

Taxonomy bar plot of top 25 Species

Female_B6_Mice

Female_NOD_Mice

F5362.S05 -

F5362.S10 -

F5362.S11 -

F5362.S06 -

F5362.S12 -

Species

- Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.590%
- Duncaniella freteri_nov_93.774%
- Acetivibrio cellulolyticus_nov_83.851%
- Lawsonibacter asaccharolyticus_nov_91.762%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_93.103%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.069%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.697%
- Duncaniella freteri_nov_90.323%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.886%
- Hathewayia proteolytica_nov_84.569%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_97.154%
- Duncaniella freteri_nov_93.208%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.267%
- Duncaniella freteri_nov_94.162%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.478%
- Lachnospiraceae_[G-11] bacterium_MOT-178_nov_94.778%
- Sporobacter termitidis_nov_83.168%
- Lachnospiraceae_[G-11] bacterium_MOT-177_nov_93.922%
- Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_84.091%
- Lawsonibacter asaccharolyticus_nov_94.798%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.450%
- Hydrogenoanaerobacterium saccharovorans_nov_89.942%
- Blautia faecicola_nov_89.749%
- Bacteroides thetaiotaomicron
- Lachnospiraceae_[G-11] bacterium_MOT-178_nov_92.293%
- Other

0.00 0.25 0.50 0.75 1.00

Abundance

0.00 0.25 0.50 0.75 1.00