

# Classification of environmental strains from order to genus levels using lipid and protein MALDI-ToF fingerprintings and chemotaxonomic network analysis

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TABLE S1. SUMMARY TABLE OF STUDIED BACTERIAL STRAINS (RKI)

<b>Names</b>	<b>Order</b>	<b>Genus</b>
<i>Achromobacter_xylosoxidans_HST_107_01</i>	Burkholderiales	<i>Achromobacter</i>
<i>Achromobacter_xylosoxidans_HST_107_02</i>	Burkholderiales	<i>Achromobacter</i>
<i>Achromobacter_xylosoxidans_HST_115_01</i>	Burkholderiales	<i>Achromobacter</i>
<i>Achromobacter_xylosoxidans_HST_115_02</i>	Burkholderiales	<i>Achromobacter</i>
<i>Acinetobacter_baumannii_DSM_24110_01</i>	Pseudomonadales	<i>Acinetobacter</i>
<i>Acinetobacter_baumannii_DSM_24110_02</i>	Pseudomonadales	<i>Acinetobacter</i>
<i>Acinetobacter_baumannii_DSM_24110_03</i>	Pseudomonadales	<i>Acinetobacter</i>
<i>Acinetobacter_baumannii_DSM_24110_04</i>	Pseudomonadales	<i>Acinetobacter</i>
<i>Aeromonas_hydrophila_subsp_hydrophila_DSM_30187_01</i>	Aeromonadales	<i>Aeromonas</i>
<i>Aeromonas_hydrophila_subsp_hydrophila_DSM_30187_02</i>	Aeromonadales	<i>Aeromonas</i>
<i>Aeromonas_hydrophila_subsp_hydrophila_DSM_30187_03</i>	Aeromonadales	<i>Aeromonas</i>
<i>Aeromonas_hydrophila_subsp_hydrophila_DSM_30187_04</i>	Aeromonadales	<i>Aeromonas</i>
<i>Bacillus_alveayuensis_DSM_19092_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_alveayuensis_DSM_19092_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_alveayuensis_DSM_19092_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_alveayuensis_DSM_19092_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_aminovorans_DSM_4337_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_aminovorans_DSM_4337_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_aminovorans_DSM_4337_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_aminovorans_DSM_4337_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_amyloliquefaciens_ssp_amyloliquefaciens_DSM_7_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_amyloliquefaciens_ssp_amyloliquefaciens_DSM_7_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_amyloliquefaciens_ssp_amyloliquefaciens_DSM_7_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_amyloliquefaciens_ssp_amyloliquefaciens_DSM_7_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_anthraxis_A1_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_anthraxis_A1_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_anthraxis_A1_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_anthraxis_A1_04</i>	Bacillales	<i>Bacillus</i>

<i>Bacillus_atrophaeus_7264_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_atrophaeus_7264_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_atrophaeus_7264_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_atrophaeus_7264_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_bataviensis_DSM_15601_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_bataviensis_DSM_15601_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_bataviensis_DSM_15601_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_bataviensis_DSM_15601_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_clarkii_DSM_9725_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_clarkii_DSM_9725_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_clarkii_DSM_9725_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_clarkii_DSM_9725_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_clausii_DSM_8716_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_clausii_DSM_8716_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_clausii_DSM_8716_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_clausii_DSM_8716_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_coagulans_DSM_1_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_coagulans_DSM_1_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_coagulans_DSM_1_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_coagulans_DSM_1_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_cohnii_DSM_6350_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_cohnii_DSM_6350_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_cohnii_DSM_6350_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_cohnii_DSM_6350_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_cucumis_DSM_101566_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_cucumis_DSM_101566_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_cucumis_DSM_101566_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_cucumis_DSM_101566_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_cytotoxicus_DSM_22905_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_cytotoxicus_DSM_22905_02</i>	Bacillales	<i>Bacillus</i>

<i>Bacillus_cytotoxicus_DSM_22905_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_cytotoxicus_DSM_22905_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_ectoiniformans_DSM_28970_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_ectoiniformans_DSM_28970_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_ectoiniformans_DSM_28970_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_ectoiniformans_DSM_28970_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_encimensis_DSM_28241_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_encimensis_DSM_28241_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_encimensis_DSM_28241_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_encimensis_DSM_28241_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_firmus_DSM_12_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_firmus_DSM_12_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_flexus_DSM_1667_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_flexus_DSM_1667_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_flexus_DSM_1667_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_flexus_DSM_1667_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_halodurans_DSM_6940_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_halodurans_DSM_6940_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_halodurans_DSM_6940_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_halodurans_DSM_6940_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_halotolerans_18911_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_halotolerans_18911_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_halotolerans_18911_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_halotolerans_18911_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_horneckiae_23495_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_horneckiae_23495_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_horneckiae_23495_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_horneckiae_23495_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_infantis_DSM_19089_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_infantis_DSM_19089_02</i>	Bacillales	<i>Bacillus</i>

<i>Bacillus_infantis_DSM_19089_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_infantis_DSM_19089_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_lentus_DSM_5221_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_lentus_DSM_5221_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_lentus_DSM_5221_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_lentus_DSM_5221_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_licheniformis_DSM_13_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_licheniformis_DSM_13_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_megaterium_DSM_90_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_megaterium_DSM_90_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_mojavensis_DSM9206_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_mojavensis_DSM9206_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_mojavensis_DSM9206_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_mojavensis_DSM9206_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_muralis_DSM16288_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_muralis_DSM16288_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_muralis_DSM16288_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_muralis_DSM16288_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_mycoides_DSM_2048_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_mycoides_DSM_2048_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_mycoides_DSM_2048_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_pseudomycoides_DSM_12442_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_pseudomycoides_DSM_12442_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_pseudomycoides_DSM_12442_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_pseudomycoides_DSM_12442_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_psychrosaccharolyticus_DSM_6_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_psychrosaccharolyticus_DSM_6_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_psychrosaccharolyticus_DSM_6_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_psychrosaccharolyticus_DSM_6_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_pumilus_DSM_13835_01</i>	Bacillales	<i>Bacillus</i>

<i>Bacillus_pumilus_DSM_13835_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_salarius_DSM16461_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_salarius_DSM16461_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_salarius_DSM16461_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_salarius_DSM16461_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_siamensis_25261_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_siamensis_25261_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_siamensis_25261_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_siamensis_25261_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_simplex_DSM_1321_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_simplex_DSM_1321_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_simplex_DSM_1321_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_simplex_DSM_1321_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_sonorensis_DSM_13779_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_sonorensis_DSM_13779_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_sonorensis_DSM_13779_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_sonorensis_DSM_13779_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_sporothermodurans_DSM_10599_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_sporothermodurans_DSM_10599_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_sporothermodurans_DSM_10599_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_sporothermodurans_DSM_10599_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis_DSM_347_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis_DSM_347_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis_DSM_347_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis_DSM_347_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_thuringiensis_DSM_6073_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_thuringiensis_DSM_6073_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_thuringiensis_DSM_6073_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_thuringiensis_DSM_6073_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_vallismortis_DSM_11031_01</i>	Bacillales	<i>Bacillus</i>

<i>Bacillus_vallismortis_DSM_11031_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_vallismortis_DSM_11031_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_vallismortis_DSM_11031_04</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_weihenstephanensis_DSM_104119_01</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_weihenstephanensis_DSM_104119_02</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_weihenstephanensis_DSM_104119_03</i>	Bacillales	<i>Bacillus</i>
<i>Bacillus_weihenstephanensis_DSM_104119_04</i>	Bacillales	<i>Bacillus</i>
<i>Brucella_abortus_S19_delta_01</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_abortus_S19_delta_02</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_abortus_S19_delta_03</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_abortus_S19_delta_04</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_canis_A183_5_01</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_canis_A183_5_02</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_canis_A183_5_03</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_canis_A183_5_04</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_melitensis_A146_13_Ringversuch_A938_3_01</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_melitensis_A146_13_Ringversuch_A938_3_02</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_melitensis_A146_13_Ringversuch_A938_3_03</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_melitensis_A146_13_Ringversuch_A938_3_04</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_microti_CCM_4915_01</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_microti_CCM_4915_02</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_neotomae_A148_7_01</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_neotomae_A148_7_02</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_neotomae_A148_7_03</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_neotomae_A148_7_04</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_ovis_A138_7_01</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_ovis_A138_7_02</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_ovis_A138_7_03</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_ovis_A138_7_04</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_pinnipedialis_A148_8_01</i>	Rhizobiales	<i>Brucella</i>

<i>Brucella_pinnipedialis_A148_8_02</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_pinnipedialis_A148_8_03</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_pinnipedialis_A148_8_04</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_vulpis_DSM_101715_01</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_vulpis_DSM_101715_02</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_vulpis_DSM_101715_03</i>	Rhizobiales	<i>Brucella</i>
<i>Brucella_vulpis_DSM_101715_04</i>	Rhizobiales	<i>Brucella</i>
<i>Burkholderia_caledonica_LMG_19076_01</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_caledonica_LMG_19076_02</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_caledonica_LMG_19076_03</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_caribensis_LMG_18531_01</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_caribensis_LMG_18531_02</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_caribensis_LMG_18531_03</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_ambifaria_DSM_16087_01</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_ambifaria_DSM_16087_02</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_ambifaria_DSM_16087_03</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_ambifaria_DSM_16087_04</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_anthina_DSM_16086_01</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_anthina_DSM_16086_02</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_anthina_DSM_16086_03</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_anthina_DSM_16086_04</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_arboris_DSM_23435_01</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_arboris_DSM_23435_02</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_arboris_DSM_23435_03</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_arboris_DSM_23435_04</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_cenocepacia_DSM_16553_01</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_cenocepacia_DSM_16553_02</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_cenocepacia_DSM_16553_03</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_cenocepacia_DSM_16553_04</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_CC_cepacia_DSM_7288_01</i>	Burkholderiales	<i>Burkholderia</i>



<i>Burkholderia</i> _CC_cepacia_DSM_7288_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_cepacia_DSM_7288_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_contaminans_DSM_22706_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_contaminans_DSM_22706_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_contaminans_DSM_22706_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_contaminans_DSM_22706_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_diffusa_DSM_23434_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_diffusa_DSM_23434_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_diffusa_DSM_23434_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_diffusa_DSM_23434_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_dolosa_DSM_16088_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_dolosa_DSM_16088_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_dolosa_DSM_16088_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_dolosa_DSM_16088_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_lata_DSM_23089_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_lata_DSM_23089_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_lata_DSM_23089_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_lata_DSM_23089_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_latens_DSM_23436_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_latens_DSM_23436_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_latens_DSM_23436_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_latens_DSM_23436_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_metallica_DSM_23519_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_metallica_DSM_23519_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_metallica_DSM_23519_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_metallica_DSM_23519_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_multivorans_ATCC_17616_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_multivorans_ATCC_17616_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_multivorans_ATCC_17616_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_pseudomultivorans_LMG_16669_01	Burkholderiales	<i>Burkholderia</i>

<i>Burkholderia</i> _CC_pseudomultivorans_LMG_16669_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_pseudomultivorans_LMG_16669_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_pseudomultivorans_LMG_16669_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_puraquae_CAMPA_565_HST_189_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_puraquae_CAMPA_565_HST_189_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_puraquae_CAMPA_565_HST_189_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_puraquae_CAMPA_565_HST_189_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_pyrrocinia_ATCC_15958_DSM_10685_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_pyrrocinia_ATCC_15958_DSM_10685_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_pyrrocinia_ATCC_15958_DSM_10685_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_pyrrocinia_ATCC_15958_DSM_10685_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_seminalis_DSM_23518_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_seminalis_DSM_23518_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_seminalis_DSM_23518_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_seminalis_DSM_23518_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_stabilis_DSM_16586_LMG_14294_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_stabilis_DSM_16586_LMG_14294_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_stabilis_DSM_16586_LMG_14294_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_stabilis_DSM_16586_LMG_14294_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_stagnalis_LMG_28156_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_stagnalis_LMG_28156_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_stagnalis_LMG_28156_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_stagnalis_LMG_28156_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_territorii_LMG_28158_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_territorii_LMG_28158_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_territorii_LMG_28158_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_territorii_LMG_28158_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_ubonensis_DSM_17311_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_ubonensis_DSM_17311_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_ubonensis_DSM_17311_03	Burkholderiales	<i>Burkholderia</i>

<i>Burkholderia</i> _CC_ubonensis_DSM_17311_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_vietnamiensis_LMG_10929_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_vietnamiensis_LMG_10929_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_vietnamiensis_LMG_10929_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _CC_vietnamiensis_LMG_10929_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _gladioli_DSM_8361_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _gladioli_DSM_8361_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _gladioli_DSM_8361_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _gladioli_DSM_8361_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _glathei_LMG_14190_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _glathei_LMG_14190_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _glathei_LMG_14190_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _glumae_LMG_1277_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _glumae_LMG_1277_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _glumae_LMG_1277_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _hospita_LMG_20574_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _hospita_LMG_20574_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _hospita_LMG_20574_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _kururiensis_DSM_13464_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _kururiensis_DSM_13464_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _kururiensis_DSM_13464_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _kururiensis_DSM_13464_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _mallei_ATCC_23344_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _mallei_ATCC_23344_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _mallei_ATCC_23344_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _mallei_ATCC_23344_04	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _oklahomensis_DSM_21774_01	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _oklahomensis_DSM_21774_02	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _oklahomensis_DSM_21774_03	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia</i> _oklahomensis_DSM_21774_04	Burkholderiales	<i>Burkholderia</i>

<i>Burkholderia_phymatum_DSM_17167_01</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_phymatum_DSM_17167_02</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_phymatum_DSM_17167_03</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_phytofirmans_DSM_17436_01</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_phytofirmans_DSM_17436_02</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_phytofirmans_DSM_17436_03</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_plantarum_LMG_10907_01</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_plantarum_LMG_10907_02</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_plantarum_LMG_10907_03</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_pseudomallei_CCUC_13790_01</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_pseudomallei_CCUC_13790_02</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_pseudomallei_CCUC_13790_03</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_thailandensis_DSM_13276_01</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_thailandensis_DSM_13276_02</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_thailandensis_DSM_13276_03</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_tuberum_LMG_21444_01</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_tuberum_LMG_21444_02</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_tuberum_LMG_21444_03</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_xenovorans_DSM_17367_01</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_xenovorans_DSM_17367_02</i>	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_xenovorans_DSM_17367_03</i>	Burkholderiales	<i>Burkholderia</i>
<i>Campylobacter_jejuni_A669_01</i>	Campylobacterales	<i>Campylobacter</i>
<i>Campylobacter_jejuni_A669_02</i>	Campylobacterales	<i>Campylobacter</i>
<i>Campylobacter_jejuni_A669_03</i>	Campylobacterales	<i>Campylobacter</i>
<i>Campylobacter_jejuni_A669_04</i>	Campylobacterales	<i>Campylobacter</i>
<i>Citrobacter_amalonaticus_04_08695_01</i>	Enterobacteriales	<i>Citrobacter</i>
<i>Citrobacter_amalonaticus_04_08695_02</i>	Enterobacteriales	<i>Citrobacter</i>
<i>Citrobacter_diversus_ATCC_25408_01</i>	Enterobacteriales	<i>Citrobacter</i>
<i>Citrobacter_diversus_ATCC_25408_02</i>	Enterobacteriales	<i>Citrobacter</i>
<i>Citrobacter_freundii_DSM_30039_01</i>	Enterobacteriales	<i>Citrobacter</i>

