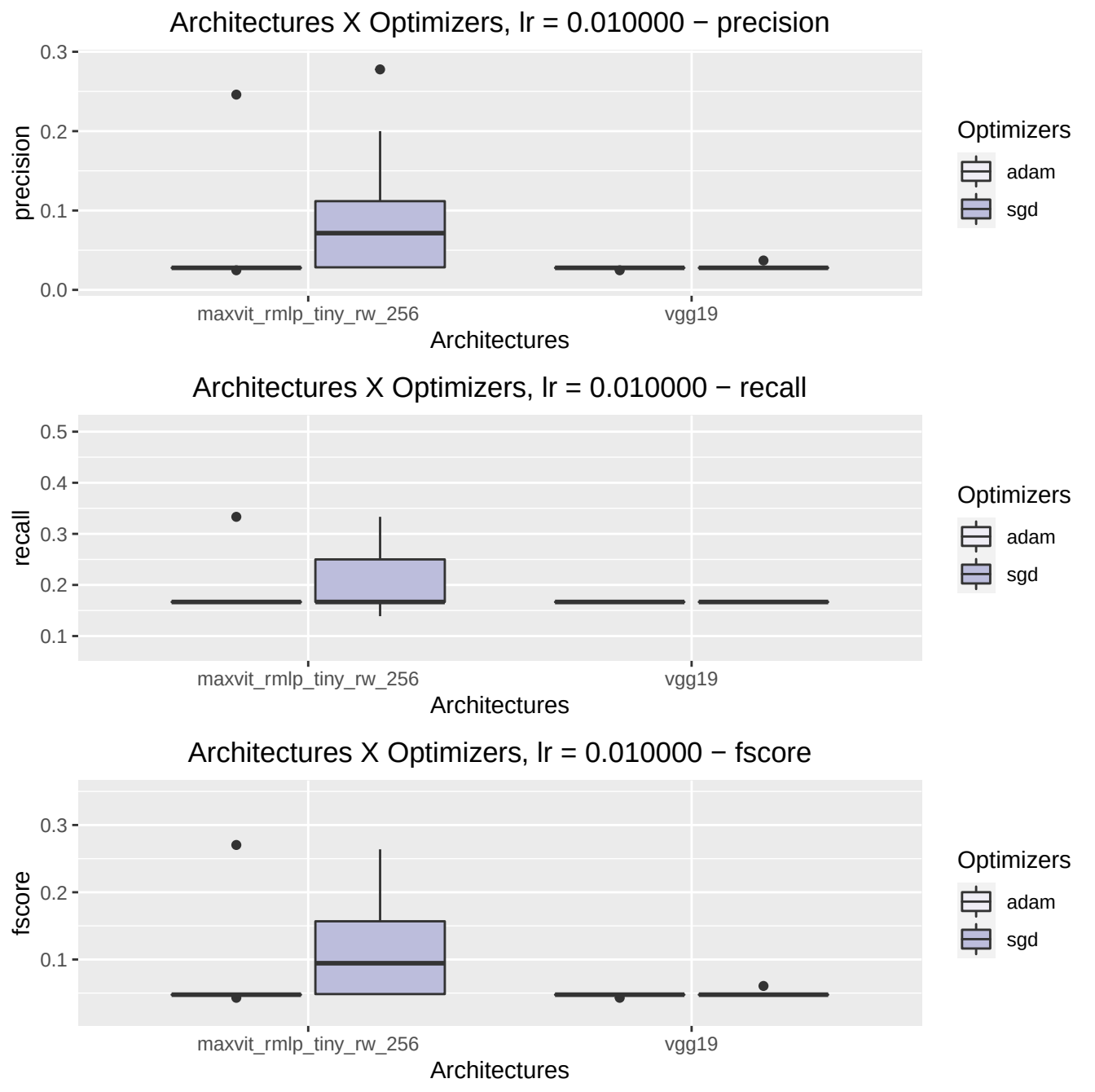
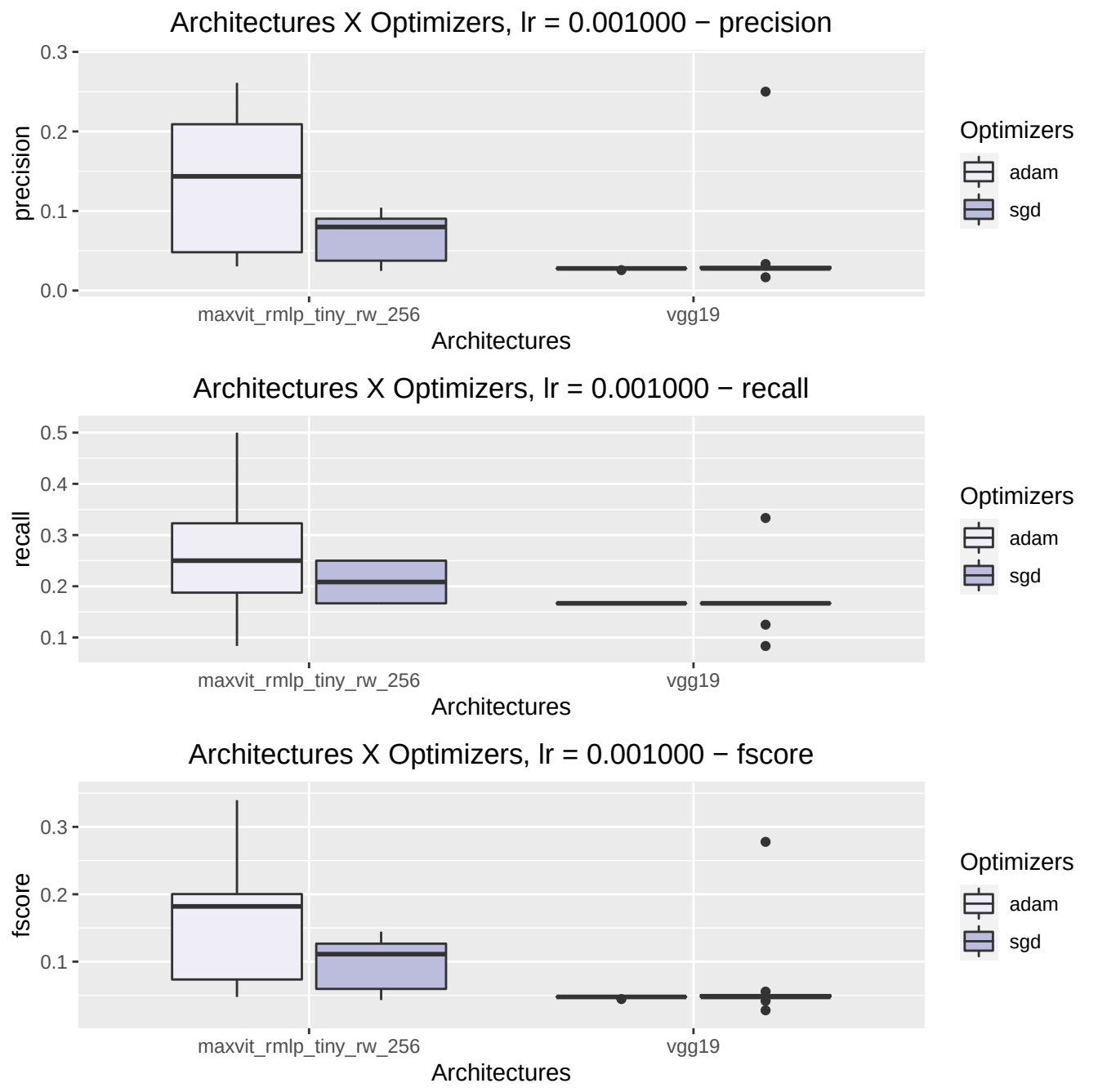
