

Microsatélites Heterólogos para *Amazona vinacea*

Espécie	Locus	Sequência de primer	Repetições	Nº de alelos	Nº de PB	GenBank	Referência	
<i>Amazona brasiliensis</i> n=15	Abra2	F: TACCCTTGCTCCTGTGATTG-6FAM R: CCACTTGCAGGATCAGAAAC	(AGAT)14	6	100-120	KX632459		
	Abra3	F: GGGGGCATTCACTTCCACTTG-ATTO532 R: TACCCTTGCTCCTGTGATTG	(ATCT)14	6	116-136	KX632460		
	Abra4	F: ACCGAGAAAGAAAATAAGCGG-ATTO550 R: AATCTTCATGCAGTGCTGGG	(GATA)12	8	116-144	KX632461		
	Abra8	F: ACCTCACATGAATGTCTTGCC-6FAM R: CGGTACAGACTGCCAGGG	(ATCT)17	7	135-167	KX632462		
	Abra9	F: CTGCATCCCCTCCTACCTTC-ATTO532 R: GTGGGGTCGGTCTGTCTATC	(TAGA)15	9	154-190	KX632463		
	Abra11	F: AGTGCTAAAAATCTTCCCATGC-ATTO550 R: ACGGATGGACGGATAGACAG	(TATC)12	12	166-220	KX632464		
	Abra12	F: TTGGAGATGTCCCAGGTTG-ATTO565 R: AGTGTGTTTTCCATTTCTCTCCC	(TGGG)14	7	150-182	KX632465		
	Abra20	F: CCCCTAAATGAGAGAATTTGGC-6FAM R: GCAGAGCAAGATGCTAAGGC	(TCTA)13	9	182-222	KX632466		
	Abra26	F: CTGAGGGCTGCTTAAGTGTG-ATTO532 R: TGGGCTCCAGGATTTAGGG	(TATC)15	8	210-242	KX632467		
	Abra32	F: ATCCTTCTGTAACCTCAACTGC-ATTO550 R: ATGCCTCAGTTTCCTCGGTG	(TATC)13	11	276-334	KX632468		
	Abra33	F: GGACCTTTCAATCCCATGTTCC-ATTO565 R: TGGTCCTTTCCTCAGCTTCC	(ATCT)14	6	265-289	KX632469		
	Abra34	F: ACAGATATCACTGCAAACAGCC-6FAM R: CCTACCTCACCACAGATGGG	(ATAG)15	6	287-311	KX632470		
	Abra35	F: GTAGCCTTCTGGAGGGGAAG-ATTO532 R: CAATGTGGTGTGGTCTAGTCTG	(AGAT)16	10	269-331	KX632471		
	Abra36	F: TGGACCTTTGGTCTGAGCTG-ATTO550 R: AGGAGTCAGGAAAGAGCTGG	(ATCT)16	5	293-309	KX632472		
	Abra37	F: ATGGGTCTCAGCGTAGATGC-ATTO565 R: GGTATGCAGGATATGCACG	(TCTA)17	11	284-326	KX632473		
	Abra41	F: CATTAAAGTCTGAGGCACCG-6FAM R: TAACCAAGCTGCTCCACAAG	(TAGA)17	7	324-352	KX632474		
	Abra42	F: GCTCTTTCAGTAGCTTCAGATTTTC-ATTO532 R: TGGCAAGATAAAGGGCTTGG	(AGAT)16	8	321-361	KX632475		
	<i>Amazona oratrix</i> n=20	Aora7	F: TACCCTTGCTCCTGTGATTG-6FAM R: GGGGCAGTTCATCCACTTG	(AGAT)16	7	109-133	KX632444	
		Aora8	F: TGGTAACACAAAAATTACCTCCTC-ATTO532 R: AGGAACTGAAATAATGAAAGATAGCC	(TATC)13	8	133-161	KX632445	

