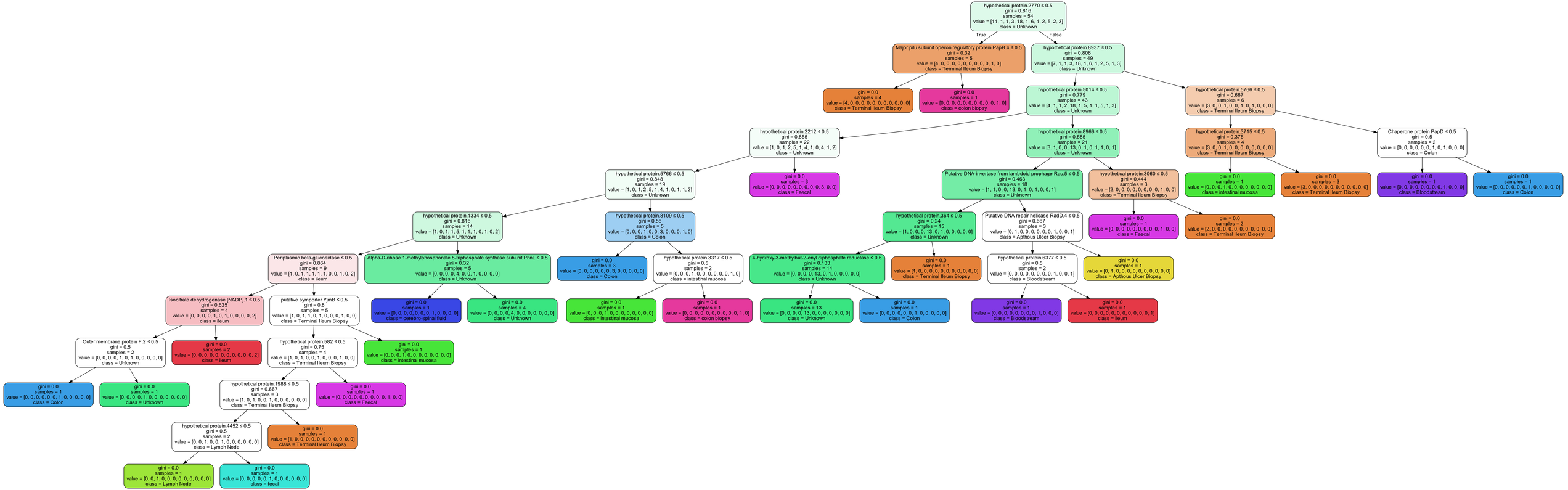
hypothetical protein.2770 ≤ 0.5
gini = 0.816
samples = 54
value = [11, 1, 1, 3, 18, 1, 6, 1, 2, 5, 2, 3]
class = Unknown
True
False
Major pilu subunit operon regulatory protein PapB.4 ≤ 0.5
gini = 0.32
samples = 5
value = [4, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0]
class = Terminal Ileum Biopsy
hypothetical protein.8937 ≤ 0.5
gini = 0.808
samples = 49
value = [7, 1, 1, 3, 18, 1, 6, 1, 2, 5, 1, 3]
class = Unknown
gini = 0.0
samples = 4
value = [4, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0]
class = Terminal Ileum Biopsy
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0]
class = colon biopsy
hypothetical protein.5014 ≤ 0.5
gini = 0.779
samples = 43
value = [4, 1, 1, 2, 18, 1, 5, 1, 1, 5, 1, 3]
class = Unknown
hypothetical protein.5766 ≤ 0.5
gini = 0.667
samples = 6
value = [3, 0, 0, 1, 0, 0, 1, 0, 1, 0, 0, 0]
class = Terminal Ileum Biopsy
hypothetical protein.2212 ≤ 0.5
gini = 0.855
samples = 22
value = [1, 0, 1, 2, 5, 1, 4, 1, 0, 4, 1, 2]
class = Unknown
hypothetical protein.8966 ≤ 0.5
gini = 0.585
samples = 21
value = [3, 1, 0, 0, 13, 0, 1, 0, 1, 1, 0, 1]
class = Unknown
hypothetical protein.3715 ≤ 0.5
gini = 0.375
samples = 4
value = [3, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0]
class = Terminal Ileum Biopsy
Chaperone protein PapD ≤ 0.5
gini = 0.5
samples = 2
value = [0, 0, 0, 0, 0, 0, 1, 0, 1, 0, 0, 0]
class = Colon
hypothetical protein.5766 ≤ 0.5
gini = 0.848
samples = 19
value = [1, 0, 1, 2, 5, 1, 4, 1, 0, 1, 1, 2]
class = Unknown
gini = 0.0
samples = 3
value = [0, 0, 0, 0, 0, 0, 0, 0, 0, 3, 0, 0]
class = Faecal
Putative DNA-invertase from lambdoid prophage Rac.5 ≤ 0.5
gini = 0.463
samples = 18
value = [1, 1, 0, 0, 13, 0, 1, 0, 1, 0, 0, 1]
class = Unknown
hypothetical protein.3060 ≤ 0.5
gini = 0.444
samples = 3
value = [2, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0]
class = Terminal Ileum Biopsy
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0]
class = colon biopsy
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0]
class = Terminal Ileum Biopsy
gini = 0.0
samples = 3
value = [3, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0]
class = Terminal Ileum Biopsy
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0]
class = Bloodstream
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1]
class = Colon
hypothetical protein.1334 ≤ 0.5
gini = 0.816
samples = 14
value = [1, 0, 1, 1, 5, 1, 1, 1, 0, 1, 0, 2]
class = Unknown
hypothetical protein.8109 ≤ 0.5
gini = 0.56
samples = 5
value = [0, 0, 0, 1, 0, 0, 3, 0, 0, 0, 1, 0]
class = Colon
hypothetical protein.364 ≤ 0.5
gini = 0.24
samples = 15
value = [1, 0, 0, 0, 13, 0, 1, 0, 0, 0, 0, 0]
class = Unknown
Putative DNA repair helicase RadD.4 ≤ 0.5
gini = 0.667
samples = 3
value = [0, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0, 1]
class = Apthous Ulcer Biopsy
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0]
class = Faecal
gini = 0.0
samples = 2
value = [2, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0]
class = Terminal Ileum Biopsy
Periplasmic beta-glucosidase ≤ 0.5
gini = 0.864
samples = 9