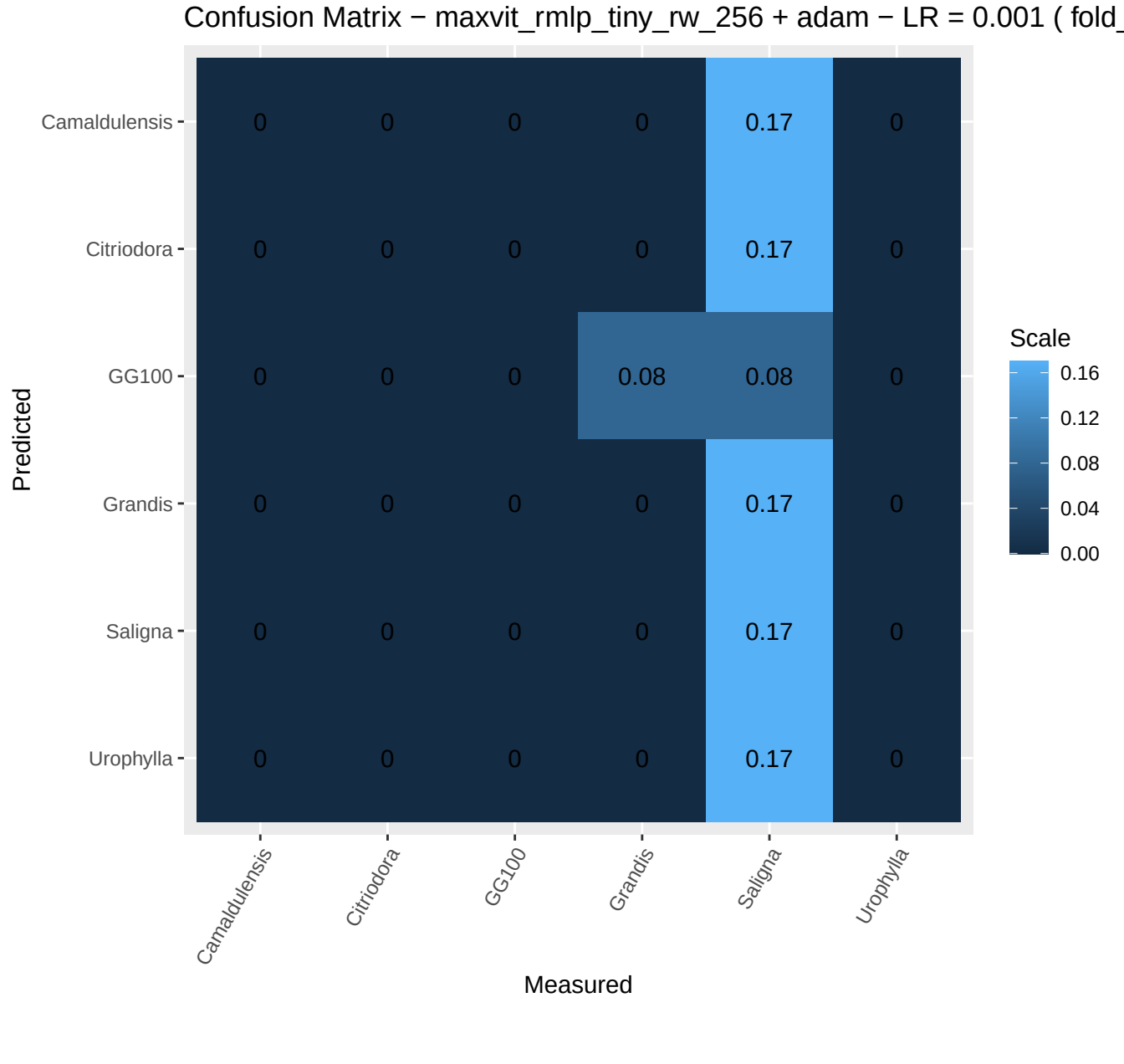
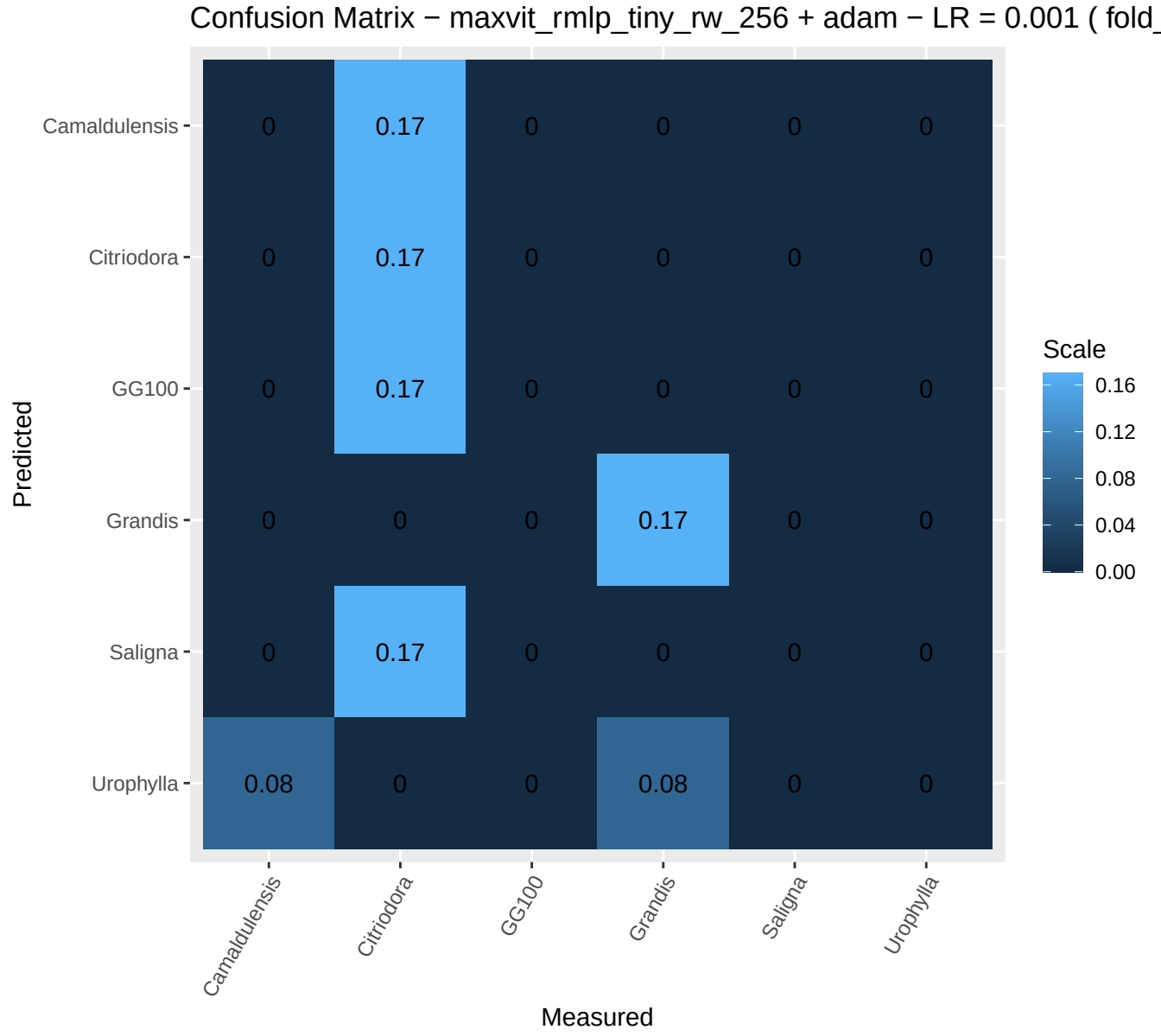