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	Aora10	F: TCAAGTGGGACGGATGGATG-ATTO550 R: TGACTCTCTCAAGGCAGGC	(AGAT)13	6	165-189	KX632446	
	Aora13	F: CGCATCTACCTAAATGAGACATTGG-ATTO565 R: TGCTTTCAGTCTGTCTATCTATCTATC	(TAGA)15	10	122-200	KX632447	
	Aora18	F: GTTCAGACGGATAGGGAGGC-6FAM R: CACTGCTCTCTGTGATAACCC	(TAGA)17	6	218-238	KX632448	
	Aora19	F: GGCTTAAACCACGCCAAC-ATTO565 R: GCCAACACGGACAGATAAGG	(TCTA)14	12	244-316	KX632449	
	Aora20	F: AAGAGCATCAGGATACTGGC-ATTO532 R: CTTTGTGATCATGCATACCTGC	(GATA)14	8	234-278	KX632450	
	Aora25	F: CACCCAGTAGCCCAGGTTC-ATTO550 R: ACTTTTGTCTGATCCATAGCC	(TATC)15	10	274-338	KX632451	
	Aora27	F: GGAAGCTGCTGATTCAGGG-ATTO565 R: GATAGGGAATGGCACAACGC	(TATC)13	7	286-314	KX632452	
	Aora28	F: GCTCTGTAAGGATGTCAGGC-6FAM R: AGTTCACCAGGCAATGAAGC	(AGAT)16	7	288-312	KX632453	
	Aora29	F: CCTGGGCTGCAATGAGTTC-ATTO532 R: ACATAGCAGGGGGAAAGGAG	(TAGA)14	9	276-312	KX632454	
	Aora30	F: AAAAGAAGACACCGCCTCAC-ATTO565 R: AAGTTTTGTGCACCCCAACC	(TAGA)16	6	300-324	KX632455	
	Aora31	F: CTCCTTTGGCTGGAGATGAG-ATTO550 R: GACAATGCTACTTTGGGGGC	(ATCT)11	8	286-338	KX632456	
	Aora33	F: AACAGAGCCCAGGAATAGCC-6FAM R: AGAAAACAAAGGGCGGTCTG	(CTAT)16	10	268-328	KX632457	
	Aora35	F: AGCTCTTCATAGACCCAGC-ATTO532 R: GGACAGATGGATAGGTAGGTAGG	(TATC)13	10	309-361	KX632458	
<i>Amazona pretrei</i> <i>n=14</i>	Apr6	F: AGGGCATAACAGGGAAGATG-6FAM R: TGCAGTTGGGTACGTTCTTG	(AAC)9	6	144-168	KX632430	
	Apr13	F: TGGTTCTGAGTTAGGGCTGG-ATTO532 R: GAGTCGATGACTACATGGAAGG	(ATCT)15	10	174-261	KX632431	
	Apr14	F: CAGTGTGAACACCATCAGCC-ATTO550 R: TCCCATCCTATGCCCATGTC	(GGAT)14	7	145-193	KX632432	
	Apr15	F: TGGAGGCTGTGCTATCTGTC-6FAM R: AGGGTTGTCAGCTCTGGAAG	(ATCT)12	10	168-224	KX632433	
	Apr18	F: CCGCATGAGATTAGCAGAAGG-ATTO565 R: ACCAGGAGGGGAAGCATAAC	(ATCT)11	6	196-224	KX632434	
	Apr19	F: ATCAAGACCTCACTGCCCTG-ATTO532 R: CAAGGTTTCATGATTTACTCTATCTC	(GATA)18	7	182-214	KX632435	

