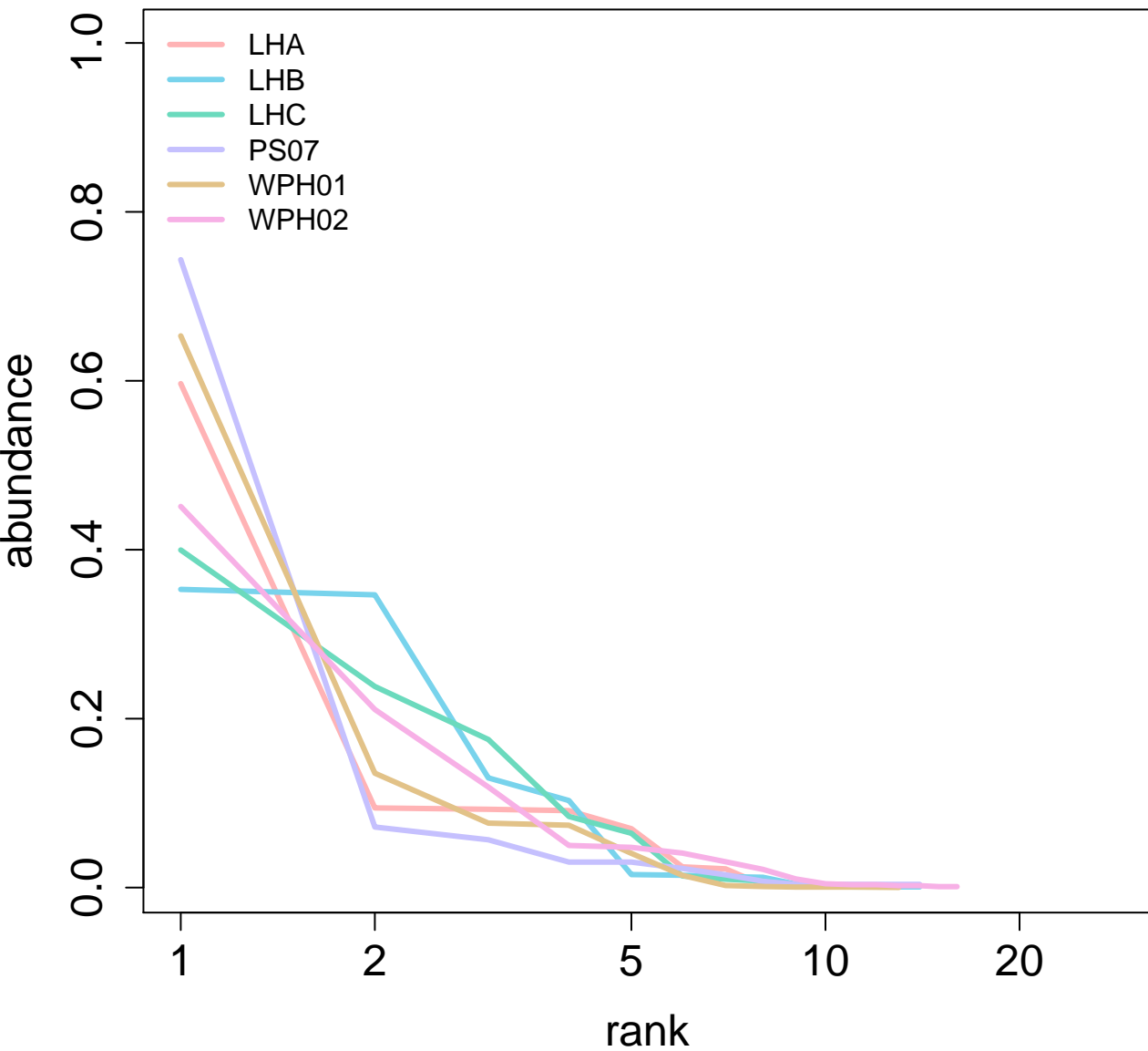
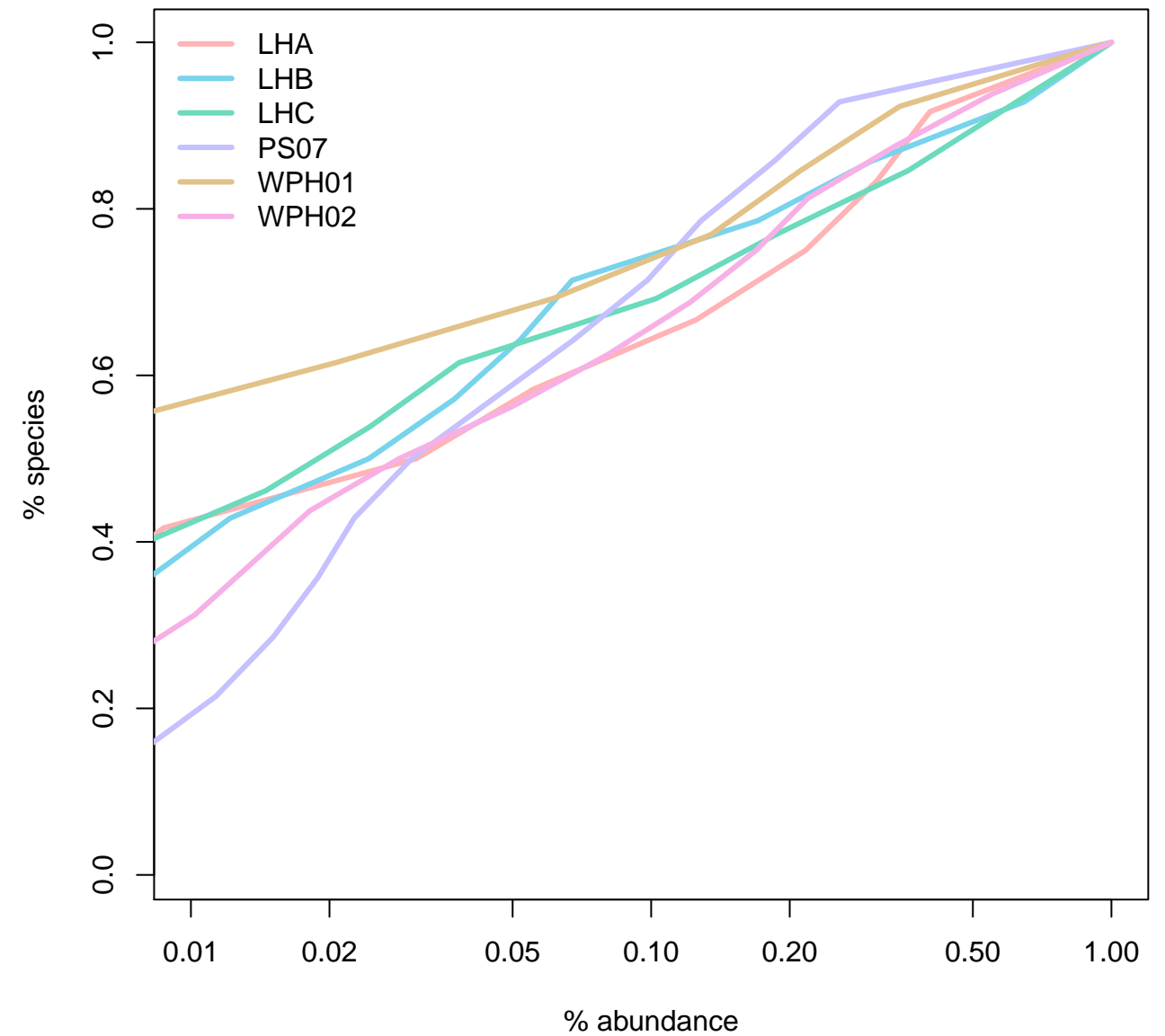
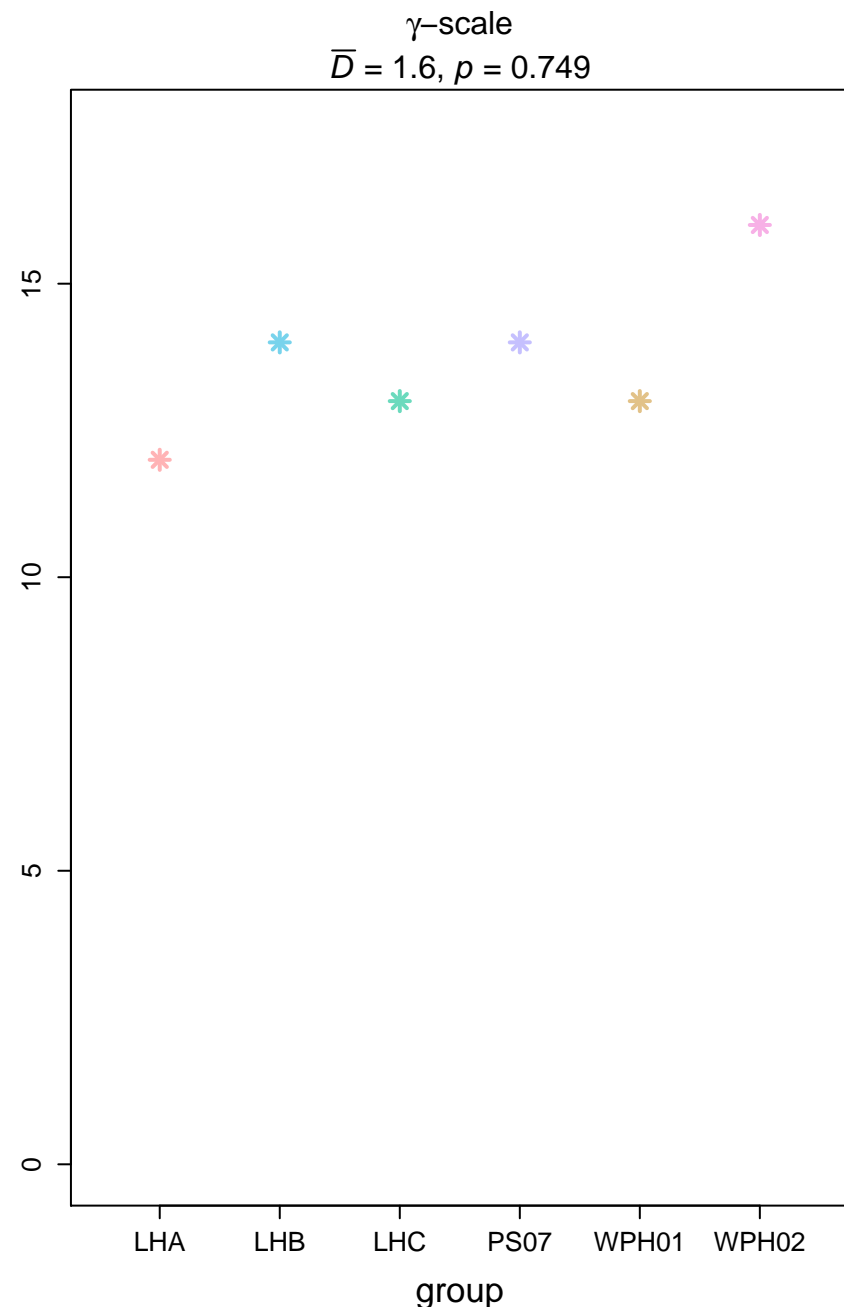
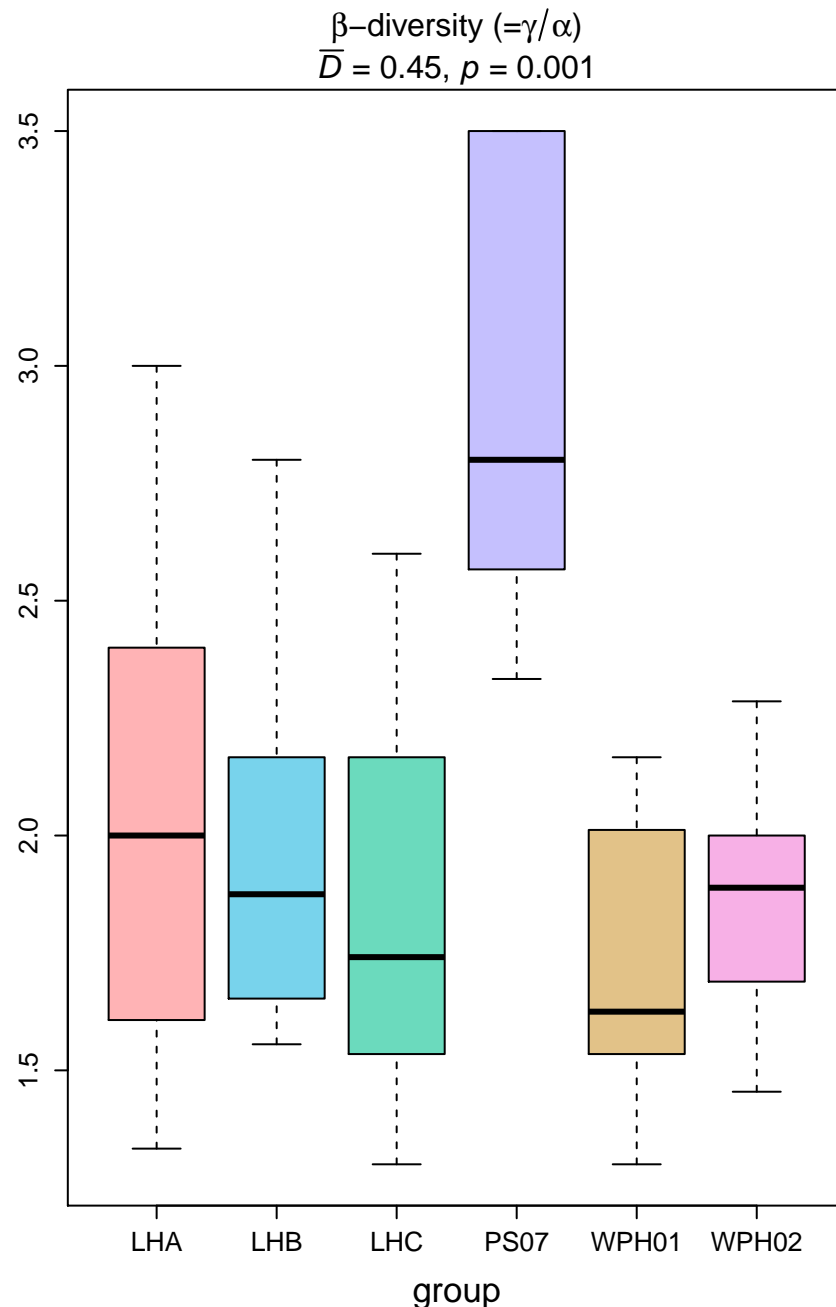
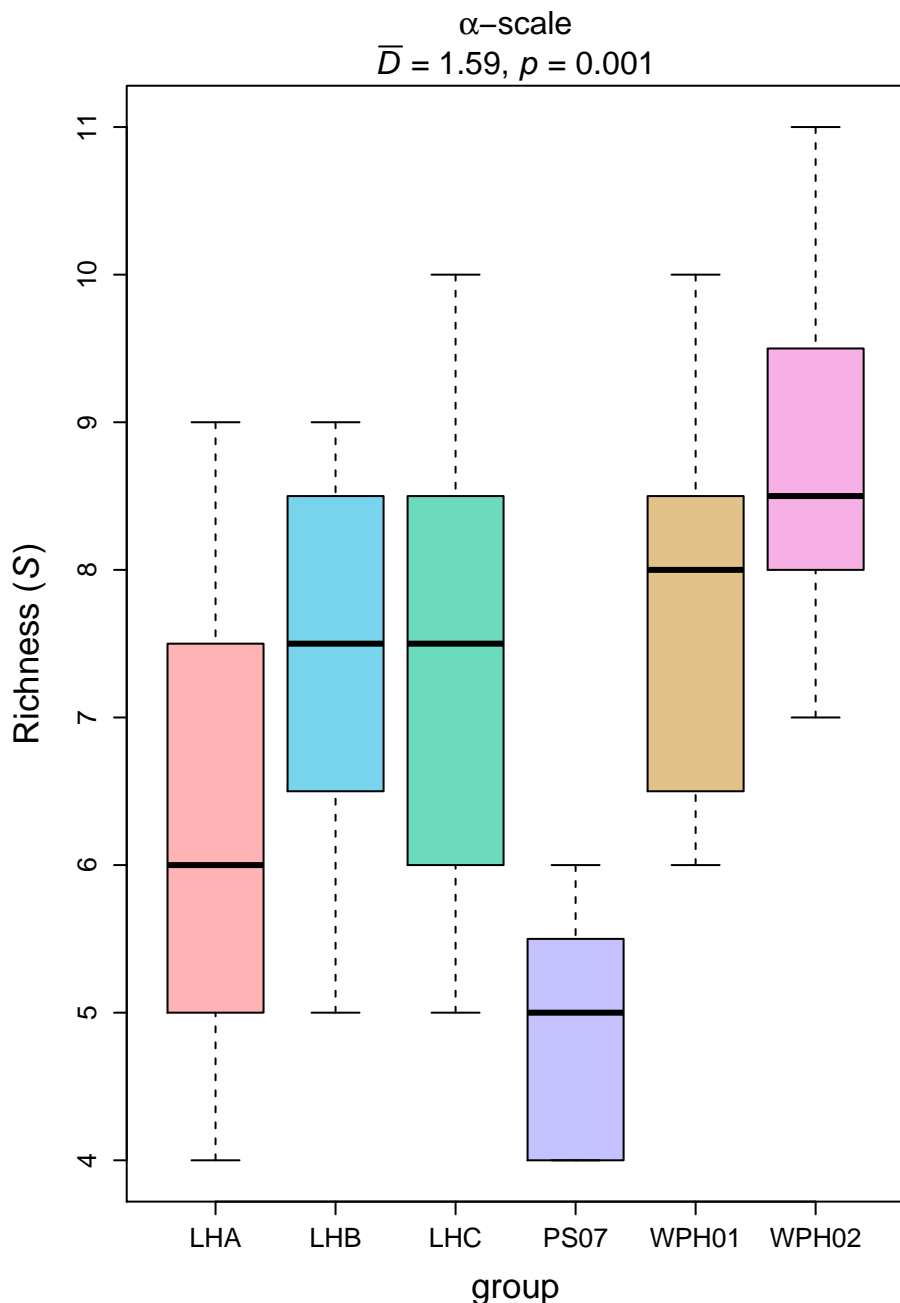