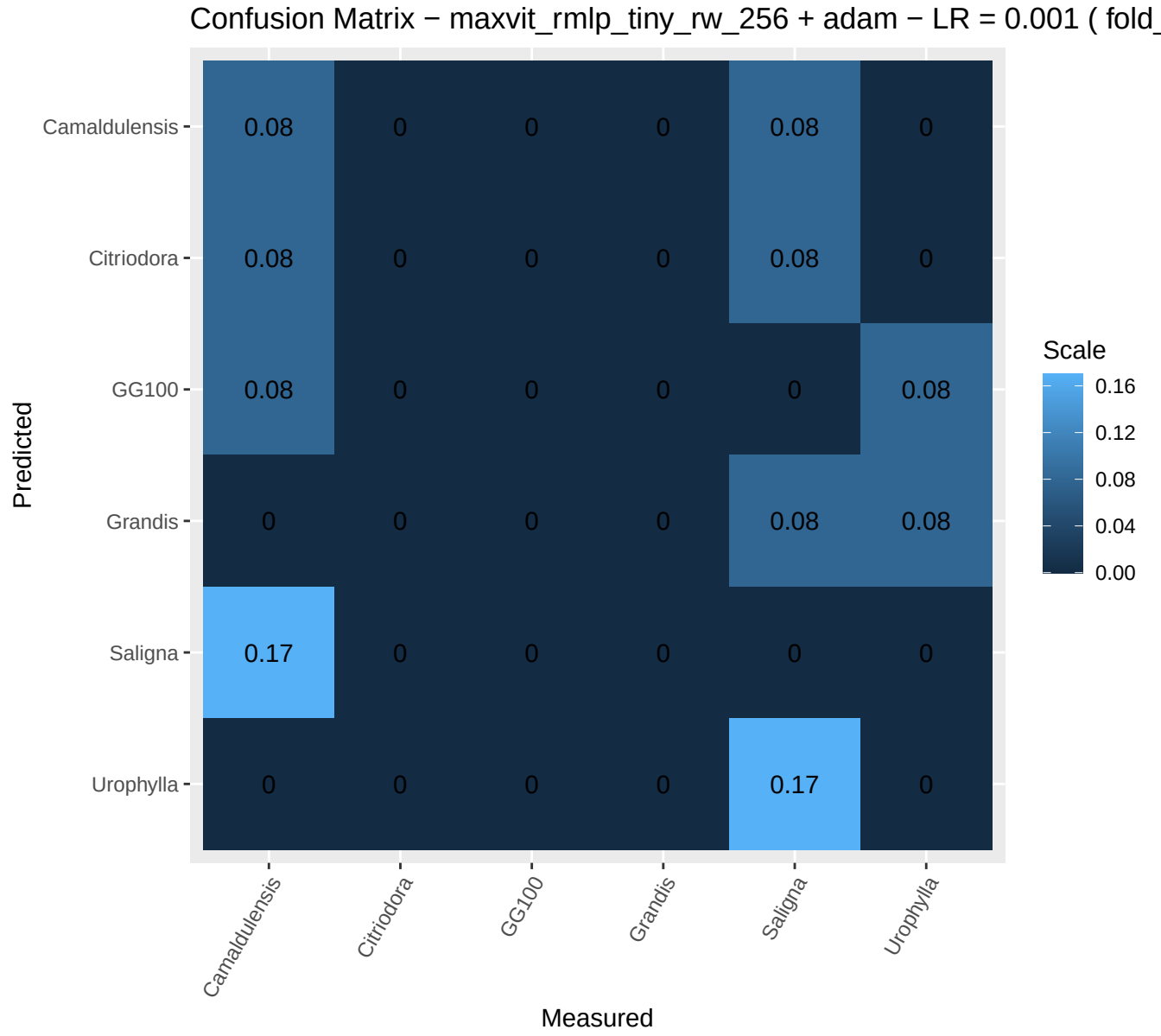
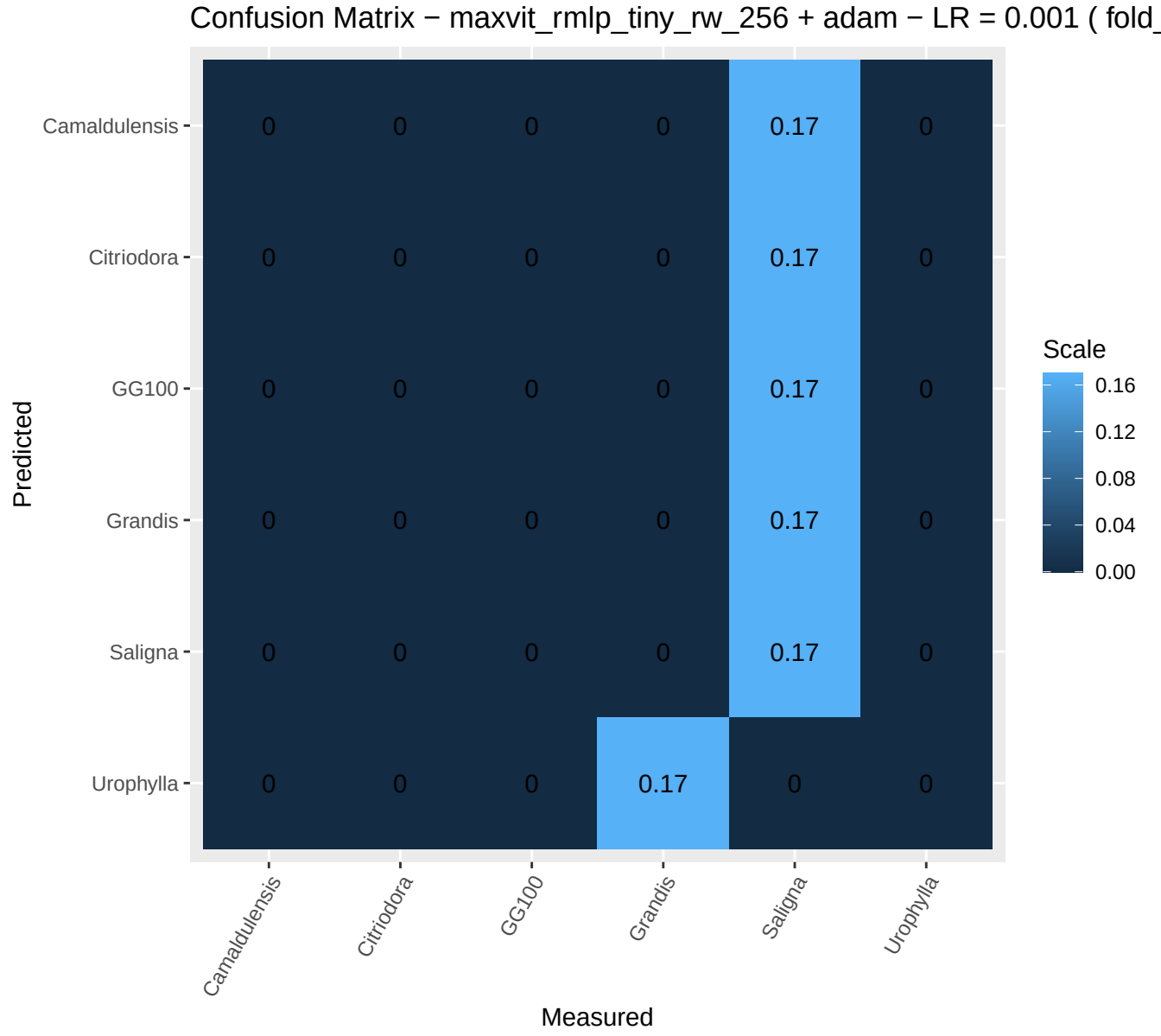

