

F0

Database

Replicates

k2\_pluspf -

k2\_standard -

Donor\_1 -

Donor\_2 -

0.00

0.25

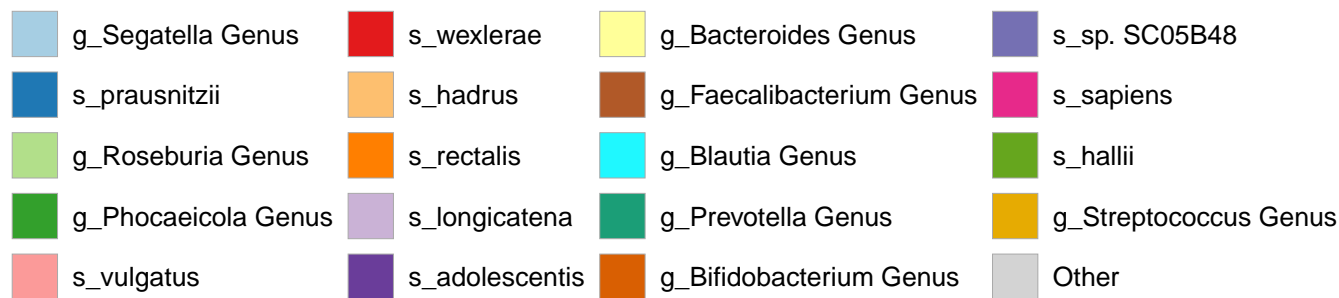
0.50

0.75

1.00

Abundance

Species



F1

Database

k2\_pluspf -

k2\_standard -

Donor\_1 -