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	Apr24	F: GGGCATGTTTTCTTGGGAGC-ATTO550 R: GCTGGTGCTAATCAGGAAGC	(GATA)12	4	220-252	KX632436	
	Apr27	F: ACAACCCCTTGTAGAGCTGG-6FAM R: AGGAGTGTGCCTAATTTACTGTC	(GATA)14	6	232-252	KX632437	
	Apr30	F: ATGCAGAAGTCTGGCACAAC-ATTO532 R: CCCTCACTAGATAGATGGATGGATAG	(ATCT)14	5	259-275	KX632438	
	Apr34	F: AAGCGAAGCTGGAGCAATTC-6FAM R: GTAGACACAGATCCACACGC	(GAT)7	5	263-281	KX632439	
	Apr35	F: GGAAACAAAGACTGCTGCCC-ATTO532 R: AACTCCTTGACTTCTTGGC	(GTAT)7	6	265-305	KX632440	
	Apr36	F: AACTCTCCCTGAGCTCCAC-ATTO550 R: GCAGCATGATCCAAGAAGGG	(CTAT)13	12	231-315	KX632441	
	Apr40	F: TGAGAGCCATTGCTTTAAGTACG-ATTO565 R: ATGCCTCAGTTTCCTCGGTG	(TATC)16	6	302-322	KX632442	
	Apr49	F: CAGCATTACTGGTAACCTTGC-6FAM R: AGGAGCAGTTCCATCTCTGC	(TTTG)8	2	342-346	KX632443	
<i>Amazona rhodocorytha</i> <i>n=19</i>	Arho2	F: AGGCAATTATGCAGGCACAG-6FAM R: CTGAACTCATCCTCACATGCAG	(TTTG)7	4	145-173	KX632416	
	Arho3	F: TCCGAACCCCTTCTTGCTAC-ATTO532 R: TGGAGGAGTATTAGTATGAAATTTGGC	(ATCT)12	6	159-183	KX632417	
	Arho4	F: GCCACCAGAGTGCCTTTTAG-ATTO550 R: GACATACCCAGAAGTCAAGAGC	(TAGA)13	5	165-181	KX632418	
	Arho5	F: AATCAAGCTCCATTGCTGCC-ATTO565 R: CAGGGTCCCAGAAGAAATCC	(AGGT)11	8	194-222	KX632419	
	Arho6	F: TCCATCTATCTATCTATCTACCTGTC-6FAM R: GTGAATGGGCATGTGGGATG	(ATCT)10	10	195-223	KX632420	
	Arho7	F: GCCCTTATCATTGACCTTCCC-ATTO532 R: AGTAGCCCAGACTGAAGAGC	(TATC)13	6	196-224	KX632421	
	Arho10	F: AGGAGCTGGATATACACACAC-ATTO550 R: CTGCCCAGCAGATGTTGTTC	(TAGA)13	12	218-254	KX632422	
	Arho13	F: ACGAAGTGGTTTGGAACAGC-ATTO565 R: AGATAAACCCAGCACCTCC	(TAGA)13	9	220-272	KX632423	
	Arho14	F: ACGGAATAAGCAAACCTTACC-6FAM R: TGAGACACCAACCCCAATCC	(ATCT)9	10	249-285	KX632424	
	Arho15	F: AGTCACTGGGAAGTTAGCTGG-ATTO532 R: TCTGGGTAGGCGGATGTTTG	(TAGA)12	6	255-275	KX632425	
	Arho23	F: ATCGTGACGGAGATTCGGG-ATTO532 R: AGCCAAAATAGAAGCCACGC	(TGC)7	4	290-322	KX632426	

Polymorphic DNA microsatellite markers for forensic individual identification and parentage analyses of seven threatened species of parrots (family Psittacidae)