<i>Citrobacter freundii</i> _DSM_30039_02	Enterobacteriales	<i>Citrobacter</i>
<i>Citrobacter freundii</i> _DSM_30039_03	Enterobacteriales	<i>Citrobacter</i>
<i>Citrobacter freundii</i> _DSM_30039_04	Enterobacteriales	<i>Citrobacter</i>
<i>Corynebacterium amycolatum</i> _RRLK02_02_01	Actinomycetales	<i>Corynebacterium</i>
<i>Corynebacterium amycolatum</i> _RRLK02_02_02	Actinomycetales	<i>Corynebacterium</i>
<i>Corynebacterium amycolatum</i> _RRLK02_02_03	Actinomycetales	<i>Corynebacterium</i>
<i>Corynebacterium amycolatum</i> _RRLK02_02_04	Actinomycetales	<i>Corynebacterium</i>
<i>Edwardsiella tarda</i> _DSM_30052_01	Enterobacteriales	<i>Edwardsiella</i>
<i>Edwardsiella tarda</i> _DSM_30052_02	Enterobacteriales	<i>Edwardsiella</i>
<i>Enterobacter aerogenes</i> _DSM_30053_01	Enterobacteriales	<i>Enterobacter</i>
<i>Enterobacter aerogenes</i> _DSM_30053_02	Enterobacteriales	<i>Enterobacter</i>
<i>Enterobacter cloacae</i> _DSM_30054_01	Enterobacteriales	<i>Enterobacter</i>
<i>Enterobacter cloacae</i> _DSM_30054_02	Enterobacteriales	<i>Enterobacter</i>
<i>Enterobacter gergoviae</i> _ATCC_33426_01	Enterobacteriales	<i>Enterobacter</i>
<i>Enterobacter gergoviae</i> _ATCC_33426_02	Enterobacteriales	<i>Enterobacter</i>
<i>Enterobacter sakazakii</i> _04_01242_01	Enterobacteriales	<i>Enterobacter</i>
<i>Enterobacter sakazakii</i> _04_01242_02	Enterobacteriales	<i>Enterobacter</i>
<i>Enterococcus faecalis</i> _DSM_20371_01	Enterobacteriales	<i>Enterococcus</i>
<i>Enterococcus faecalis</i> _DSM_20371_02	Enterobacteriales	<i>Enterococcus</i>
<i>Enterococcus faecalis</i> _DSM_20371_03	Enterobacteriales	<i>Enterococcus</i>
<i>Enterococcus faecalis</i> _DSM_20371_04	Enterobacteriales	<i>Enterococcus</i>
<i>Escherichia coli</i> _DSM_3871_01	Enterobacteriales	<i>Escherichia</i>
<i>Escherichia coli</i> _DSM_3871_02	Enterobacteriales	<i>Escherichia</i>
<i>Escherichia coli</i> _DSM_3871_03	Enterobacteriales	<i>Escherichia</i>
<i>Escherichia coli</i> _DSM_3871_04	Enterobacteriales	<i>Escherichia</i>
<i>Francisella guangzhouensis</i> _DSM_102975_01	Thiotrichales	<i>Francisella</i>
<i>Francisella guangzhouensis</i> _DSM_102975_02	Thiotrichales	<i>Francisella</i>
<i>Francisella guangzhouensis</i> _DSM_102975_03	Thiotrichales	<i>Francisella</i>
<i>Francisella guangzhouensis</i> _DSM_102975_04	Thiotrichales	<i>Francisella</i>
<i>Francisella hispaniensis</i> _DSM_22475_01	Thiotrichales	<i>Francisella</i>

<i>Francisella_hispaniensis_DSM_22475_02</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_hispaniensis_DSM_22475_03</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_hispaniensis_DSM_22475_04</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_naotunensis_ssp_orientalis_DSM_21254_01</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_naotunensis_ssp_orientalis_DSM_21254_02</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_naotunensis_ssp_orientalis_DSM_21254_03</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_naotunensis_ssp_orientalis_DSM_21254_04</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_tularensis_novicida_Ft26_01</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_tularensis_novicida_Ft26_02</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_tularensis_novicida_Ft26_03</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_tularensis_novicida_Ft26_04</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_tularensis_ssp_mediasiatica_Ft31_01</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_tularensis_ssp_mediasiatica_Ft31_02</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_tularensis_ssp_mediasiatica_Ft31_03</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_tularensis_ssp_mediasiatica_Ft31_04</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_tularensis_subsp_holarctica_A466_1_01</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_tularensis_subsp_holarctica_A466_1_02</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_tularensis_subsp_holarctica_A466_1_03</i>	Thiotrichales	<i>Francisella</i>
<i>Francisella_tularensis_subsp_holarctica_A466_1_04</i>	Thiotrichales	<i>Francisella</i>
<i>Inquilinus_limosus_DSM_16000_01</i>	Rhodospirillales	<i>Inquilinus</i>
<i>Inquilinus_limosus_DSM_16000_02</i>	Rhodospirillales	<i>Inquilinus</i>
<i>Inquilinus_limosus_DSM_16000_03</i>	Rhodospirillales	<i>Inquilinus</i>
<i>Inquilinus_limosus_DSM_16000_04</i>	Rhodospirillales	<i>Inquilinus</i>
<i>Klebsiella_oxytoca_ATCC_13182_01</i>	Enterobacteriales	<i>Klebsiella</i>
<i>Klebsiella_oxytoca_ATCC_13182_02</i>	Enterobacteriales	<i>Klebsiella</i>
<i>Klebsiella_pneumoniae_ssp_ozeanae_DSM_68_01</i>	Enterobacteriales	<i>Klebsiella</i>
<i>Klebsiella_pneumoniae_ssp_ozeanae_DSM_68_02</i>	Enterobacteriales	<i>Klebsiella</i>
<i>Kocuria_palustris_RRLK06_02_01</i>	Actinomycetales	<i>Kocuria</i>
<i>Kocuria_palustris_RRLK06_02_02</i>	Actinomycetales	<i>Kocuria</i>
<i>Kocuria_palustris_RRLK06_02_03</i>	Actinomycetales	<i>Kocuria</i>

<i>Kocuria_palustris_RRLK06_02_04</i>	Actinomycetales	<i>Kocuria</i>
<i>Lysinibacillus_fusiformis_B207_01</i>	Bacillales	<i>Lysinibacillus</i>
<i>Lysinibacillus_fusiformis_B207_02</i>	Bacillales	<i>Lysinibacillus</i>
<i>Lysinibacillus_fusiformis_B207_03</i>	Bacillales	<i>Lysinibacillus</i>
<i>Lysinibacillus_fusiformis_B207_04</i>	Bacillales	<i>Lysinibacillus</i>
<i>Lysinibacillus_sphaericus_DSM_396_01</i>	Bacillales	<i>Lysinibacillus</i>
<i>Lysinibacillus_sphaericus_DSM_396_02</i>	Bacillales	<i>Lysinibacillus</i>
<i>Micrococcus_luteus_RRLK07_01</i>	Actinomycetales	<i>Micrococcus</i>
<i>Micrococcus_luteus_RRLK07_02</i>	Actinomycetales	<i>Micrococcus</i>
<i>Micrococcus_luteus_RRLK07_03</i>	Actinomycetales	<i>Micrococcus</i>
<i>Micrococcus_luteus_RRLK07_04</i>	Actinomycetales	<i>Micrococcus</i>
<i>Ochrobactrum_anthropi_DSM_20150_01</i>	Rhizobiales	<i>Ochrobactrum</i>
<i>Ochrobactrum_anthropi_DSM_20150_02</i>	Rhizobiales	<i>Ochrobactrum</i>
<i>Ochrobactrum_anthropi_DSM_20150_03</i>	Rhizobiales	<i>Ochrobactrum</i>
<i>Ochrobactrum_anthropi_DSM_20150_04</i>	Rhizobiales	<i>Ochrobactrum</i>
<i>Oligella_urethralis_A691_2_01</i>	Burkholderiales	<i>Oligella</i>
<i>Oligella_urethralis_A691_2_02</i>	Burkholderiales	<i>Oligella</i>
<i>Paenibacillus_alvei_B210_01</i>	Bacillales	<i>Paenibacillus</i>
<i>Paenibacillus_alvei_B210_02</i>	Bacillales	<i>Paenibacillus</i>
<i>Paenibacillus_alvei_B210_03</i>	Bacillales	<i>Paenibacillus</i>
<i>Paenibacillus_alvei_B210_04</i>	Bacillales	<i>Paenibacillus</i>
<i>Paenibacillus_polymyxa_DSM_365_01</i>	Bacillales	<i>Paenibacillus</i>
<i>Paenibacillus_polymyxa_DSM_365_02</i>	Bacillales	<i>Paenibacillus</i>
<i>Paenibacillus_polymyxa_DSM_365_03</i>	Bacillales	<i>Paenibacillus</i>
<i>Paenibacillus_polymyxa_DSM_365_04</i>	Bacillales	<i>Paenibacillus</i>
<i>Pandoraea_apista_DSM_16535_01</i>	Burkholderiales	<i>Pandoraea</i>
<i>Pandoraea_apista_DSM_16535_02</i>	Burkholderiales	<i>Pandoraea</i>
<i>Pandoraea_apista_DSM_16535_03</i>	Burkholderiales	<i>Pandoraea</i>
<i>Pandoraea_apista_DSM_16535_04</i>	Burkholderiales	<i>Pandoraea</i>
<i>Pandoraea_pnomenusa_DSM_16536_01</i>	Burkholderiales	<i>Pandoraea</i>

<i>Pandoraea_pnomenusa_DSM_16536_02</i>	Burkholderiales	<i>Pandoraea</i>
<i>Pandoraea_pnomenusa_DSM_16536_03</i>	Burkholderiales	<i>Pandoraea</i>
<i>Pandoraea_pnomenusa_DSM_16536_04</i>	Burkholderiales	<i>Pandoraea</i>
<i>Pandoraea_pulmonicola_DSM_16583_01</i>	Burkholderiales	<i>Pandoraea</i>
<i>Pandoraea_pulmonicola_DSM_16583_02</i>	Burkholderiales	<i>Pandoraea</i>
<i>Pandoraea_pulmonicola_DSM_16583_03</i>	Burkholderiales	<i>Pandoraea</i>
<i>Pandoraea_pulmonicola_DSM_16583_04</i>	Burkholderiales	<i>Pandoraea</i>
<i>Pantoea_agglomerans_ATCC_27988_01</i>	Enterobacteriales	<i>Pantoea</i>
<i>Pantoea_agglomerans_ATCC_27988_02</i>	Enterobacteriales	<i>Pantoea</i>
<i>Paracoccus_yeei_RRLK05_01_01</i>	Rhodobacterales	<i>Paracoccus</i>
<i>Paracoccus_yeei_RRLK05_01_02</i>	Rhodobacterales	<i>Paracoccus</i>
<i>Paracoccus_yeei_RRLK05_01_03</i>	Rhodobacterales	<i>Paracoccus</i>
<i>Paracoccus_yeei_RRLK05_01_04</i>	Rhodobacterales	<i>Paracoccus</i>
<i>Proteus_inconstans_ATCC_25827_01</i>	Enterobacteriales	<i>Proteus</i>
<i>Proteus_inconstans_ATCC_25827_02</i>	Enterobacteriales	<i>Proteus</i>
<i>Proteus_mirabilis_SM_788_01</i>	Enterobacteriales	<i>Proteus</i>
<i>Proteus_mirabilis_SM_788_02</i>	Enterobacteriales	<i>Proteus</i>
<i>Proteus_morganii_DSM_30117_01</i>	Enterobacteriales	<i>Proteus</i>
<i>Proteus_morganii_DSM_30117_02</i>	Enterobacteriales	<i>Proteus</i>
<i>Proteus_vulgaris_ATCC_33420_01</i>	Enterobacteriales	<i>Proteus</i>
<i>Proteus_vulgaris_ATCC_33420_02</i>	Enterobacteriales	<i>Proteus</i>
<i>Pseudomonas_aeruginosa_ATCC_27853_01</i>	Pseudomonadales	<i>Pseudomonas</i>
<i>Pseudomonas_aeruginosa_ATCC_27853_02</i>	Pseudomonadales	<i>Pseudomonas</i>
<i>Pseudomonas_aeruginosa_ATCC_27853_03</i>	Pseudomonadales	<i>Pseudomonas</i>
<i>Pseudomonas_aeruginosa_ATCC_27853_04</i>	Pseudomonadales	<i>Pseudomonas</i>
<i>Pseudomonas_chlororaphis_ATCC_17809_01</i>	Pseudomonadales	<i>Pseudomonas</i>
<i>Pseudomonas_chlororaphis_ATCC_17809_02</i>	Pseudomonadales	<i>Pseudomonas</i>
<i>Pseudomonas_chlororaphis_ATCC_17809_03</i>	Pseudomonadales	<i>Pseudomonas</i>
<i>Pseudomonas_chlororaphis_ATCC_17809_04</i>	Pseudomonadales	<i>Pseudomonas</i>
<i>Ralstonia_insidiosa_DSM_17714_01</i>	Burkholderiales	<i>Ralstonia</i>



<i>Ralstonia_insidiosa_DSM_17714_02</i>	Burkholderiales	<i>Ralstonia</i>
<i>Ralstonia_insidiosa_DSM_17714_03</i>	Burkholderiales	<i>Ralstonia</i>
<i>Ralstonia_insidiosa_DSM_17714_04</i>	Burkholderiales	<i>Ralstonia</i>
<i>Ralstonia_mannitolilytica_DSM_17512_01</i>	Burkholderiales	<i>Ralstonia</i>
<i>Ralstonia_mannitolilytica_DSM_17512_02</i>	Burkholderiales	<i>Ralstonia</i>
<i>Ralstonia_mannitolilytica_DSM_17512_03</i>	Burkholderiales	<i>Ralstonia</i>
<i>Ralstonia_mannitolilytica_DSM_17512_04</i>	Burkholderiales	<i>Ralstonia</i>
<i>Ralstonia_pickettii_DSM_6297_01</i>	Burkholderiales	<i>Ralstonia</i>
<i>Ralstonia_pickettii_DSM_6297_02</i>	Burkholderiales	<i>Ralstonia</i>
<i>Ralstonia_pickettii_DSM_6297_03</i>	Burkholderiales	<i>Ralstonia</i>
<i>Ralstonia_pickettii_DSM_6297_04</i>	Burkholderiales	<i>Ralstonia</i>
<i>Salmonella_enterica_serovar_minnesota_SF_1111_01</i>	Enterobacteriales	<i>Salmonella</i>
<i>Salmonella_enterica_serovar_minnesota_SF_1111_02</i>	Enterobacteriales	<i>Salmonella</i>
<i>Salmonella_enteritidis_LT21_01</i>	Enterobacteriales	<i>Salmonella</i>
<i>Salmonella_enteritidis_LT21_02</i>	Enterobacteriales	<i>Salmonella</i>
<i>Salmonella_hadar_01</i>	Enterobacteriales	<i>Salmonella</i>
<i>Salmonella_hadar_02</i>	Enterobacteriales	<i>Salmonella</i>
<i>Salmonella_typhimurium_SH_9178_01</i>	Enterobacteriales	<i>Salmonella</i>
<i>Salmonella_typhimurium_SH_9178_02</i>	Enterobacteriales	<i>Salmonella</i>
<i>Serratia_grimesii_DSM_30063_01</i>	Enterobacteriales	<i>Salmonella</i>
<i>Serratia_grimesii_DSM_30063_02</i>	Enterobacteriales	<i>Salmonella</i>
<i>Serratia_marcescens_DSM_30121_01</i>	Enterobacteriales	<i>Salmonella</i>
<i>Serratia_marcescens_DSM_30121_02</i>	Enterobacteriales	<i>Salmonella</i>
<i>Shigella_boydii_01</i>	Enterobacteriales	<i>Salmonella</i>
<i>Shigella_boydii_02</i>	Enterobacteriales	<i>Salmonella</i>
<i>Shigella_flexneri_3A_01</i>	Enterobacteriales	<i>Salmonella</i>
<i>Shigella_flexneri_3A_02</i>	Enterobacteriales	<i>Salmonella</i>
<i>Staphylococcus_aureus_DSM_20231_01</i>	Bacillales	<i>Staphylococcus</i>
<i>Staphylococcus_aureus_DSM_20231_02</i>	Bacillales	<i>Staphylococcus</i>
<i>Staphylococcus_aureus_DSM_20231_03</i>	Bacillales	<i>Staphylococcus</i>

