

TABLE S2 | Characterization of the *Rhinoptera bonasus* five loci. Na: number of alleles per locus; polymorphic information content (PIC) values, H_o and H_e : observed and expected heterozygosity, respectively; fixation indices F_{IS} , F_{IT} , F_{ST}

Locus	Repeat Motif	Primer sequence 5'-3'	Size (pb)	Fluorescence	Na	PIC	H_o	H_e	F_{IS}	F_{IT}	F_{ST}
Rbon_30	(AC) ₁₈	F:GGGCATAAATTCAGTCTCTC R:GTGAATCCTCTCTGCAACC	143	FAM	4	0.376	0.200	0.313	0.361	0.507	0.229
Rbon_38	(ACTC) ₁	F:TGAACGATGCAGTAGGAGGG R:TGTGATGATCGAGTGTGAG	141	FAM	3	0.386	0.351	0.407	0.137	0.323	0.216
Rbon_41	(AAGT) ₆	F:GCCTGTTTCTGCGCTGTAG R:GGGAGGGTGAAGAGGAAGTTC	126	VIC	6	0.108	0.110	0.137	0.199	0.276	0.096
Rbon_52	(AC) ₁₆	F:ACAAGGTTCAAGGTTCTTGG R:TATGTGTGCCGGTGTGCC	180	FAM	7	0.468	0.337	0.322	-0.048	0.369	0.397
Rbon_56	(AC) ₁₃	F:AGGGAGAGACAGGTCAATGC R:TCCAAGTCCAACAGAGGTCC	169	VIC	12	0.711	0.680	0.733	0.072	0.129	0.061
					Mean	0.409	0.336	0.382	0.144	0.321	0.200



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