# Supplemental Appendix in support of PAPER TITLE

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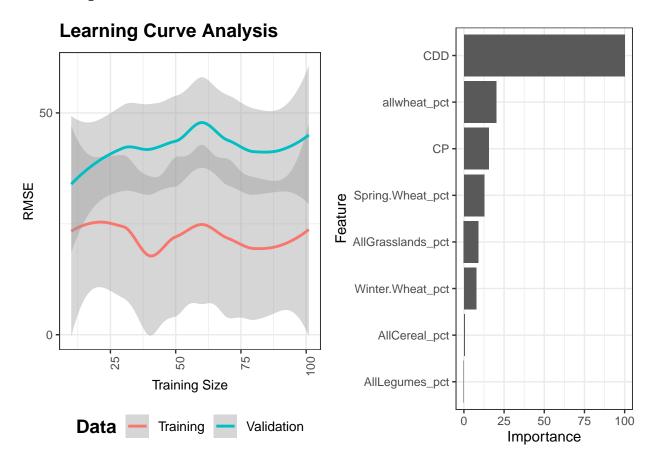
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#### **Total Aphids Model**



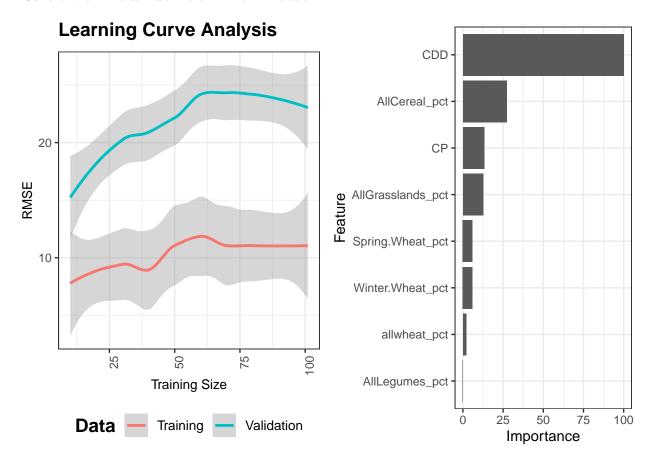
```
## Random Forest
##
## 101 samples
##
     8 predictors
##
## No pre-processing
## Resampling: Cross-Validated (10 fold, repeated 1 times)
## Summary of sample sizes: 89, 90, 92, 91, 91, 90, ...
## Resampling results across tuning parameters:
##
##
     mtry
           RMSE
                     Rsquared
                                 MAE
     2
                     0.1811534
                                 47.38910
##
           63.73568
     5
           65.99397
                     0.1735441
                                 48.38993
##
##
     8
           68.40480
                     0.1574900
                                 49.77404
##
## RMSE was used to select the optimal model using the smallest value.
## The final value used for the model was mtry = 2.
```

#### Total Aphids No Mfc Model



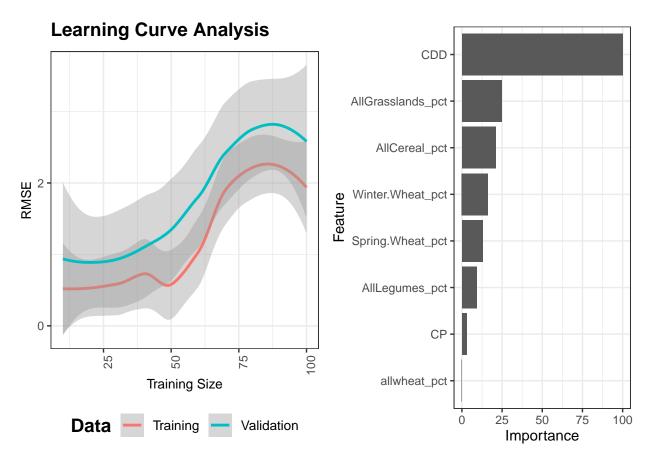
```
## Random Forest
##
## 101 samples
##
     8 predictors
##
## No pre-processing
## Resampling: Cross-Validated (10 fold, repeated 1 times)
## Summary of sample sizes: 89, 90, 92, 91, 91, 90, ...
## Resampling results across tuning parameters:
##
##
     mtry
           RMSE
                     Rsquared
                                 MAE
     2
                     0.2563659
                                 36.33553
##
           49.33028
     5
           49.26247
                     0.2559647
                                 35.33321
##
##
     8
           49.22627
                     0.2541812
                                 34.64219
##
## RMSE was used to select the optimal model using the smallest value.
## The final value used for the model was mtry = 8.
```