<i>Staphylococcus_aureus_DSM_20231_04</i>	Bacillales	<i>Staphylococcus</i>
<i>Staphylococcus_epidermidis_DSM_1798_01</i>	Bacillales	<i>Staphylococcus</i>
<i>Staphylococcus_epidermidis_DSM_1798_02</i>	Bacillales	<i>Staphylococcus</i>
<i>Staphylococcus_epidermidis_DSM_1798_03</i>	Bacillales	<i>Staphylococcus</i>
<i>Staphylococcus_epidermidis_DSM_1798_04</i>	Bacillales	<i>Staphylococcus</i>
<i>Staphylococcus_hominis_RRLK01_03_01</i>	Bacillales	<i>Staphylococcus</i>
<i>Staphylococcus_hominis_RRLK01_03_02</i>	Bacillales	<i>Staphylococcus</i>
<i>Staphylococcus_hominis_RRLK01_03_03</i>	Bacillales	<i>Staphylococcus</i>
<i>Staphylococcus_hominis_RRLK01_03_04</i>	Bacillales	<i>Staphylococcus</i>
<i>Stenotrophomonas_maltophilia_Sm_36_01</i>	Xanthomonadales	<i>Stenotrophomonas</i>
<i>Stenotrophomonas_maltophilia_Sm_36_02</i>	Xanthomonadales	<i>Stenotrophomonas</i>
<i>Stenotrophomonas_maltophilia_Sm_36_03</i>	Xanthomonadales	<i>Stenotrophomonas</i>
<i>Stenotrophomonas_rhizophila_RRLK03_01_01</i>	Xanthomonadales	<i>Stenotrophomonas</i>
<i>Stenotrophomonas_rhizophila_RRLK03_01_02</i>	Xanthomonadales	<i>Stenotrophomonas</i>
<i>Stenotrophomonas_rhizophila_RRLK03_01_03</i>	Xanthomonadales	<i>Stenotrophomonas</i>
<i>Stenotrophomonas_rhizophila_RRLK03_01_04</i>	Xanthomonadales	<i>Stenotrophomonas</i>
<i>Streptococcus_pyogenes_DSM_20565_01</i>	Lactobacillales	<i>Streptococcus</i>
<i>Streptococcus_pyogenes_DSM_20565_02</i>	Lactobacillales	<i>Streptococcus</i>
<i>Streptococcus_pyogenes_DSM_20565_03</i>	Lactobacillales	<i>Streptococcus</i>
<i>Streptococcus_pyogenes_DSM_20565_04</i>	Lactobacillales	<i>Streptococcus</i>
<i>Vibrio_alginolyticus_DSM_2171_01</i>	Enterobacteriales	<i>Vibrio</i>
<i>Vibrio_cholerae_A220_01</i>	Enterobacteriales	<i>Vibrio</i>
<i>Vibrio_cholerae_A220_02</i>	Enterobacteriales	<i>Vibrio</i>
<i>Vibrio_cholerae_A220_03</i>	Enterobacteriales	<i>Vibrio</i>
<i>Vibrio_cholerae_A220_04</i>	Enterobacteriales	<i>Vibrio</i>
<i>Vibrio_diazotrophicus_174_2_01</i>	Enterobacteriales	<i>Vibrio</i>
<i>Vibrio_mimicus_A177_1_01</i>	Enterobacteriales	<i>Vibrio</i>
<i>Vibrio_parahaemolyticus_177_2_01</i>	Enterobacteriales	<i>Vibrio</i>
<i>Vibrio_vulnificus_A177_3_02</i>	Enterobacteriales	<i>Vibrio</i>
<i>Xenophilus_sp_A395_01</i>	Burkholderiales	<i>Xenophilus</i>

<i>Xenophilus_sp_A395_02</i>	Burkholderiales	<i>Xenophilus</i>
<i>Yersinia_aldovae_DSM_18303_01</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_aldovae_DSM_18303_02</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_aldovae_DSM_18303_03</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_aleksiciae_DSM_14987_01</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_aleksiciae_DSM_14987_02</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_aleksiciae_DSM_14987_03</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_bercovieri_DSM_18528_01</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_bercovieri_DSM_18528_02</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_bercovieri_DSM_18528_03</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_enterocolitica_DSM_9676_01</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_enterocolitica_DSM_9676_02</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_enterocolitica_DSM_9676_03</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_entomophaga_DSM_22339_01</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_entomophaga_DSM_22339_02</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_entomophaga_DSM_22339_03</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_entomophaga_DSM_22339_04</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_frederiksenii_DSM_18490_01</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_frederiksenii_DSM_18490_02</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_frederiksenii_DSM_18490_03</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_intermedia_DSM_18517_01</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_intermedia_DSM_18517_02</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_intermedia_DSM_18517_03</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_kristensenii_DSM_18543_01</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_kristensenii_DSM_18543_02</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_kristensenii_DSM_18543_03</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_massiliensis_DSM_21859_01</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_massiliensis_DSM_21859_02</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_massiliensis_DSM_21859_03</i>	Enterobacteriales	<i>Yersinia</i>
<i>Yersinia_massiliensis_DSM_21859_04</i>	Enterobacteriales	<i>Yersinia</i>

<i>Yersinia_mollaretii_DSM_18520_01</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_mollaretii_DSM_18520_02</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_mollaretii_DSM_18520_03</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_nurmii_DSM_22296_01</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_nurmii_DSM_22296_02</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_nurmii_DSM_22296_03</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_nurmii_DSM_22296_04</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_pekkanenii_DSM_22769_01</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_pekkanenii_DSM_22769_02</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_pekkanenii_DSM_22769_03</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_pekkanenii_DSM_22769_04</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_pestis_10329_01</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_pestis_10329_04</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_pseudotuberculosis_DSM_8992_01</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_pseudotuberculosis_DSM_8992_02</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_pseudotuberculosis_DSM_8992_03</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_rohdei_DSM_18270_01</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_rohdei_DSM_18270_02</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_rohdei_DSM_18270_03</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_ruckeri_DSM_18506_01</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_ruckeri_DSM_18506_02</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_ruckeri_DSM_18506_03</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_similis_DSM_18211_01</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_similis_DSM_18211_02</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_similis_DSM_18211_03</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_similis_DSM_18211_04</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_wautersii_DSM_27350_01</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_wautersii_DSM_27350_02</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_wautersii_DSM_27350_03</i>	Enterobacterales	<i>Yersinia</i>
<i>Yersinia_wautersii_DSM_27350_04</i>	Enterobacterales	<i>Yersinia</i>

TABLE S2. SUMMARY TABLE OF STUDIED BACTERIAL STRAINS (BSNB)

<b>Names</b>	<b>ID</b>	<b>BSNB</b>	<b>Order</b>	<b>Genus</b>
<i>Bacillus_cereus</i>	GTC2112	BSNB_0016	Bacillales	<i>Bacillus</i>
<i>Brevibacillus_sp</i>	GCYPB01	BSNB_0034	Bacillales	<i>Brevibacillus</i>
<i>Bacillus_sp</i>	GTC2802	BSNB_0053	Bacillales	<i>Bacillus</i>
<i>Serratia_sp</i>	CN1	BSNB_0139	Enterobacteriales	<i>Serratia</i>
<i>Serratia_marcescens</i>	CN4	BSNB_0142	Enterobacteriales	<i>Serratia</i>
<i>Burkholderia_seminalis</i>	CN5A	BSNB_0143	Burkholderiales	<i>Burkholderia</i>
<i>Pseudomonas_beteli</i>	CN12	BSNB_0150	Pseudomonadales	<i>Pseudomonas</i>
<i>Serratia_marcescens</i>	CN14	BSNB_0152	Enterobacteriales	<i>Serratia</i>
<i>Achromobacter_sp</i>	CN15	BSNB_0153	Burkholderiales	<i>Achromobacter</i>
<i>Burkholderia_sp</i>	CN20	BSNB_0158	Burkholderiales	<i>Burkholderia</i>
<i>Serratia_marcescens</i>	CN27	BSNB_0165	Enterobacteriales	<i>Serratia</i>
<i>Burkholderia_sp</i>	CN28	BSNB_0166	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_pyrrocinia</i>	CN32	BSNB_0170	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_sp</i>	CN33	BSNB_0171	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_cepacia</i>	CN34	BSNB_0172	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_sp</i>	CN38	BSNB_0178	Burkholderiales	<i>Burkholderia</i>
<i>Lactococcus_garvieae</i>	CN39	BSNB_0179	Lactobacillales	<i>Lactococcus</i>
<i>Lactococcus_garvieae</i>	CN40	BSNB_0180	Lactobacillales	<i>Lactococcus</i>
<i>Burkholderia_arboris</i>	CN41	BSNB_0181	Burkholderiales	<i>Burkholderia</i>
<i>Enterobacter_tabaci</i>	CN42A	BSNB_0182	Enterobacteriales	<i>Enterobacter</i>
<i>Enterobacter_tabaci</i>	CN51	BSNB_0191	Enterobacteriales	<i>Enterobacter</i>
<i>Enterobacter_tabaci</i>	CN52	BSNB_0192	Enterobacteriales	<i>Enterobacter</i>
<i>Pseudomonas_sp</i>	CN54	BSNB_0194	Pseudomonadales	<i>Pseudomonas</i>
<i>Kitasatospora_sp</i>	CN61	BSNB_0202	Actinomycetales	<i>Kitasatospora</i>
<i>Bacillus_cereus</i>	CN89	BSNB_0230	Bacillales	<i>Bacillus</i>
<i>Bacillus_cereus</i>	CN91	BSNB_0232	Bacillales	<i>Bacillus</i>
<i>Lysinibacillus_xylanilyticus</i>	CN92	BSNB_0233	Bacillales	<i>Lysinibacillus</i>
<i>Streptomyces_sp</i>	CN93	BSNB_0234	Actinomycetales	<i>Streptomyces</i>

<i>Bacillus_firmus</i>	CN94	BSNB_0235	Bacillales	<i>Bacillus</i>
<i>Bacillus_sp</i>	CN95	BSNB_0236	Bacillales	<i>Bacillus</i>
<i>Lysinibacillus_sp</i>	CN98	BSNB_0239	Bacillales	<i>Lysinibacillus</i>
<i>Enterobacter_tabaci</i>	CN99	BSNB_0240	Enterobacteriales	<i>Enterobacter</i>
<i>Serratia_marcescens</i>	CN101	BSNB_0242	Enterobacteriales	<i>Serratia</i>
<i>Lysinibacillus_xylanilyticus</i>	CN104	BSNB_0245	Bacillales	<i>Lysinibacillus</i>
<i>Burkholderia_sp</i>	CN105	BSNB_0246	Burkholderiales	<i>Burkholderia</i>
<i>Streptomyces_malaysiense</i>	CN115	BSNB_0256	Actinomycetales	<i>Streptomyces</i>
<i>Paraburkholderia_tropica</i>	A1F1 2	BSNB_0493	Burkholderiales	<i>Paraburkholderia</i>
<i>Klebsiella_michiganensis</i>	A1F1 3	BSNB_0494	Enterobacteriales	<i>Klebsiella</i>
<i>Enterobacter_asburiae</i>	A2F_1 1.1	BSNB_0517	Enterobacteriales	<i>Enterobacter</i>
<i>Burkholderia_tropica</i>	A2F_1 3	BSNB_0519	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_sp</i>	A2F_1 5.2	BSNB_0521	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_tropica</i>	A2F11 4	BSNB_0547	Burkholderiales	<i>Burkholderia</i>
<i>Bacillus_thuringiensis</i>	A3F_1 2	BSNB_0561	Bacillales	<i>Bacillus</i>
<i>Paraburkholderia_tropica_</i>	A3F_1 3	BSNB_0562	Burkholderiales	<i>Paraburkholderia</i>
<i>Paraburkholderia_tropica_</i>	A3F_1 6	BSNB_0565	Burkholderiales	<i>Paraburkholderia</i>
<i>Bacillus_subtilis</i>	A3F1 1	BSNB_0566	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	A3F1 8.2	BSNB_0567	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	A3F1 3	BSNB_0568	Bacillales	<i>Bacillus</i>
<i>Pantoea_dispersa</i>	A3F1 4.1	BSNB_0569	Enterobacteriales	<i>Pantoea</i>
<i>Burkholderia_tropica</i>	A3F1 5	BSNB_0570	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_tropica</i>	A3F1 7	BSNB_0572	Burkholderiales	<i>Burkholderia</i>
<i>Bacillus_subtilis</i>	A3F1 8.111	BSNB_0573	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	A3F1 2	BSNB_0576	Bacillales	<i>Bacillus</i>
<i>Klebsiella_variicola</i>	A4F1 1	BSNB_0591	Enterobacteriales	<i>Klebsiella</i>
<i>Klebsiella_variicola</i>	A4F1 2	BSNB_0592	Enterobacteriales	<i>Klebsiella</i>
<i>Burkholderia_tropica</i>	A4F1 3	BSNB_0593	Burkholderiales	<i>Burkholderia</i>
<i>Klebsiella_variicola</i>	A4F1 4	BSNB_0594	Enterobacteriales	<i>Klebsiella</i>
<i>Klebsiella_variicola</i>	A4F1 5	BSNB_0595	Enterobacteriales	<i>Klebsiella</i>

<i>Burkholderia_plantarum</i>	A4F1 6	BSNB_0596	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_plantarum</i>	A4F1 7	BSNB_0597	Burkholderiales	<i>Burkholderia</i>
<i>Klebsiella_variicola</i>	A4F1 9	BSNB_0598	Enterobacteriales	<i>Klebsiella</i>
<i>Burkholderia_sp</i>	A5F_1 2	BSNB_0616	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_arvi</i>	A5F_1 4	BSNB_0617	Burkholderiales	<i>Burkholderia</i>
<i>Bacillus_cereus</i>	A5F_1 6	BSNB_0618	Bacillales	<i>Bacillus</i>
<i>Burkholderia_sp</i>	A5F_1 3	BSNB_0619	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_tropica</i>	A5F1 8	BSNB_0626	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_tropica</i>	A5F10 2	BSNB_0630	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_tropica</i>	A5F10 4	BSNB_0632	Burkholderiales	<i>Burkholderia</i>
<i>Paraburkholderia_guayanaensis</i>	A5F10 7	BSNB_0635	Burkholderiales	<i>Paraburkholderia</i>
<i>Burkholderia_sp</i>	A5F10 8	BSNB_0636	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_tropica</i>	A5F10 11	BSNB_0639	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_oxypila</i>	A5F10 12	BSNB_0640	Burkholderiales	<i>Burkholderia</i>
<i>Paenibacillus_glucanolyticus</i>	A6F_1 1	BSNB_0643	Bacillales	<i>Paenibacillus</i>
<i>Bacillus_sp</i>	A6F1 4	BSNB_0648	Bacillales	<i>Bacillus</i>
<i>Burkholderia_tropica</i>	A6F16 4	BSNB_0656	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_tropica</i>	A6F16 6	BSNB_0658	Burkholderiales	<i>Burkholderia</i>
<i>Bacillus_subtilis</i>	A6F16 7.1	BSNB_0659	Bacillales	<i>Bacillus</i>
<i>Burkholderia_tropica</i>	A6F16 8	BSNB_0661	Burkholderiales	<i>Burkholderia</i>
<i>Burkholderia_tropica</i>	A6F16 10	BSNB_0663	Burkholderiales	<i>Burkholderia</i>
<i>Enterobacter_amnigenus</i>	A7F_1 1	BSNB_0669	Enterobacteriales	<i>Enterobacter</i>
<i>Pantoea_stewartii</i>	A7F_1 3	BSNB_0671	Enterobacteriales	<i>Pantoea</i>
<i>Pantoea_stewartii</i>	A7F_1 4	BSNB_0672	Enterobacteriales	<i>Pantoea</i>
<i>Enterobacter_mori</i>	A7F_1 7	BSNB_0675	Enterobacteriales	<i>Enterobacter</i>
<i>Bacillus_subtilis</i>	A7F1 5.1	BSNB_0683	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	A7F1 8	BSNB_0685	Bacillales	<i>Bacillus</i>
<i>Stenotrophomonas_maltophilia</i>	A8F_1 1	BSNB_0704	Xanthomonadales	<i>Stenotrophomonas</i>
<i>Enterobacter_mori</i>	A8F_1 3	BSNB_0706	Enterobacteriales	<i>Enterobacter</i>
<i>Enterobacter_mori</i>	A8F_1 5.1	BSNB_0708	Enterobacteriales	<i>Enterobacter</i>