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	Arho52	F: CTTATTGTGACCGTGCTGG-ATTO550 R: TATGCTGAGGCAAGGTTACG	(TAGA)14	10	228-288	KX632427	
	Arho55	F: GGCCTGGTAGTTACCCTCTC-6FAM R: CCAGTCAGCACAGCATTAGC	(GATG)12	9	276-320	KX632428	
	Arho57	F: AGAAAACAGACAGATAAAGTCAGGAG-ATTO565 R: CCTTCTGCACAGCCATTCTG	(TAGA)15	10	305-337	KX632429	
<i>Anodorhynchus leari</i> n=26	Alea2	F: GGCTGGGCTAAGACTCTCTC-6FAM R: TGCCTTTCAGAGCAAACAAGG	(ATAC)9	3	84-110	KX632400	
	Alea4	F: TCACGACTGAAGCTTTTATGTCC-ATTO532 R: GACAAAGCTAAATGAAAGGGCAG	(GATA)10	4	139-167	KX632401	
	Alea5	F: GCCTAATGCAGTGAGCTGTG-ATTO550 R: AGACAGTAGATTGGAGGTAGGTAG	(CTAT)8	7	122-154	KX632402	
	Alea10	F: CTGACAGGTAGATAGGTAGGTAGG-ATTO565 R: TCTTCCAGGGCTTTTTGTGTCAG	(GATA)10	5	128-160	KX632403	
	Alea11	F: CCGTGTCAACGACAATGATG-6FAM R: ACTGCATATTTTATCTGTCTGTCTATC	(GATA)11	10	128-168	KX632404	
	Alea14	F: CCAGTCAGCACAGCATTAGC-ATTO532 R: GCATCCTGTAAACCTCTGGC	(ATCC)10	4	176-190	KX632405	
	Alea20	F: GTGAGTCATCACGTGTCAGC-ATTO550 R: TGCTTGTGACCCTCATTTTGG	(TATC)12	7	182-206	KX632406	
	Alea23	F: ACCACCCACATTTCTGCTTC-ATTO565 R: ACCCTCAGTTTCTCATCTATG	(GATA)12	7	197-225	KX632407	
	Alea24	F: CAACCTTCATGTTCCAGGGC-ATTO532 R: AGGCATTCTAGTTCTCACAGG	(TATC)12	10	204-230	KX632408	
	Alea25	F: ACTCCTGTGTGAATTTATTCTTGC-6FAM R: CAGCCCCGTCTATGGATACC	(AGAT)10	7	190-230	KX632409	
	Alea27	F: AGTGAAAGAGCCCTGCAAAG-ATTO565 R: GGACATTCATTCCCTGCAGATTAG	(GATA)12	5	213-241	KX632410	
	Alea28	F: TGAACCTCCTACCATAGCCG-ATTO532 R: TGTATTAACATCTGCCTGGATAGAC	(CATC)13	8	210-250	KX632411	
	Alea31	F: CAATGAAACACGGGAAGGGG-ATTO550 R: TTCTGTGTTGTGCAGAGGTG	(CATC)12	8	216-244	KX632412	
	Alea35	F: GCTTGCAGAAATGGGAGAGG-6FAM R: TGATTCAACCTCAGCTGCAC	(GGAT)11	3	259-271	KX632413	
	Alea46	F: AGATTTCTCCAACTGCCTC-6FAM R: AGACCTACTTTCCAATTCCTATGC	(GGAT)12	8	314-346	KX632414	
	Alea49	F: ACGTTACATGAACTGCGCTC-ATTO550 R: CCATGAAACATCATCAGCAGG	(GATA)11	7	299-347	KX632415	

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<i>Ara rubrogenys</i> n=22	Arub1	F: ACAGGCATCTGCAAGGAGAG-6FAM R: CACAGAGGAAACAGTTCCTAGC	(AGAT)12	7	92-125	KX632385	
	Arub4	F: GGCCAAAGAAGTACAGGTAG-ATTO532 R: GGGCTCTGCATAGAAAAGCTG	(GATA)13	5	130-150	KX632386	
	Arub5	F: TCAAGACGGCTGAAAATCGC-ATTO550 R: GTGCTAGAAAAGGTATATAAGTAGTGG	(ATAG)14	6	140-160	KX632387	
	Arub7	F: TGGTCTTTCTTTTCAGGCATCTC-ATTO565 R: AATTTACAGCCAAAAGACAAC	(GATG)11	6	170-190	KX632388	
	Arub12	F: TCACTTTGGTAGGGCCTGTG-ATTO532 R: AGAAAGAAAGCATTGTAGGGC	(TAAC)13	7	159-199	KX632389	
	Arub13	F: TTTAATGTTGGTGGTCTCAGC-ATTO550 R: GTCCACAAGTGAAAAGTGG	(CTAT)14	7	187-211	KX632390	
	Arub14	F: TCGAGAAGTTTGAAGTTCAG-ATTO565 R: CTGCTGCCATGATCCACATC	(TATC)16	8	180-208	KX632391	
	Arub16	F: GCTCCAAGAGTCAAAGAGGTC-6FAM R: AGAAGGGGTTTTGTATGATTCCTC	(ATCT)14	5	201-217	KX632392	
	Arub17	F: TGGGCCATTTGTCAAGCAAC-ATTO532 R: TGCAACCTGCTTTGGATCAC	(TCTA)11	7	191-215	KX632393	
	Arub19	F: GAGACACTGCCTACCCAG-ATTO550 R: AAGGGCAGCATTGGTTTCAG	(TAGA)14	10	206-250	KX632394	
	Arub25	F: ATTGGGATCACCTGGCAGAC-ATTO565 R: CATGCAGAGAAGAGGTGCTTG	(GATA)14	6	215-245	KX632395	
	Arub26	F: AAGTGACACACCCAGAGGTC-6FAM R: TGTTGTGCTGCTTGAGAAAC	(TATC)15	5	225-245	KX632396	
	Arub29	F: TTCCAGGATGGGATTGAC-ATTO532 R: ACATGTCAGTGTGCCACTTC	(TAGG)11	9	229-285	KX632397	
	Arub31	F: AGCCAGGGAGATTCCAGTTC-ATTO550 R: GAGCCTACTGGCTGGAAGAG	(ATCT)13	6	272-292	KX632398	
	Arub32	F: GACCAGGGAAGGGTTTTAC-ATTO565 R: AGCCATTCTCACTTCACTGC	(TCTA)14	6	241-273	KX632399	
	<i>Primolius couloni</i> n=19	Pcou1	F: TGCACAGCTTGTTGTATCTGTGTC-6FAM R: CTGTTCACTTCATTTCAGCTATGG	(TCTA)14	6	96-120	KX632370
Pcou4		F: TGAAGTGTCAAACACCAACG-ATTO532 R: TCCAATTCAAAGTACGTTTTTC	(GATA)13	6	131-151	KX632371	
Pcou15		F: GGGCAACTGGAGAAAAGTCAG-ATTO550 R: GGGTGGCTCAAAAATCTGGG	(TAGA)14	10	179-215	KX632372	
Pcou16		F: ATGGATGGATGGACAGACGG-ATTO565 R: TGCAACTGGGAAGAAAAGTCC	(TAGA)11	7	187-211	KX632373	