0.00

0.25

0.50

0.75

1.00

Abundance

Species



F2

Database

Replicates

k2\_pluspf-

k2\_standard-

Donor\_1-

Donor\_2-

0.00

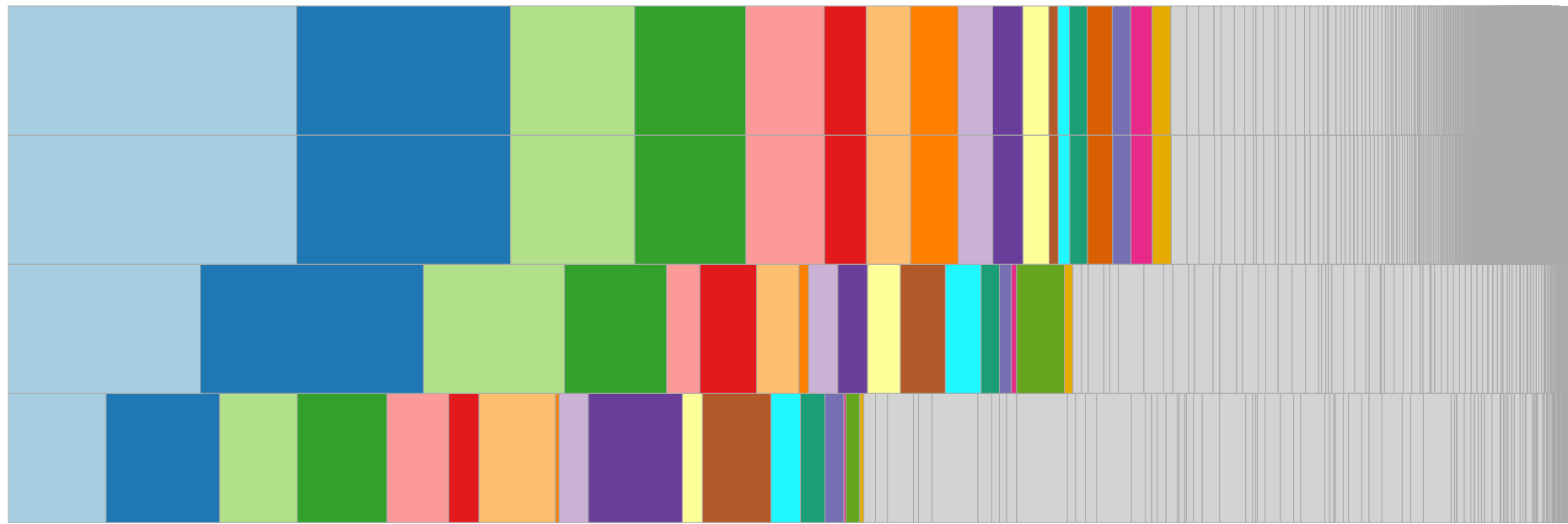
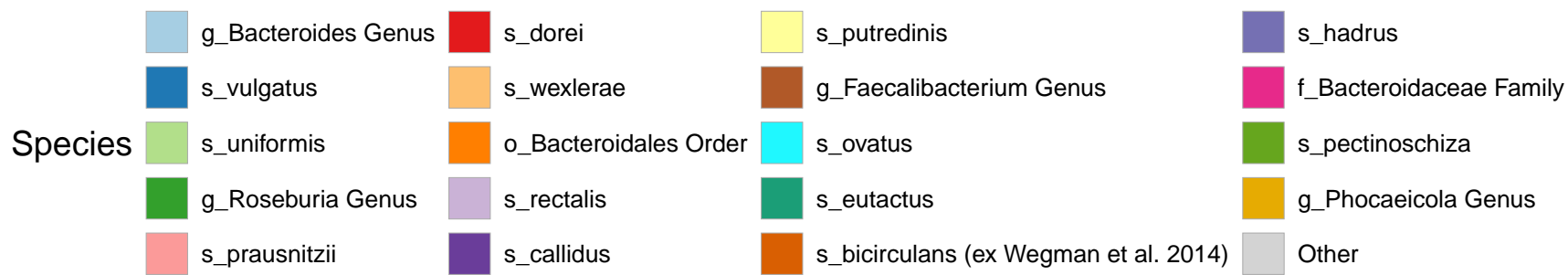
0.25

