Contradicts

Case: s2523728.1

Case: s238462.1

Contradicts

Supports

Case: s457952.1

Contradicts

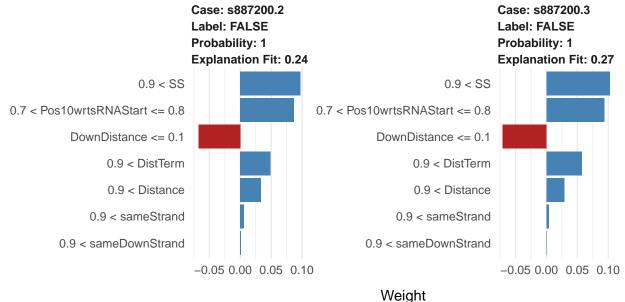
Supports

Case: s852175



Weight

Case: s852175.1





-0.05 0.00 0.05 0.10

Case: s1195937

0.9 < sameDownStrand



Weight

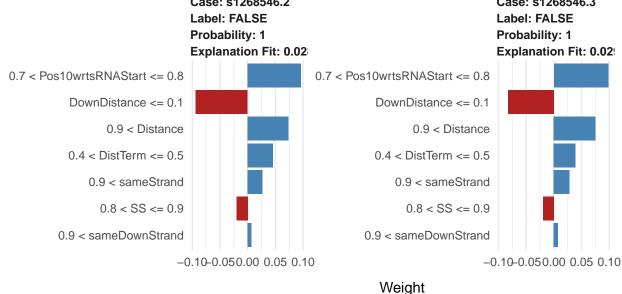
0.9 < sameDownStrand

Case: s1195937.1

-0.05 0.00 0.05 0.10

Case: s1268546

Feature





Case: s1268546.1

Contradicts

Case: s1403676.1

Explanation Fit: 0.08

Label: FALSE

Probability: 1

Case: s1403676

Explanation Fit: 0.09

Label: FALSE

Probability: 1

Contradicts

Case: s1407387.1

Case: s1489467.1

Label: FALSE

Probability: 1

Contradicts

Supports

Case: s1489467

Label: FALSE

Probability: 1

Case: B_cenocepacia

Label: TRUE

0.1 < DownDistance <= 0.2

0.6 < SS <= 0.7

0.0

0.1

0.2



Weight

0.1 < DownDistance <= 0.2

0.6 < SS <= 0.7

0.0

0.1

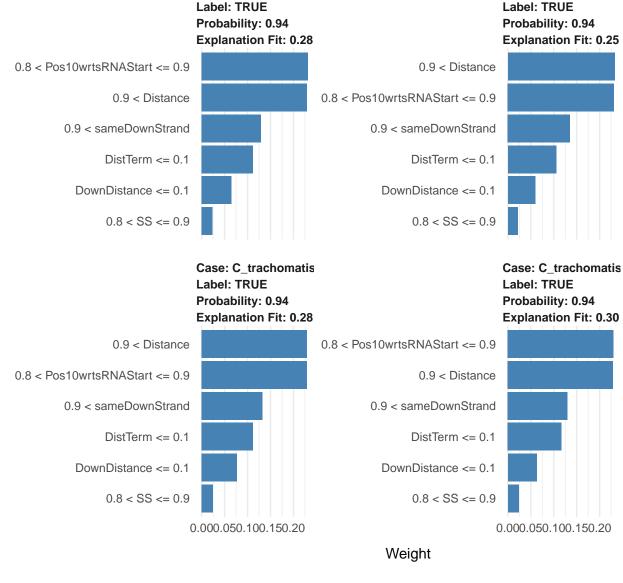
0.2

Case: B_cenocepacia

Contradicts

Case: B_cenocepacia

Case: B_cenocepacia



Contradicts

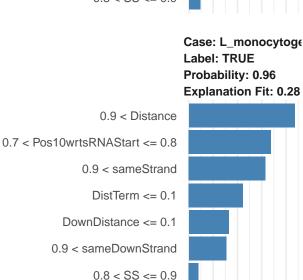
Case: C trachomatis

Case: C trachomatis

Case: L_monocytoge

Label: TRUE

Probability: 0.96



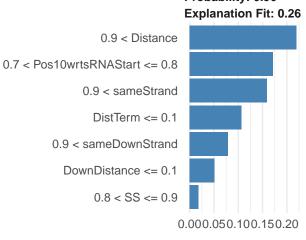
0.000.050.100.150.20

Case: L_monocytoge

Explanation Fit: 0.26

Label: TRUE