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	Pcou17	F: TGCCACCAGAGTGTCTTTTAG-6FAM R: AAGTGACACACCCAGAGGTC	(GATA)12	5	190-206	KX632374	
	Pcou18	F: ACAGCGTCCATCTAGAAACG-ATTO532 R: CCTCTCACATGTTGGGGAAAG	(ATCT)16	11	182-216	KX632375	
	Pcou25	F: GGAACAGTCATCAACAGTGGG-ATTO550 R: TAGAGTGGATGTCAAGGCGG	(GATA)12	5	218-240	KX632376	
	Pcou28	F: GCCACGTAACCTACAGGCTTC-ATTO565 R: ATCTCAGGCACTGAAAGCTC	(AGAT)16	7	210-234	KX632377	
	Pcou35	F: TCAGAATATCACATTAAGACTGAAGC-6FAM R: GTAAACTGTCTCCAGCGTGC	(TATC)15	4	240-252	KX632378	
	Pcou42	F: CCTGCTCTTCTCCAGTCAG-ATTO532 R: GCCATACAAATCTTGCTCTGC	(GATA)14	10	297-349	KX632379	
	Pcou44	F: CCCTAACAAGGCAGACATACC-ATTO550 R: CCGAATTAGGTCTTTCATCTGC	(ATCT)12	6	294-318	KX632380	
	Pcou45	F: TGGTCCTTTCCTCAGCTTCC-ATTO565 R: ACTTGAGGACCTTCCAATCCC	(GATA)13	6	290-310	KX632381	
	Pcou46	F: GGGCTTCCAAGTCCAATATG-6FAM R: ACGGAAAGTTCCTCCCTCAC	(AGAT)12	6	285-313	KX632382	
	Pcou47	F: GGGGCTGTTCCATGAATCG-ATTO532 R: GAAGCAGTGCAGCTTTAGGG	(TATC)13	6	284-316	KX632383	
	Pcou49	F: AGCAGTTGAGGGGGAGTTTC-ATTO550 R: CCTGCACCTGCATCTGTTTC	(TCTA)13	6	315-335	KX632384	
<i>Amazona guildingii</i> n=9	AgGT02	F: GATCTCATTTTGATGCCAGT R: GACTTCTGACAAGCGTGCTG	(GT)28	5	320–328	AF339754	
	AgGT04	F: CTTGCTCCACTGAAAGG R: GCGATGAAGATAACCC	(GT)11	3	264–272	AF339755	
	AgGT07	F: CAAACCATTTACACCC R: GCTCTTGAGTTTTCCC	(GT)2	6	271–284	AF339756	
	AgGT08	F: GTGGCCTAACCTGAGAGTGG R: ACATGTGCACACCTGATGG	(GT)6(GCGT)7	3	321–330	AF339757	
	AgGT12	F: ACTCATGCAGGGTTCTCAG R: TTGTGGCTGGTAGAGGTGTG	(GT)17	2	296–303	AF339758	
	AgGT17	F: TGAGTAAGGGCTGTGCAATG R: GCCTCAAGTTCTCCATTTCC	(GT)11	3	421–427	AF339759	
	AgGT21	F: TCCAGGCCAACACATTTAC R: GCTTAGTGCATATCCCAAGCTA	(GT)21	3	307–318	AF339760	
	AgGT81	F: GGGGAACATCATTCTCCAG R: AGAAGGAGGGGAAGCACATGA	(GT)20	8	330–344	AF339761	