value = [1, 0, 1, 1, 1, 1, 1, 0, 0, 1, 0, 2]
class = ileum
Alpha-D-ribose 1-methylphosphonate 5-triphosphate synthase subunit PhnL ≤ 0.5
gini = 0.32
samples = 5
value = [0, 0, 0, 0, 4, 0, 0, 1, 0, 0, 0, 0]
class = Unknown
gini = 0.0
samples = 3
value = [0, 0, 0, 0, 0, 0, 3, 0, 0, 0, 0, 0]
class = Colon
hypothetical protein.3317 ≤ 0.5
gini = 0.5
samples = 2
value = [0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 1, 0]
class = intestinal mucosa
4-hydroxy-3-methylbut-2-enyl diphosphate reductase ≤ 0.5
gini = 0.133
samples = 14
value = [0, 0, 0, 0, 13, 0, 1, 0, 0, 0, 0, 0]
class = Unknown
gini = 0.0
samples = 1
value = [1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0]
class = Terminal Ileum Biopsy
hypothetical protein.6377 ≤ 0.5
gini = 0.5
samples = 2
value = [0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 1]
class = Bloodstream
gini = 0.0
samples = 1
value = [0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0]
class = Apthous Ulcer Biopsy
Isocitrate dehydrogenase [NADP].1 ≤ 0.5
gini = 0.625
samples = 4
value = [0, 0, 0, 0, 1, 0, 1, 0, 0, 0, 0, 2]
class = ileum
putative symporter YjmB ≤ 0.5
gini = 0.8
samples = 5
value = [1, 0, 1, 1, 0, 1, 0, 0, 0, 1, 0, 0]
class = Terminal Ileum Biopsy
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0]
class = cerebro-spinal fluid
gini = 0.0
samples = 4
value = [0, 0, 0, 0, 4, 0, 0, 0, 0, 0, 0, 0]
class = Unknown
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0]
class = intestinal mucosa
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0]
class = colon biopsy
gini = 0.0
samples = 13
value = [0, 0, 0, 0, 13, 0, 0, 0, 0, 0, 0, 0]
class = Unknown
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0]
class = Colon
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0]
class = Bloodstream
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1]
class = ileum
Outer membrane protein F.2 ≤ 0.5
gini = 0.5
samples = 2
value = [0, 0, 0, 0, 1, 0, 1, 0, 0, 0, 0, 0]
class = Unknown
gini = 0.0
samples = 2
value = [0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 2]
class = ileum
hypothetical protein.582 ≤ 0.5
gini = 0.75
samples = 4
value = [1, 0, 1, 0, 0, 1, 0, 0, 0, 1, 0, 0]
class = Terminal Ileum Biopsy
gini = 0.0
samples = 1
value = [0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0]
class = intestinal mucosa
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0]
class = Colon
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0]
class = Unknown
hypothetical protein.1988 ≤ 0.5
gini = 0.667
samples = 3
value = [1, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0]
class = Terminal Ileum Biopsy
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0]
class = Faecal
hypothetical protein.4452 ≤ 0.5
gini = 0.5
samples = 2
value = [0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0]
class = Lymph Node
gini = 0.0
samples = 1
value = [1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0]
class = Terminal Ileum Biopsy
gini = 0.0
samples = 1
value = [0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0]
class = Lymph Node
gini = 0.0
samples = 1
value = [0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0]
class = fecal