<i>Enterobacter_mori</i>	A8F_1 6.1	BSNB_0709	Enterobacteriales	<i>Enterobacter</i>
<i>Bacillus_subtilis</i>	A8F_1 8	BSNB_0712	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	A8F1 1	BSNB_0716	Bacillales	<i>Bacillus</i>
<i>Luteibacter_yeojuensis</i>	A8F1 6.1	BSNB_0719	Xanthomonadales	<i>Luteibacter</i>
<i>Luteibacter_yeojuensis</i>	A8F1 7	BSNB_0721	Xanthomonadales	<i>Luteibacter</i>
<i>Bacillus_subtilis</i>	A8F1 8	BSNB_0722	Bacillales	<i>Bacillus</i>
<i>Enterobacter_mori</i>	A8F11 1	BSNB_0723	Enterobacteriales	<i>Enterobacter</i>
<i>Enterobacter_mori</i>	A8F11 3	BSNB_0725	Enterobacteriales	<i>Enterobacter</i>
<i>Enterobacter_mori</i>	A8F11 4	BSNB_0726	Enterobacteriales	<i>Enterobacter</i>
<i>Enterobacter_mori</i>	A8F11 5	BSNB_0727	Enterobacteriales	<i>Enterobacter</i>
<i>Bacillus_subtilis</i>	A8F11 8	BSNB_0730	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	C1F9 10	BSNB_0762	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	P1F1 1	BSNB_0803	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	P1F1 2	BSNB_0804	Bacillales	<i>Bacillus</i>
<i>Methylobacterium_mesophilicum</i>	P1F1 3	BSNB_0805	Rhizobiales	<i>Methylobacterium</i>
<i>Bacillus_subtilis</i>	P1F5 8	BSNB_0816	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	P1F5 9	BSNB_0817	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	W2F_2 1	BSNB_0955	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	W2F_2 3	BSNB_0956	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	W2F_2 9	BSNB_0959	Bacillales	<i>Bacillus</i>
<i>Acidisoma_sp</i>	W3F_3 8	BSNB_0978	Rhodospirillales	<i>Acidosoma</i>
<i>Burkholderia_tropica</i>	AC A2 F11 9.1B	BSNB_1020	Burkholderiales	<i>Burkholderia</i>
<i>Methylobacterium_populi</i>	VECD13D	BSNB_1022	Rhizobiales	<i>Methylobacterium</i>
<i>Curtobacterium_sp</i>	VECD14A	BSNB_1023	Actinomycetales	<i>Curtobacterium</i>
<i>Pantoea_dispersa</i>	VECD14B	BSNB_1024	Enterobacteriales	<i>Pantoea</i>
<i>Bacillus_megaterium</i>	VECD14C	BSNB_1025	Bacillales	<i>Bacillus</i>
<i>Methylobacterium_sp</i>	VECD14G	BSNB_1026	Rhizobiales	<i>Methylobacterium</i>
<i>Asaia_sp</i>	VECD15A	BSNB_1027	Rhodospirillales	<i>Asaia</i>
<i>Pseudomonas_sp</i>	ST5C	BSNB_1030	Pseudomonadales	<i>Pseudomonas</i>



<i>Arthrobacter_sp</i>	ST8C	BSNB_1032	Actinomycetales	<i>Arthrobacter</i>
<i>Pantoea_agglomerans</i>	ST16C	BSNB_1034	Enterobacteriales	<i>Pantoea</i>
<i>Methylobacterium_mesophilicum</i>	LR_FP43	BSNB_1116	Rhizobiales	<i>Methylobacterium</i>
<i>Methylobacterium_mesophilicum</i>	LR_FP51	BSNB_1123	Rhizobiales	<i>Methylobacterium</i>
<i>Methylobacterium_mesophilicum</i>	LR_FP67	BSNB_1136	Rhizobiales	<i>Methylobacterium</i>
<i>Methylobacterium_mesophilicum</i>	LR_FP68	BSNB_1137	Rhizobiales	<i>Methylobacterium</i>
<i>Methylobacterium_mesophilicum</i>	LR_FP74	BSNB_1141	Rhizobiales	<i>Methylobacterium</i>
<i>Rhodococcus_kroppenstedtii</i>	LR_FP135	BSNB_1179	Actinomycetales	<i>Rhodococcus</i>
<i>Rhodococcus_kroppenstedtii</i>	LR_FP150	BSNB_1190	Actinomycetales	<i>Rhodococcus</i>
<i>Microbacterium_testaceum</i>	LR_FP151	BSNB_1191	Actinomycetales	<i>Microbacterium</i>
<i>Microbacterium_testaceum</i>	LR_FP152	BSNB_1192	Actinomycetales	<i>Microbacterium</i>
<i>Bacillus_subtilis</i>	LR_FP171	BSNB_1207	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	LR_FP172	BSNB_1208	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	LR_FP174	BSNB_1209	Bacillales	<i>Bacillus</i>
<i>Bacillus_subtilis</i>	LR_FP175	BSNB_1210	Bacillales	<i>Bacillus</i>
<i>Rhodococcus_cerastii</i>	LR_EL13	BSNB_1220	Actinomycetales	<i>Rhodococcus</i>
<i>Bacillus_subtilis</i>	LR_EL109	BSNB_1238	Bacillales	<i>Bacillus</i>
<i>Rhodococcus_cerastii</i>	LR_EL133	BSNB_1246	Actinomycetales	<i>Rhodococcus</i>
<i>Klebsiella_michiganensis</i>	A1F2A1	NO_BSNB	Enterobacteriales	<i>Klebsiella</i>
<i>Klebsiella_michiganensis</i>	A4F1_12	NO_BSNB	Enterobacteriales	<i>Klebsiella</i>
<i>Methylobacterium_mesophilicum</i>	LRFP_069	NO_BSNB	Rhizobiales	<i>Methylobacterium</i>

TABLE S3. SUMMARY TABLE OF STUDIED FUNGAL STRAINS (BSNB)

<b>Names</b>	<b>ID</b>	<b>BSNB</b>	<b>Order</b>	<b>Genus</b>
<i>Epicoccum_nigrum</i>	GTG01	BSNB_0001	Pleosporales	<i>Epicoccum</i>
<i>Colletotrichum_gloeosporioides</i>	GTC2001	BSNB_0002	Glomerellales	<i>Colletotrichum</i>
<i>Pestalotiopsis_sp</i>	GTC2102	BSNB_0006	Xylariales	<i>Pestalotiopsis</i>
<i>Pestalotiopsis_sp</i>	GTC2104	BSNB_0008	Xylariales	<i>Pestalotiopsis</i>
<i>Chaetomium_globosum</i>	GTC2114	BSNB_0018	Sordariales	<i>Chaetomium</i>
<i>Diaporthe_phaseolorum</i>	GTC2201	BSNB_0020	Diaporthales	<i>Diaporthe</i>
<i>Xylaria_cubensis</i>	GPP02	BSNB_0022	Xylariales	<i>Xylaria</i>
<i>Mycoleptodiscus_sp</i>	GTC2304	BSNB_0024	Magnaporthales	<i>Mycoleptodiscus</i>
<i>Colletotrichum_siamense</i>	GTC2303	BSNB_0025	Glomerellales	<i>Colletotrichum</i>
<i>Mycoleptodiscus_sp</i>	GTC2304	BSNB_0026	Magnaporthales	<i>Mycoleptodiscus</i>
<i>Colletotrichum_gloeosporioides</i>	GTC2401	BSNB_0029	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_boninense</i>	GTC2502	BSNB_0032	Glomerellales	<i>Colletotrichum</i>
<i>Diaporthe_phaseolorum</i>	GTC2503	BSNB_0033	Diaporthales	<i>Diaporthe</i>
<i>Colletotrichum_boninense</i>	GTC2601	BSNB_0035	Glomerellales	<i>Colletotrichum</i>
<i>Xylaria_cubensis</i>	GTC2604	BSNB_0038	Xylariales	<i>Xylaria</i>
<i>Colletotrichum_gloeosporioides</i>	GTC2605	BSNB_0039	Glomerellales	<i>Colletotrichum</i>
<i>Xylaria_sp</i>	GTC2702	BSNB_0041	Xylariales	<i>Xylaria</i>
<i>Guignardia_mangiferae</i>	GTC2703	BSNB_0042	Botryosphaerales	<i>Guignardia</i>
<i>Pestalotiopsis_sp</i>	GTC2704	BSNB_0043	Xylariales	<i>Pestalotiopsis</i>
<i>Diaporthe_sp</i>	GTC2705	BSNB_0044	Diaporthales	<i>Diaporthe</i>
<i>Diaporthe_sp</i>	GTC2708	BSNB_0047	Diaporthales	<i>Diaporthe</i>
<i>Xylariales_sp</i>	GTC2709	BSNB_0048	Xylariales	<i>Xylaria</i>
<i>Xylariales_sp</i>	GTC2711	BSNB_0050	Xylariales	<i>Xylaria</i>
<i>Penicillium_sp</i>	GVL02	BSNB_0051	Eurotiales	<i>Penicillium</i>
<i>Phomopsis_sp</i>	GTC2801	BSNB_0052	Diaporthales	<i>Phomopsis</i>
<i>Phomopsis_phyllanthicola</i>	GTC2804	BSNB_0055	Diaporthales	<i>Phomopsis</i>
<i>Xylaria_cubensis</i>	GTC2805	BSNB_0056	Xylariales	<i>Xylaria</i>
<i>Phomopsis_sp</i>	GTC2806	BSNB_0057	Diaporthales	<i>Phomopsis</i>

<i>Diaporthe_phaseolorum</i>	GTC2808	BSNB_0059	Diaporthales	<i>Diaporthe</i>
<i>Diaporthe_phaseolorum</i>	GTC2809	BSNB_0060	Diaporthales	<i>Diaporthe</i>
<i>Phomopsis_sp</i>	GTC2812	BSNB_0063	Diaporthales	<i>Phomopsis</i>
<i>Phomopsis_sp</i>	GTC2820	BSNB_0071	Diaporthales	<i>Phomopsis</i>
<i>Chaetomium_sp</i>	GTC2901	BSNB_0074	Sordariales	<i>Chaetomium</i>
<i>Xylaria_sp</i>	GTC2902	BSNB_0075	Xylariales	<i>Xylaria</i>
<i>Acremonium_sp</i>	GTC2903	BSNB_0076	Hypocreales	<i>Acremonium</i>
<i>Pestalotiopsis_microspora</i>	GTC2904	BSNB_0077	Xylariales	<i>Pestalotiopsis</i>
<i>Chaetomium_sp</i>	GTC2905	BSNB_0078	Sordariales	<i>Chaetomium</i>
<i>Colletotrichum_gloeosporioides</i>	GTC3003	BSNB_0085	Glomerellales	<i>Colletotrichum</i>
<i>Guignardia_mangiferae</i>	GSS01	BSNB_0087	Botryosphaeriales	<i>Guignardia</i>
<i>Xylaria_cubensis</i>	GSS02	BSNB_0088	Xylariales	<i>Xylaria</i>
<i>Xylaria_cubensis</i>	GSS04	BSNB_0089	Xylariales	<i>Xylaria</i>
<i>Xylaria_cubensis</i>	GSS05	BSNB_0090	Xylariales	<i>Xylaria</i>
<i>Xylaria_cubensis</i>	GSS06	BSNB_0091	Xylariales	<i>Xylaria</i>
<i>Colletotrichum_sp</i>	GSS08	BSNB_0093	Glomerellales	<i>Colletotrichum</i>
<i>Diaporthe_phaseolorum</i>	GSS09	BSNB_0094	Diaporthales	<i>Diaporthe</i>
<i>Diaporthe_sp</i>	GSS11	BSNB_0096	Diaporthales	<i>Diaporthe</i>
<i>Colletotrichum_sp</i>	GSS13	BSNB_0098	Glomerellales	<i>Colletotrichum</i>
<i>Diaporthe_sp</i>	GSS15	BSNB_0100	Diaporthales	<i>Diaporthe</i>
<i>Pestalotiopsis_sp</i>	GCI01	BSNB_0102	Xylariales	<i>Pestalotiopsis</i>
<i>Xylaria_cubensis</i>	GCI02	BSNB_0103	Xylariales	<i>Xylaria</i>
<i>Xylaria_cubensis</i>	LD1.6	BSNB_0106	Xylariales	<i>Xylaria</i>
<i>Muscodor_sp</i>	LD1.5	BSNB_0107	Xylariales	<i>Muscodor</i>
<i>Guignardia_mangiferae</i>	LD2.13	BSNB_0115	Botryosphaeriales	<i>Guignardia</i>
<i>Colletotrichum_gloeosporioides</i>	LD3.4	BSNB_0118	Glomerellales	<i>Colletotrichum</i>
<i>Xylaria_cubensis</i>	LD3.6.2	BSNB_0121	Xylariales	<i>Xylaria</i>
<i>Cystobasidium_minutum</i>	LR01	BSNB_0122	Saccharomycetales	<i>Cystobasidium</i>
<i>Xylaria_sp</i>	LD4.11	BSNB_0123	Xylariales	<i>Xylaria</i>
<i>Colletotrichum_sp</i>	LD4.3.2	BSNB_0124	Glomerellales	<i>Colletotrichum</i>

