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Corrigendum

The authors (Hill *et al*.) of the study 'Novel microsatellite markers for the saltmarsh sharp-tailed sparrow, *Ammodramus caudacutus* (Aves: Passeriformes)' have alerted us to an error in Table 1 of their work.

The corrected table is as follows:

Table 1 Characteristics of nine new microsatellite loci from the saltmarsh sparrow, *Ammodramus caucadutus*. Size ranges, number of alleles (N_A), observed (H_O) and expected (H_E) heterozygosity from 250–350 individuals from Connecticut. Clone column lists size (in bp) of PCR product in each clone

Locus GenBank Accession no.	Primer sequences (5′–3′) 5′ fluorescent tag as indicated	Repeat motif	Clone (bp)	Range (bp)	N_A	H_O	H_E	Amplification conditions ([$MgCl_2$], AT, no. of cycles)
Aca01 EF447093	6FAM-AGCCCACTAATGGGTTTTCC TGAGTGTTCAAAGTTGCCAGA	$(TCTA)_{14}(TCA)_2$- $TCTATCA(TCTA)_{13}$	229	165–229	15	0.867	0.877	2.5 mM, 56C, 32
Aca04†	6FAM-CAATGGTGACTGCAATCCTG	$(TCTA)_{11}(TC)_3TATCC$-	211	157–291	32	0.858	0.902	2.5 mM, 56C, 32
EF447094	CCTCAGCCATTTCTGTTGTCT	$(TCTA)_{13}$						
Aca05	HEX-CCTGCTAGGCTGCATCTTCT	$(TGTC)_2(TATG)_7(TATC)_{14}$	203	193–239	12	0.823	0.860	2.5 mM, 56C, 32
EF447095	GAGTGTCATCACATTTGTACTTTGG							
Aca08	6FAM-TAGCCACAAGCAAGACCTGA	$(AC)_{18}$	175	165–185	10	0.767	0.790	3.0 mM, 56C, 32
EF447096	CTGTGACAGGAAGGGCAGTT							
Aca10	NED-GACCGGGATTTCCATTCATA	$(TATC)(TGTC)(TATC)_{11}$	190	158–194	10	0.795	0.775	2.5 mM, 56C, 32
EF447097	TGCTATTAGGTCTTTTCATTGTCC							
Aca11	NED-AGCTTCCCATACCTGAATGC	$(AGAT)_{12}$	142	130–166	14	0.789	0.832*	2.5 mM, 56C, 32
EF447098	GAAAGGCATGAGTTTACAGTGG							
Aca12‡	HEX-GCTTGTTTCCTGTTCCCAAA	$(GATA)_9G(GATA)_5(GACA)_2$-	243	213–257	20	0.785	0.902	2.5 mM, 56C, 32
EF447099	AATCGGATCCATAGACTTCAAA	$(GACG)(GATA)_2GATG$						
Aca17	NED-GGAGCATGTGACAATGGAGT	$(TCTA)_{13}(TC)_9$	242	234–258	8	0.752	0.771	2.5 mM, 50C, 40
EF447100	TCTGTGCTGTTCCAAGCAGA							
Aca21§	NED-CCCTCTCCCTGGCTACTCTT	$(ATCT)_{14}(TC)_8$	247	223–259	11	0.595	0.773*	2.5 mM, 50C, 40
EF447101	CTGGTGCTTCTGGCTCAGT							

*Significant departure from HWE.
†Although a tetranucleotide repeat with typical 4 bp stutter, Aca04 has alleles spaced every 2 bp for much of its range.
‡Although a tetranucleotide repeat, Aca12 has two series of alleles differing by a single bp, so that spacing between allelic sizes in bp goes 3, 1, 3, 1 etc.
§Aca21 has known, fairly common null alleles, with frequency estimated at 0.125.

Reference

Hill CE, Tomko S, Hagen C, Schable NA, Glenn TC (2007) Novel microsatellite markers for the saltmarsh sharp-tailed sparrow, *Ammodramus Caudacutus* (Aves: Passeriformes). *Molecular Ecology Notes*, doi: 10.1111/j.1471-8286.2007.01885.x

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