Architectures X Optimizers, lr = 0.010000 − precision
precision
0.3
0.2
0.1
0.0
maxvit_rmlp_tiny_rw_256
vgg19
Architectures
Optimizers
adam
sgd
Architectures X Optimizers, lr = 0.010000 − recall
recall
0.5
0.4
0.3
0.2
0.1
maxvit_rmlp_tiny_rw_256
vgg19
Architectures
Optimizers
adam
sgd
Architectures X Optimizers, lr = 0.010000 − fscore
fscore
0.3
0.2
0.1
maxvit_rmlp_tiny_rw_256
vgg19
Architectures
Optimizers
adam
sgd

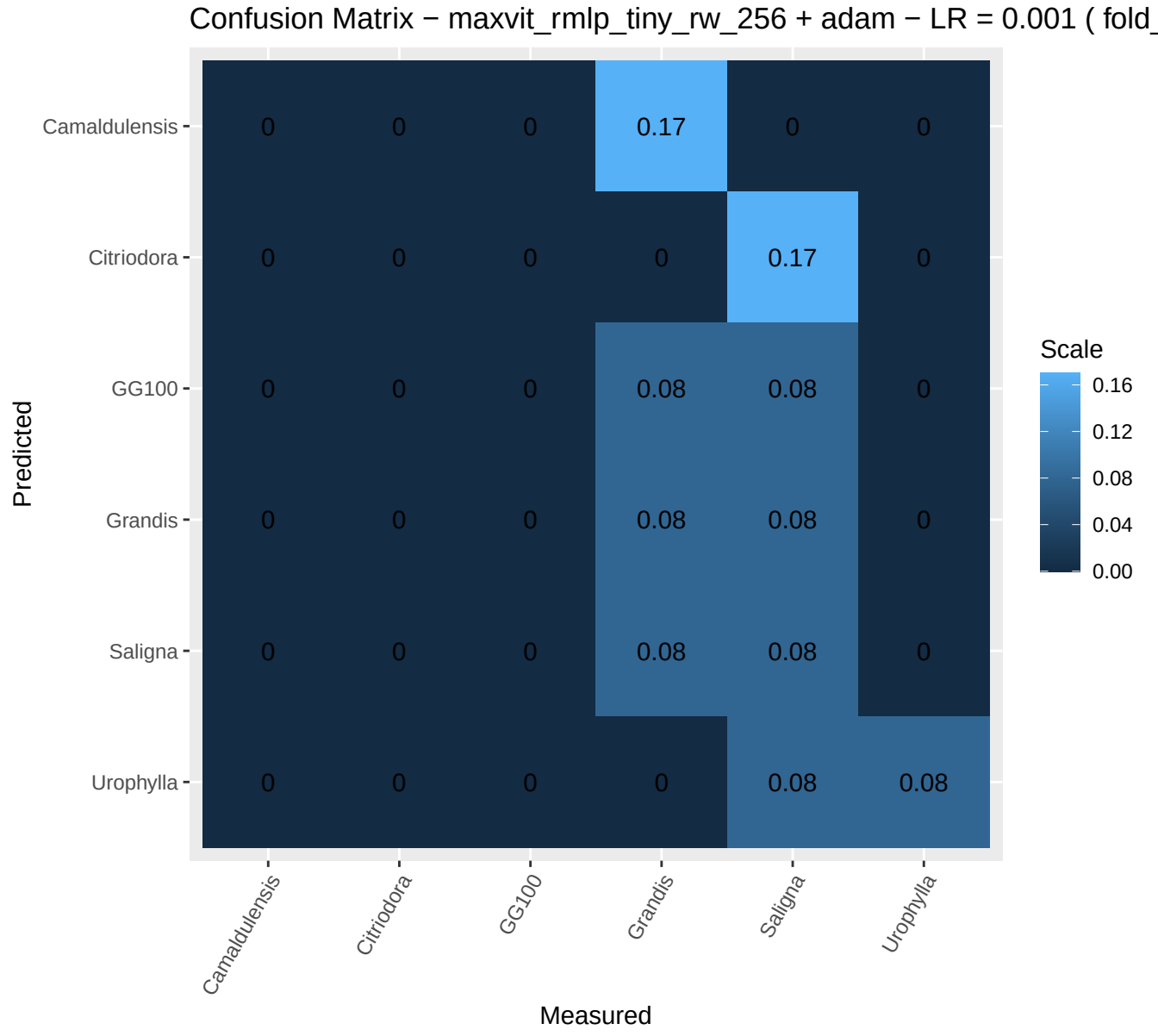

Architectures X Optimizers, lr = 0.001000 − precision
precision
0.3
0.2
0.1
0.0
maxvit_rmlp_tiny_rw_256
vgg19
Architectures
Optimizers
adam
sgd
Architectures X Optimizers, lr = 0.001000 − recall
recall
0.5
0.4
0.3
0.2
0.1
maxvit_rmlp_tiny_rw_256
vgg19
Architectures
Optimizers
adam
sgd
Architectures X Optimizers, lr = 0.001000 − fscore
fscore
0.3
0.2
0.1
maxvit_rmlp_tiny_rw_256
vgg19
Architectures
Optimizers
adam
sgd