<i>Candida_etchellsii</i>	LD5.7	BSNB_0125	Saccharomycetales	<i>Candida</i>
<i>Colletotrichum_gloeosporioides</i>	LD8.10	BSNB_0132	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	LD8.6	BSNB_0133	Glomerellales	<i>Colletotrichum</i>
<i>Guignardia_mangiferae</i>	LD8.9	BSNB_0135	Botryosphaeriales	<i>Guignardia</i>
<i>Rhizomucor_variabilis</i>	CN16	BSNB_0154	Mucorales	<i>Rhizomucor</i>
<i>Rhizomucor_variabilis</i>	CN17	BSNB_0155	Mucorales	<i>Rhizomucor</i>
<i>Penicillium_shearrii</i>	CN29	BSNB_0167	Eurotiales	<i>Penicillium</i>
<i>Penicillium_simplicissimum</i>	CN30	BSNB_0168	Eurotiales	<i>Penicillium</i>
<i>Candida_tropicalis</i>	CN36A	BSNB_0174	Saccharomycetales	<i>Candida</i>
<i>Rhizomucor_variabilis</i>	CN36B	BSNB_0175	Mucorales	<i>Rhizomucor</i>
<i>Rhizomucor_variabilis</i>	CN37B	BSNB_0177	Mucorales	<i>Rhizomucor</i>
<i>Cladosporium_sp</i>	CN56	BSNB_0196	Capnodiales	<i>Cladosporium</i>
<i>Penicillium_sanguifluum</i>	CN57	BSNB_0197	Eurotiales	<i>Penicillium</i>
<i>Exophiala_xenobiotica</i>	CN60bis	BSNB_0201	Chaetothyriales	<i>Exophiala</i>
<i>Pseudallescheria_boydii</i>	CN71	BSNB_0212	Microascales	<i>Pseudallescheria</i>
<i>Scedosporium_boydii</i>	CN75	BSNB_0216	Microascales	<i>Scedosporium</i>
<i>Penicillium_citrinum</i>	CN77	BSNB_0218	Eurotiales	<i>Penicillium</i>
<i>Paecilomyces_formosus</i>	CN79	BSNB_0220	Eurotiales	<i>Paecilomyces</i>
<i>Scedosporium_boydii</i>	CN81	BSNB_0222	Microascales	<i>Scedosporium</i>
<i>Paecilomyces_formosus</i>	CN83	BSNB_0224	Eurotiales	<i>Paecilomyces</i>
<i>Pestalotiopsis_microspora</i>	CN84	BSNB_0225	Xylariales	<i>Pestalotiopsis</i>
<i>Pseudallescheria_boydii</i>	CN85	BSNB_0226	Microascales	<i>Pseudallescheria</i>
<i>Aspergillus_aculeatus</i>	CN86	BSNB_0227	Eurotiales	<i>Aspergillus</i>
<i>Penicillium_rubidurum</i>	CN87	BSNB_0228	Eurotiales	<i>Penicillium</i>
<i>Penicillium_araracuarensense</i>	CN102	BSNB_0243	Eurotiales	<i>Penicillium</i>
<i>Penicillium_araracuarensense</i>	CN103	BSNB_0244	Eurotiales	<i>Penicillium</i>
<i>Aspergillus_sp</i>	CN109	BSNB_0250	Eurotiales	<i>Aspergillus</i>
<i>Penicillium_sp</i>	CN112	BSNB_0253	Eurotiales	<i>Penicillium</i>

<i>Paecilomyces_formosus</i>	CN122	BSNB_0263	Eurotiales	<i>Paecilomyces</i>
<i>Rhizomucor_variabilis</i>	CN125	BSNB_0266	Mucorales	<i>Rhizomucor</i>
<i>Acrodictys_sp</i>	A3F8 11.1	BSNB_0283	Hypocreales	<i>Acrodictys</i>
<i>Ochroconis_sp</i>	A5F_1 3	BSNB_0284	Venturiales	<i>Ochroconis</i>
<i>Ochroconis_sp</i>	A5F_1 5	BSNB_0285	Venturiales	<i>Ochroconis</i>
<i>Sordariomycetes_sp</i>	A5F1 13	BSNB_0287	Sordariales	<i>Sordariomycetes</i>
<i>Paramicrothyrium_sp</i>	A5F1 9	BSNB_0289	Microthyriales	<i>Paramicrothyrium</i>
<i>Colletotrichum_gloeosporioides</i>	A5F1 12	BSNB_0290	Glomerellales	<i>Colletotrichum</i>
<i>Apioclypea_sp</i>	A5F1 15	BSNB_0292	Xylariales	<i>Apioclypea</i>
<i>Penicillium_sp</i>	A5F1 10.1	BSNB_0293	Eurotiales	<i>Penicillium</i>
<i>Chaetomium_sp</i>	A6F1 9	BSNB_0295	Sordariales	<i>Chaetomium</i>
<i>Chaetomium_sp</i>	A6F1 12	BSNB_0296	Sordariales	<i>Chaetomium</i>
<i>Penicillium_sp</i>	A6F1 13	BSNB_0298	Eurotiales	<i>Penicillium</i>
<i>Muscodor_sp</i>	A6F1 19	BSNB_0299	Xylariales	<i>Muscodor</i>
<i>Xylaria_sp</i>	A6F1 8.2	BSNB_0300	Xylariales	<i>Xylaria</i>
<i>Albonectria_rigidiuscula</i>	A8F_1 9.1	BSNB_0301	Hypocreales	<i>Albonectria</i>
<i>Fusarium_sp</i>	A8F_1 9.2	BSNB_0302	Hypocreales	<i>Fusarium</i>
<i>Fusarium_sp</i>	A8F_1 13	BSNB_0303	Hypocreales	<i>Fusarium</i>
<i>Sordariomycetes_sp</i>	A8F1 2	BSNB_0304	Sordariales	<i>Sordariomycetes</i>
<i>Chaetomium_sp</i>	A8F1 5	BSNB_0305	Sordariales	<i>Chaetomium</i>
<i>Oxydothis_sp</i>	A1F12 12	BSNB_0516	Xylariales	<i>Oxydothis</i>
<i>Diaporthe_sp</i>	A2F_1 4.1	BSNB_0520	Diaporthales	<i>Diaporthe</i>
<i>Stereum_hirsutum</i>	A2F_1 7	BSNB_0523	Russulales	<i>Stereum</i>
<i>Pseudopestalotiopsis_sp</i>	A2F_1 10	BSNB_0526	Xylariales	<i>Pseudopestalotiopsis</i>
<i>Endomelanconiopsis_endophytica</i>	A2F1 1	BSNB_0528	Botryosphaerales	<i>Endomelanconiopsis</i>
<i>Colletotrichum_gloeosporioides</i>	A2F1_2	BSNB_0529	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	A2F1 3.1	BSNB_0530	Glomerellales	<i>Colletotrichum</i>
<i>Stereum_hirsutum</i>	A2F1 3.2	BSNB_0531	Russulales	<i>Stereum</i>
<i>Stereum_hirsutum</i>	A2F1 4.1	BSNB_0532	Russulales	<i>Stereum</i>
<i>Nodulisporium_sp</i>	A2F1 6	BSNB_0534	Xylariales	<i>Nodulisporium</i>

<i>Colletotrichum_gloeosporioides</i>	A2F1 8	BSNB_0536	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	A2F1 9	BSNB_0537	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	A2F1 10.22	BSNB_0538	Glomerellales	<i>Colletotrichum</i>
<i>Nemania_sp</i>	A2F1 11	BSNB_0539	Xylariales	<i>Nemania</i>
<i>Colletotrichum_gloeosporioides</i>	A2F1 12	BSNB_0540	Glomerellales	<i>Colletotrichum</i>
<i>Diaporthe_sp</i>	A2F1 13	BSNB_0541	Diaporthales	<i>Diaporthe</i>
<i>Clonostachys_sp</i>	A2F11 3	BSNB_0546	Hypocreales	<i>Clonostachys</i>
<i>Stereum_hirsutum</i>	A2F11 6.11	BSNB_0532	Russulales	<i>Stereum</i>
<i>Colletotrichum_sp</i>	A2F11 6.21	BSNB_0549	Glomerellales	<i>Colletotrichum</i>
<i>Stereum_hirsutum</i>	A2F11 7	BSNB_0550	Russulales	<i>Stereum</i>
<i>Colletotrichum_gloeosporioides</i>	A2F11_8	BSNB_0551	Glomerellales	<i>Colletotrichum</i>
<i>Neopestalotiopsis_sp</i>	A2F11 12.1	BSNB_0556	Xylariales	<i>Neopestalotiopsis</i>
<i>Colletotrichum_vietnamense</i>	A2F11 13	BSNB_0557	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	A2F11 15	BSNB_0559	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	A3F1 8.121	BSNB_0574	Glomerellales	<i>Colletotrichum</i>
<i>Fusarium_concolor</i>	A3F1 8.13	BSNB_0575	Hypocreales	<i>Fusarium</i>
<i>Diaporthe_sp</i>	A3F1 9	BSNB_0577	Diaporthales	<i>Diaporthe</i>
<i>Diaporthe_sp</i>	A3F1 10	BSNB_0578	Diaporthales	<i>Diaporthe</i>
<i>Cladosporium_cladosporioides</i>	A3F1 11	BSNB_0579	Capnodiales	<i>Cladosporium</i>
<i>Colletotrichum_gloeosporioides</i>	A3F8 1	BSNB_0580	Glomerellales	<i>Colletotrichum</i>
<i>Fusarium_sp</i>	A3F8 2	BSNB_0581	Hypocreales	<i>Fusarium</i>
<i>Endomelanconiopsis_endophytica</i>	A3F8 3	BSNB_0582	Botryosphaerales	<i>Endomelanconiopsis</i>
<i>Colletotrichum_sp</i>	A3F8 4	BSNB_0583	Glomerellales	<i>Colletotrichum</i>
<i>Endomelanconiopsis_endophytica</i>	A3F8 6	BSNB_0584	Botryosphaerales	<i>Endomelanconiopsis</i>
<i>Xylaria_sp</i>	A3F8 5.2	BSNB_0586	Xylariales	<i>Xylaria</i>
<i>Endomelanconiopsis_sp</i>	A3F8 8	BSNB_0587	Botryosphaerales	<i>Endomelanconiopsis</i>
<i>Endomelanconiopsis_endophytica</i>	A3F8 9	BSNB_0588	Botryosphaerales	<i>Endomelanconiopsis</i>
<i>Cladosporium_sp</i>	A3F8 10	BSNB_0589	Capnodiales	<i>Cladosporium</i>
<i>Colletotrichum_gloeosporioides</i>	A3F8 12	BSNB_0590	Glomerellales	<i>Colletotrichum</i>

<i>Colletotrichum_fruticola</i>	A5F_1 1	BSNB_0615	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	A5F1 3	BSNB_0622	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	A5F1 3	BSNB_0623	Glomerellales	<i>Colletotrichum</i>
<i>Thanatephorus_cucumeris</i>	A5F1 5	BSNB_0624	Cantharellales	<i>Thanatephorus</i>
<i>Colletotrichum_gloeosporioides</i>	A5F1 6.1	BSNB_0625	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_sp</i>	A5F1 11	BSNB_0627	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_theobromicola</i>	A5F1 17	BSNB_0628	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_theobromicola</i>	A5F10 9	BSNB_0637	Glomerellales	<i>Colletotrichum</i>
<i>Cladosporium_cladosporioides</i>	A5F10 10	BSNB_0638	Capnodiales	<i>Cladosporium</i>
<i>Colletotrichum_gloeosporioides</i>	A5F10 13	BSNB_0641	Glomerellales	<i>Colletotrichum</i>
<i>Hypoxyton_investiens</i>	A5F10 14.1	BSNB_0642	Xylariales	<i>Hypoxyton</i>
<i>Diaporthe_sp</i>	A6F_1 2	BSNB_0644	Diaporthales	<i>Diaporthe</i>
<i>Colletotrichum_theobromicola</i>	A6F_1 3	BSNB_0645	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	A6F1 1	BSNB_0646	Glomerellales	<i>Colletotrichum</i>
<i>Fusarium_sp</i>	A6F1 2	BSNB_0647	Hypocreales	<i>Fusarium</i>
<i>Colletotrichum_gloeosporioides</i>	A6F1 5.1	BSNB_0649	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	A6F1 7	BSNB_0650	Glomerellales	<i>Colletotrichum</i>
<i>Fusarium_sp</i>	A6F1 10	BSNB_0651	Hypocreales	<i>Fusarium</i>
<i>Colletotrichum_theobromicola</i>	A6F1 11	BSNB_0652	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	A6F1 15.2	BSNB_0653	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	A6F1 16.1	BSNB_0654	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	A6F16 2	BSNB_0655	Glomerellales	<i>Colletotrichum</i>
<i>Thanatephorus_cucumeris</i>	A6F16 7.2	BSNB_0660	Cantharellales	<i>Thanatephorus</i>
<i>Colletotrichum_gloeosporioides</i>	A6F16 9.1	BSNB_0662	Glomerellales	<i>Colletotrichum</i>
<i>Acutodesmus_obliquus</i>	A6F16 7.2	BSNB_0664	Sphaeropleales	<i>Acutodesmus</i>
<i>Nodulisporium_sp</i>	A6F16 13	BSNB_0666	Xylariales	<i>Nodulisporium</i>
<i>Cladosporium_herbarum</i>	A6F16 15	BSNB_0668	Capnodiales	<i>Cladosporium</i>
<i>Fusarium_lateritium</i>	A7F_1 10	BSNB_0678	Hypocreales	<i>Fusarium</i>
<i>Diaporthe_sp</i>	A7F1 1	BSNB_0679	Diaporthales	<i>Diaporthe</i>

<i>Endomelanconiopsis_endophytica</i>	A7F1 2	BSNB_0680	Botryosphaerales	<i>Endomelanconiopsis</i>
<i>Endomelanconiopsis_endophytica</i>	A7F1 3	BSNB_0681	Botryosphaerales	<i>Endomelanconiopsis</i>
<i>Colletotrichum_siamense</i>	A7F1 4	BSNB_0682	Glomerellales	<i>Colletotrichum</i>
<i>Endomelanconiopsis_endophytica</i>	A7F1 6	BSNB_0684	Botryosphaerales	<i>Endomelanconiopsis</i>
<i>Fusarium_avenaceum</i>	A7F1 9	BSNB_0686	Hypocreales	<i>Fusarium</i>
<i>Diaporthe_sp</i>	A7F1 10	BSNB_0687	Diaporthales	<i>Diaporthe</i>
<i>Colletotrichum_gloeosporioides</i>	A7F1 11	BSNB_0688	Glomerellales	<i>Colletotrichum</i>
<i>Penicillium_verrucosum</i>	A7F10 1	BSNB_0691	Eurotiales	<i>Penicillium</i>
<i>Endomelanconiopsis_endophytica</i>	A7F10 2	BSNB_0692	Botryosphaerales	<i>Endomelanconiopsis</i>
<i>Endomelanconiopsis_endophytica</i>	A7F10 3	BSNB_0693	Botryosphaerales	<i>Endomelanconiopsis</i>
<i>Colletotrichum_theobromicola</i>	A7F10 6.1	BSNB_0694	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_theobromicola</i>	A7F10 7.1	BSNB_0695	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_gloeosporioides</i>	A7F10 8	BSNB_0696	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_theobromicola</i>	A7F10 11	BSNB_0699	Glomerellales	<i>Colletotrichum</i>
<i>Apioclypea_sp</i>	A7F10 12	BSNB_0700	Xylariales	<i>Apioclypea</i>
<i>Colletotrichum_gloeosporioides</i>	A7F10 13.1	BSNB_0701	Glomerellales	<i>Colletotrichum</i>
<i>Fusarium_lateritium</i>	A7F10 13.2	BSNB_0702	Hypocreales	<i>Fusarium</i>
<i>Colletotrichum_theobromicola</i>	A7F10 14.2	BSNB_0703	Glomerellales	<i>Colletotrichum</i>
<i>Fusarium_lateritium</i>	A8F_1 4	BSNB_0707	Hypocreales	<i>Fusarium</i>
<i>Harknessia_australiensis</i>	A8F1 3	BSNB_0717	Xylariales	<i>Harknessia</i>
<i>Xylaria_sp</i>	A8F1 4	BSNB_0718	Xylariales	<i>Xylaria</i>
<i>Oxydothis_sp</i>	A8F1 6.2	BSNB_0720	Xylariales	<i>Oxydothis</i>
<i>Endomelanconiopsis_endophytica</i>	A8F11 9	BSNB_0731	Botryosphaerales	<i>Endomelanconiopsis</i>
<i>Akanthomyces_attenuatus</i>	A8F11 10	BSNB_0732	Hypocreales	<i>Akanthomyces</i>
<i>Endomelanconiopsis_endophytica</i>	A8F11 11	BSNB_0733	Botryosphaerales	<i>Endomelanconiopsis</i>
<i>Colletotrichum_gloeosporioides</i>	AC A2 F11 9.21	BSNB_1019	Glomerellales	<i>Colletotrichum</i>
<i>Colletotrichum_theobromicola</i>	VECD4B	BSNB_1021	Glomerellales	<i>Colletotrichum</i>
<i>Pestalotiopsis_sp</i>	VECD4A	BSNB_1036	Xylariales	<i>Pestalotiopsis</i>
<i>Pestalotiopsis_sp</i>	VECD4C	BSNB_1037	Xylariales	<i>Pestalotiopsis</i>



