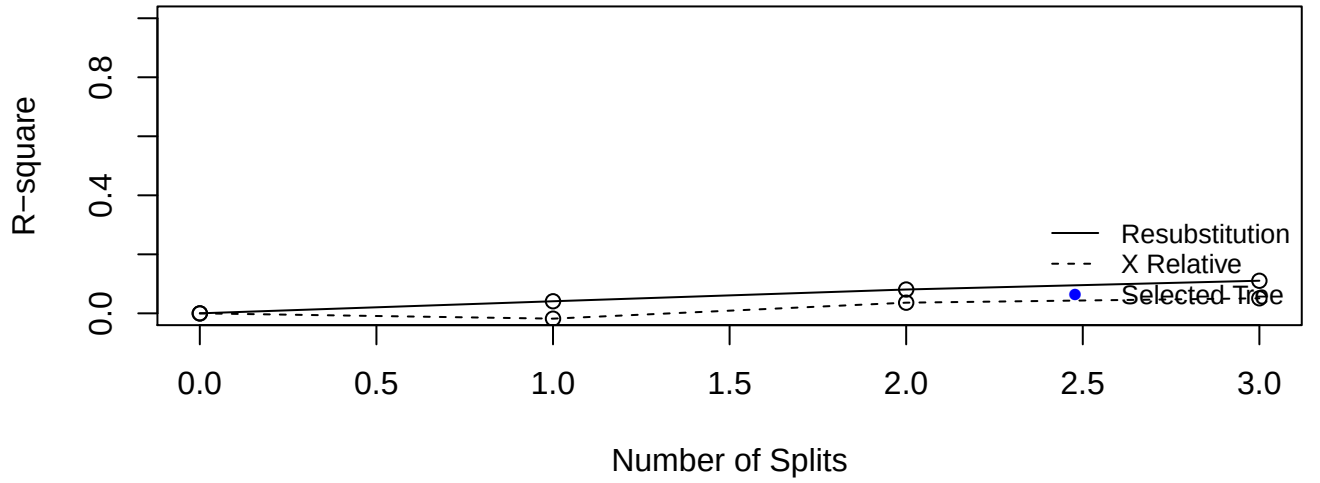
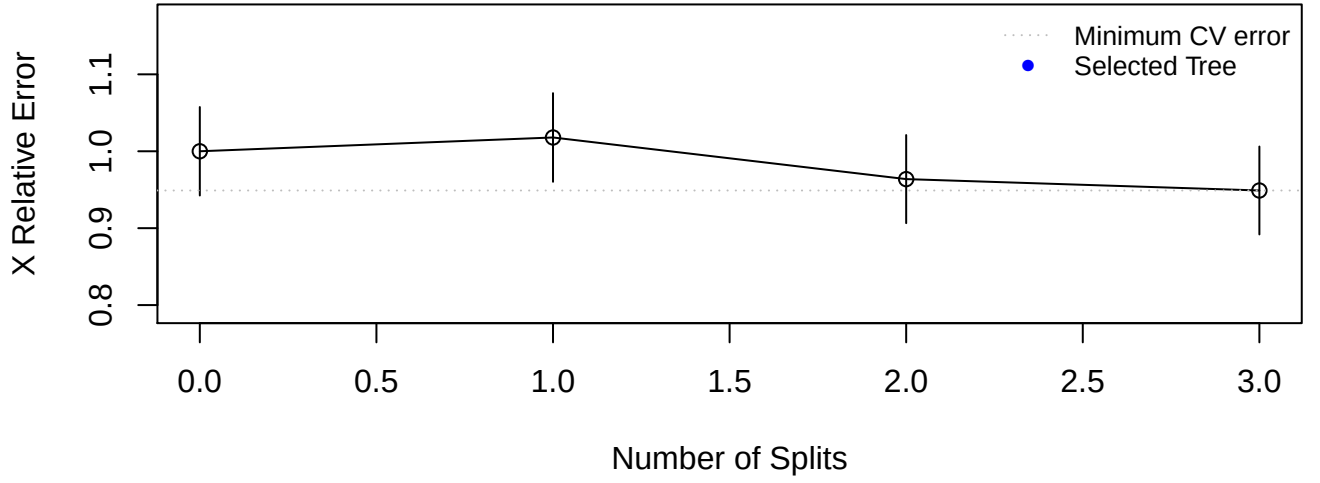
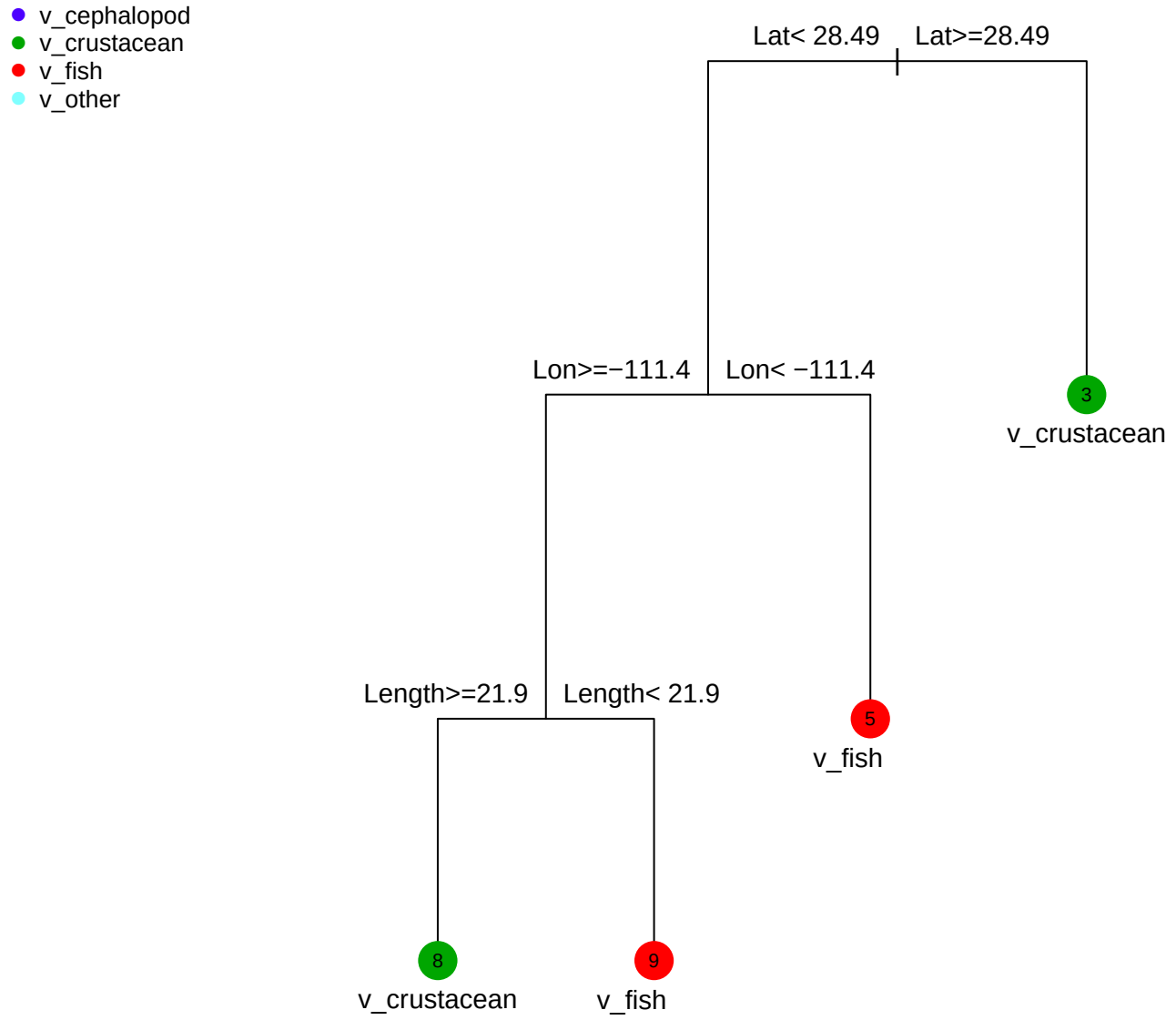

R−square
0.0
0.4
0.8
Resubstitution
X Relative
Selected Tree
0.0
0.5
1.0
1.5
2.0
2.5
3.0
Number of Splits


X Relative Error
0.8
0.9
1.0
1.1
Minimum CV error
Selected Tree
0.0
0.5
1.0
1.5
2.0
2.5
3.0
Number of Splits


v_cephalopod
v_crustacean
v_fish
v_other
Lat< 28.49
Lat>=28.49
3
v_crustacean
Lon>=−111.4
Lon< −111.4
5
v_fish
Length>=21.9
Length< 21.9
8
v_crustacean
9
v_fish

Variable Importance
1.0
0.8
0.6
0.4
0.2
0.0
Lat
Lon
Length
Year