0.50

0.75

1.00

Abundance



F3

Database

Replicates

k2\_pluspf -

k2\_standard -

Donor\_1 -

Donor\_2 -

0.00

0.25

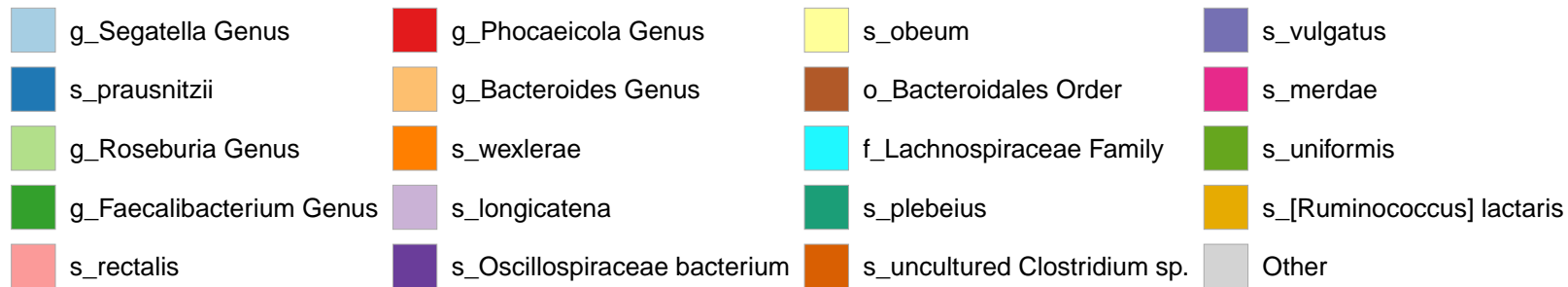
0.50

0.75

1.00

Abundance

Species



M4

Database

Replicates

k2\_pluspf -

k2\_standard -

Donor\_1 -

Donor\_2 -

0.00

0.25

0.50

0.75

1.00

Abundance



M5

Database

k2\_pluspf -

k2\_standard -

