

J . i f G t , V . 97, S 1, D
/10.1007/s12041-018-0935-7

2018, O R 61 65

I A S .



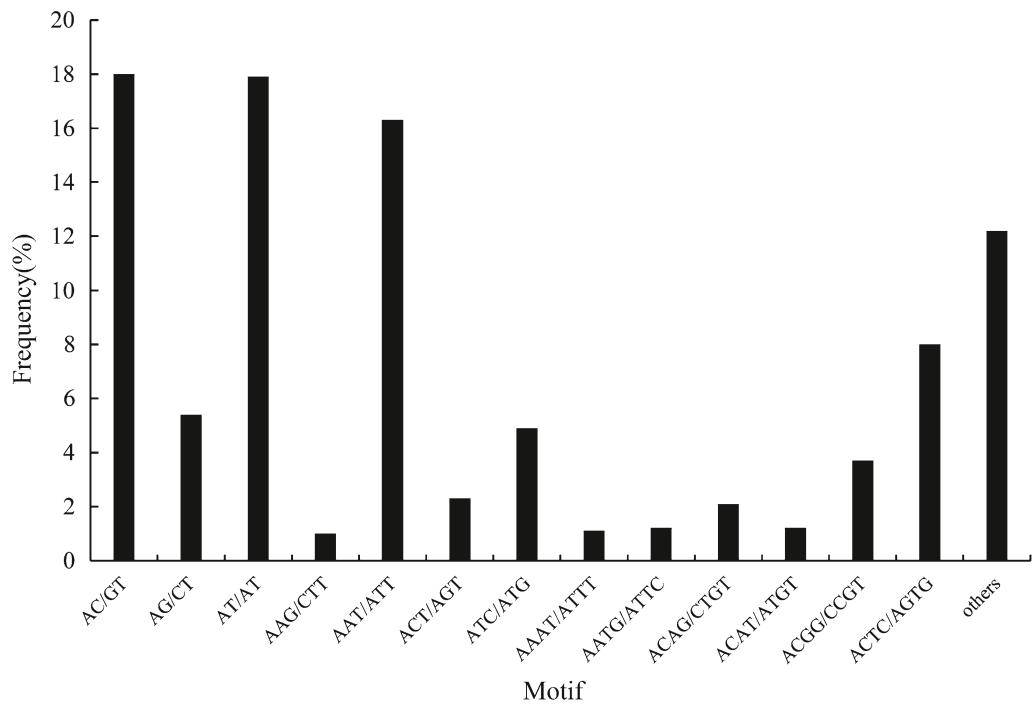
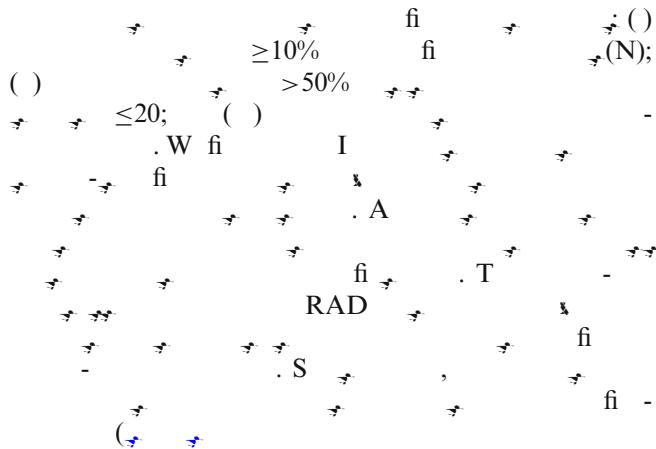


Fig. 1. D.

C.



