

Taxonomy bar plot of top 25 Species

Male Disorder

Male Normal

F30682.S01 -

F30682.S06 -

F30682.S07 -

F30682.S02 -

F30682.S08 -

Species

- Acetivibrio cellulolyticus_nov_83.153%
- Hathewayia proteolytica_nov_83.297%
- Lachnospiraceae_[G-14] bacterium_MOT-183_nov_93.348%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
- Hathewayia proteolytica_nov_84.233%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.719%
- Lachnospiraceae_[G-11] bacterium_MOT-177_nov_96.066%
- Anaerotignum faecicola_nov_85.287%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.466%
- Anaerotignum lactatifermentans_nov_84.040%
- Acetivibrio cellulolyticus_nov_83.190%
- Agrococcus versicolor_nov_83.227%
- Turicibacter sanguinis_nov_95.923%
- Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_91.775%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.489%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_93.348%
- Lachnospiraceae_[G-1] bacterium_MOT-166
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.407%
- Lachnospiraceae_[G-14] bacterium_MOT-182_nov_90.042%
- Phoceia massiliensis_nov_89.722%
- Eubacterium ventriosum_nov_93.320%
- Acetivibrio cellulolyticus_nov_85.560%
- Lachnoclostridium phytofermentans_nov_91.458%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.125%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_97.210%
- Other

0.00 0.25 0.50 0.75 1.00

Abundance

0.00 0.25 0.50 0.75 1.00