<i>Beauveria_bassiana</i>	VECD6B	BSNB_1038	Hypocreales	<i>Beauveria</i>
<i>Trichoderma_sp</i>	VECD6D	BSNB_1040	Hypocreales	<i>Trichoderma</i>
<i>Beauveria_bassiana</i>	VECD6E	BSNB_1041	Hypocreales	<i>Beauveria</i>
<i>Beauveria_bassiana</i>	VECD11A	BSNB_1042	Hypocreales	<i>Beauveria</i>
<i>Mucor_sp</i>	VECD11D	BSNB_1043	Mucorales	<i>Mucor</i>
<i>Penicillium_sp</i>	VECD11E	BSNB_1044	Eurotiales	<i>Penicillium</i>
<i>Penicillium_sp</i>	VECD11F	BSNB_1045	Eurotiales	<i>Penicillium</i>
<i>Penicillium_simplicissimum</i>	VECD11G	BSNB_1046	Eurotiales	<i>Penicillium</i>
<i>Neosartorya_sp</i>	VECD13B	BSNB_1048	Eurotiales	<i>Neosartorya</i>
<i>Penicillium_oxalicum</i>	VECD13C	BSNB_1049	Eurotiales	<i>Penicillium</i>
<i>Beauveria_bassiana</i>	VECD16B	BSNB_1053	Hypocreales	<i>Beauveria</i>
<i>Pestalotiopsis_theae</i>	VECD14E	BSNB_1054	Xylariales	<i>Pestalotiopsis</i>
<i>Fusarium_sp</i>	ST1C	BSNB_1055	Hypocreales	<i>Fusarium</i>
<i>Lecanicillium_sp</i>	ST4C	BSNB_1056	Hypocreales	<i>Lecanicillium</i>
<i>Fusarium_sp</i>	ST7C	BSNB_1057	Hypocreales	<i>Fusarium</i>
<i>Fusarium_sp</i>	ST9C	BSNB_1058	Hypocreales	<i>Fusarium</i>
<i>Arthrinium_sp</i>	ST10C	BSNB_1059	Sordariales	<i>Arthrinium</i>
<i>Penicillium_sp</i>	ST12C	BSNB_1061	Hypocreales	<i>Penicillium</i>
<i>Mucor_hiemalis</i>	ST15C	BSNB_1063	Mucorales	<i>Mucor</i>
<i>Fusarium_sp</i>	ST18C	BSNB_1065	Hypocreales	<i>Fusarium</i>
<i>Penicillium_sp</i>	ST21C	BSNB_1067	Eurotiales	<i>Penicillium</i>
<i>Beauveria_sp</i>	CLL14070	BSNB_1069	Hypocreales	<i>Beauveria</i>
<i>Isaria_farinosa</i>	CLL10038	BSNB_1250	Hypocreales	<i>Isaria</i>
<i>Cordyceps_militaris</i>	CLL10030	BSNB_1252	Hypocreales	<i>Cordyceps</i>
<i>Ophiocordyceps_forquignonii</i>	CLL11056	BSNB_1253	Hypocreales	<i>Ophiocordyceps</i>

TABLE S4. 16S OF STUDIED BACTERIAL STRAINS (BSNB)

>BSNB\_0016\_ *Bacillus\_cereus*

CCCGCGTCGCATTAGCTAGTTGGTGAGGTAACGGCTCACCAAGGCAACGATGCGTAGCCGA  
CCTGAGAGGGTGATCGGCCACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCA  
GCAGTAGGGAATCTTCCGCAATGGACGAAAGTCTGACGGAGCAACGCCGCGTGAGTGATGA  
AGGCTTTCGGGTCGTAAACTCTGTTGTTAGGGAAGAACAAGTGCTAGTTGAATAAGCTGGC  
ACCTTGACGGTACCTAACCAGAAAGCCACGGCTAACTACGTGCCAGCAGCCGCGGTAAATAC  
GTAGGTGGCAAGCGTTATCCGGAATTATTGGGCGTAAAGCGCGCGCAGGTGGTTTCTTAAGT  
CTGATGTGAAAGCCCACGGCTCAACCGTGGAGGGTCATTGGAAACTGGGAGACTTGAGTGC  
AGAAGAGGAAAGTGGAATTCATGTGTAGCGGTGAAATGCGTAGAGATATGGAGGAACAC  
CAGTGGCGAAGGCGACTTCTGGTCTGTAAGTACACTGAGGCGCGAAAGCGTGGGGAGCA  
AACAGGATTAGATACCCTGGTAGTCCACGCCGTAAACGATGAGTGCTAAGTGTTAGGGGGT  
TTCCGCCCTTTAGTGCTGAAGTTAACGCATTAAGCACTCCGCC

>BSNB\_0034\_ *Brevibacillus\_sp*

GGCGGGCTGCTAATACATGCAAGTCGAGCGAGTCTCTTCGGAGGCTAGCGGCGGACGGGTG  
AGTAACACGTAGGCAACCTGCCTCTCAGACTGGGATAACATAGGGAACTTATGCTAATAC  
CGGATAGGTTTTTGGACCGCATGGTCCGAAAAGAAAAGATGGCTTCGGCTATCACTGGGAG  
ATGGGCCTGCGGCGCATTAGCTAGTTGGTGGGGTAACGGCCTACCAAGGCGACGATGCGTA  
GCCGACCTGAGAGGGTGACCGGCCACACTGGGACTGAGACACGGCCCAGACTCCTACGGG  
AGGCAGCAGTAGGGAATTTCCACAATGGACGAAAGTCTGATGGAGCAACGCCGCGTGAAC  
GATGAAGGTCTTCGATTGTAAAGTTCTGTTGTTAGGGACGAATAAGTACCGTTCTGAATAGG  
GCGGTACCTTGACGGTACCTGACGAGAAAGCCACGGCTAACTACGTGCCAGCAGCCGCGGT  
AATACA

>BSNB\_0053\_ *Bacillus\_sp*

GACAGTGGCGGGGTGCTTACACATGCAAGTCGAACGGAAAGGCCCTGCTTTTGTGGGGTGT  
CGTGTGGAGAAGGGGTGTGTAAACCGTGTGTACCCTGCCCTTGTCTTTGGGATAACTTCGGG  
AAACTGGGGCTAATACCGGATAAGAGCTCCTGCTGCGTGGTGGGGGTGGAAAGTTTCTGCG  
GGGGGATGGACTCGCGGCTTATCTATTTGTTGGTGGGGTAGTGGCTTACCAAGGCTTTGAC  
GGGTAGCCGGCCTGACAGGGTGGCCGACCACTTTGGGACTGAGATACAGCCCACACTCATA  
CCGGAGGCAGCAGTGGGGAATATTTATAGATGGGCGGAAGCCTGATGCACTGCGCCGCGTG  
CGGGATGACGGCCTTAAGGTTGTAAACCCCTTTACCTGTGACTAAGCGTGAATGACGGGTA  
TGGGTAAATAAGCACCGCCTAACTACATGCCATCAGACGCCCTAATATAACACCGTGCCAG  
CAGCCGCGGTAATACAAAAAAGGGGGGT

>BSNB\_0139\_ *Serratia\_sp*

GCGGCGGCTTACACATGCAAGTCGAGCGGTAGCACAAAGGAGCTTGCTCYCTGGGTGACGA  
GCGGCGGACGGGTGAGTAATGTCTGGGAACTGCCTGATGGAGGGGGATAACTACTGGAAA  
CGGTAGCTAATACCGCATAACGTCGCARGACCAAAGAGGGGGACCTTCGGGCCTCTTGCCA  
TCAGATGTGCCAGATGGGATTAGCTAGTAGGTGGGGTAATGGCTCACCTAGGCAACRATSC  
CTAGMTGGTCTGAGAGGATGACCAGCCACACTGGAAGTGGAGACACGGTCCAKACTCCTACG  
GGAGGCAGCAGTGGGGAATATTGCACAATGGGCGCAAGCCTGATGCAGCCRTGCCGCGTGT  
GTGAAGAAGGCCTTCGGGTTGTAAAGCACTTTCAGCGAGRAGGAAGGTGGTGAAGTTAATA  
CGTTCATCAATTGACGTTACTCGAAAAAAGCACCGGCCAACTCCGTGCCAGCASCSSGR  
GTAATAMAGCGGCGGCTTACACATGCAAGTCGAGCGGTAGCACAAAGGAGCTTGCTCYCTG  
GGTGACGAGCGGCGGACGGGTGAGTAATGTCTGGGAACTGCCTGATGGAGGGGGATAACT  
ACTGGAAACGGTAGCTAATACCGCATAACGTCGCARGACCAAAGAGGGGGACCTTCGGGCC  
TCTTGCCATCAGATGTGCCAGATGGGATTAGCTAGTAGGTGGGGTAATGGCTCACCTAGGC  
AACRATSCCTAGMTGGTCTGAGAGGATGACCAGCCACACTGGAAGTGGAGACACGGTCCAKA

CTCCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGCAAGCCTGATGCAGCCRTG  
CCGCGTGTGTGAAGAAGGCCTTCGGGTTGTAAAGCACTTTCAGCGAGRAGGAAGGTGGTGA  
ACTTAATACGTTTCATCAATTGACGTTACTCGCAAAAAAAGCACCGGCCAACTCCGTGCCAGC  
ASCCGSGRGTAAATAMA

>BSNB\_0140\_ *Serratia\_marcescens*

GCGGGAGGCCTAACACATGCAAGCCGAGCGGTAGAGATCTTTCGGGATCTTGAGAGCGGCG  
TACGGGTGCGGAACACGTGTGCAACCTGCCTTTATCTGGGGGATAGCCTTTCGAAAGGAAG  
ATTAATACCCATAATATATCGACTGGCATCAGTTGATATTGAAAACCTCCGGTGGATAGAGA  
TGGGCACGCGCAAGATTAGATAGTTGGTGAGGTAACGGCTCACCAAGTCTGCGATCTTTAGG  
GGGCCTGAGAGGGTGATCCCCCACACTGGTACTGAGACACGGACCAGACTCCTACGGGAGG  
CAGCAGTGAGGAATATTGGACAATGGGTGAGAGCCTGATCCAGCCATCCCGCGTGAAGGAC  
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>BSNB\_0143\_ *Burkholderia\_seminalis*

TGCTCTTGGGTGACGAGTGGCGGACGGGTGAGTAATGTCTGGGGATCTGCCCGATAGAGGG  
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CGGTCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGCAAGCCTGAT  
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AGGCGACGGGGTTAATAACCTGTGATTGACGTTACCCGCAGAAGAAGCACCGGCTAACT  
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CTCAGGTGCGAAAGCGTGGGGAGCAAACAGGATTAGATACCCTGGTAGTCCACGCCGTAAA  
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>BSNB\_0150\_ *Pseudomonas\_beteli*

GAGTGAACGCTGGCGGTAGGCCTAACACATGCAAGTCGAACGGCAGCACAGGAGAGCTTG  
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GGTCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGGACAATGGGCGCAAGCCTGATC  
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ATCCAGCCGGCTAATACCTGGTTGGGATGACGGTACCCAAAGAATAAGCACCGGCTAACTT  
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GATGCGAACTGGATGTTGGGTGCAATTTGGCACGCAGTATCGAAGCTAACGCGTTAAGTTCCG  
CCG

>BSNB\_0152\_ *Serratia\_marcescens*

CGGCAGGCTTAACACATGCAAGTCGAGCGGTAGCACAAGGGAGCTTGCTCYCTGGGTGACG  
AGCGGCGGACGGGTGAGTAATGTCTGGGAACTGCCTGATGGAGGGGGATAACTACTGGAA  
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GTGTGAAGAAGGCCTTCGGGTTGTAAAGCACTTTCAGCGAGGAGGAAGGTGGTGAACCTAA  
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>BSNB\_0153\_ *Achromobacter\_sp*

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GCGGCCGATATCGGATTAGCTAGTTGGTGGGGTAACGGCTCACCAAGGCGACGATCCGTAG  
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GCAGCAGTGGGGAATTTTGGACAATGGGGGAAACCCTGATCCAGCCATCCCGCGTGTGCGA  
TGAAGGCCTTCGGGTTGTAAAGCACTTTTGGCAGGAAAGAAACGTCATGGGTAAATACCCC  
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>BSNB\_0158\_ *Burkholderia\_sp*

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GCAGTGGGGAATTTTGGACAATGGGCGAAAGCCTGATCCAGCAATGCCGCGTGTGTGAAGA  
AGGCCTTCGGGTTGTAAAGCACTTTTGTCCGGAAGAAATCCTTGCCCTAATACGGTCGGG  
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ACA

>BSNB\_0165\_ *Serratia\_marcescens*

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GTGTGAAGAAGGCCTTCGGGTTGTAAAGCACTTTCAGCGAGGAGGAAGGTGGTGARCTTAA  
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GGGGTAATACA

>BSNB\_0166\_ *Burkholderia\_sp*

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GCTGGTCTGAGAGGACGACCAGCCACACTGGGACTGAGACACGGCCCAGACTCCTACGGG  
AGGCAGCAGTGGGGAATTTTGGACAATGGGCGAAAGCCTGATCCAGCAATGCCGCGTGTGT  
GAAGAAGGCCTTCGGGTTGTAAAGCACTTTTGTCCGGAAAGAAATCCTTGGCCCTAATACGG  
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TAATACA

>BSNB\_0170\_ Burkholderia\_pyrrocinia

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TAATACA

>BSNB\_0171\_ Burkholderia\_sp

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GCTGGTCTGAGAGGACGACCAGCCACACTGGGACTGAGACACGGCCCAGACTCCTACSGGA  
GGCAGCAGTGGGGAATTTTGGACAATGGGCGAAAGCCTGATCCAGCMATGCCKCGTGTGTG  
AAGAAGGCCTTCGGGTTGTAAAGCACTTTTGTCCGGAAAGAAATCCTTGRCTCTAATACAGT  
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ATACA

>BSNB\_0172\_ Burkholderia\_cepacia

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CCGATGGCTGATTAGCTAGTTGGTGGGGTAAAGGCCTACCAAGGCGACGATCAGTAGCTGG  
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GCAGTGGGGAATTTTGGACAATGGGCGAAAGCCTGATCCAGCAATGCCGCGTGTGTGAAGA  
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CAA

>BSNB\_0178\_ Burkholderia\_sp

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GAGGCAGCAGTGGGGAATTTTGGACAATGGGCGAAAGCCTGATCCAGCAATGCCGCGTGTG  
TGAAGAAGGCCTTCGGGTTGTAAAGCACTTTTGTCCGGAAAGAAATCCTTGCTCTAATACA  
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GTAATACA

>BSNB\_0179\_ *Lactococcus\_garvieae*

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ATTTAAAAGAAGCAATTGCTTCACTACTTGATGATCCCGCGTTGTATTAGCTAGTTGGTAGTG  
TAAAGGACTACCAAGGCGATGATACATAGCCGACCTGAGAGGGTGATCGGCCACACTGGG  
ACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTAGGGAATCTTCGGCAATGGGGG  
CAACCCTGACCGAGCAACGCCGCGTGAGTGAAGAAGGTTTTCGGATCGTAAAACTCTGTTGT  
TAGAGAAGAACGTTAAGTAGAGTGGAATACTTAAGTGACGGTATCTAACCAGAAAGGG  
ACGGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGTCCCAAGCGTTGTCCGGATTTAT  
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GCGGTGAAATGCGTAGATATATGGAGGAACACCGGAGGCGAAAGCGGCTCTCTGGCCTGTA  
ACTGACACTGAGGCTCGAAAGCGTGCGGAGCAAACAGGATTAGATACCCTGGTAGTCCACG  
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AAGCACTCCGCCT