#### Relative Abundance Mfc Model



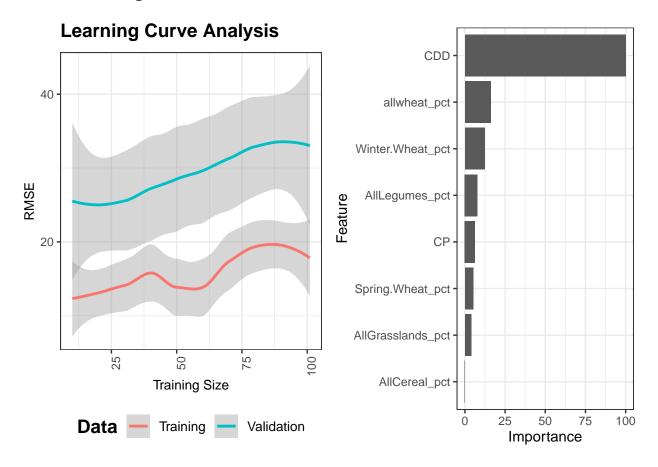
```
## Random Forest
##
## 101 samples
     8 predictors
##
##
## No pre-processing
## Resampling: Cross-Validated (10 fold, repeated 1 times)
## Summary of sample sizes: 89, 90, 92, 91, 91, 90, ...
## Resampling results across tuning parameters:
##
##
     mtry
           RMSE
                     Rsquared
                                 MAE
##
     2
           23.75841
                     0.2415788
                                 19.97434
##
     5
           24.29576
                     0.2339301
                                 20.53495
##
     8
           24.53101
                     0.2530602
                                20.70043
##
## RMSE was used to select the optimal model using the smallest value.
## The final value used for the model was mtry = 2.
```

# Individual Aphids: Rp



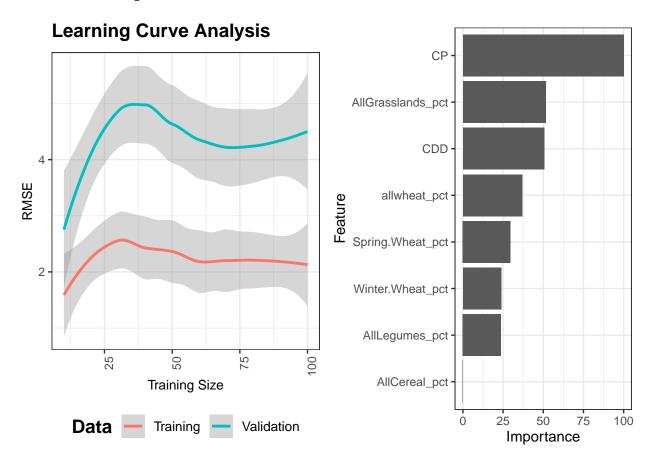
```
## Random Forest
##
##
  100 samples
     8 predictors
##
## No pre-processing
## Resampling: Cross-Validated (10 fold, repeated 1 times)
## Summary of sample sizes: 89, 91, 90, 89, 91, 90, ...
## Resampling results across tuning parameters:
##
##
     mtry
           RMSE
                     Rsquared
                                  {\tt MAE}
     2
                      0.09381127
                                  1.541037
##
           2.431611
     5
           2.575098
##
                      0.18266845
                                  1.536897
##
     8
           2.872344
                     0.16819091
                                  1.625897
##
## RMSE was used to select the optimal model using the smallest value.
## The final value used for the model was mtry = 2.
```