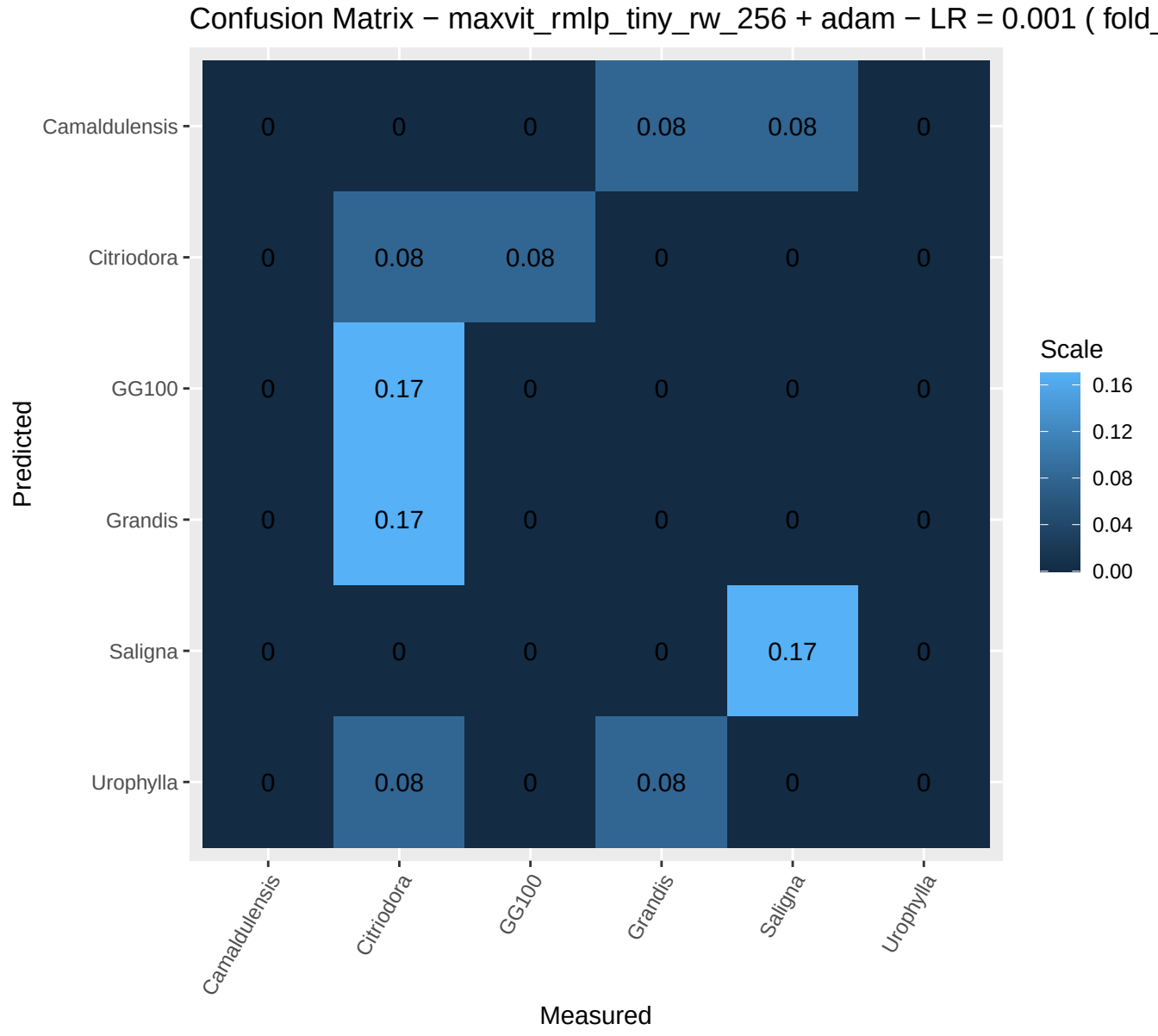

Confusion Matrix − maxvit_rmlp_tiny_rw_256 + adam − LR = 0.001 (fold_
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Predicted
Measured
Scale
0.16
0.12
0.08
0.04
0.00
0 0 0 0 0.17 0
0 0 0 0 0.17 0
0 0 0 0.08 0.08 0
0 0 0 0 0.17 0
0 0 0 0 0.17 0
0 0 0 0 0.17 0