Replicates

Donor\_1 -

Donor\_2 -

0.00

0.25

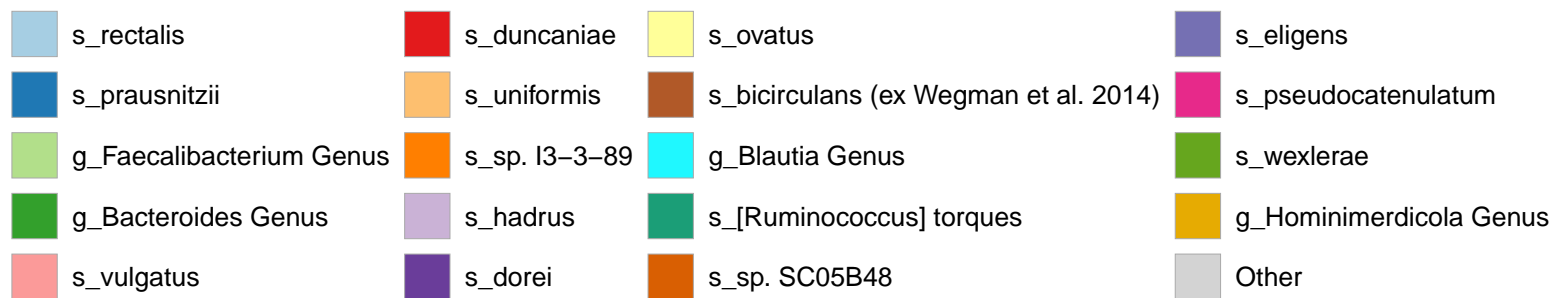
0.50

0.75

1.00

Abundance

Species



F6

Database

Replicates

k2\_pluspf -

k2\_standard -

Donor\_1 -

Donor\_2 -

0.00

0.25

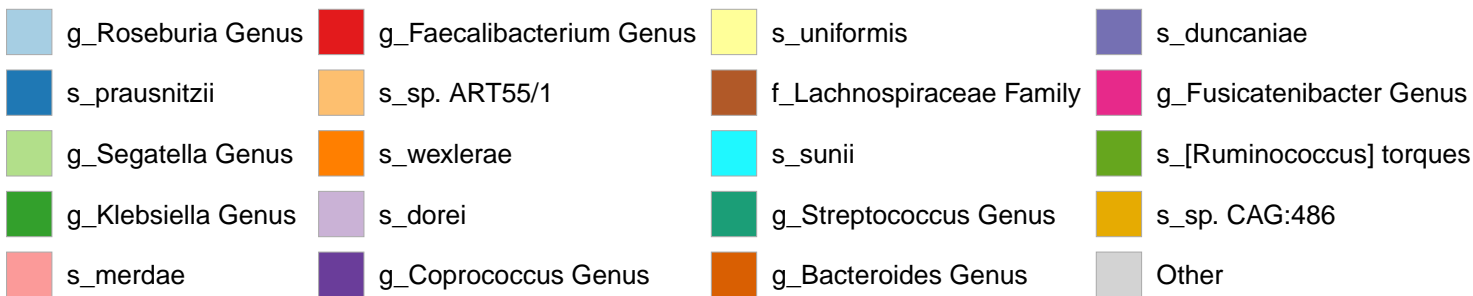
0.50

0.75

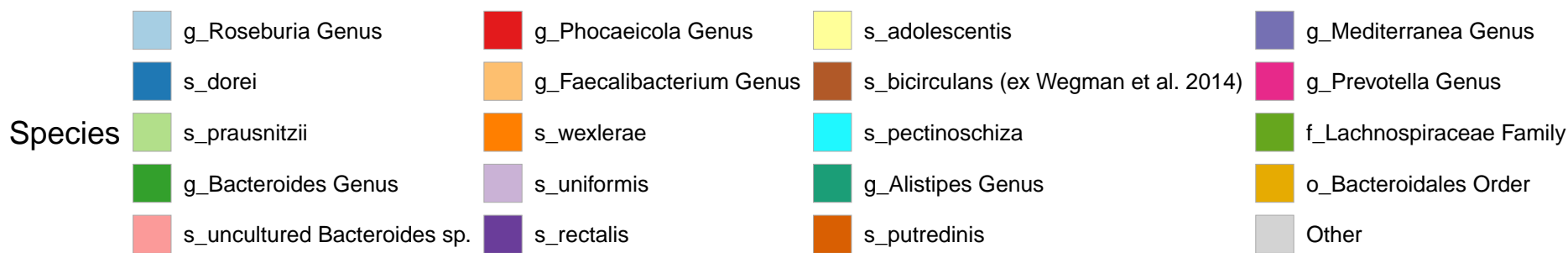
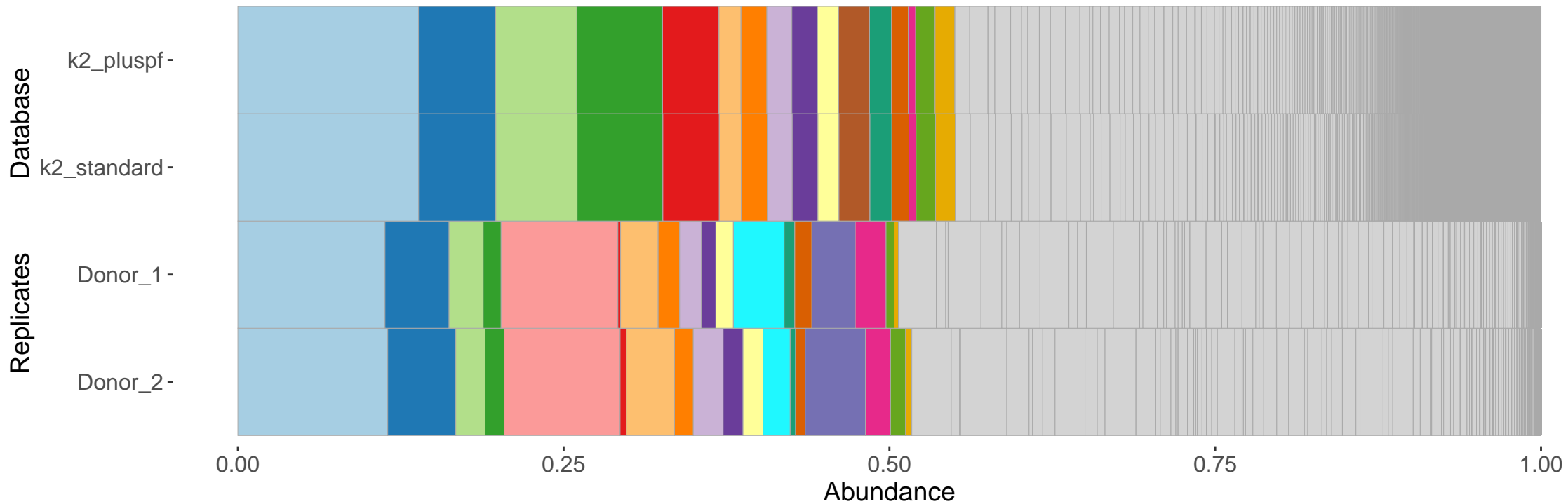
1.00