Probability: 0.96







Contradicts

Case: S coelicolor F

Label: TRUE

Case: S coelicolor F

Label: TRUE

Contradicts

Case: V cholerae R

Label: TRUE

Case: V cholerae R/

Label: TRUE

DownDistance <= 0.1

0.8 < SS <= 0.9

0.000.050.100.150.200.25

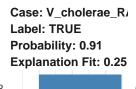


Weight

DownDistance <= 0.1

0.8 < SS <= 0.9

0.000.050.100.150.200.25



Case: V_cholerae_R/

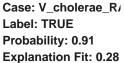
Label: TRUE

0.0

0.1

0.2

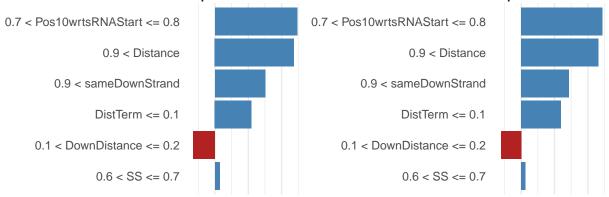
0.6 < SS <= 0.7

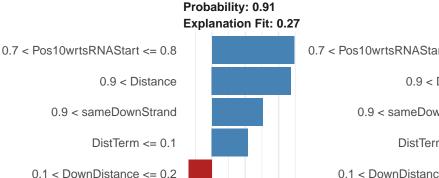


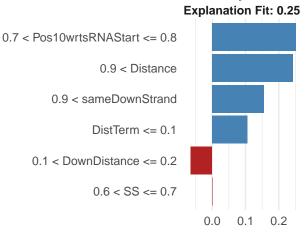
Case: V_cholerae_R/

Label: TRUE

Probability: 0.91







Weight



Label: TRUE
Probability: 0.91
Explanation Fit: 0.26

0.8 < Pos10wrtsRNAStart <= 0.9

once

0.9 < Distance

0.9 < sameStrand

DistTerm <= 0.1

DownDistance <= 0.1



Case: V_cholerae_R/ Label: TRUE Probability: 0.91

Case: V cholerae R

Probability: 0.91
Explanation Fit: 0.28

0.8 < Pos10wrtsRNAStart <= 0.9

0.9 < Distance

0.9 < sameStrand

DistTerm <= 0.1

DownDistance <= 0.1

0.8 < SS <= 0.9

0.8 < SS <= 0.9

Case: V cholerae R/

Case: V_cholerae_R/

Explanation Fit: 0.29

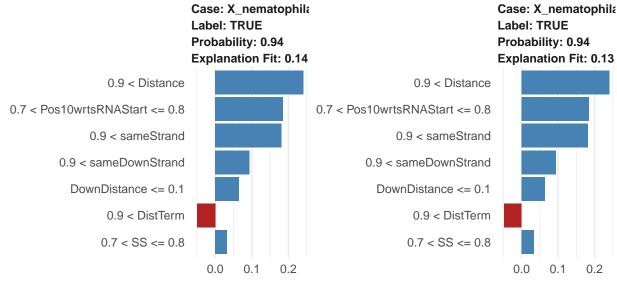
0.000.050.100.150.20

Label: TRUE

Probability: 0.91

Weight







Weight