Original Group	AP_A	AP_B	WGA_A	WGA_B	SH_A	SH_B
Taxonomy level phylum $(m = 2, \delta_1^2 = 0.97, P = 0.0001)$						
% correct	100	100	100	0	33	33
correct/total	3/3	3/3	3/3	0/3	1/3	1/3
N.C. 1 (C. 1)				SH_A		
Misclassified to group	n/a	n/a	n/a	SH_B	SH_B	SH_A
group				WGA_A		
Taxonomy level	class $(m = 5, \delta_1^2 = 0.98, P = 0.0001)$					
% correct	100	100	100	0	67	67
correct/total	3/3	3/3	3/3	0/3	2/3	2/3
N.C. 1 (C. 1)				SH_B		
Misclassified to	n/a	n/a	n/a	WGA_A	SH_B	WGA_B
group						
Taxonomy level	order $(m = 3, \delta_1^2 = 0)$	0.98, P = 0.0	001)			
% correct	100	100	100	0	33	67
correct/total	3/3	3/3	3/3	0/3	1/3	2/3
3.61 1 107 1				SH_B		
Misclassified to	n/a	n/a	n/a	WGA_A	SH_B	SH_A
group						
Taxonomy level	family $(m = 9, \delta_1^2 = 0.99, P = 0.0081)$					
% correct	100	100	100	0	67	67
correct/total	3/3	3/3	3/3	0/3	2/3	2/3
M: 1 :C 14				SH_B		
Misclassified to group	n/a	n/a	n/a	WGA_A	SH_B	WGA_B
				SH_A		
Taxonomy level	genus $(m = 10, \delta_1^2 =$	= 0.99, P = 0	.0078)			
% correct	100	100	100	0	67	67
correct/total	3/3	3/3	3/3	0/3	2/3	2/3
M. 1 'C 1.				SH_B		
Misclassified to group	n/a	n/a	n/a	WGA_A	SH_B	SH_A
group				SH_A		
Taxonomy level	species $(m = 10, \delta_1^2 = 0.99, P = 0.0621)$					
% correct	100	100	100	0	67	67
correct/total	3/3	3/3	3/3	0/3	2/3	2/3
AC 1 10 1				SH_B		
Misclassified to	n/a	n/a	n/a	WGA_A	SH_B	SH_A
group				SH_A		