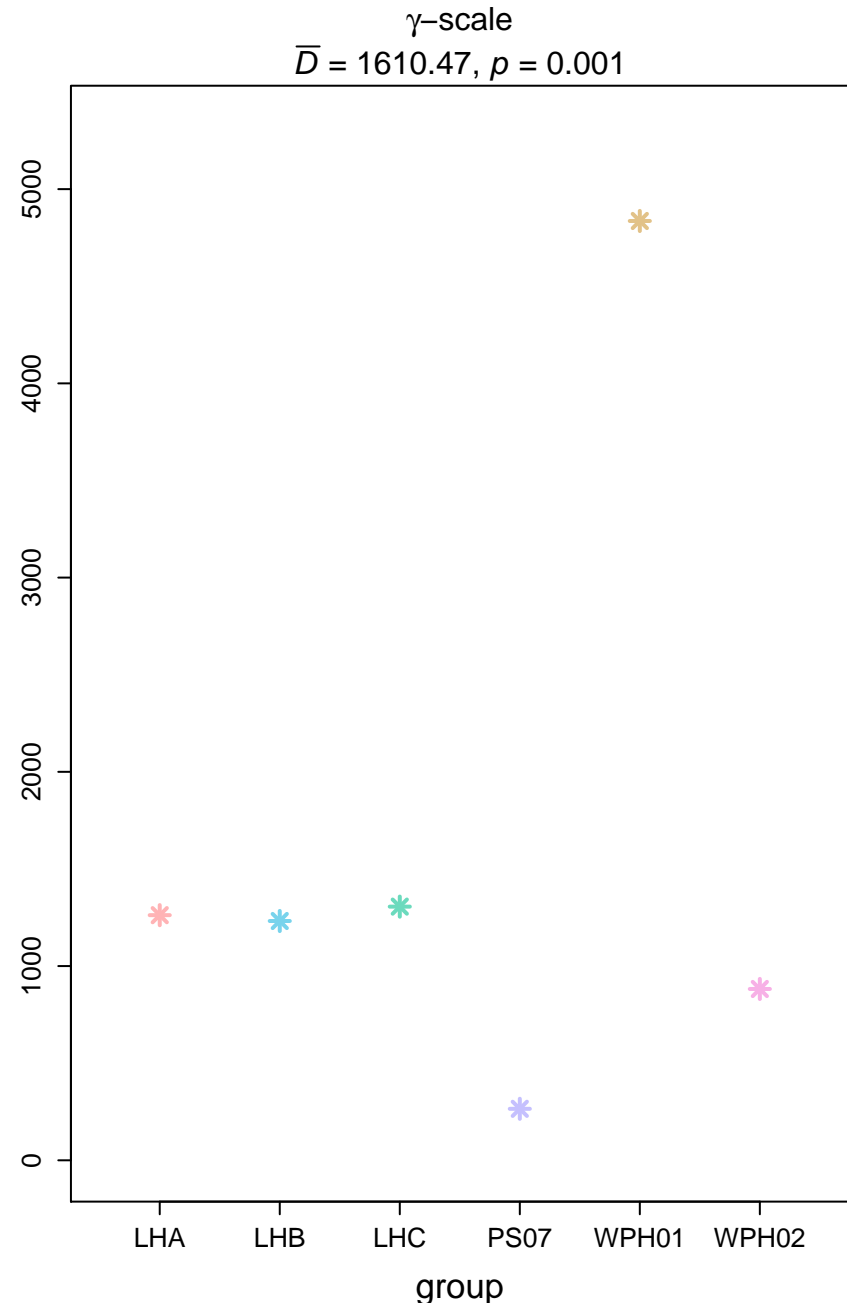
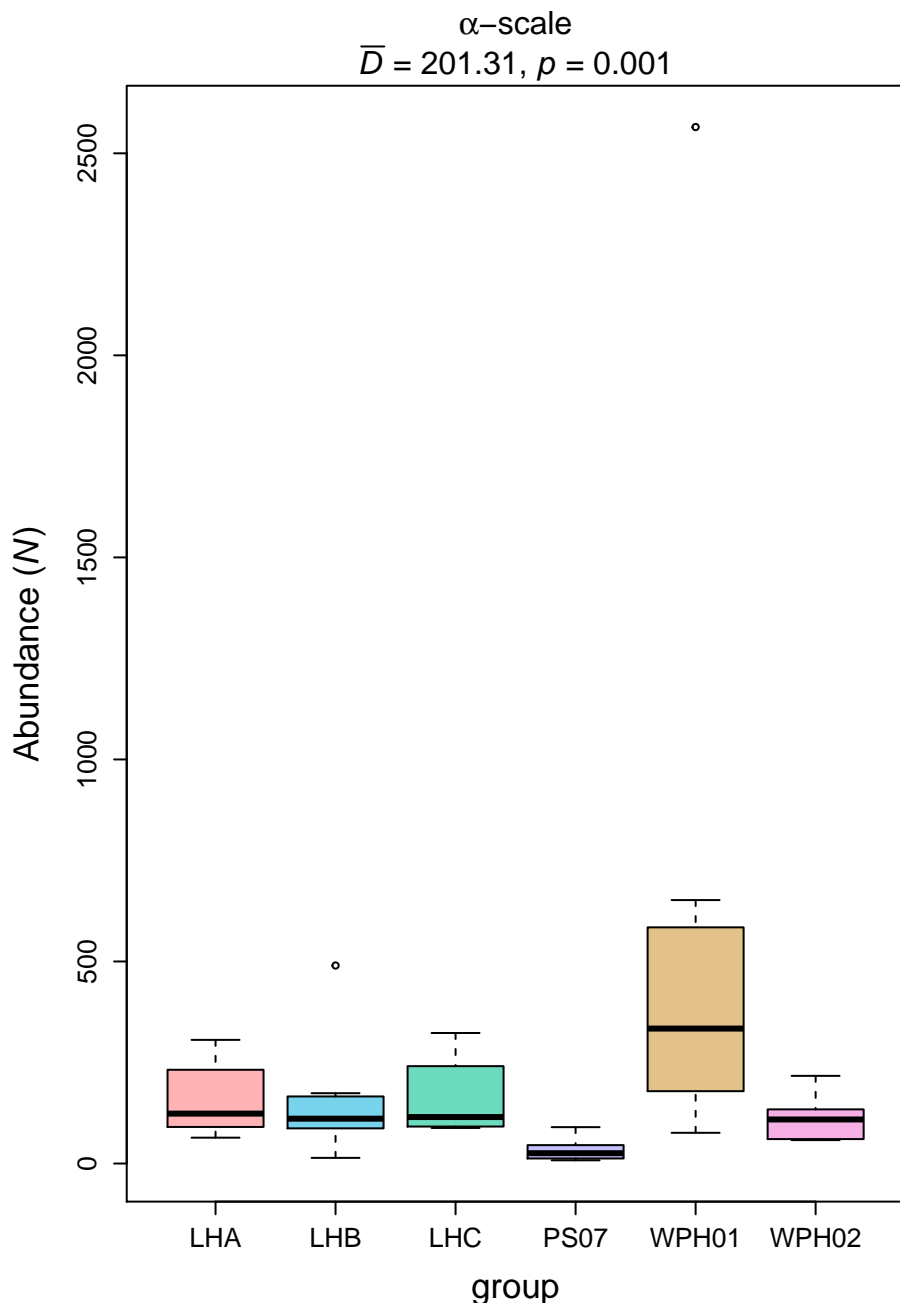
Group Scale

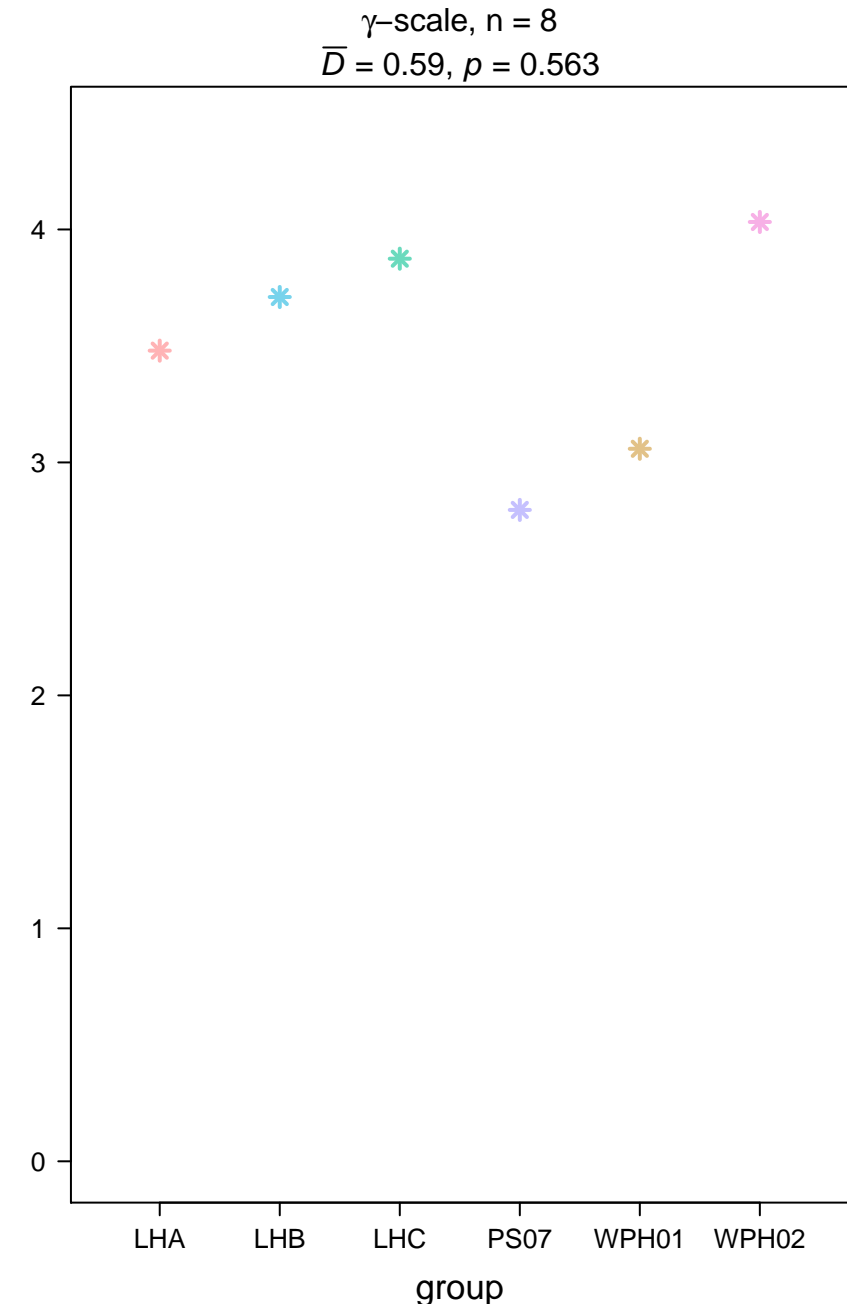
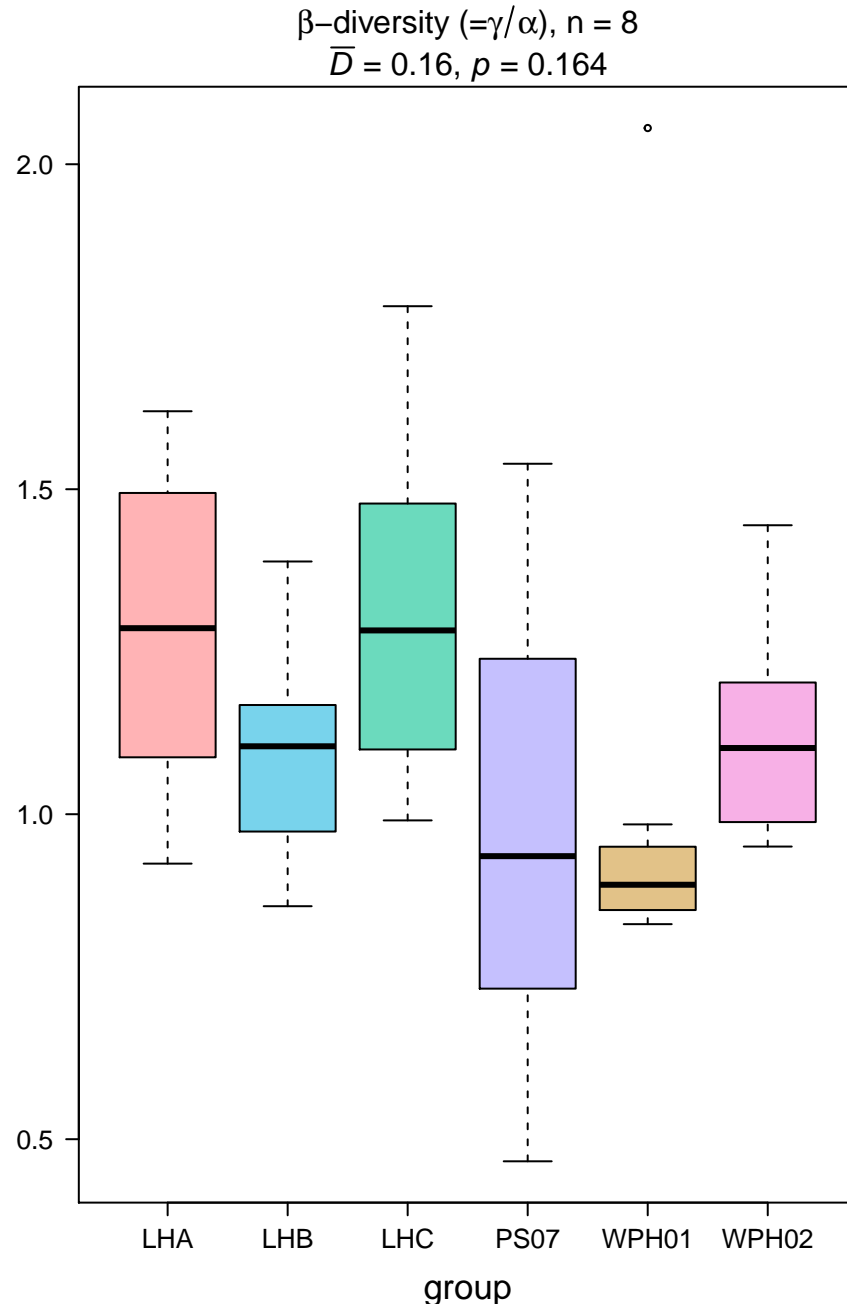