Abundance

Species



F7



F8

Database

Replicates

k2\_pluspf -

k2\_standard -

Donor\_1 -

Donor\_2 -

0.00

0.25

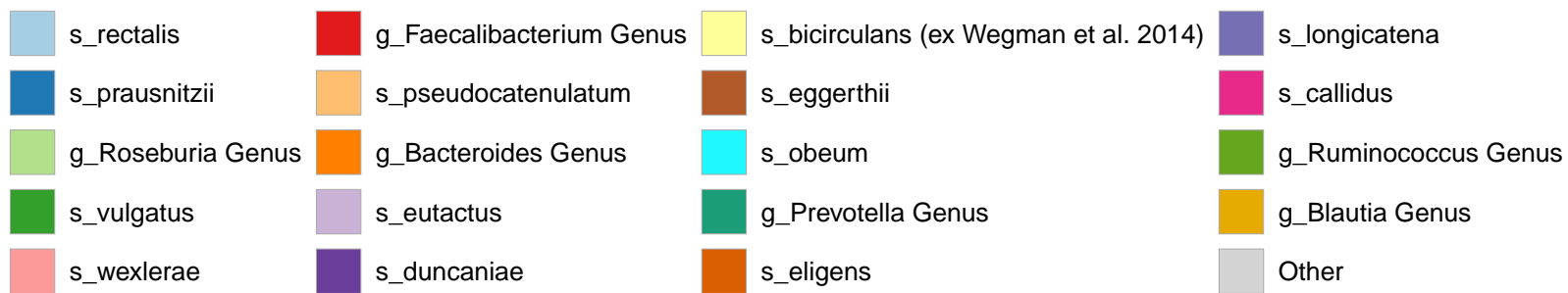
0.50

0.75

1.00

Abundance

Species



F9

Database

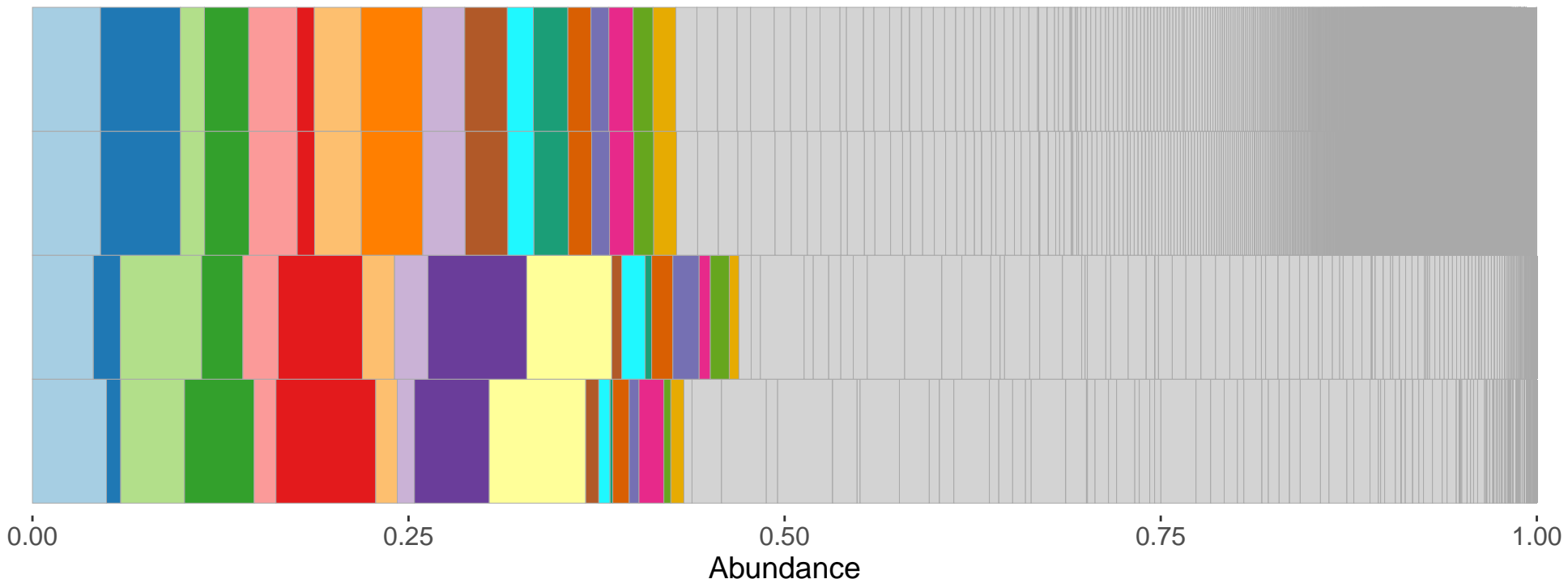
Replicates

k2\_pluspf -

k2\_standard -

Donor\_1 -

Donor\_2 -



0.00

0.25





















0.50

0.75

1.00

Abundance

Species

- |   |  |   |   |
|---|--|---|---|
|  s_vulgatus      |  g_Coprococcus Genus        |  g_Ruminococcoides Genus |  g_Faecalibacterium Genus  |
|  s_prausnitzii   |  g_Roseburia Genus          |  g_Bacteroides Genus     |  s_[Ruminococcus] torques  |
|  g_Blautia Genus |  s_sp. ART55/1              |  s_hallii                |  g_Faecalibacillus Genus   |
|  s_eggerthii    |  s_hadrus                  |  s_obeum                |  f_Lachnospiraceae Family |
|  s_wexlerae    |  g_Fusicatenibacter Genus |  s_eligens             |  Other                   |