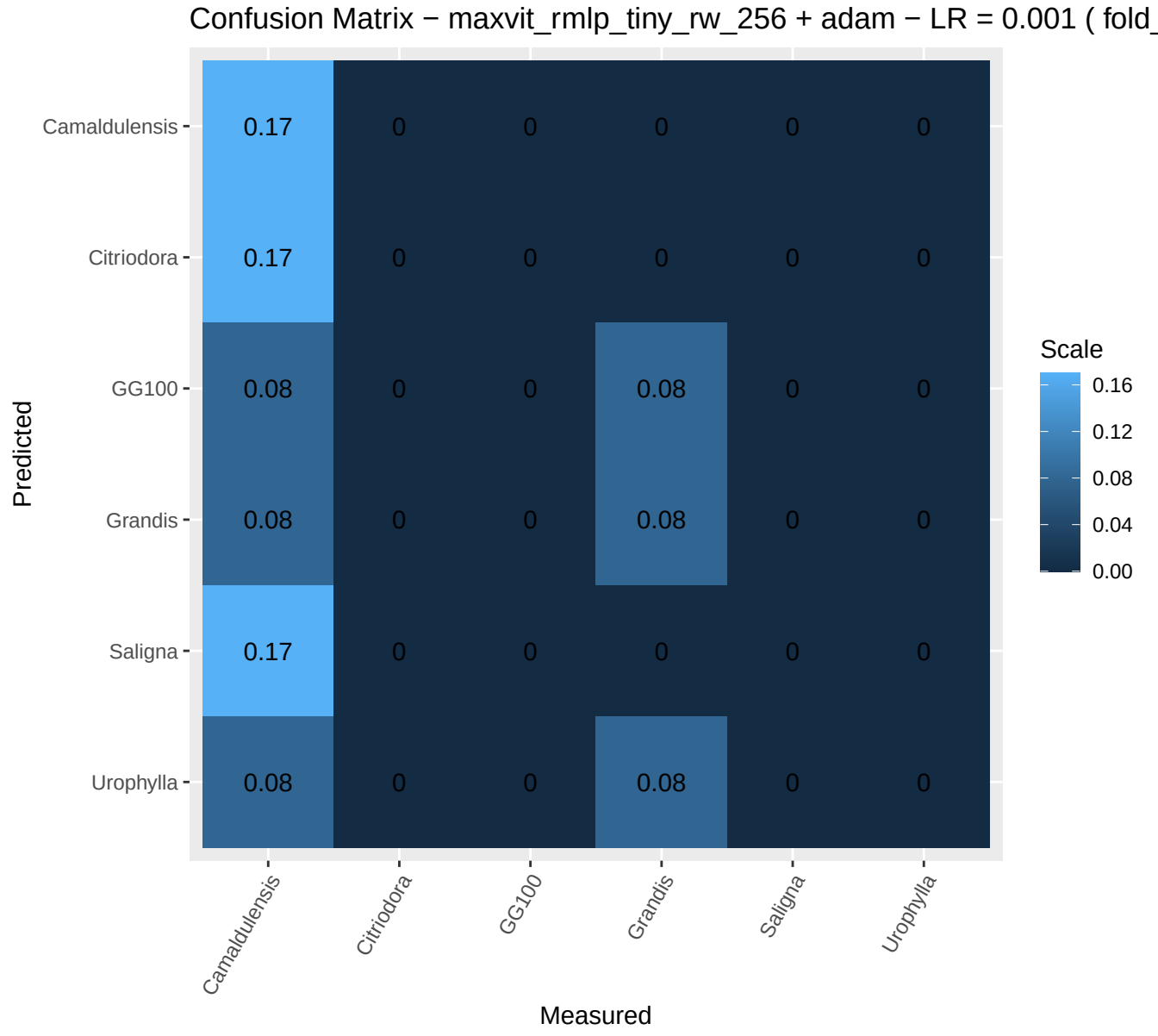

Confusion Matrix − maxvit_rmlp_tiny_rw_256 + adam − LR = 0.001 (fold_
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Predicted
Measured
Scale
0.16
0.12
0.08
0.04
0.00
0 0.17 0 0 0 0
0 0.17 0 0 0 0
0 0.17 0 0 0 0
0 0 0 0.17 0 0
0 0.17 0 0 0 0
0.08 0 0 0.08 0 0

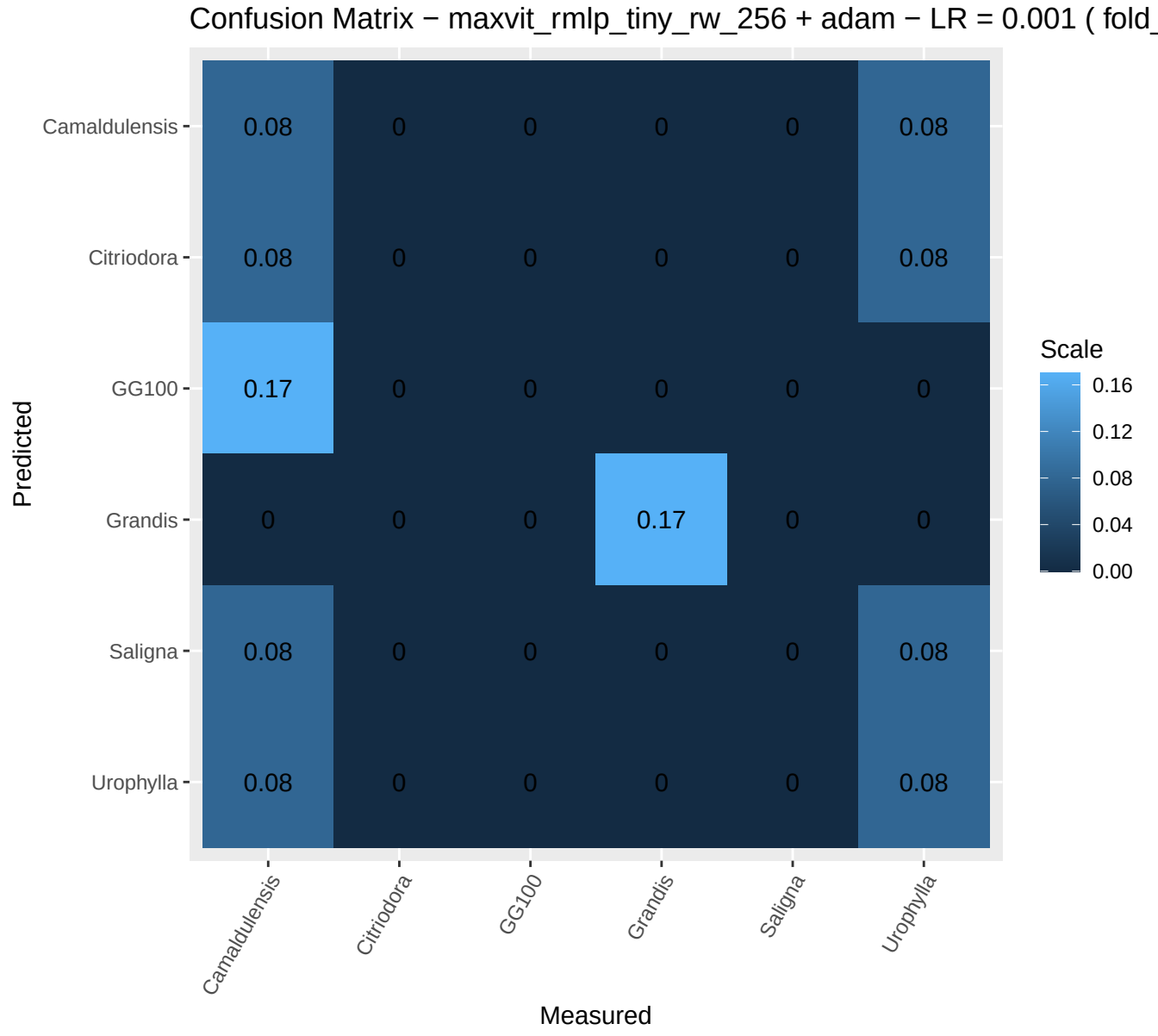

Confusion Matrix − maxvit_rmlp_tiny_rw_256 + adam − LR = 0.001 (fold_
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Predicted
0.08 0 0 0 0.08 0
0.08 0 0 0 0.08 0
0.08 0 0 0 0 0.08
0 0 0 0 0.08 0.08
0.17 0 0 0 0 0
0 0 0 0 0.17 0
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Measured
Scale
0.16
0.12
0.08
0.04
0.00

