



Supplementary Material

Multi-gene Phylogenetic Analysis and Genetic Diversity of Discrete Elytral Color Phenotypes in *Menochilus sexmaculatus* (Coleoptera: Coccinellidae)

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Supplementary Table S1. GenBank accession numbers.

Specimen	Code	COI	COII	CAD	H3
<i>M. sexmaculatus</i>	A1	MH589162	MH589194	MH589228	MH589128
<i>M. sexmaculatus</i>	A2	MH589163	MH589195	MH589229	MH589129
<i>M. sexmaculatus</i>	B1	MH589164	MH589196	MH589230	MH589130
<i>M. sexmaculatus</i>	B2	MH589165	MH589197	MH589231	MH589131
<i>M. sexmaculatus</i>	C1	MH589166	MH589198	MH589232	MH589132
<i>M. sexmaculatus</i>	C2	MH589161	MH589199	MH589233	MH589133
<i>M. sexmaculatus</i>	D1	MH589167	MH589200	MH589234	MH589134
<i>M. sexmaculatus</i>	D2	MH589168	MH589201	MH589235	MH589135
<i>M. sexmaculatus</i>	E1	MH589169	MH589202	MH589236	MH589136
<i>M. sexmaculatus</i>	E2	MH589170	MH589203	MH589237	MH589137
<i>M. sexmaculatus</i>	F1	MH589171	MH589204	MH589238	MH589138
<i>M. sexmaculatus</i>	F2	MH589172	MH589205	MH589239	MH589139
<i>M. sexmaculatus</i>	G1	MH589173	MH589206	MH589240	MH589140
<i>M. sexmaculatus</i>	G2	MH589174	MH589207	MH589241	MH589141
<i>M. sexmaculatus</i>	H1	MH589175	MH589208	MH589242	MH589142
<i>M. sexmaculatus</i>	H2	MH589176	MH589209	MH589243	MH589143
<i>M. sexmaculatus</i>	I1	MH589177	MH589210	MH589244	MH589144
<i>M. sexmaculatus</i>	I2	MH589178	MH589211	MH589245	MH589145
<i>M. sexmaculatus</i>	J1	MH589178	MH589212	MH589246	MH589145
<i>M. sexmaculatus</i>	J2	MH589179	MH589213	MH589247	MH589146
<i>M. sexmaculatus</i>	K1	MH589180	MH589214	MH589248	MH589147
<i>M. sexmaculatus</i>	K2	MH589181	MH589215	MH589249	MH589148
<i>M. sexmaculatus</i>	L1	MH589182	MH589216	MH589250	MH589149
<i>M. sexmaculatus</i>	L2	MH589183	MH589217	MH589251	MH589150
<i>M. sexmaculatus</i>	M1	MH589184	MH589218	MH589252	MH589151
<i>M. sexmaculatus</i>	M2	MH589185	MH589219	MH589253	MH589152
<i>M. sexmaculatus</i>	N1	MH589186	MH589220	MH589254	MH589153
<i>M. sexmaculatus</i>	N2	MH589187	MH589221	MH589255	MH589154
<i>M. sexmaculatus</i>	O1	MH589188	MH589222	MH589256	MH589155
<i>M. sexmaculatus</i>	O2	MH589189	MH589223	MH589257	MH589156
<i>M. sexmaculatus</i>	P1	MH589190	MH589224	MH589258	MH589157
<i>M. sexmaculatus</i>	P2	MH589191	MH589225	MH589259	MH589158
<i>M. sexmaculatus</i>	Q1	MH589192	MH589226	MH589260	MH589159
<i>M. sexmaculatus</i>	Q2	MH589193	MH589227	MH589261	MH589160

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Supplementary Table S2, The genetic distance of the different elytra forms in *M.sexmaculatus* based on combined dataset.