### Individual Aphids: Sa



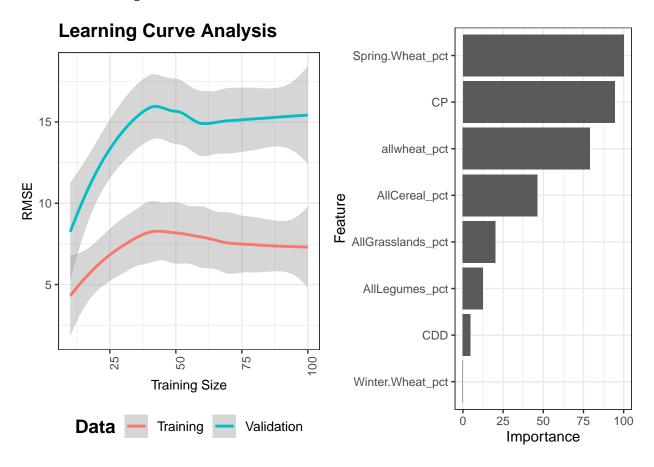
```
## Random Forest
##
## 101 samples
##
     8 predictors
##
## No pre-processing
## Resampling: Cross-Validated (10 fold, repeated 1 times)
## Summary of sample sizes: 89, 90, 92, 91, 91, 90, ...
## Resampling results across tuning parameters:
##
##
     mtry
           RMSE
                     Rsquared
                                 MAE
     2
           32.39298
                     0.4948489
                                 21.99353
##
     5
           31.82151
                     0.5220429
                                 21.25240
##
##
     8
           33.24165
                     0.5102561
                                 21.71563
##
## RMSE was used to select the optimal model using the smallest value.
## The final value used for the model was mtry = 5.
```

### Individual Aphids: Md



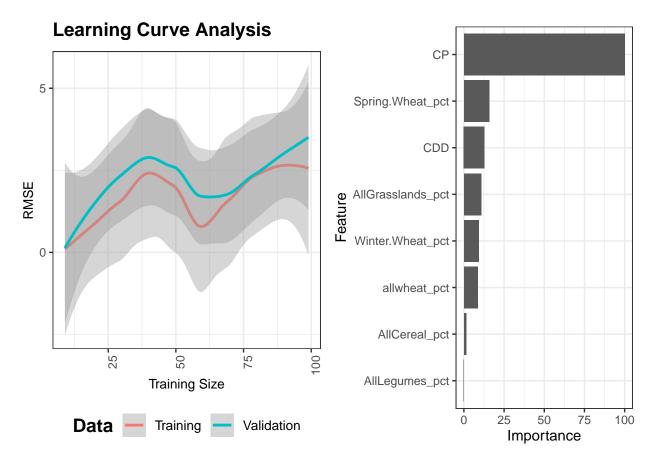
```
## Random Forest
##
## 100 samples
     8 predictors
##
## No pre-processing
## Resampling: Cross-Validated (10 fold, repeated 1 times)
## Summary of sample sizes: 90, 89, 89, 89, 91, 90, ...
## Resampling results across tuning parameters:
##
##
     mtry
           RMSE
                     Rsquared
                                 MAE
     2
           4.277962
                     0.2618058
                                 3.343674
##
     5
           4.335352
                     0.2462028
                                 3.387907
##
##
     8
           4.410895
                     0.2265757
                                 3.438736
##
## RMSE was used to select the optimal model using the smallest value.
## The final value used for the model was mtry = 2.
```

### Individual Aphids: Mfc



```
## Random Forest
##
##
  100 samples
     8 predictors
##
## No pre-processing
## Resampling: Cross-Validated (10 fold, repeated 1 times)
## Summary of sample sizes: 90, 89, 89, 90, 90, 91, ...
## Resampling results across tuning parameters:
##
##
     mtry
           RMSE
                     Rsquared
                                 MAE
     2
           14.57525
                     0.1221381
                                 11.71662
##
     5
           15.00601
                     0.1049100
                                 12.11249
##
##
     8
           15.07874
                     0.1190409
                                 12.22297
##
## RMSE was used to select the optimal model using the smallest value.
## The final value used for the model was mtry = 2.
```

# Individual Aphids: Rp



```
## Random Forest
##
## 99 samples
   8 predictors
##
## No pre-processing
## Resampling: Cross-Validated (10 fold, repeated 1 times)
## Summary of sample sizes: 89, 89, 89, 89, 89, 89, ...
## Resampling results across tuning parameters:
##
##
     mtry
           RMSE
                     Rsquared
                                 MAE
     2
           3.267462
                     0.2959596
                                 1.431210
##
     5
                     0.2503583
##
           3.406572
                                 1.420783
##
     8
           3.694826
                     0.2584216
                                 1.461916
##
## RMSE was used to select the optimal model using the smallest value.
## The final value used for the model was mtry = 2.
```

