

Table S1. Allele-specific primer pairs used for *S*-genotyping.

S-allele	Accession no.	Sequence (5'-3')	Consensus PCR product [bp]^a	PCR product [bp]	[μM]	Reference
PcS ₁₀₃	B39S3F1	AGAACATTGATAATTAGGACCAT	1.582	430	0.3	Sanzol and Robbins 2008
	B40S3R1	GGTTTCTTATAGTCGATACCTG				
PcS ₁₀₅	A55S5F1	ACGGTATGTGGCCTTCAAACAGCTC	651	352	0.3	Sanzol and Robbins 2008
	A57S5R2	GATTGTCGTTCTGTATTGCGGGTCG				
PcS ₁₀₆	PycomC1F	ATTTTCAATTTACGCAGCAATATCAGC	670	462	0.3	Sanzol and Robbins 2008
	PycomS6R	GCGTGTTCGTTATTATTCGC				
PcS ₁₀₇	A60S7F1	CGACCCGGAATATTGCAAGATAAGG	653	443	0.4	Sanzol and Robbins 2008
	A63S7R3	GGCATTTCATGTCCAGCACT				
PcS ₁₀₈	PycomS8F	CTTGTAACGATCGTCCTGAACAA	680	613	0.3	Sanzol and Robbins 2008
	PycomS8R	CCTCAACTAATTCAGTCGTCGTC				
PcS ₁₀₉	B47S9F2	TTTGAATGGACCTCACCTGC	646	437	0.4	Sanzol and Robbins 2008
	B48S9R3	TATCTACCAGTGGCCTTTTCAA				
PcS ₁₁₁	A68S11F1	CAGAAAAATGTAAGACTACAGCTC	678	431	0.6	Sanzol and Robbins 2008
	PycomS11R	TTGTCCCGTTCGGTTGAATAGC				
PcS ₁₁₄	B36S14F2	TGGTTTGTGGCCTTCAGAGTC	642	490	0.3	Sanzol and Robbins 2008
	A71S14R4	GTCGCTTATCATCGGTACTTGG				
PcS ₁₁₅	A83SmF1	TGGCCTTCAAACCTGGTACTTATCC	658	380	0.6	Sanzol and Robbins 2008
	B37SmR2	ATGTACATTTTGATAACAGTTCCG				
PcS ₁₁₆	PcS16 forward	GCCAGCTGTCTGCAACTCC	670	430	0.4	Nikzad et al., 2014
	PcS16 reverse	GTCGTTATTATTGCGGGATAC				
PcS ₁₁₇	PpS9 forward	CTCAATCGAAGGGCAAATGT	1.600	400	0.4	Nikzad et al., 2014
	PpS9 reverse	TTATTGGTGGGGCAGAAAAA				
PcS ₁₂₁	B52S21F2	TCACCCAGAAAATTGCACGGAC	661	483	0.3	Sanzol and Robbins 2008
	B53S21R2	TTTGGTTTCTTATTGATGCTC				
PcS ₁₂₂	A84S22F1	CACGCTAGGACCTGACCCGAGT	683	418	0.6	Sanzol and Robbins 2008
	A89S22R1	CTAGAGACGTTTTGTTTCTTGACAG				
PcS ₁₂₃	A88S23F1	CAGCACAGGAAATGACCCAAGAT	653	523	0.3	Sanzol and Robbins 2008
	B38S23R2	AATTCAGTCCTCTGAGTATAGT				
PcS ₁₂₄	A85S24F1	TCTGGTGGCCTTCAAACCTGGATACT	654	489	0.6	Sanzol and Robbins 2008
	A86S24F2	GTTGGTACTAATGAGTATGGCAGC				
PcS ₁₂₅	PycomC1F	ATTTTCAATTTACGCAGCAATATCAGC	-	1095	0.4	Sanzol and Robbins 2008
	S125r1 (B61)	TGCACCGAAATGTTAGTACCA				

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PcS ₁₂₆	PcS26 forward	TCCGGTTCAATCTGCGCTTT	680	100	0.5	This study
	PcS26 reverse	TCCCGCAATAACGAACGAGG			0.5	This study
PcS ₁₂₇	PcS27 forward	CTGGCGTAAACAGTGGGAGA	850	277	0.5	This study
	PcS27 reverse	GCAAAGACCGACCTCAACCA			0.5	This study

^a Size of the amplicons obtained by using *consensus* primers PycomC1F and PycomC5R for each S-RNase alleles

Table S2. S-genotypes assigned to the accessions in analysis combining *consensus* and allele-specific PCR primers.