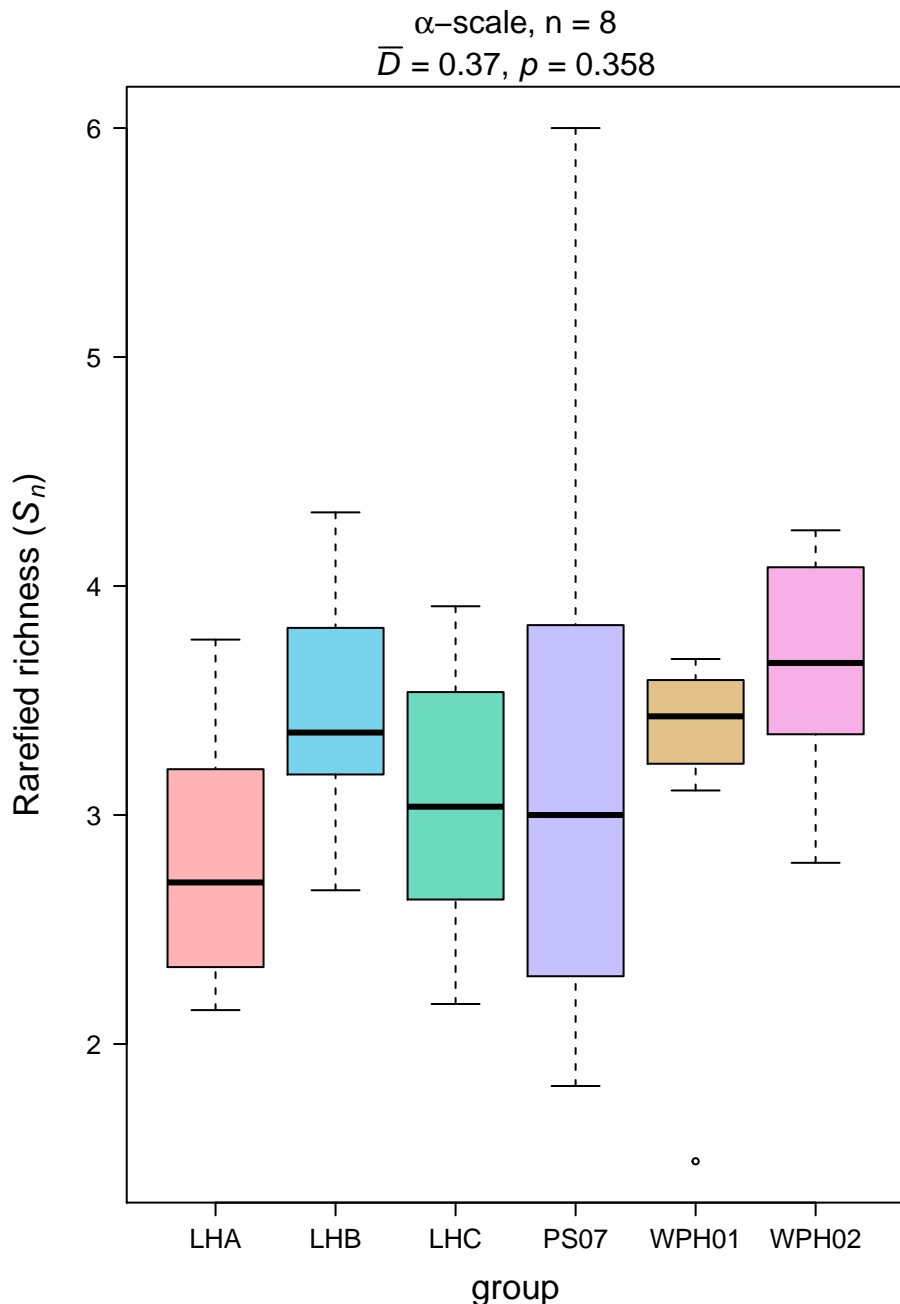


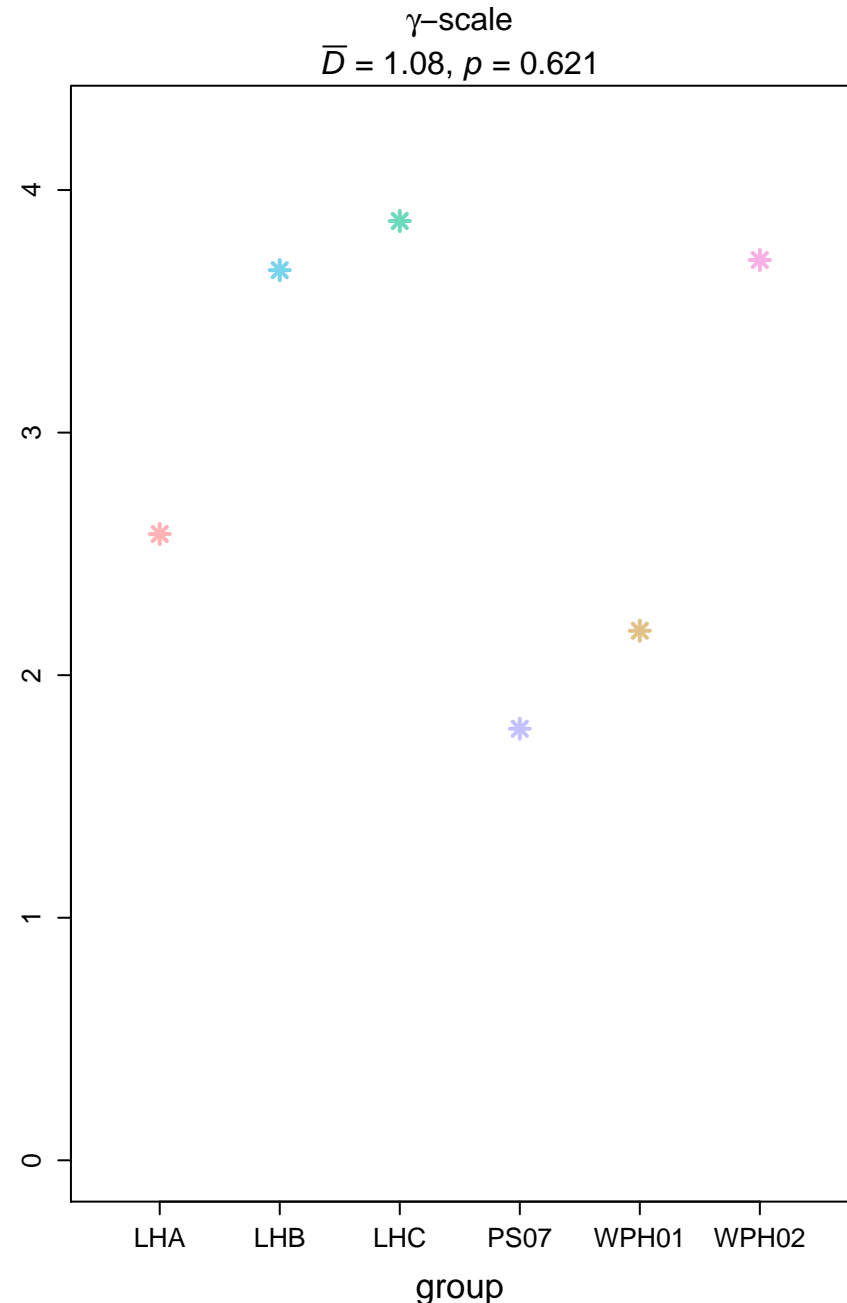
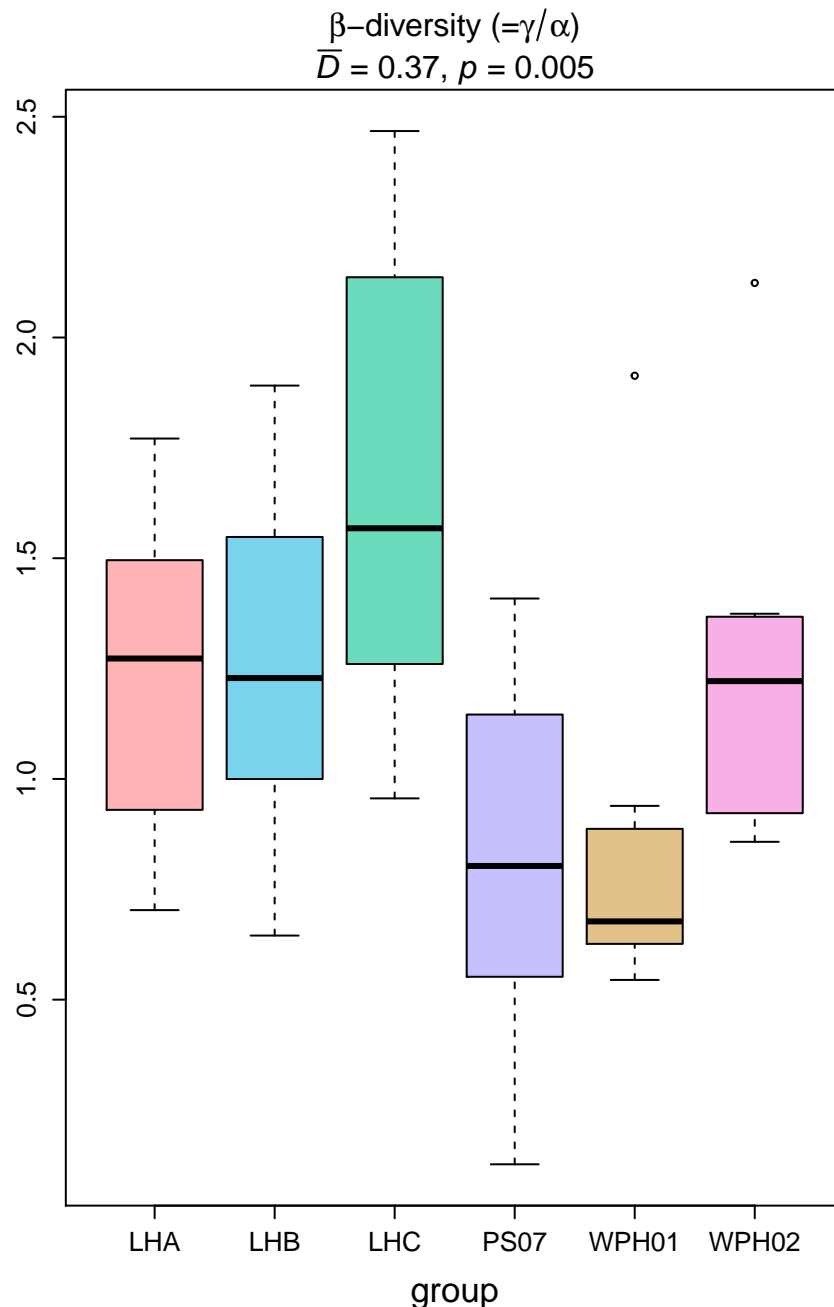
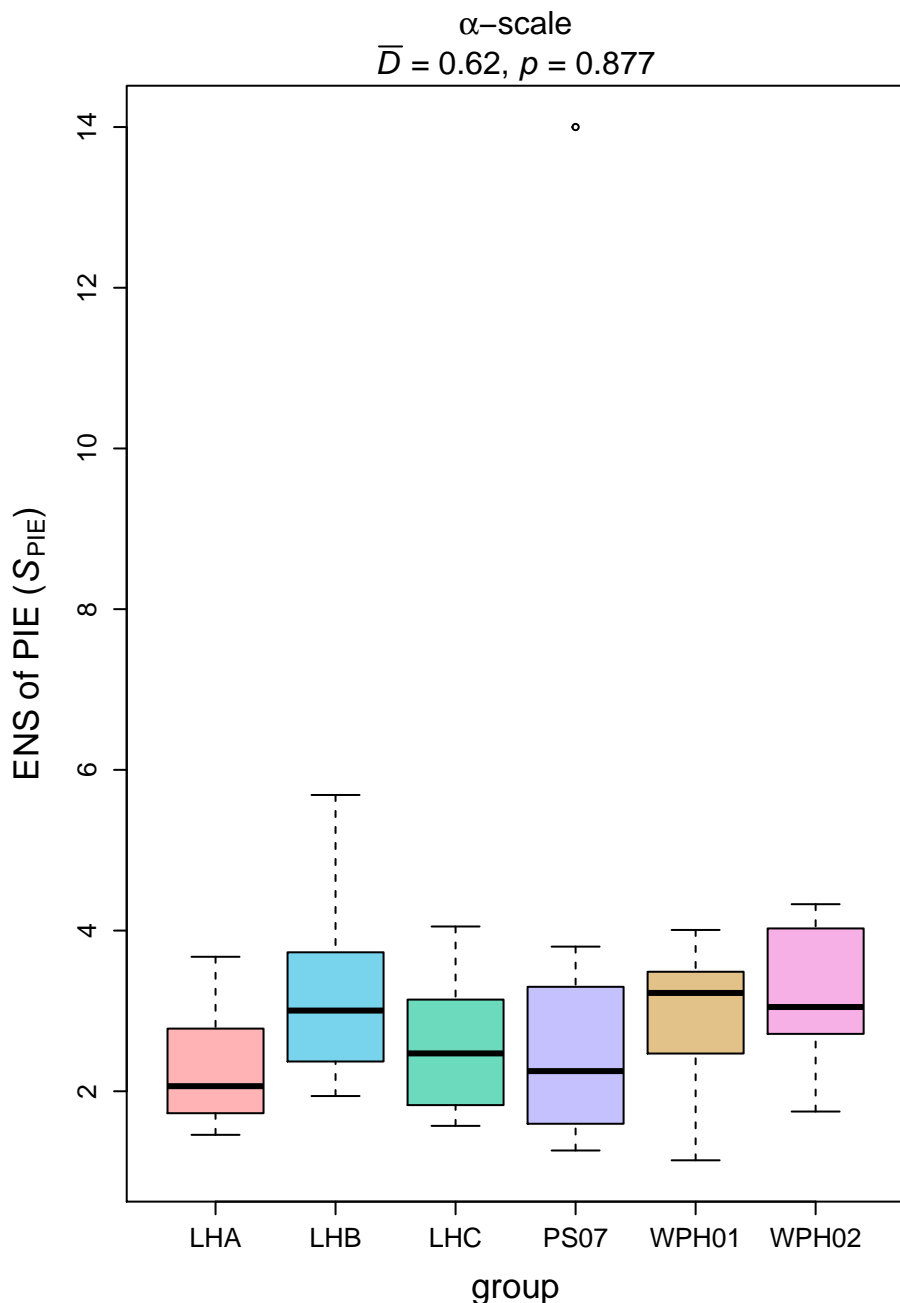
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