Characterization of microsatellite loci
in the endangered St. Vincent Parrot
(*Amazona guildingii*)

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	AgGT83	F: GGCCTGGGTAACAAAAAGGT R: ACCTAAGAGGCCATCCCAGT	(GT)2	9	247–271	AF339762	
<i>Cacatua moluccensis</i>	Cmol01	F: CTGCCTCAGAGAAGCAAAC R: ATGGGTGGATTGATACGAAG	(CCAT)8	4	191–203	DQ354125	Characterisation of 13 microsatellite loci for the Moluccan Cockatoo, <i>Cacatua moluccensis</i> , and Cuban Amazon, <i>Amazona leucocephala</i> , and their conservation and utility in other parrot species (Psittaciformes)
	Cmol02	F: GGGTGAGGGAGAAAGGGAAG R:AGCAGGTATGGGGACAGCAG	(CCAT)9	5	236–252	DQ354126	
	Cmol03	F: TGGGGTGCTTTACCACTGAG R: AGAGGAACAGAATCAACAGAAG	(CTAT)9CCAT (CTAT)5	2	158–162	DQ354127	
	Cmol07	F: AGTCCCAGCCATAAGCAGAG R: CAGCATGGTGTGGGTTGAG	(CCAT)3CTAT (CCAT)5CCCT GTCC(CCAT)4 CCAC(CCAT)2 CCCTCCCCAC AT(CCCT)4 (CCAT)3 (CCCT)3	3	358–370	DQ354130	
	Cmol08	F: TGCCAGGCACAGGAGACAG R: (CCCC)GATCATCTGCTCTTG	(CCTT)8CCGT (CCTT)4	1	253	DQ354131	
<i>Amazona leucocephala</i>	Alel01	F: TCCTACCCGATCCCTAAAAG R: AGGCAGGTCATACTTCAAATC	(GTTT)5	1	218	DQ354133	
	Alel02	F: CCAGTTCAGCAGTCCTATTG R: ATTCTCTCCCTTCTCAGTTAC	(CAAA)5AA (CAAA)3	1	228	DQ354134	
	Alel03	F: CATGGTCTTCTTGCTTAC R: GTTTCAACGTCCATATCCAG	(CAAA)6	1	201	DQ354135	
	Alel04	F: GAAGGAAGGTCTGGTAGTCA R: ATCCCTCCAATGTCCACAAG	(GGAA)12	4	148–160	DQ354136	
	Alel05	F: ATCCTGTCCTTTGTCCTG R: AATGCCACCTGATGATTGATG	(CCCT)4	1	188	DQ354137	
	Alel07	F: CTGTGGTGCTGGGTGCTTG R: GGATGGGAGGATAAATGGAC	(CCCT)4TCCT (CCATCCCT)2 CTATCCAC (CCAT)5CATC (CCAT)6	1	226	DQ354139	
	Alel08	F: CATTCTTCTCTGCTCTACTAC R: GTTTTTCTAGCTGCACTCTTG	(GGAA)7GGGA (GGAA)2	1	220	DQ354140	

