Original Group	WGA_A	WGA_B	SH_A	SH_B
Taxonomy level	*phylum* ($m = 9$, $\delta_1^2 = 1$, P = 0.207)			
% correct	100	33	33	0
correct/total	3/3	1/3	1/3	0/3
Misclassified to group	n/a	SH_B	SH_B	SH_A WGA_B
Taxonomy level	*class* ($m = 5$, $\delta_1^2 = 0.936$, P = 0.039)			
% correct	33	0	33	100
correct/total	1/3	0/3	1/3	3/3
Misclassified to group	SH_B	SH_A SH_B	SH_B	n/a
Taxonomy level	*order* ($m = 4$, $\delta_1^2 = 0.928$, P = 0.009)			
% correct	66	0	66	100
correct/total	2/3	0/3	2/3	3/3
Misclassified to group	SH_B	SH_A SH_B	SH_B	n/a
Taxonomy level	*family* ($m = 6$, $\delta_1^2 = 0.99$, P = 0.0108)			
% correct	66	0	66	66
correct/total	2/3	0/3	2/3	2/3
Misclassified to group	SH_B	SH_B	SH_B	WGA_B
Taxonomy level	*genus* ($m = 8$, $\delta_1^2 = 099$, P = 0.0008)			
% correct	100	0	100	33
correct/total	3/3	0/3	3/3	1/3
Misclassified to group	n/a	SH_A SH_B	n/a	SH_A WGA_B
Taxonomy level	*species* ($m = 8$, $\delta_1^2 = 0.99$, P = 0.003)			
% correct	100	0	66	33
correct/total	3/3	0/3	2/3	1/3
Misclassified to group	n/a	SH_B	SH_B	SH_A WGA_B