485	166/186	219/275	342/379
40	165/168	216/226	250/250	372/372	485/485	166/186	219/275	342/379
41	165/168	216/226	246/246	364/364	489/489	166/186	219/275	342/379
42	165/168	216/226	246/246	368/368	406/406	166/186	219/275	342/379
43	165/168	216/226	246/246	368/368	477/477	166/186	219/275	342/379
44	165/168	216/226	250/250	368/368	493/493	166/186	219/275	342/379
45	165/168	216/226	250/250	368/368	null	166/186	219/275	342/379
46	165/168	216/226	250/250	368/368	485/485	166/186	219/277	342/379
47	165/168	216/226	250/250	368/368	489/489	166/186	219/277	342/379
48	165/168	216/226	250/250	368/368	485/485	166/186	219/279	342/379
49	165/168	216/226	250/250	376/376	485/485	166/166	219/279	342/379
50	165/168	216/226	254/254	360/360	485/485	166/186	219/271	342/379
51	165/168	216/226	254/254	364/364	481/481	166/186	219/275	342/379
52	165/168	216/226	254/254	372/372	485/485	166/186	219/275	342/379
53	165/168	216/226	254/254	368/368	485/485	186/186	219/275	342/379

***NA2 lineage***

1	165/168	216/216	145/151	157/170	null	null	221/221	344/360
2	165/168	216/216	145/151	161/174	null	null	221/221	344/360

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