Accessions	Consensus PCR (pb)	S-alleles by <i>consensus</i> PCR						S-alleles by allele-specific PCR															S-genotype			
		PcS101	PcS102	PcS104	PcS110	PcS113	PcS120	PcS103	PcS105	PcS106	PcS107	PcS108	PcS109	PcS111	PcS114	PcS115	PcS116	PcS117	PcS121	PcS122	PcS123	PcS124		PcS125	PcS126	PcS127
Adamo	650/680	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-	PcS105/PcS126
Alessio	650	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS107/-
Angelico	750/1650	-	-	+	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS104/PcS103
Azzone di Cassone	850/1650	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	PcS103/PcS127
Bella di Giugno	750/800	-	-	+	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS104/PcS120 ^a
Bergamotto	680/1650	-	-	-	-	-	-	+	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	PcS103/PcS111
Bianchetto	650/1650	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	+	-	-	-	-	-	-	-	PcS109/PcS117
Bianchettone	650/750	-	-	+	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS104/PcS105
Bruttu Beddu	1650	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS103/-
Buona Luisa	1300/1650	+	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS101/PcS103
Butirra	1300/1650	+	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS101/PcS103
Campana	700/1650	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	PcS103/PcS122
Catanese	680/750	-	-	+	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS104/PcS108
Cavaliere	650/750	-	-	+	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	PcS104/PcS115
Chiuzzu	850/1650	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	PcS103/PcS127
Coscia	750/1650	-	-	+	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS104/PcS103 ^b
Duchessa d'Angio'	650/1650	-	-	-	-	-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS103/PcS105
Faccia Donna	640	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	PcS124/-
Faccibedda	680/850	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	+	PcS126/PcS127
Franconello	680/850	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	+	PcS126/PcS127
Garibaldi	650/750	-	-	+	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	PcS104/PcS115
Garofalo	640	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	PcS124/-
Gentile	670/1300	+	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS101/PcS106 ^b
Ialufaru	640	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	PcS124/-
Ianculiddu	850	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-	+	PcS125/PcS127
Iazzuleddu	850/1650	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	PcS103/PcS127
Mezza Campana	680/750	-	-	+	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS104/PcS108
Moscateello 1	650/750	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	PcS104/PcS123
Moscateello 2	680/1650	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	PcS103/PcS116
Moscateello Maiolino	680/850	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	+	PcS126/PcS127
Moscateello Nero	670/1300	+	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS101/PcS106
Paradiso/Confittaru	680/1650	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-	PcS103/PcS126

<i>P. pyraister</i> 2	680	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS111/-
<i>P. pyraister</i> 3	700	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	PcS122/-
<i>P. pyraister</i> 4	850	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	PcS127/-
<i>P. pyraister</i> 5	650	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	PcS105/PcS123
<i>P. pyraister</i> 7	680/1300	+	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	PcS101/PcS111
<i>P. pyraister</i> 8	680/850	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	+	PcS108/PcS127
<i>P. pyraister</i> 9	650	-	-	-	-	-	-	-	+	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	PcS105/PcS109
<i>P. pyraister</i> 10	750/2200	-	-	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS104/PcS110
<i>P. pyraister</i> 11	680	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS108/-
Abbé Fétel	650/750	-	-	+	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS104/PcS105 ^{a, d}
Beurre Hardy	640/680	-	-	-	-	-	-	-	-	-	-	+	-	-	+	-	-	-	-	-	-	-	-	-	-	PcS108/PcS114 ^{a, e}
Cascade	750/1300	+	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS101/PcS104 ^{a, d}
Dr. Jules Guyot	650/1300	+	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS101/PcS105 ^b
Kaiser	650	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	+	-	PcS107/PcS125 ^{a, c}
Max Red Bartlett	1300/1700	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS101/PcS102 ^d
Old Home	1300/2000	+	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS101/PcS113 ^e
Harrow Sweet	650/1700	-	+	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS102/PcS105 ^d
Williams	1300/1700	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PcS101/PcS102 ^{d, f}

For every accession, the table shows the *consensus* PCR products (bp), the positive (+) and negative (-) amplification for each S-RNase allele tested and the S-genotype.

Reference genotype: (a) Zuccherelli et al., 2002; (b) Zisovich et al., 2004; (c) Tassinari 2005; (d) Takasaki et al., 2006; (e) Moriya et al., 2007; (f) Mota et al., 2007; (g) Sanzol 2009.