



- Map_10_002
- Map_10_001
- Map_175_001
- Map_31_004
- Map_37_008
- Map_45_004
- Map_54_005
- Map_80_004
- 630129.13
- Map_154_001
- M264_MCI
- Map_59_012
- 630129.8
- 28026.754
- DianaHT_2018_SRR6795021_bin.13
- ERR7738228_bin.16
- Bj155_06_4B1
- MGYG00000756
- Map_36_006
- Map_72_005
- N4G05
- ShaoY_2019_ERR3404927_bin.4
- Map_129_001
- Map_151_001
- Map_168_002
- Map_47_002
- Map_138_001
- 1447716.4
- Map_20_007
- BP223_mabbin.011_sub
- YIT13052
- MurphyR_2019_SRR7351817_bin.7
- MurphyR_2019_SRR7352076_bin.1
- MurphyR_2019_SRR7351942_bin.8
- MurphyR_2019_SRR7352002_bin.12
- MurphyR_2019_SRR7351642_bin.8
- Map_63_012
- ShaoY_2019_ERR3405657_bin.2
- MurphyR_2019_SRR7352101_bin.11
- MGYG000264346
- MurphyR_2019_SRR7351965_bin.4
- Y150460.7
- YIT13051
- YIT13061
- APCKJ1
- YIT13053
- YIT13060
- YIT13054
- YIT13056
- ShaoY_2019_ERR3406028_bin.3
- YIT13057
- ShaoY_2019_ERR3406125_bin.1
- YIT13052
- YIT13055
- WumpachL_2018_SRR5575423_bin.7
- YIT13056
- ShaoY_2019_ERR3405065_bin.8
- ShaoY_2019_ERR3406102_bin.6
- ShaoY_2019_ERR3405688_bin.7
- ShaoY_2019_ERR3406253_bin.8
- ShaoY_2019_ERR3405918_bin.15
- ShaoY_2019_ERR3406290_bin.8
- ShaoY_2019_ERR3405915_bin.14
- 630129.10
- DianaHT_2018_SRR6795059_bin.13
- MurphyR_2019_SRR7351718_bin.6
- ShaoY_2019_ERR3405871_bin.8
- 630129.38
- ShaoY_2019_ERR3404970_bin.5
- ShaoY_2019_ERR3405119_bin.12
- ParnanenK_2018_SRR5723876_bin.5
- 1686.6
- FerretP_2018_SRR5273974_bin.15
- ShaoY_2019_ERR3405896_bin.18
- ShaoY_2019_ERR3405246_bin.4
- ShaoY_2019_ERR3405190_bin.3
- MurphyR_2019_SRR7351904_bin.13
- ChuDM_2017_SRR8942351_bin.4
- ShaoY_2019_ERR3405965_bin.2
- MGYG00013873
- ShaoY_2019_ERR3404734_bin.10
- VatanenT_2016_SRR4408034_bin.6
- BackheF_2015_ERR526045_bin.2
- ChuDM_2017_SRR8942415_bin.5
- ParnanenK_2018_SRR5723804_bin.1
- MurphyR_2019_SRR7352095_bin.8
- ShaoY_2019_ERR3406266_bin.7
- VatanenT_2016_SRR4305363_bin.2
- CokerMO_2019_SRR8692185_bin.21
- CasaburiG_2019_SRR6277129_bin.6
- 56652.4
- BackheF_2016_ERR525997_bin.2
- ShaoY_2019_ERR3405651_bin.7
- VatanenT_2016_SRR4305474_bin.9
- ShaoY_2019_ERR3406144_bin.2
- Bifidobacterium_catenulatum_BIOML_A1
- SPMP38_bin.25
- YIT13059
- VatanenT_2016_SRR4408139_bin.2
- TM115-191
- MurphyR_2019_SRR7411410_bin.1
- MurphyR_2019_SRR7351968_bin.3
- MurphyR_2019_SRR7351945_bin.4
- MurphyR_2019_SRR7351807_bin.6
- MGYG00031974
- MGYG00011443
- MGYG000110618
- MGYG000008610
- KY225_44
- CokerMO_2019_SRR8692222_bin.7
- ChuDM_2017_SRR8942384_bin.1
- BackheF_2016_ERR525762_bin.5
- Baumann-DudenhoeflerAM_2018_SRR7217889_bin.9
- MGYG000152456
- MGYG000091537
- ShaoY_2019_ERR3405213_bin.4
- MurphyR_2019_SRR7351990_bin.1
- MGYG000143087
- MurphyR_2019_SRR7351743_bin.7
- ShaoY_2019_ERR3406198_bin.10
- YassourM_2018_SRR7280800_bin.4
- KU_genome_26637
- Bj84_11_2010
- ShaoY_2019_ERR3405802_bin.7
- MurphyR_2019_SRR7351639_bin.4
- ShaoY_2019_ERR3406025_bin.10
- 1686.302
- DianaHT_2018_SRR6795047_bin.4
- MGYG000279952
- ShaoY_2019_ERR3405650_bin.3
- CLA-AA-H311
- SPMP78_bin.95
- AG465_3
- KU_genome_18534
- NSG01

Predicted phenotype

- 0
- 1

Taxonomy

- Bifidobacterium catenulatum subsp. catenulatum
- Bifidobacterium catenulatum subsp. kashiwanohense
- Bifidobacterium catenulatum subsp. kashiwanohense_A
- Bifidobacterium hominis

type

- di_and_oligosaccharides
- monosaccharides_and_derivatives
- polysaccharides

origin

- bacterial
- fungal
- host (animal)
- plant
- universal