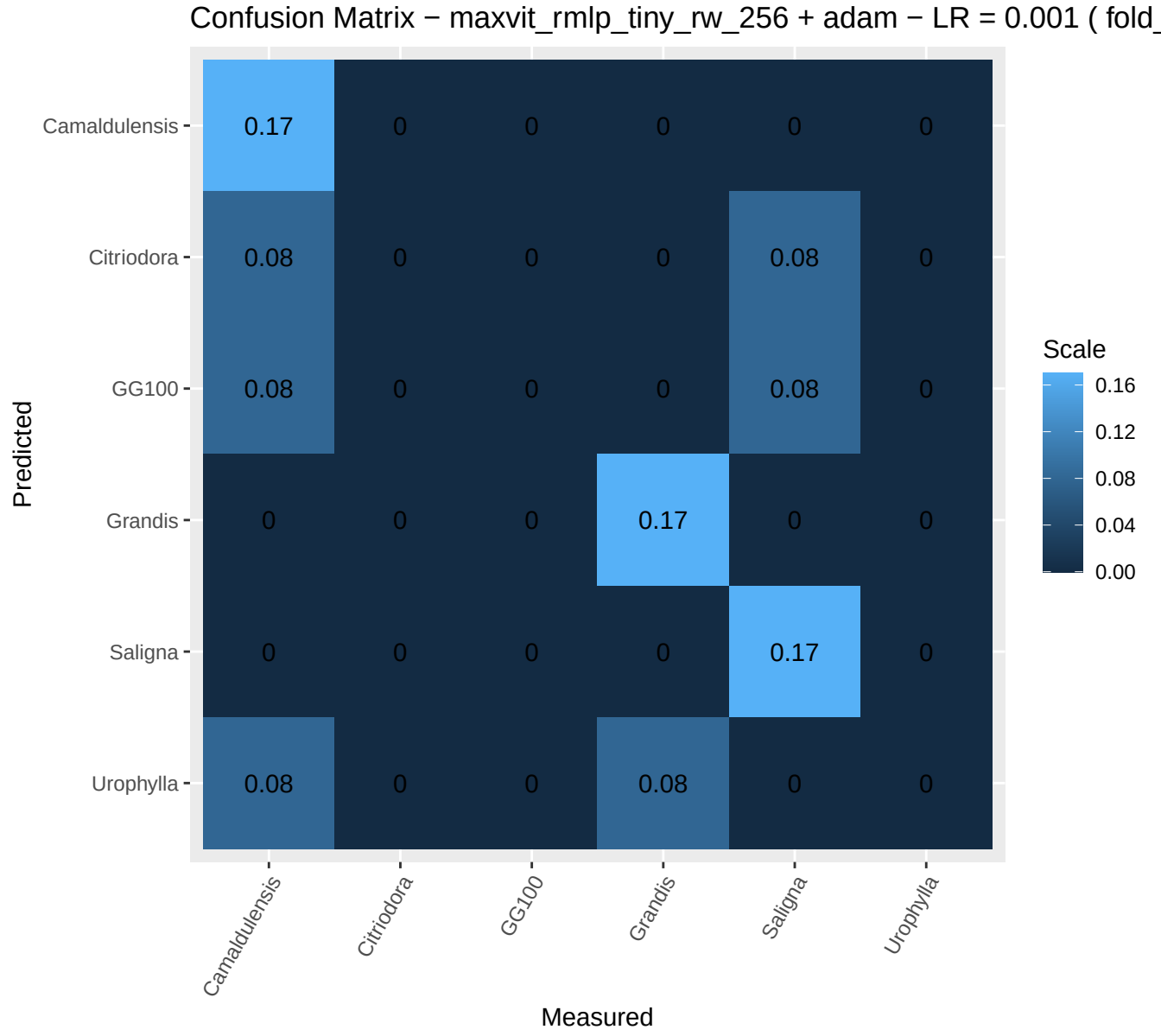

Confusion Matrix − maxvit_rmlp_tiny_rw_256 + adam − LR = 0.001 (fold_
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Predicted
Measured
Scale
0.16
0.12
0.08
0.04
0.00

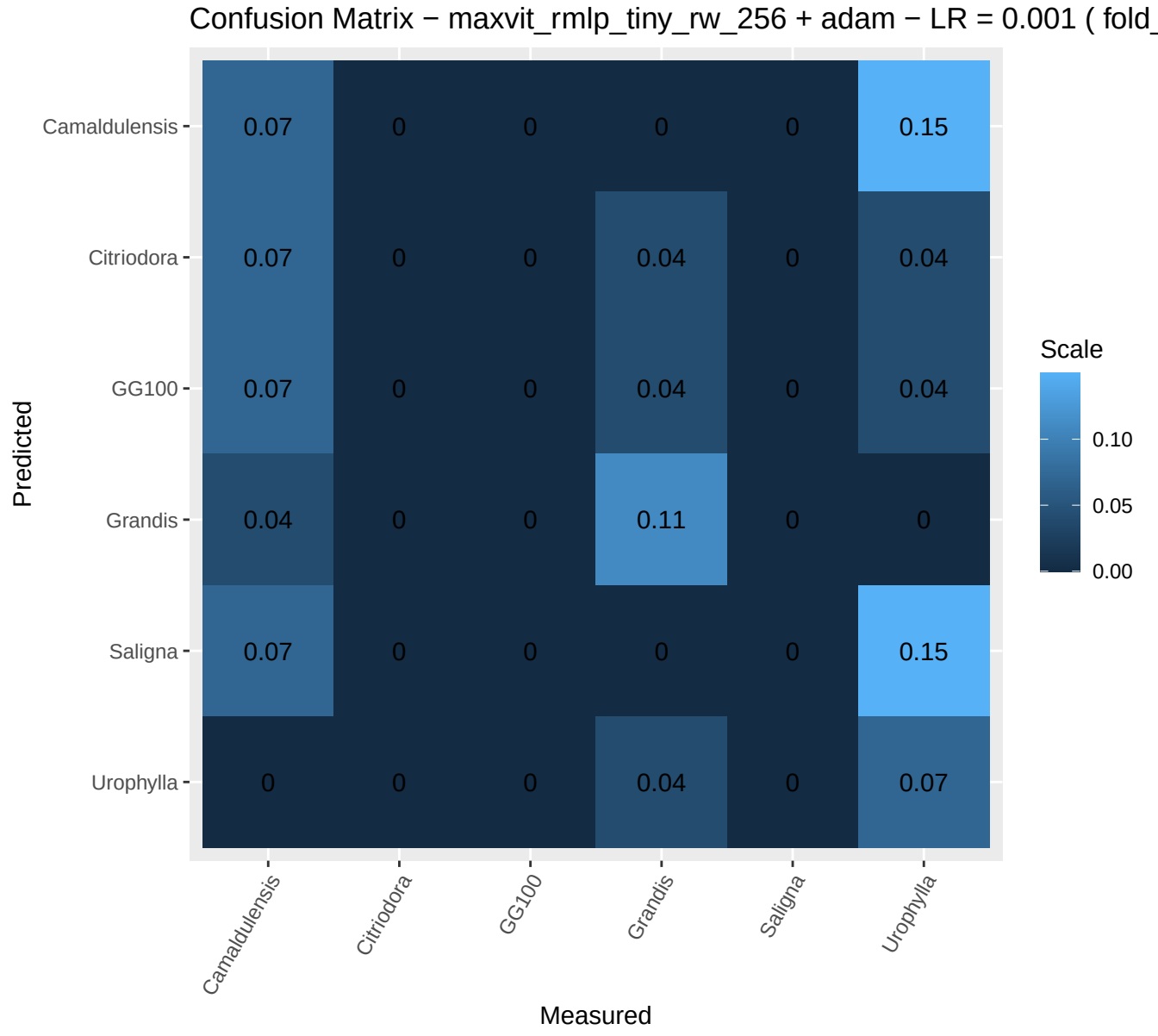

Confusion Matrix − maxvit_rmlp_tiny_rw_256 + adam − LR = 0.001 (fold_
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Predicted
Measured
0 0 0 0.17 0 0
0 0 0 0 0.17 0
0 0 0 0.08 0.08 0
0 0 0 0.08 0.08 0
0 0 0 0.08 0.08 0
0 0 0 0 0.08 0.08
Scale
0.16
0.12
0.08
0.04
0.00

