

Correction. In the article titled “Ant Diversity and Distribution (Hymenoptera: Formicidae) Throughout Maine Lowbush Blueberry Fields in Hancock and Washington Counties” by Choate and Drummond in the April issue, 41 (2): 222–232, Figures 2 and 3 were reversed although their captions were correct. This has been corrected in the online version of the article.

Correction. In the article titled “Polymorphic Microsatellite Loci From an Indigenous Asian Fungus-Growing Termite, *Macrotermes gilvus* (Blattodea: Termitidae) and Cross Amplification in Related Taxa” by Singham et al in the April issue, 41 (2): 426–431, Table 1 was missing the gene accession numbers. The correct table is:

Table 1. Characteristics of 11 polymorphic microsatellite loci in the fungus-growing termite, *Macrotermes gilvus*, isolated using 454 pyrosequencing

Locus	Repeat motif	Primer sequence (5'-3')	Ta (°C)	N	N _A	Expected allele size (bp)	H _O	H _E	P _{HWE}	GenBank accession no.
Mg 18	(GAT) ₈	F:GGAAGAGGCTGTGGACCTG R:GTCCGTGCGTGATATTTAAACG	58.5	22	4	184	0.363	0.325	1.000	JQ065703
Mg 34	(AACT) ₆	F:TACCCGTAGCCAGAAACGC R:CCTCGTTCTGGGTGAGGAC	58.5	22	3	328	0.091	0.090	1.000	JQ065706
Mg 2	(ATC) ₈	F:GCATCATCAGTGAGAAAGGTTTG R:TGACACAGCAGGTTCCITTTG	58.5	22	4	185	0.636	0.538	0.017	JQ065695
Mg 1	(AAT) ₉	F:TTCCGCTTACTTCTGCACAC R:TTGTGACCCCGTAAGTCTGA	58.5	22	4	272	0.318	0.285	1.000	JQ065694
Mg 30	(AACT) ₁₃	F:GCTGCAAAACCAGTTAGCTTC R:TTGACTCGTTACACAATCCGTA	58.5	22	2	227	0.091	0.089	1.000	JQ065704
Mg 37	(ACTT) ₁₁	F:TTTGGCACCTGTTCGTACC R:AGGACTACTGCCACAGAC	58.5	22	3	196	0.363	0.540	0.088	JQ065708
Mg 3	(AAT) ₈	F:TAGATAGGGGTGGTGGTCA R:CCGAAAGGCGACTATGAAA	58.5	22	3	158	0.091	0.090	1.000	JQ065696
Mg 7	(ATT) ₈	F:CGAATATTTGCATCCTGGAA R:ACCCCGTAGCAGGTCTCTAT	58.5	22	2	226	0.545	0.444	0.360	JQ065699
Mg 5	(AAT) ₉	F:AAGCGATCTTGAACGTCTCTC R:GCACATTTCTTAGCCCTGGT	58.5	22	2	154	0.227	0.206	1.000	JQ065697
Mg 11	(ATT) ₉	F:CCTCCATTGGTATAAAGGCAC R:GGATAGGACAGGTGGCC	57.0	22	2	201	0.227	0.206	1.000	JQ065702
Mg 8	(AAT) ₁₁	F:CGTTCCCTCCITCCACTAAC R:ACGTCGTAGAACTCAGCGG	57.0	22	3	193	0.727	0.489	0.017	JQ065700

T_a, Primer annealing temperature; N, sample size for genetic analysis; N_A, number of alleles; H_O, observed heterozygosity, H_E, expected heterozygosity; P_{HWE}, Hardy-Weinberg equilibrium value.

Primer sequence and GenBank accession numbers of the four other monomorphic loci can be obtained from Supplemental Material (Table S1).

Correction. In the article titled “Prevalence of *Candidatus* Erwinia dacicola in Wild and Laboratory Olive Fruit Fly Populations and Across Developmental Stages” by Estes et al in the April issue 41 (2): 265–274, the corresponding author affiliations footnotes were incorrect. The correct footnotes are:

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