

**Table S2. Mating type, and banding patterns for the allozyme *glucose-6-phosphate isomerase* and for a restriction fragment length polymorphism (RFLP) assay using the RG57 probe for the 20 unique NYS-2010/11 multilocus genotypes.** Polymorphic sites for the RFLP assay using the RG57 probe are highlighted in grey. Seven of the 24 possible combinations are observed within the 20 NYS-2010/11 genotypes.

MLG <sup>a</sup>	MT <sup>b</sup>	GPI <sup>c</sup>	MH <sup>d</sup>	RG57 <sup>e</sup>																									
				1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	24a	25
US-1	A1	86/100	H22	1	0	1	1	1	0	1	0	1	1	0	0	1	1	0	1	0	0	0	1	1	0	0	1	0	1
US-6	A1	100/100	-	1	0	1	1	1	1	1	0	0	1	0	0	1	1	0	0	0	1	0	1	1	0	0	1	0	1
US-7	A2	100/111	-	1	0	0	1	1	0	0	0	0	1	0	0	1	1	0	1	0	1	0	1	1	0	0	1	0	1
US-8	A2	100/111/122	H20	1	0	0	1	1	0	0	0	0	1	0	0	1	1	0	1	0	0	0	1	1	0	1	1	0	1
US-11	A1	100/100/111	H1	1	0	1	0	1	1	1	0	0	1	0	0	1	1	0	1	0	1	0	1	1	0	0	1	0	1
US-12	A1	100/111	-	1	0	0	0	1	0	0	0	0	1	0	0	1	1	0	0	0	1	0	1	1	0	0	1	0	1
US-14	A2	100/122	-	1	0	0	0	1	0	0	0	0	1	0	0	1	1	0	1	0	0	0	1	1	0	1	1	0	1
US-16	A1	100/111	-	1	0	0	0	1	1	0	0	0	1	0	0	1	1	0	1	0	1	0	1	1	0	0	1	0	1
US-17	A1	100/122	H28	1	0	1	0	0	0	1	0	0	0	0	0	1	1	0	1	0	1	0	1	1	0	0	1	0	1
US-19	A2	100/100	-	1	0	1	0	1	0	1	0	0	0	0	0	1	1	0	1	0	0	0	1	1	0	0	1	0	1
US-20	A2	100/100	-	1	0	1	0	1	0	1	0	0	1	0	0	1	1	0	1	0	1	0	1	1	0	0	1	0	1
US-21	A2	100/122	H20	1	0	0	0	1	0	0	0	0	1	0	0	1	1	0	0	0	0	0	1	1	0	1	1	0	1
US-22	A2	100/122	H20	1	0	0	0	1	0	0	0	0	0	0	0	1	1	0	1	0	0	0	1	1	0	0	1	0	1
US-23 <sup>f</sup>	A1	100/100	H25	1	1	0	0	1	1	0	0	0	1	0	0	1	1	0	0	1	0	1*	1	1	1*	0	1	1	1
US-24	A1	100/100	H13	1	0	1	0	1	0	1	0	0	1	0	0	1	1	0	1	0	0	0	1	1	0	1	1	0	1
GDT-01	A1	100/100	H20	1	0	0	0	1	0	0	0	0	0	0	0	1	1	0	1	0	0	0	1	1	1	0	1	0	1
GDT-02	A2	111/122	H20	1	0	0	0	1	0	0	0	0	0	0	0	1	1	0	1	0	0	1	1	1	1	0	1	0	1
GDT-03	A2	100/111/122	H20	1	0	0	0	1	0	0	0	0	0	0	0	1	1	0	1	0	0	1	1	1	1	0	1	0	1
GDT-04	A2	100/111	H20	1	0	0	0	1	0	0	0	0	0	0	0	1	1	0	1	0	0	0	1	1	0	1	0	1	1
GDT-05	A1	100/111	H20	1	0	0	0	1	0	0	0	0	0	0	0	1	1	0	1	0	0	0	1	1	0	0	1	0	1

GDT-06	A2	100/111	H20	1	0	0	0	1	0	0	0	0	0	0	0	1	1	0	1	0	0	0	1	0	1		
GDT-07	A2	100/111	H20	1	0	0	0	1	0	0	0	0	0	0	1	1	0	0	0	1	1	1	0	1	0	1	
GDT-08	A1	100/111	H20	1	0	0	0	1	0	0	0	0	0	0	1	1	0	1	1	1	1	0	0	1	0	1	
GDT-09	A2	100/122	H20	1	0	0	0	1	0	0	0	0	0	0	1	1	0	1	0	1	1	0	0	1	0	1	
GDT-10	A1	100/122	H20	1	0	0	0	1	0	0	0	0	0	0	1	1	0	1	0	1	1	1	0	1	0	1	
GDT-11	A1	111/111	H20	1	0	0	0	0	0	0	0	0	0	0	1	1	0	1	0	1	1	0	0	1	0	1	
GDT-12	A1	100/111	H20	1	0	0	0	1	0	0	0	0	0	0	1	1	0	1	0	1	1	1	0	0	1	0	1
GDT-13	A2	100/111	H20	1	0	0	0	1	0	0	0	0	0	0	1	1	0	1	0	1	1	1	0	0	1	0	1
GDT-14	A1	111/122	H20	1	0	0	0	1	0	0	0	0	0	0	1	1	0	1	0	0	1	1	0	0	1	0	1
GDT-15	A2	111/122	H20	1	0	0	0	1	0	0	0	0	0	0	1	1	0	1	0	1	1	0	0	1	0	1	
GDT-16	A2	111/122	H20	1	0	0	0	1	0	0	0	0	0	0	1	1	0	1	0	1	1	0	0	1	0	1	
GDT-17	A2	100/122	H20	1	0	0	0	1	0	0	0	0	0	0	1	1	0	1	0	1	1	1	0	1	0	1	
GDT-18	A2	100/111	H20	1	0	0	0	1	0	0	0	0	0	0	1	1	0	0	0	1	1	1	0	1	0	1	
GDT-19	A1	100/100	H20	1	0	0	0	1	0	0	0	0	0	0	1	1	0	0	0	1	1	0	0	1	0	1	
GDT-20	A1	100/111	H20	1	0	0	0	1	0	0	0	0	0	0	1	1	0	0	0	1	1	0	0	1	0	1	

<sup>a</sup> Multilocus genotype

<sup>b</sup> Mating type

<sup>c</sup> Glucose-6-phosphate isomerase

<sup>d</sup> Mitochondrial haplotype

<sup>e</sup> Restriction Fragment Length Polymorphism (RFLP) bands using the RG57 probe

<sup>f</sup> Isolate BL2009 belonging to clonal lineage US-23 lacks band 17

\* Bands 19 and 22 have not been described in previous reports for US-23 isolate. We decided to include them here because these two bands are seen on all US-23 isolates analyzed.