>BSNB\_0180\_ *Lactococcus\_garvieae*

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ATTGCTTCACTACTTGATGATCCCGCGTTGTATTAGCTAGTTGGTAGTGTAAGGACTACCAA  
GGCGATGATACATAGCCGACCTGAGAGGGTGATCGGCCACACTGGGACTGAGACACGGCCC  
AGACTCCTACGGGAGGCAGCAGTAGGGAATCTTCGGCAATGGGGGCAACCCTGACCGAGC  
AACGCCGCGTGAGTGAAGAAGGTTTTCGGATCGTAAAACTCTGTTGTTAGAGAAGAACGTT  
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>BSNB\_0181\_ *Burkholderia\_arboris*

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CCAGACTCCTACGGGAGGCAGCAGTGGGGAATTTTGGACAATGGGCGAAAGCCTGATCCAG  
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>BSNB\_0182\_ *Enterobacter*\_tabaci

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GATGCAATTGCATCACTCAAAGATGATCCCGCGTTGTATTAGCTAGTTGGTGAGGTAAAGGC  
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ACGGCCCAAACCTCCTACGGGAGGCAGCAGTAGGGAATCTTCGGCAATGGACGAAAGTCTGA  
CCGAGCAACGCCGCGTGAGTGAAGAAGGTTTTTCGGATCGTAAAACCTCTGTTGGTAGAGAAG  
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ATGCGTAGATATATGGAGGAACACCGGTGGCGAAAGCGGCTCTCTGGCCTGTAAGTACAC  
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>BSNB\_0191\_ *Enterobacter*\_tabaci

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GCCTAGGTCCGATTAGCTAGTTGGTGAGGTAATGGCTCACCAAGGCGACGATCCGTAAGT  
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AGGTCTTCGGATTGTAAAGCACTTTAAGTTGGGAGGAAGGGCAGTAAATTAATACTTTGCTG  
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A

>BSNB\_0192\_ *Enterobacter*\_tabaci

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ATCCCTAGCTGGTCTGAGAGGATGACCAGCCACACTGGAAGTGAAGACACGGTCCAGACTCC  
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GTGTATGAAGAAGGCCTTCGGGTTGTAAAGTACTTTCAGCGGGGAGGAAGGTGTTGAGGTTA  
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GCGGTAATACGGAGGGTGCAAGCGTTAATCGGAATTACTGGGCGTAAAGCGCACGCAGGCG  
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>BSNB\_0194\_ *Pseudomonas*\_sp

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AGGTCTTCGGATTGTAAAGCACTTTAAGTTGGGAGGAAGGGCATTAAACCTAATACGTTAGTG

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A

>BSNB\_0202\_ *Kitasatospora\_sp*

GGCGTGCTTACACATGCAAGTCGAACGGTGAAGCCCTTCGGGGTGGATCAGTGGCGAACGG  
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CAGTGGGGAATATTGCACAATGGGCGGAAGCCTGATGCAGCGACGCCGCGTGAGGGATGA  
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>BSNB\_0230\_ *Bacillus\_cereus*

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CAACGCCGCGTGAGTGATGAAGGCTTTCGGGTCGTAAAACCTCTGTNGTTAGGGAAGAACAA  
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TGCCAGCAGCCGCRGTAATACA

>BSNB\_0232\_ *Bacillus\_cereus*

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>BSNB\_0233\_ *Lysinibacillus\_xylanilyticus*

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>BSNB\_0234\_ *Streptomyces\_sp*

TGGSGGGCTGCTTACACATGCAAGTCGAACGATGAAGCCCTTCGGGGTGGATTAGTGGCGA  
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GAGCCCCGCGCCTATCAGCTTGTTGGTGAGGTAATGGCTCACCAAGGCGACGACGGGTAGC  
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GCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGGG  
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>BSNB\_0235\_ *Bacillus\_firmus*

GGCGGCGTGCTAATACATGCAAGTCGAGCGGAYTKMTGGGAGCTTGCTCCCTGAAGTCAG  
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GCGACGATGCGTAGCCGACCTGAGAGGGTGATCGGCCACACTGGGACTGAGACACGGCCC  
AGACTCCTACGGGAGGCAGCAGTAGGGAATCTTCCGCAATGGACGAAAGTCTGACGGAGC  
AACGCCGCGTGAGTGATGAAGGTTTTCGGATCGTAAACTCTGTTGTTAGGGAAGAACAAG  
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>BSNB\_0236\_ *Bacillus\_sp*

TGCGGCGTGCTAATACATGCAAGTCGAGCGAATGGATTAAGAGCTTGCTCTTATGAAGTTA  
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AACCGGGGCTAATACCGGATAACATTTTGAACYGCATGGTTGAAATTGAAAGGCGGCTTC  
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CAGACTCCTACGGGAGGCAGCAGTAGGGAATCTTCCGCAATGGACGAAAGTCTGACGGAGC  
AACGCCGCGTGAGTGATGAAGGCTTTCGGGTCGTAAACTCTGTTGTTAGGGAAGAACAAG  
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GCCAGCAGCCGCGGGTAATACA

>BSNB\_0239\_ *Lysinibacillus\_sp*

GGCGGCGTGCTAATACATGCAAGTCGAGCGAACAGAAAAGGAGCTTGCTCCTTTGACGTT  
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AACCGGGGCTAATACCGAATAATCTATTTCACTTCATGGTGAAATACTGAAAGACGGCATCT  
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CCAGACTCCTACGGGAGGCAGCAGTAGGGAATCTTCCACAATGGGCGAAAGCCTGATGGAG  
CAACGCCCCGTGAGTGAAGAAGGTTTTCGGATCGTAAACTCTGTTGTAAGGGAAGAACA  
GTACAGTAGTAACTGGCTGTACCTTGACGGTACCTTATTAGAAAGCCACGGCTAACTACGTG  
CCAGCAGCCGCGGKTAATACA

>BSNB\_0240\_ *Enterobacter\_tabaci*

GTGCTAACAATGCAAGTCGAAGCGGTATYACATAGKCCTTGCTTCCCTGGGAGTMRAGGGG

GGGAGGGGTGACCATGTGTGGGCCCACCTTTAAGATAACGGAAGCTTCCGGAATGTTGTCTM  
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GATCCATGACCGGACTSAGARGAAGGTTGGGCACAGGGTCACTGAAATAACAAGCCTACCT  
TAGWAWGAGGTCATACCAAMCAATTTTGAACATGACCSAGYCATTGATCCCSCTGGAGG  
CAACAGGAGGGAATCTTTATRGTATCCCTCTGGGCTAAGGAGGATAATGACATTACCTCTTG  
ATTAATCCCGGGTTTTCTRSTGCCAAAMTCCGSKATGAAGGGGGAAAACCTTRTKACRTTCATC  
ATTTGMCTCTACTCGCAGCCCAAKCACCAAGCCCACTCCGTGCCACCKGCCGCRRAACCCC  
AG

>BSNB\_0242\_ *Serratia\_marcescens*

GGCGGCTGCCTAATACATGCAAGTCGAGCGAACAGAAAAGGAGCTTGCTCCTTTGACGTTA  
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GCAACAATCCKTASCCAACCTGARAGGGTGATCGGCCACMCTGGGACTGAAACMCGGCCC  
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MCSCCCCGGGAKTRAAAAAGGTTTTCGGATCGWAAAACCTCTTTTGTARGGRAAAAAACA  
ACRGTAKTAMCTGGYTKTACCTTGACGGTMCCTTATWAAAARSCCMCGGCWAAAYTACTTGC  
CASCACCGCGGGTAATACA

>BSNB\_0245\_ *Lysinibacillus\_xylanilyticus*

ACTATCATCAGCGGAGGAGYATMTCAATAAGCGGAGGAKSTWTCRMKAWGCGGRGGAKYG  
YCTAAAKATKAGRGGGARAGGWCTTTAGGATGTTAACCTAGCTAAAGTCAGGCTTAGGCCT  
GGTATCCTAATTCATTATTTACCAAAAGAATTCAGAATTAATTATTGTAACATAAGCGTAAA  
AAACTTATAAAACAACCTTTTAACAACGGATCTCTTGTTCTCGCATCGATGAARAACGTAGC  
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GCGCTCAATGGTATTCCATTGAGCACGCCTGTTTCAGTATCAACAACAACCCACATCCRCAA  
TTTTGTTGTGAATGGAATGAGAGTAATCGACGTTAAAATTGAACTCTTTAAAATTATTAGG  
CCTGAACTATTGTTCTTTTASCCTGAACATTAATTTTAATATAAAGGAATGCTCTAGTTATTAA  
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>BSNB\_0246\_ *Burkholderia\_sp*

TGGGCGGCATGCCTTACACATGCAAGTCGAACGGCAGCACGGGGGCAACCCTGGTGCGGAG  
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CCGGATTAATACCGCATACGATCTACGGAAGAAAGCGGGGGATCTTCGGACCTCGCGCTGC  
AGGGGCGGCCGATGGCAGATTAGCTAGTTGGTGGGGTAAAGGCCTACCAAGGCGACGATCT  
GTAGCTGGTCTGAGAGGACGACCAGCCACACTGGGACTGAGACACGGCCCAGACTCCTACG  
GGAGGCAGCAGTGGGGAATTTTGACAATGGGGGCAACCCTGATCCAGCAATGCCGCGTGT  
GTGAAGAAGGCCTTCGGGTGTAAAGCACTTTTGTCCGGAAAGAAAACCTTACCCGCTAATAT  
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GGTAATACA

>BSNB\_0256\_ *Streptomyces\_malaysiense*

ACGTACGCTGGCGGCGTGCTTAACACATGCAAGTCGAACGATGAAGCCCTTCGGGGTGGAT  
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ACGGGTAGCCGGCCTGAGAGGGGCGACCGGCCACACTGGGACTGAGACACGGCCCAGACTC  
CTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCC  
GCGTGAGGGATGACGGCCTTCGGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGA  
CGGTACCTGCAGAAGAAGCGCCGGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGGC  
GCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTAGGCGGCTTGTACGTTCGGTTGT  
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GAGATCGGAATTCCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCG  
AAGGCGGATCTCTGGGCCGATACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGA  
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>BSNB\_0493\_ *Paraburkholderia\_tropica*

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CCGCGTGTGTGAAGAAGGCCTTCGGGTTGTAAAGCACTTTTGTCCGGAAAGAAATCCTTGGT  
CCTAATATGGTCGGGGGATGACGGTACCGGAAGAATAAGCACCGGCTAACTACGTGCCAGC  
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GCATCGCTTGAGTATGGCAGAGGGGGGTAGAATTCCACGTGTAGCAGTGAAATGCGTAGAG  
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>BSNB\_0494\_ *Klebsiella\_michiganensis*

TTCATGGCTCAGATTGAACGCTGGCGGCAGGCCTAACACATGCAAGTCGAGCGGTAGCACG  
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GGGGGACCTTCGGGCCTCTTGCCATCAGATGAACCCAGATGGGATTAGCTAGTAGGTGGGG  
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CTGAGACACGGTCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGCA  
AGCCTGATGCAGCCATGCCGCGTGTATGAAGAAGGCCTTCGGGTTGTAAAGTACTTTCAGCG  
AGGAGGAAGGCGTTAAGGTTAATAACCTTGCGGATTGACGTTACTCGCAGAAGAAGCACCG  
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GGCGTAAAGCGCACGCAGGCGGTTTGTAAAGTCGGATGTGAAATCCCCGGGCTCAACCTGG  
GAACTGCATTCGAAACTGGCAAGCTTGAGTCTTGTAGAGGGGGGTAGAATTCCAGGTGTAG  
CGGTGAAATGCGTAGAGATCTGGAGGAATACCGGTGGCGAAGGCGGCCCCCTGGACAAAG  
ACTGACGCTCAGGTGCGAAAGCGTGGGGAGCAAACAGGATTAGATACCCTGGTAGTCCACG  
CCGTAAACGATGTCGACTTGGAGGTTGTGCCCTTGAGGCGTGGCTTCGGGAGCTAACGCGTT  
AAGTCGACCGCCTG

>BSNB\_0517\_ *Enterobacter\_asburiae*

GGTCGACTTAACGCGTTAGCTCCGGAAGCCACGCCTCAAGGGCACAACCTCCAAGTCGACA  
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GCGTCAGTCTTTGTCCAGGGGGCCGCTTCGCCACCGGTATTCCTCCAGATCTCTACGCATTT  
CACCGCTACACCTGGAATTCTACCCCCCTCTACAAGACTCTAGCCTGCCAGTTTCGAATGCA  
GTTCCCAGGTTGAGCCCCGGGGATTTCACATCCGACTTGACAGACCGCCTGCGTGCGCTTTAC  
GCCCAGTAATTCCGATTAACGCTTGACCCCTCCGTATTACCGCGGCTGCTGGCACGGAGTTA  
GCCGGTGCTTCTTCTGCGGGTAACGTCAATCGACAAGGTTATTAACCTTATCGCCTTCCTCCC

CGCTGAAAGTACTTTACAACCCGAAGGCCTTCTTCATACACGCGGCATGGCTGCATCAGGCT  
TGCGCCCATTTGTGCAATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGACCGTGTCTCAGTT  
CCAGTGTGGCTGGTCATCCTCTCAGACCAGCTAGGGATCGTCGCCTAGGTGAGCCGTTACCC  
CACCTACTAGCTAATCCCATCTGGGCACATCTGATGGCAAGAGGCCCGAAGGTCCCCCTCTT  
TGGTCTTGCGACGTTATGCGGTATTAGCTACCGTTTCCAGTAGTTATCCCCCTCCATCAGGCA  
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>BSNB\_0519\_ Burkholderia\_tropica

TGATTCATGGCTCAGATTGAACGCTGGCGGCATGCCTTACACATGCAAGTCGAACGGCAGC  
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GTAGTGGGGGATAGCCCGGCGAAAGCCGGATTAATACCGCATAACGATCTACGGATGAAAGC  
GGGGGATCTTCGGACCTCGCGCTATAGGGGGCGGCCGATGGCGGATTAGCTAGTTGGTGAGG  
TAAAGGCTCACCAAGGCGACGATCCGTAGCTGGTCTGAGAGGACGACCAGCCACACTGGG  
ACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTGGGGAATTTTGGACAATGGGCG  
AAAGCCTGATCCAGCAATGCCGCGTGTGTGAAGAAGGCCTTCGGGTGTAAAGCACTTTTGT  
CCGGAAGAAATCCTTGATCCTAATATGGTTCGGGGGATGACGGTACCGGAAGAATAAGCAC  
CGGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGGTGCAAGCGTTAATCGGAATTAC  
TGGGCGTAAAGCGTGCGCAGGCGGTGATGTAAGACCGATGTGAAATCCCCGGGCTCAACCT  
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AGCAGTGAAATGCGTAGAGATGTGGAGGAATACCGATGGCGAAGGCAGCCCCCTGGGTCA  
ATACTGACGCTCATGCACGAAAGCGTGGGGAGCAAACAGGATTAGATACCCTGGTAGTCCA  
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GAAGTGACCGCCTGG