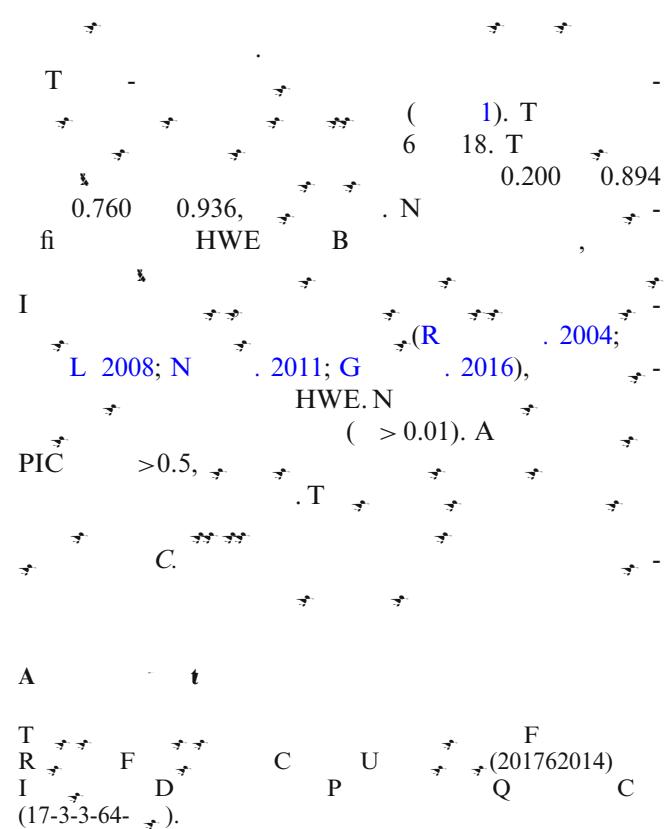
L	r	P	r	(5' 3')	R	N . r	S ()	({°C})	H	H	PIC	G	B
												r	r
X 04	F: CATGTTTCTTGCTTTTCACG R: AACTGGTTGGCTGGAGTAA	(TG) ₂₃	11	270 320	60	0.744	0.851	0.117	0.822	MF460397			
X 08	F: GCTGGACTGCTGGAGACT R: TTGGTAGTTGTGCTGGTCCCCG	(TG) ₁₃	9	350 380	60	0.698	0.823	0.148	0.790	MF460398			
X 11	F: TTACAAAACCTTGTGTGCG R: GTTTGATTACGACACGCC	(AC) ₃₀	10	250 290	60	0.600	0.882	0.001*	0.857	MF460399			
X 17	F: CGTTTGAATACCTTTGAA R: CGGTGTTCAAAAGATGCTGAA	(AG) ₁₅	11	180 220	60	0.382	0.843	0.030	0.810	MF460400			
X 23	F: CGTCGTGACCTCTGGCTAT R: GTAGTATGGGGAGGACAA	(AT) ₁₂	13	240 310	60	0.682	0.919	0.000*	0.902	MF460401			
X 47	F: TTGATGACAACAATAAGGCG R: ACAAAACCCCTCCCAAC	(AT) ₈	7	188 208	60	0.467	0.763	0.000*	0.713	MF460402			
X 48	F: TTCTACATGGCCGGATTG R: CTGATATCAAACCCACGCA	(AT) ₁₀	6	262 278	60	0.500	0.760	0.004	0.709	MF460403			
X 51	F: GTTATCTGCATGCTGGC R: GTGCAGAAAAAAGTGGAA	(TG) ₁₅	16	236 276	60	0.591	0.873	0.000*	0.850	MF460404			
X 07	F: TGACTGGAAACAAAACGGAA R: ACGCACAACTGCACACATT	(AAT) ₁₃	10	150 186	60	0.870	0.868	0.137	0.843	MF460405			
X 09	F: GGACTGTAAAGCGCTATGAGCA R: GAAGAAAACGCTAAATCTCCA	(TAT) ₁₇	10	191 09	60	0.870	0.869	0.168	0.844	MF460406			
X 12	F: TCATCAAACGCCGTAATCA R: GCTGAAAAGCGCTCCCTAAA	(TGA) ₁₂	12	189 228	60	0.650	0.833	0.001*	0.859	MF460407			
X 17	F: CGCTCAAGAACATCTGTCCA R: TTGTGACTTAATTTGACGGC	(TAA) ₉	13	199 238	60	0.711	0.868	0.030	0.843	MF460408			
X 19	F: CATGCAGTCAGTCAGTTCAT R: TGGCGTGTCAAGCAACGAT	(AAT) ₁₃	17	169 220	60	0.783	0.936	0.003	0.921	MF460409			
X 22	F: CCTGTTCTCCAGCAACGAT R: TGGCAGTCAGTCAGTTCAT	(TAC) ₉	18	223 289	60	0.763	0.893	0.839	0.871	MF460410			
X 27	F: CTTCTGGGTGTGGAAATTGG R: TCTCCTACAGCGATGACGTG	(TAT) ₉	10	226 253	60	0.609	0.860	0.004	0.834	MF460411			
X 29	F: CAAGTAGCAAATGCAAAATCCC R: TATTCTGTCACGGTTATTGC	(TTA) ₉	14	172 215	60	0.500	0.919	0.000*	0.899	MF460412			
X 31	F: TCATCAAACGCCGTAATCA R: GATTGCGCGTGAATAGG	(TAC) ₂₈	15	166 217	60	0.894	0.928	0.147	0.912	MF460413			