	A1	A2	B1	B2	C1	C2	D1	D2	E1	E2	F1	F2	G1	G2	H1	H2	I1	I2	J1	J2	K1	K2	L1	L2	M1	M2	N1	N2	O1	O2	P1	P2	Q1	Q2
A1																																		
A2	0.001																																	
B1	0.002	0.002																																
B2	0.001	0.001	0.001																															
C1	0.008	0.008	0.009	0.008																														
C2	0.010	0.010	0.010	0.010	0.002																													
D1	0.000	0.001	0.001	0.000	0.008	0.009																												
D2	0.007	0.007	0.007	0.008	0.007	0.001	0.003	0.006																										
E1	0.000	0.001	0.001	0.000	0.008	0.009	0.000	0.006																										
E2	0.007	0.008	0.007	0.008	0.002	0.004	0.008	0.001	0.008																									
F1	0.007	0.007	0.008	0.007	0.001	0.003	0.006	0.000	0.006	0.001																								
F2	0.008	0.008	0.008	0.008	0.002	0.003	0.008	0.001	0.008	0.002	0.001																							
G1	0.000	0.001	0.001	0.000	0.008	0.009	0.000	0.006	0.000	0.008	0.006	0.008																						
G2	0.008	0.009	0.009	0.008	0.003	0.004	0.008	0.002	0.008	0.002	0.002	0.003	0.008																					
H1	0.002	0.002	0.002	0.001	0.008	0.011	0.002	0.008	0.002	0.009	0.008	0.009	0.002	0.010																				
H2	0.008	0.008	0.009	0.008	0.002	0.004	0.008	0.001	0.008	0.002	0.001	0.002	0.008	0.001	0.009																			
I1	0.007	0.008	0.008	0.007	0.002	0.003	0.007	0.000	0.007	0.002	0.000	0.002	0.000	0.002	0.002	0.001																		
I2	0.001	0.001	0.001	0.000	0.007	0.006	0.000	0.005	0.000	0.007	0.005	0.007	0.000	0.007	0.002	0.006	0.005																	
J1	0.001	0.001	0.002	0.001	0.007	0.009	0.000	0.006	0.000	0.007	0.006	0.007	0.000	0.008	0.002	0.007	0.006	0.000																
J2	0.008	0.009	0.009	0.008	0.002	0.004	0.008	0.002	0.008	0.003	0.002	0.003	0.008	0.003	0.007	0.003	0.002	0.008	0.007															
K1	0.007	0.008	0.008	0.007	0.002	0.003	0.007	0.000	0.007	0.002	0.000	0.002	0.000	0.002	0.002	0.001	0.005	0.006	0.002															
K2	0.007	0.008	0.008	0.007	0.002	0.003	0.007	0.000	0.007	0.002	0.000	0.002	0.000	0.002	0.007	0.002	0.001	0.006	0.001	0.001														
L1	0.001	0.002	0.002	0.001	0.008	0.009	0.001	0.007	0.001	0.008	0.007	0.008	0.001	0.009	0.002	0.008	0.008	0.001	0.009	0.008	0.008													
L2	0.008	0.009	0.009	0.008	0.002	0.003	0.008	0.002	0.008	0.003	0.002	0.002	0.008	0.003	0.010	0.003	0.002	0.008	0.003	0.002	0.007	0.008	0.002	0.009										
M1	0.002	0.002	0.002	0.002	0.008	0.010	0.001	0.008	0.001	0.009	0.008	0.009	0.001	0.009	0.000	0.009	0.008	0.002	0.007	0.008	0.008	0.002	0.009											
M2	0.000	0.001	0.001	0.000	0.008	0.009	0.000	0.006	0.000	0.008	0.006	0.008	0.000	0.008	0.002	0.008	0.007	0.000	0.008	0.007	0.007	0.001	0.008	0.001										
N1	0.007	0.008	0.008	0.007	0.002	0.003	0.007	0.000	0.007	0.002	0.000	0.002	0.000	0.002	0.002	0.001	0.005	0.006	0.001	0.001	0.000	0.008	0.002	0.008	0.007									
N2	0.007	0.008	0.008	0.007	0.002	0.003	0.007	0.000	0.007	0.002	0.000	0.002	0.000	0.002	0.007	0.002	0.005	0.006	0.002	0.001	0.001	0.008	0.002	0.008	0.007									
O1	0.002	0.002	0.003	0.002	0.008	0.011	0.002	0.008	0.002	0.009	0.008	0.009	0.002	0.010	0.001	0.009	0.008	0.002	0.007	0.008	0.008	0.002	0.010	0.000	0.002	0.008								
O2	0.008	0.008	0.009	0.008	0.002	0.004	0.008	0.001	0.008	0.002	0.001	0.002	0.008	0.003	0.007	0.002	0.002	0.007	0.000	0.002	0.002	0.008	0.003	0.006	0.008	0.002	0.002	0.007						
P1	0.000	0.001	0.001	0.000	0.008	0.009	0.000	0.006	0.000	0.008	0.006	0.008	0.000	0.008	0.002	0.008	0.007	0.000	0.008	0.007	0.007	0.001	0.008	0.001	0.000	0.007	0.007	0.002	0.008					
P2	0.007	0.007	0.008	0.007	0.001	0.003	0.006	0.000	0.006	0.001	0.000	0.001	0.000	0.002	0.008	0.001	0.000	0.005	0.006	0.002	0.000	0.007	0.002	0.008	0.006	0.000	0.000	0.008	0.001	0.006				
Q1	0.008	0.009	0.009	0.008	0.002	0.004	0.008	0.002	0.008	0.003	0.002	0.003	0.008	0.003	0.007	0.003	0.002	0.008	0.000	0.002	0.001	0.009	0.003	0.007	0.008	0.001	0.002	0.007	0.000	0.008	0.002			
Q2	0.001	0.002	0.001	0.001	0.008	0.010	0.001	0.007	0.001	0.008	0.007	0.008	0.001	0.009	0.002	0.008	0.008	0.001	0.009	0.008	0.008	0.002	0.009	0.002	0.001	0.008	0.008	0.002	0.008	0.001	0.007	0.009		

*The code in the table corresponds to code in the table!