

PERMANENT GENETIC RESOURCES NOTE

Permanent Genetic Resources added to Molecular Ecology
Resources Database 1 June 2011–31 July 2011

MOLECULAR ECOLOGY RESOURCES PRIMER DEVELOPMENT CONSORTIUM¹ F. KEITH BARKER,^{2,3} JAMES J. BELL,⁴ STEVEN M. BOGDANOWICZ,⁵ S. L. BONATTO,⁶ FRANK CEZILLY,⁷ SARAH M. COLLINS,⁵ CHRISTINE DUBREUIL,⁷ MATTHEW J. DUFORT,² CYRIL ERAUD,⁸ REIKO FUSEYA,⁹ E. A. HEAP,^{10,11} N. JACOBSEN,¹² M. MADDERS,¹¹ R. McEWING,¹⁰ ANDREW P. MICHEL,¹³ F. MOUGEOT,^{14,15} R. S. OGDEN,¹⁰ LUCIA C. ORANTES,¹³ A. S. OTHMAN,¹⁶ ÉRIC PARENT,¹⁷ P. PULIDO-SANTACRUZ,⁶ RACHEL RIOUX-PARÉ,¹⁷ M. F. ROBERTS,¹⁰ R. ROSAZLINA,¹⁶ TAKASHI SAKAMOTO,¹⁸ PELAYO SALINAS DE-LEÓN,⁴ JEAN-MARIE SÉVIGNY,¹⁷ PHILIPPE ST-ONGE,¹⁹ J. TERRAUBE,^{14,11} R. E. TINGAY,¹¹ RÉJEAN TREMBLAY,¹⁹ SEIICHI WATANABE¹⁸ and RÉMI A. WATTIER⁷

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Abstract

This article documents the addition of 112 microsatellite marker loci and 24 pairs of single nucleotide polymorphism (SNP) sequencing primers to the Molecular Ecology Resources Database. Loci were developed for the following species: *Agelaius phoeniceus*, *Austrolittorina cincta*, *Circus cyaneus*, *Circus macrourus*, *Circus pygargus*, *Cryptocoryne × purpurea* Ridl. nothovar. *purpurea*, *Mya arenaria*, *Patagioenas squamosa*, *Prochilodus mariae*, *Scylla serrata* and *Scyphalopus speluncae*. These loci were cross-tested on the following species: *Cryptocoryne × purpurea* nothovar. *purpurea*, *Cryptocoryne affinis*, *Cryptocoryne ciliata*, *Cryptocoryne cordata* var. *cordata*, *Cryptocoryne elliptica*, *Cryptocoryne griffithii*, *Cryptocoryne minima*, *Cryptocoryne nurii* and *Cryptocoryne schulzei*. This article also documents the addition of 24 sequencing primer pairs and 24 allele-specific primers or probes for *Aphis glycines*.

This article documents the addition of 112 microsatellite marker loci and 24 pairs of single nucleotide polymorphism (SNP) genotyping primers to the Molecular Ecology Resources Database. Table 1 contains information

on the focal species, the number of loci developed, any other species the loci were tested in and the accession numbers for the loci in both the Molecular Ecology Resources Database and GenBank. The authors

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responsible for each set of loci are listed in the final column. Table 2 presents information on SNP genotyping resources added to the MER database and presents data on the focal species, the number of sequencing primer pairs, the observed number of SNPs, other species the loci were tested in and the number of allele-specific

primers or probes. The MER database and GenBank accession numbers and the authors responsible are also listed. A full description of the development protocol for the loci presented here can be found on the Molecular Ecology Resources Database (<http://tomato.biol.trinity.edu/>).

Table 1 Information on the focal species, the number of loci developed, any other species the loci were tested in and the accession numbers for the loci in both the Molecular Ecology Resources Database and GenBank. The authors responsible for each set of loci are listed in the final column

Species	No. of primers developed	Other species tested	MER database no.	GenBank accession no.	Authors
<i>Agelaius phoeniceus</i>	10	n/a	46739–46748	JF907495–JF907504	Dufort, M. J.; Barker, F. K.
<i>Austrolittorina cincta</i>	8	n/a	46832–46825	HQ171471–HQ171478	Salinas-de-León, Pelayo; Bell, James J.
<i>Circus macrourus</i> , <i>Circus pygargus</i> , <i>Circus cyaneus</i>	19	n/a	46833–46913	AF469499–AF469502, AF469504, AF469506, AF469508, AJ620419, AJ620437, AY312455, AY312457, AY312459–AY312461, AY631063, AY631070, AY817040, AY817052, AY817053	Heap, E. A.; McEwing, R.; Roberts, M. F.; Tingay, R. E.; Mougeot, F.; Terraube, J.; Madders, M.; Ogden, R. S.
<i>Cryptocoryne</i> × <i>purpurea</i> Ridl. nothovar. <i>purpurea</i>	16	<i>C. × purpurea</i> nothovar. <i>purpurea</i> , <i>C. cordata</i> var. <i>cordata</i> , <i>C. griffithii</i> , <i>C. affinis</i> , <i>C. ciliata</i> , <i>C. elliptica</i> , <i>C. minima</i> , <i>C. nurii</i> , <i>C. schulzei</i>	46723–46738	HQ150007–HQ150022	Rosazlina, R.; Jacobsen, N.; Othman, A. S.
<i>Mya arenaria</i>	8	n/a	46786–46793	JN191327–JN191334	St-Onge, Philippe; Parent, Éric; Sévigny, Jean-Marie; Tremblay, Réjean; Rioux-Paré, Rachel
<i>Patagioenas squamosa</i>	15	n/a	46771–46785	GF111033–GF111047	Cezilly, Frank; Dubreuil, Christine; Eraud, Cyril; Wattier, Rémi A.
<i>Prochilodus mariae</i>	11	n/a	46760–46770	JF832395–JF832405	Collins, Sarah M.; Bogdanowicz, Steven M.
<i>Scylla serrata</i>	13	n/a	47010–47022	AB622868–AB622880	Fuseya, Reiko; Sakamoto, Takashi; Watanabe, Seiichi
<i>Scytalopus speluncae</i>	12	n/a	46710–46722	JF827281–JF827292	Pulido-Santacruz, P.; Bonatto, S. L.

Table 2 Information on the focal species, the sequencing primer pairs developed, the number of single nucleotide polymorphisms observed and any other species the loci were tested in. The next columns contain the number of allele-specific primers and probes developed, and the Molecular Ecology Resources database and GenBank accession numbers, respectively. The authors responsible for each set of loci are listed in the final column

Species	No. of primer pairs	No. of SNPs in sequence	Other species tested	No. of allele-specific primers/probe	Target gene(s)	MER database numbers	Genbank accession no	Authors
<i>Aphis glycines</i>	24	24	n/a	24	See Table 3 in manuscript	47024–47047	JN122354–JN122375	Orantes, Lucia C.; Michel, Andrew P.