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	Alel09	F: CTTTTGAAGGGTTCCTATTC R: TCCATCCTGTCCCACATAC	(GGAA)4 TAAAGGAAG GCA(GGAA)6	1	190	DQ354141	
<i>Amazona aestiva</i>	AaesA	F: TAATCCTCACGTCCTATTCC R: GAAGGCAAAGAGAAAGAAGA	(CTCCTT)21	12	233–305	944212521	Development and characterization of novel microsatellite loci for the Blue-fronted Amazon (<i>Amazona aestiva</i> , Psittaciformes, Aves) and cross-species amplification for other two threatened <i>Amazona</i> species
	AaesB	F: TCTTGTGAGTGTGTTGCAG R: AACCAAACAGAAACCAATGC	(TGTGTA)15	7	270–306	944206582	
	AaesC	F: AGTCCAACACTGCACACTCT R: TATCATTACGTCATGCTCA	(TCTATA)15	14	302–422	944212093	
	AaesD	F: TGAGCAGTTCTTAGGACAGG R: AGGCACTCCAGAAAATAT	(TTCTC)25	24	343–543	944296203	
	AaesE	F: CTCATTGCTTATTCCTGTGG R: CAGGGATAGAGGAAAGGATT	(AGGAT)23	11	279–334	944212765	
	AaesF	F: AACTGCCTGAGACACACTG R: GCTGTTCTAAAGTGCGTAA	(ACGGC)21	13	277–367	944228742	
	AaesG	F: AACTTGTGTTCCCAAGGTAT R: TCTGCCAGAGTGAATTGGTC	(AGGAT)21	17	224–324	944207741	
	AaesH	F: ACTTGCATCTTTGTTAGGG R: GGCAGTTGGACTAGGTGATT	(AGGAT)17	13	315–405	944212964	
	AaesI	F: GCAGAACAGAACAGATTGGA R: GAACAAGCTCCCTACAGAGG	(AGGAT)16	13	248–338	944218382	
	AaesJ	F: CTCCCAGAAGTTTAGCCATT R: CAATACAGACCACCTTCTCG	(CTCTT)16	5	260–290	944213706	
	AaesK	F: AGAGGCTACACCTTGTCTCC R: ACCTCTCAACCACATCATTG	(AAGAG)16	16	260–375	944206688	
	AaesL	F: TGGTGAAACTTTTTGTCAGC R: ACAGGCAGTTAATGTGACCA	(TCCTA)16	18	295–410	944207207	
	AaesM	F: GAGGGCAAGTGGGTATATTT R: ACAGCTTGTGCAGATCAACT	(GGTGT)15	11	338–393	944215733	
	AaesN	F: GGTAGTGATATGAGGCAGCA R: TTTGGAGAGCAGAGGTACAA	(AGAGC)15	15	328–413	944214338	
	AaesO	F: TTCCTACGCTGATTACCTCA R: TATCTCGGTTTTATGCCTGA	(AGATG)15	15	250–340	944210789	
	AaesP	F: ATGTGTTCCAGTGAGACAGC R: AGACATGGAAGCACAGTCAG	(AACAT)15			944211022	
	AaesQ	F: TCAGAGTGCTTGCCTCATT R: TCTGCAAAGACTTAGCCAAC	(AAGAG)14	11	235–305	944218360	
	AaesR	F: GAAACACCCAGCCTTAACTG	(CCTGT)14	12	305–360	944216362	

Microsatélites Heterólogos para Amazona vinacea

Espécie	Locus	Sequência de primer	Repetições	Nº de alelos	Nº de PB	GenBank	Referência
		R: CTATGACGGAAATCAAGCAG					
	AaesS	F: TCCTCAGCCTTTTGATGTA	(ACTCT)14	12	333–403	944214087	
		R: TGGTGATAGCAGGGAATTTA					
	AaesT	F: AACTGGCAGATTAAGCACAA	(GGAGT)14	9	253–308	944211765	
		R: CTCGTCTCGTCTTCTCCT					
	AaesU	F: AATCAGGTGTGGCAGGTACT	(AACT)23	20	292–413	944297128	
		R: CTCGTTTGCTTCTAGCAATG					
	AaesV	F: CTGGGCTTAGGTTTTGTTTT	(GGAT)26	13	257–329	944216972	
		R: TGCTCATAAGGGGTTGAGTA					
	AaesW	F: CACCTTTCGTGCTTGACAGC	(AGGC)19	10	175–211	944216243	
		R: GTGGGTGCTCCGTGTGTC					
	AaesX	F: GATGGAGTTTGCAGGATACA	(CATT)20	9	266–302	944211341	
		R: TTCAACCATCAAGGAATGAC					