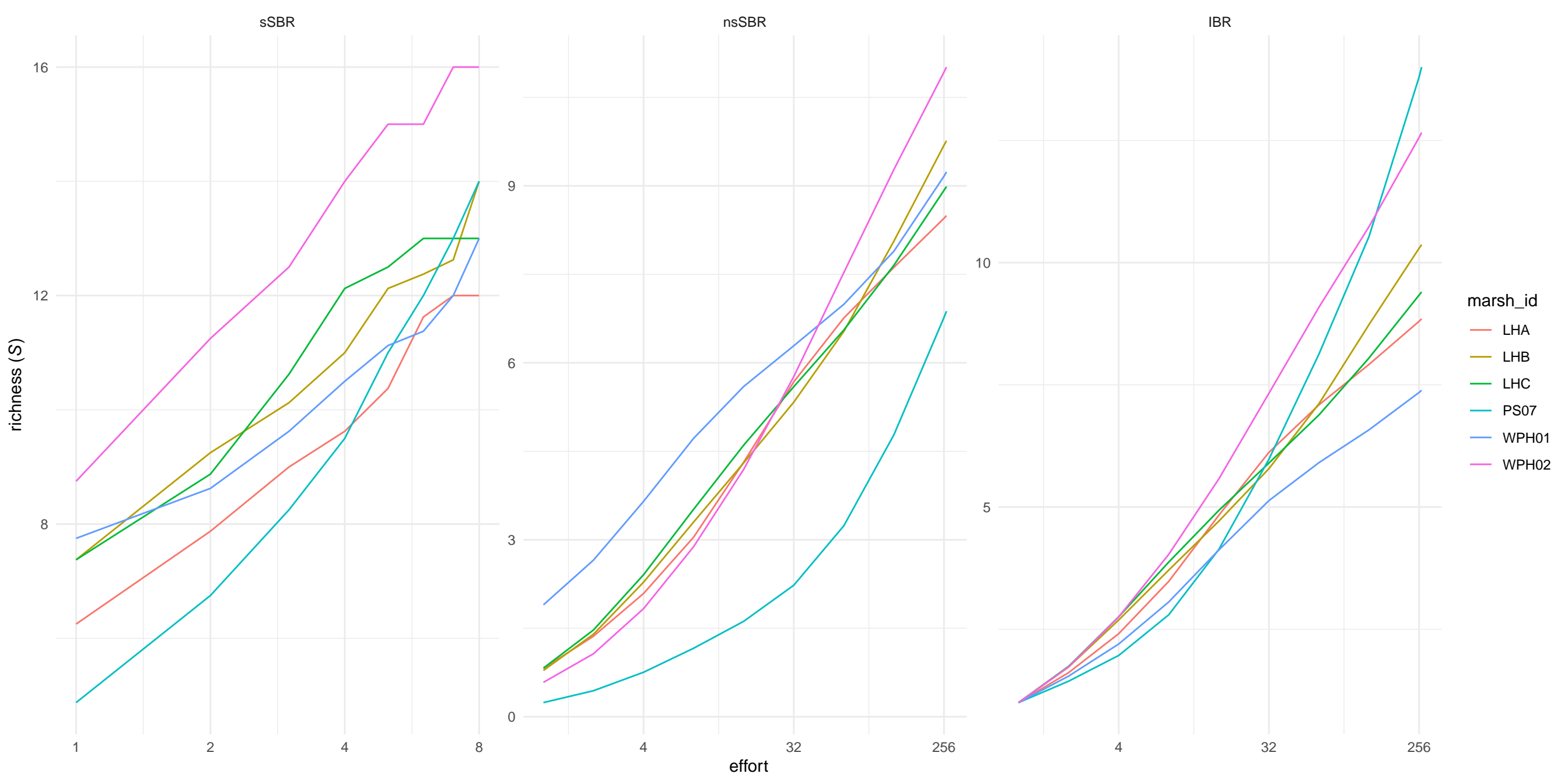


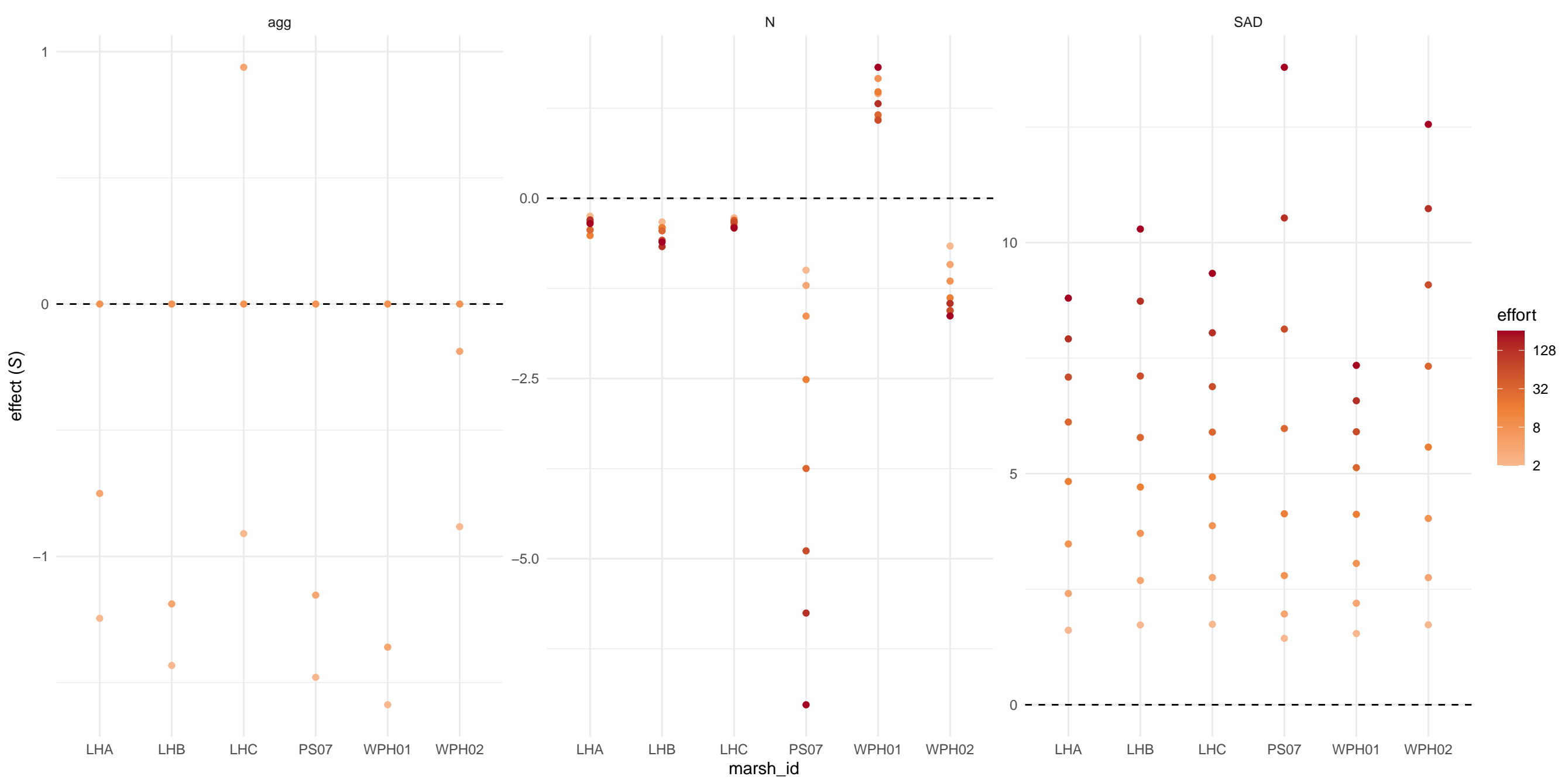


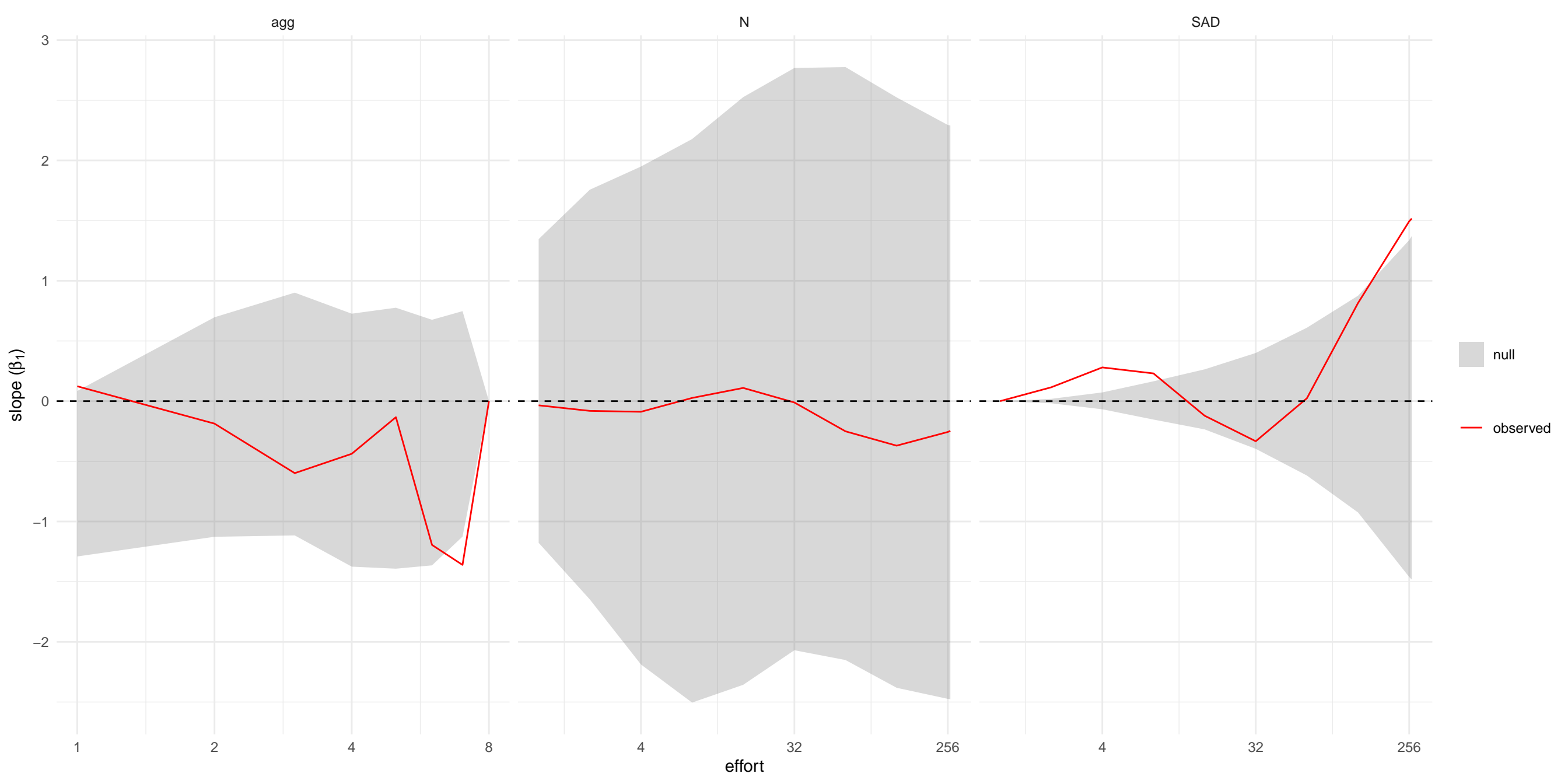




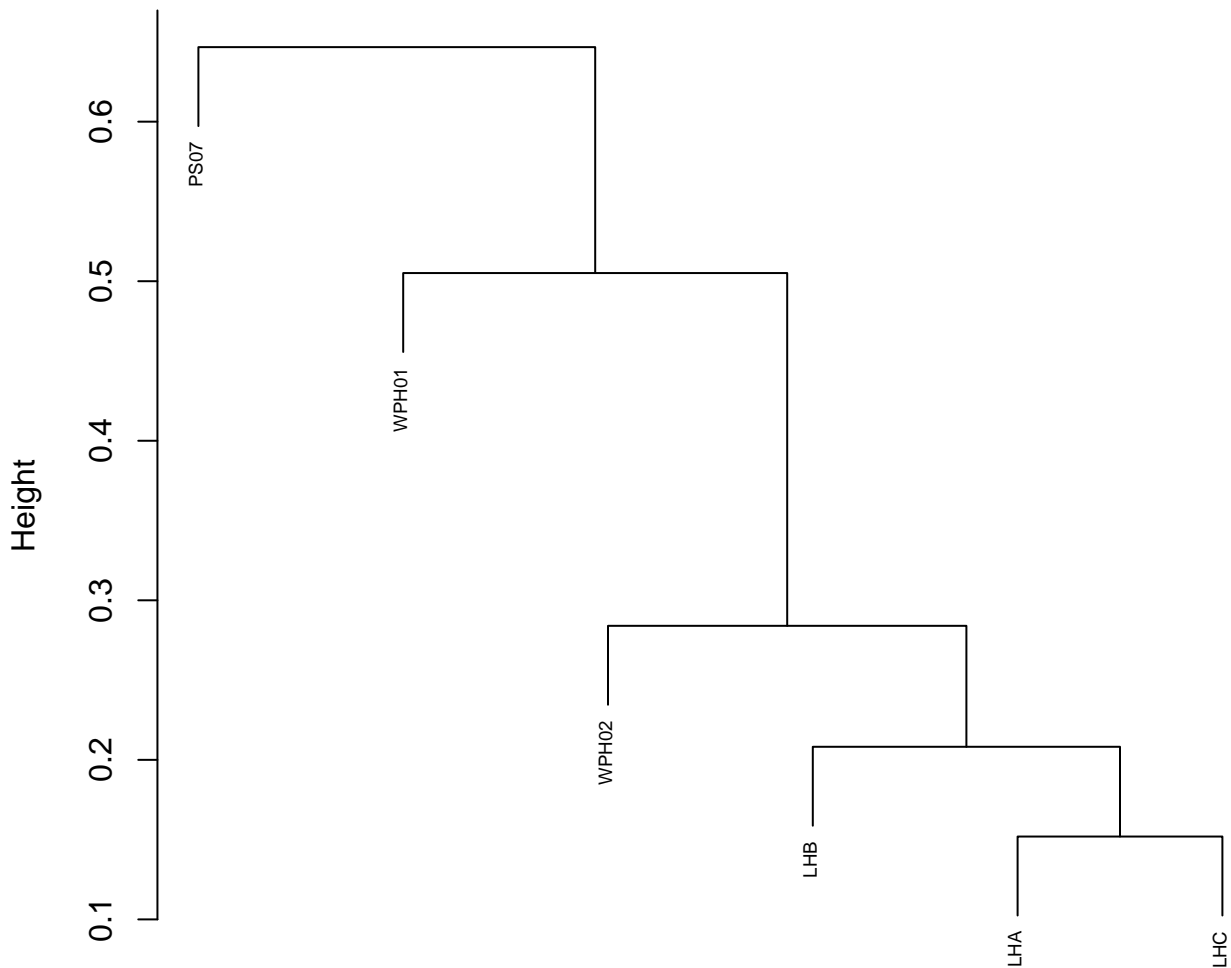








Cluster Dendrogram



	pairs	R2	p.value	p.adjusted
1	LHA vs LHB	0.08182420	0.226773227	1.00000000
2	LHA vs LHC	0.05933934	0.495504496	1.00000000
3	LHA vs PS07	0.31579191	0.001998002	0.02997003
4	LHA vs WPH01	0.22352236	0.005994006	0.08991009
5	LHA vs WPH02	0.19274518	0.009990010	0.14985015
6	LHB vs LHC	0.06120527	0.491508492	1.00000000
7	LHB vs PS07	0.22193295	0.008991009	0.13486513
8	LHB vs WPH01	0.15523083	0.019980020	0.29970030
9	LHB vs WPH02	0.11039332	0.083916084	1.00000000
10	LHC vs PS07	0.33211747	0.000999001	0.01498501
11	LHC vs WPH01	0.18456885	0.029970030	0.44955045
12	LHC vs WPH02	0.16195992	0.045954046	0.68931069
13	PS07 vs WPH01	0.32057027	0.001998002	0.02997003
14	PS07 vs WPH02	0.29778415	0.002997003	0.04495504
15	WPH01 vs WPH02	0.29798847	0.002997003	0.04495504

bray_cent_dist[[i]]

hclust (*, "ward.D")