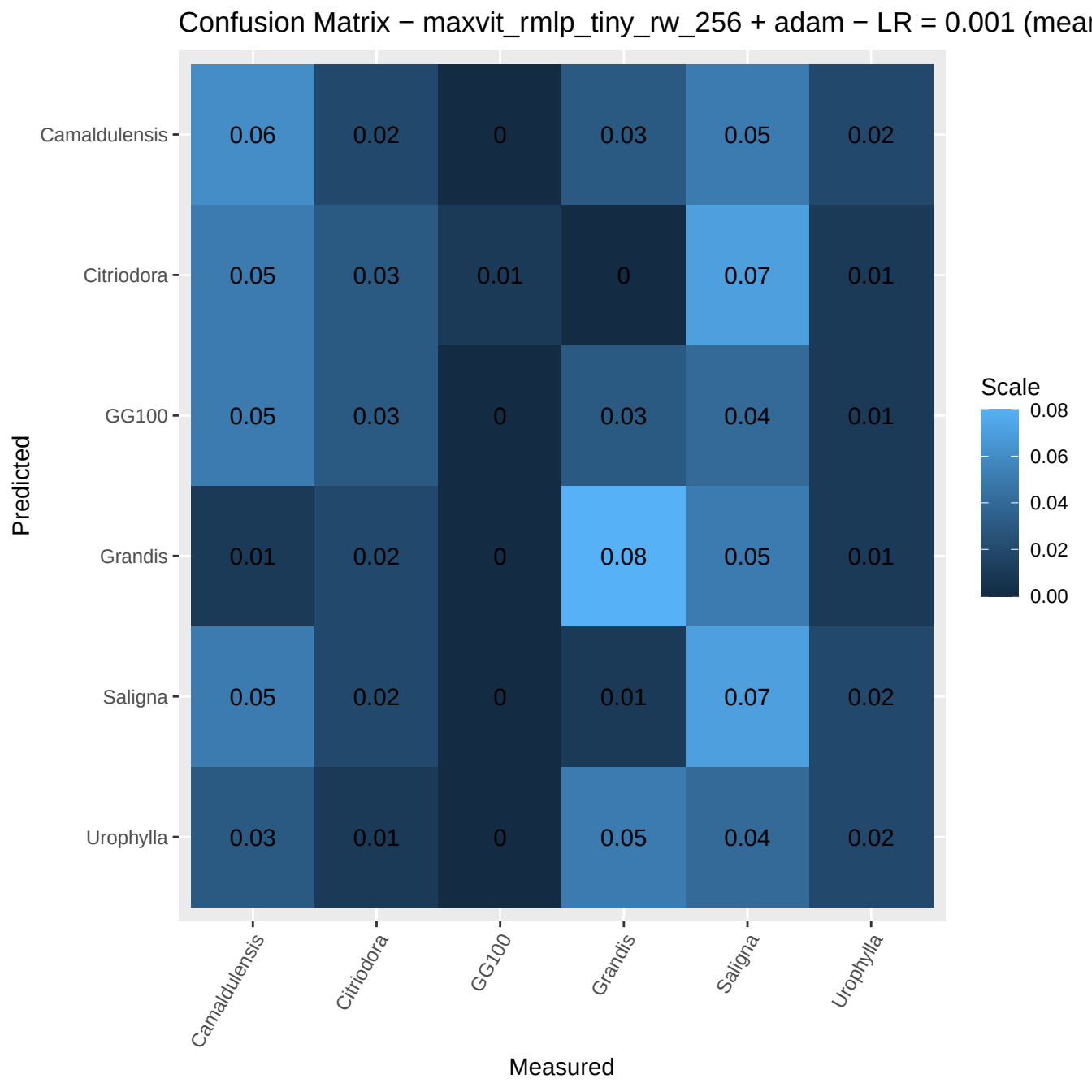

Confusion Matrix − maxvit_rmlp_tiny_rw_256 + adam − LR = 0.001 (fold_
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Predicted
Measured
Scale
0.16
0.12
0.08
0.04
0.00
0 0 0 0.08 0.08 0
0 0.08 0.08 0 0 0
0 0.17 0 0 0 0
0 0.17 0 0 0 0
0 0 0 0 0.17 0
0 0.08 0 0.08 0 0


Confusion Matrix − maxvit_rmlp_tiny_rw_256 + adam − LR = 0.001 (fold_
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Predicted
Measured
0.17
0.08
0
Scale
0.16
0.12
0.08
0.04
0.00


Confusion Matrix − maxvit_rmlp_tiny_rw_256 + adam − LR = 0.001 (fold_
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Predicted
Measured
Scale
0.16
0.12
0.08
0.04
0.00


Confusion Matrix − maxvit_rmlp_tiny_rw_256 + adam − LR = 0.001 (fold_
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Predicted
Measured
Scale
0.16
0.12
0.08
0.04
0.00
0.17 0 0 0 0 0
0.08 0 0 0 0.08 0
0.08 0 0 0 0.08 0
0 0 0 0.17 0 0
0 0 0 0 0.17 0
0.08 0 0 0.08 0 0


Confusion Matrix − maxvit_rmlp_tiny_rw_256 + adam − LR = 0.001 (fold_
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Predicted
0.07 0 0 0 0 0.15
0.07 0 0 0.04 0 0.04
0.07 0 0 0.04 0 0.04
0.04 0 0 0.11 0 0
0.07 0 0 0 0 0.15
0 0 0 0.04 0 0.07
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Measured
Scale
0.10
0.05
0.00


Confusion Matrix − maxvit_rmlp_tiny_rw_256 + adam − LR = 0.001 (mean
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Predicted
0.06 0.02 0 0.03 0.05 0.02
0.05 0.03 0.01 0 0.07 0.01
0.05 0.03 0 0.03 0.04 0.01
0.01 0.02 0 0.08 0.05 0.01
0.05 0.02 0 0.01 0.07 0.02
0.03 0.01 0 0.05 0.04 0.02
Camaldulensis
Citriodora
GG100
Grandis
Saligna
Urophylla
Measured
Scale
0.08
0.06
0.04
0.02
0.00