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L	P	(5' 3')	R	N.	S	()	(°C)	H	H	PIC	G B PIC
X 33	F: GGGAAATCATCGCCAAAGTTA R: ACGTGAACTGTTCATTCATTGCAG	(ATD) ₇	11	172	202	60	0.439	0.872	0.000*	0.846	MF460414
X 37	F: AATTCACTGCAGCCGTITAGG R: CAACTGTGAGATGACACGGG	(AAT) ₁₁	11	248	281	60	0.200	0.857	0.000*	0.828	MF460415
X 41	F: CATGCAGTCGTTAGGAGCA R: TACATGAIGGCCGTTTICA	(ATD) ₉	12	198	234	60	0.717	0.895	0.020*	0.874	MF460416
X 45	F: CATTGTTGTGCTGGACTGGA R: CGGTACCGCACCATCTACTCA	(TAD) ₉	16	166	211	60	0.783	0.919	0.004	0.902	MF460417
X 47	F: ACCATCGGGATTTACG R: CCTGTCTGTGGTAAAGTG	(ATG) ₈	13	243	288	60	0.761	0.880	0.480	0.858	MF460418
X 54	F: CATATCATGCAAGTCATTG R: GGCCCTGTCTAAATG	(TCA) ₁₃	8	184	208	60	0.745	0.826	0.451	0.793	MF460420

*S ;H , * ;H , * ;PIC,
(< 0.05/23).

L W



B N. A., E P. D., A T. S., C M. C., S A.
L., L A. J E. A. 2008 R SNP RAD L
3, 3376.

B M., A C. M. M A. D. 2014 C 13
1179 1187. L M B N C 41,
D D. S C. 2003 M
(MSA): X., X., B Q. M E 31, 167 169.
G X., X., B Q. M E L Q. 2016 P (S
1920) (C O B E C 66, 129 136.
K S. T., T M. L. M T. C. 2007 R
CERVUS M
E 16, 1099 1106.
M M., H M. P W. 2002 M DNA
N L, L Q. G K L 2011 M fi
(M B M C M N C M
E 32, 488 497.

R K. S., R W. L., G P. M., C R. B.
A S. K. J. 2004 M

J. H. 95, 346 352.
 R. F. 2008 G 007:
 T. 8, 103 106.
 EST T., M W, V R. K. G A. 2003

(C -M L.): fi - W L.). A G SSR-
 W (M J. 1994 E 106, 411 422.
 H. Z Q. 2008 E C F 13, 38 44 (C).
 EST EST-SSR P fi
 (C). J. H. 99, 208 214.

C : INDRAJIT NANDA