>BSNB\_0521\_ Burkholderia\_sp

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AGTATTGACCCAGGGGGCTGCCTTCGCCATCGGTATTTCCTCCACATCTCTACGCATTTCACTG  
CTACACGTGGAATTCTACCCCCCTCTGCCATACTCAAGCCTTGCAAGTACCAATGCAGTTCC  
CAGGTAAAGCCCGGGGATTTACATCGGTCTTACAAAACCGCCTGCGCACGCTTTACGCCCA  
GTAATTCCGATTAACGCTCGCACCCCTACGTATTACCGCGGCTGCTGGCACGTAGTTAGCCGG  
TGCTTATTCTTCCGGTACCGTCATCCCCGCGGGATATTATCCAACAGGATTTCTTTCCGGACA  
AAAGTGCTTTACAACCCGAAGGCCTTCTTACACACGCGGCATTGCTGGATCAGGCTTTTCGC  
CCATTGTCCAAAATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCCAGT  
GTGGCTGGTTCCTCTCAGACCAGCTACGGATCGTCGCCTTGGTAGGCCTTTACCCACCA  
ACTAGCTAATCCGCCATCGGCCACCCCAATAGCGCGAGGTCTTGCGATCCCCCGCTTTTCATC  
CAAAGATCGTATGCGGTATTAATCCGGCTTTCGCCGGGCTATCCCCCACTACTGGACATGTT  
CCGATGTATTACTACCCGTTTCGCCACTCGCCACCAGGTGCAAGCACCCGTGCTGCCGTTTCG  
ACTTGCAATGTGTAAGGCAT

>BSNB\_0547\_ Burkholderia\_tropica

ACACATGCAAGTCGAACGGCAGCACGGGTGCTTGACCTGGTGGCGAGTGGCGAACGGGTG  
AGTAATACATCGGAACGTGTCCTGTAGTGGGGGATAGCCCGGCGAAAGCCGGATTAATACC  
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GGCGGATTAGCTAGTTGGTGGGGTAAAGGCTCACCAAGGCGACGATCCGTAGCTGGTCTGA  
GAGGACGACCAGCCACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGT  
GGGGAATTTTGGACAATGGGCGAAAGCCTGATCCAGCAATGCCGCGTGTGTGAAGAAGGCC  
TTCGGGTGTAAAGCACTTTTGTCCGGAAAGAAATCCTTGATCCTAATATGGTCGGGGGATG  
ACGGTACCGGAAGAATAAGCACCGGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGG

GTGCAAGCGTTAATCGGAATTACTGGGCGTAAAGCGTGCGCAGGCGGTGATGTAAGACCGA  
TGTGAAATCCCCGGGCTCAACCTGGGAACTGCATTGGTGAAGTGCATCGCTTGAGTATGGCAG  
AGGGGGGTAGAATTCCACGTGTAGCAGTGAAATGCGTAGAGATGTGGAGGAATACCGATGG  
CGAAGGCAGCCCCCTGGGTCAATACTGACGCTCATGCACGAAAGCGTGGGGAGCAAACAG  
GATTAGATACCCTGGTAGTCCACGCCCTAAACGATGTCAACTGGTTGTGCGGTCTTCATTGA  
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>BSNB\_0561\_ *Bacillus\_thuringiensis*

GTGCTTAATGCGTTAACTTCAGCACTAAAGGGCGGAAACCCTCTAACACTTAGCACTCATCG  
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TCAGTTACAGACCAGAAAGTCGCCCTTCGCCACTGGTGTTCCTCCATATCTCTACGCATTTAC  
CGCTACACATGGAATTCCACTTTCCTCTTCTGCACTCAAGTCTCCCAGTTTCCAATGACCTC  
CACGGTTGAGCCGTGGGCTTTCACATCAGACTTAAGAAACCACCTGCGCGCGCTTTACGCCC  
AATAATTCCGGATAACGCTTGCCACCTACGTATTACCGCGGCTGCTGGCACGTAGTTAGCCG  
TGGCTTTCCTGGTTAGGTACCGTCAAGGTGCCAGCTTATTCAACTAGCACTTGTTCCTCCCTAA  
CAACAGAGTTTTACGACCCGAAAGCCTTCATCACTCACGCGGCGTTGCTCCGTCAGACTTTC  
GTCCATTGCGGAAGATTCCCTACTGCTGCCTCCCGTAGGAGTCTGGGCGGTGTCTCAGTCCC  
AGTGTGGCCGATCACCTCTCAGGTGGCTACGCATCGTTGCCTTGGTGAGCCGTTACCTCA  
CCAAGTACTAGCTAATGCGACGCGGGTCCATCCATAAGTGACAGCCGAAGCCGCTTTCAATTC  
GAACCATGCAGTTCAAAATGTTATCCGGTATTAGCCCCGGTTTCCCGGAGTTATCCCAGTCTT  
ATGGGCAGGTTACCCACGTGTTACTACCCGTCGCGCGCTAACTTCATAAGAGCAAGCTCTT  
AATCCATTGCTCGACTTGCATGTATTAGGCACGC

>BSNB\_0562\_ *Paraburkholderia\_tropica*

ACTTCACGCGTTAGCTACGTTACTAAGGAAATGAATCCCCAACAACCAGTTGACATCGTTTA  
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GTATTGGCCCAGGGGGCTGCCCTTCGCCATCGGTATTCTCCACATCTCTACGCATTTCACTGC  
TACACGTGGAATTCTACCCCCCTCTGCCATACTCCAGCGATGCAGTCACCAATGCAGTTCCC  
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TAATTCCGATTAACGCTTGACCCCTACGTATTACCGCGGCTGCTGGCACGTAGTTAGCCGGT  
GCTTATTCCTCCGGTACCGTCATCCCCGACGGATATTAGCCATCAGGATTTCTTTCCGGACAA  
AAGTGCTTTACAACCCGAAGGCCTTCTTCACACACGCGGCATTGCTGGATCAGGGTTTCCCC  
CATTGTCCAAAATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCGGTGTCTCAGTCCCAGTG  
TGGCTGGTCGTCCTCTCAGACCAGCTACGGATCGTCGCCTTGGTGGGCCTTTACCCACCAA  
CTAGCTAATCCGCCATCGGCCGCCCTATAGCGCGAGGTCCGAAGATCCCCCGCTTTCATCC  
GTAGATCGTATGCGGTATTAATCCGGCTTTCGCCGGGCTATCCCCCACTACAGGACACGTTT  
CGATGTATTACTACCCGTTTCGCCACTCGCCACCAGGTGCAAGCACCCGTGCTGCCGTTTCA  
CTTGCATGTGTAAGGCATGCCGCCAGCGTTCAATC

>BSNB\_0565\_ *Paraburkholderia\_tropica*

CATGCCTTACACATGCAAGTACGAACGGCAGCACGGGTGCTTGCACCTGGTGGCGAGTGGC  
GAACGGGTGAGTAATACATCGGAACGTGTCTGTAGTGGGGGATAGCCCGGCGAAAGCCGG  
ATTAATACCGCATACGATCTACGGATGAAAGCGGGGGATCTTCGGACCTCGCGCTATAGGG  
GCGGCCGATGGCGGATTAGCTAGTTGGTGGGGTAAAGGCCACCAAGGCGACGATCCGTAG  
CTGGTCTGAGAGGACGACCAGCCACACTGGGACTGAGACACGGCCCAGACTCCTACGGGA  
GGCAGCAGTGGGGAATTTGGACAATGGGGGAAACCCTGATCCAGCAATGCCGCGTGTGTG  
AAGAAGGCCTTCGGGTGTAAAGCACTTTTGTCCGGAAAGAAATCCTGATGGCTAATATCCG  
TCGGGGATGACGGTACCGGAAGAATAAGCACCGGCTAACTACGTGCCAGCAGCCGCGGTA  
ATACGTAGGGTGCAAGCGTTAATCGGAATTACTGGGCGTAAAGCGTGCGCAGGCGGTGATG

TAAGACCGATGTGAAATCCCCGGGCTCAACCTGGGAACTGCATTGGTGACTGCATCGCTGG  
AGTATGGCAGAGGGGGGTAGAATTCCACGTGTAGCAGTGAAATGCGTAGAGATGTGGAGG  
AATACCGATGGCGAAGGCAGCCCCCTGGGCCAATACTGACGCTCATGCACGAAAGCGTGGG  
GAGCAAACAGGATTAGATACCCTGGTAGTCCACGCCCTAAACGATGTCAAGGGGTGTGG  
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>BSNB\_0566\_ *Bacillus\_subtilis*

CGGAGTGCTTAATGCGTTAGCTGCAGCACTAAGGGGCGGAAACCCCCTAACACTTAGCACT  
CATCGTTTACGGCGTGGACTACCAGGGTATCTAATCCTGTTTCGCTCCCCACGCTTTCGCTCCT  
CAGCGTCAGTTACAGACCAGAGAGTCGCCTTCGCCACTGGTGTTCCTCCACATCTCTACGCA  
TTTCACCGCTACACGTGGAATTCCACTCTCCTCTTCTGCACTCAAGTTCCCCAGTTTCCAATG  
ACCCTCCCCGGTTGAGCCGGGGGCTTTCACATCAGACTTAAGAAACCGCCTGCGAGCCCTTT  
ACGCCCAATAATTCCGGACAACGCTTGCCACCTACGTATTACCGCGGCTGCTGGCACGTAGT  
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CCCTAACAACAGAGCTTTACGATCCGAAAACCTTCATCACTCACGCGGCGTTGCTCCGTCAG  
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GTCCCAGTGTGGCCGATCACCTCTCAGGTCGGCTACGCATCGTTGCCTTGTTGAGCCGTTAC  
CTCACCAACTAGCTAATGCGCCGCGGGTCCATCTGTAAGTGGTAGCCGAAGCCACCTTTTAT  
GTTTGAACCATGCGGTTCAAACAACCATCCGGTATTAGCCCCGGTTTCCCGGAGTTATCCCA  
GTCTTACAGGCAGGTTACCCACGTGTTACTCACCCGTCCGCGGCTAACATCAGGGAGCAAGC  
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>BSNB\_0567\_ *Bacillus\_subtilis*

CGGAGTGCTTAATGCGTTAGCTGCAGCACTAAGGGGCGGAAACCCCCTAACACTTAGCACT  
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CAGCGTCAGTTACAGACCAGAGAGTCGCCTTCGCCACTGGTGTTCCTCCACATCTCTACGCA  
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ACGCCCAATAATTCCGGACAACGCTTGCCACCTACGTATTACCGCGGCTGCTGGCACGTAGT  
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CCCTAACAACAGAGCTTTACGATCCGAAAACCTTCATCACTCACGCGGCGTTGCTCCGTCAG  
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GTCCCAGTGTGGCCGATCACCTCTCAGGTCGGCTACGCATCGTTGCCTTGTTGAGCCGTTAC  
CTCACCAACTAGCTAATGCGCCGCGGGTCCATCTGTAAGTGGTAGCCGAAGCCACCTTTTAT  
GTTTGAACCATGCGGTTCAAACAACCATCCGGTATTAGCCCCGGTTTCCCGGAGTTATCCCA  
GTCTTACAGGCAGGTTACCCACGTGTTACTCACCCGTCCGCGGCTAACATCAGGGAGCAAGC  
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>BSNB\_0568\_ *Bacillus\_subtilis*

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CATCGTTTACGGCGTGGACTACCAGGGTATCTAATCCTGTTTCGCTCCCCACGCTTTCGCTCCT  
CAGCGTCAGTTACAGACCAGAGAGTCGCCTTCGCCACTGGTGTTCCTCCACATCTCTACGCA  
TTTCACCGCTACACGTGGAATTCCACTCTCCTCTTCTGCACTCAAGTTCCCCAGTTTCCAATG  
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ACGCCCAATAATTCCGGACAACGCTTGCCACCTACGTATTACCGCGGCTGCTGGCACGTAGT  
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CCCTAACAACAGAGCTTTACGATCCGAAAACCTTCATCACTCACGCGGCGTTGCTCCGTCAG  
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CTCACCAACTAGCTAATGCGCCGCGGGTCCATCTGTAAGTGGTAGCCGAAGCCACCTTTTAT  
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GTCTTACAGGCAGGTTACCCACGTGTTACTACCCCGTCCGCCGCTAACATCAGGGAGCAAGC  
TCCCATCTGTCCGCTCGACTTGCATGTATTAGGCACGCCGCCAGCGTTCGTC

>BSNB\_0569\_ *Pantoea\_dispersa*

GGTCGACTTAACGCGTTAGCTCCGGAAGCCACGCCTCAAGGGCACAACCTCCAAGTCGACA  
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GCCCAGTAATTCCGATTAAACGCTTGCACCCTCCGTATTACCGCGGCTGCTGGCACGGAGTTA  
GCCGGTGCTTCTTCTGCGGGTAACGTCAATCGGCAAGGTTATTAACCTCACCGCCTTCCTCCC  
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TGCGCCCATTTGTGCAATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGACCGTGTCTCAGTT  
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CACCTACTAGCTAATCCCATCTGGGCACATCCGATGGTGTGAGGCCCGAAGGTCCCCCACTT  
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GTTTCCAGACATTACTACCCCGTCCGCCACTCGCCACCCAAAGAGCAAGCTCTTCTGTGCT  
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>BSNB\_0570\_ *Burkholderia\_tropica*

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CACCAAGGCGACGATCCGTAGCTGGTCTGAGAGGACGACCAGCCACACTGGGACTGAGAC  
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GCCTG

>BSNB\_0572\_ *Burkholderia\_tropica*

ACTTCACGCGTTAGCTACGTTACCAAGTCAATGAAGACCCGACAACCAGTTGACATCGTTTA  
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AAGTGCTTTACAACCCGAAGGCCTTCTTCACACACGCGGCATTGCTGGATCAGGCTTTCGCC  
CATTGTCCAAAATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCCAGTG  
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GTAGATCGTATGCGGTATTAATCCGGCTTTCGCCGGGCTATCCCCACTACAGGACACGTTC  
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CTTGCATGTGT

>BSNB\_0573\_ *Bacillus\_subtilis*

CGGAGTGCTTAATGCGTTAGCTGCAGCACTAAGGGGCGGAAACCCCCTAACACTTAGCACT  
CATCGTTTACGGCGTGGACTACCAGGGTATCTAATCCTGTTTCGCTCCCCACGCTTTCGCTCCT  
CAGCGTCAGTTACAGACCAGAGAGTCGCCTTCGCCACTGGTGTTCCTCCACATCTCTACGCA  
TTTCACCGCTACACGTGGAATTCCACTCTCCTCTTCTGCACTCAAGTTCCCCAGTTTCCAATG  
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CTCACCAACTAGCTAATGCGCCGCGGGTCCATCTGTAAGTGGTAGCCGAAGCCACCTTTTAT  
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>BSNB\_0576\_ *Bacillus\_subtilis*

CGGAGTGCTTAATGCGTTAGCTGCAGCACTAAGGGGCGGAAACCCCCTAACACTTAGCACT  
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ACGCCCAATAATTCCGGACAACGCTTGCCACCTACGTATTACCGCGGCTGCTGGCACGTAGT  
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>BSNB\_0591\_ *Klebsiella\_variicola*

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CGGGTGAATGCGTAGAGATCTGGAGGAATACCGGTGGCGAAGGCGGCCCCCTGGACAAA



GACTGACGCTCAGGTGCGAAAGCGTGGGGAGCAAACAGGATTAGATACCCTGGTAGTCCAC  
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>BSNB\_0592\_ *Klebsiella\_variicola*

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GCGGTGAAATGCGTAGAGATCTGGAGGAATACCGGTGGCGAAGGCGGCCCCCTGGACAAA  
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>BSNB\_0593\_ *Burkholderia\_tropica*

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GATCTTCGGACCTCGCGCTATAGGGGCGGCCGATGGCGGATTAGCTAGTTGGTGAGGTA  
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>BSNB\_0594\_ *Klebsiella\_variicola*

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GTGGGGGACCTTCGGGCCTCATGCCATCAGATGTGCCCAGATGGGATTAGCTGGTAGGTGGG  
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TAAATCGACCGC  
>BSNB\_0595\_ *Klebsiella\_variicola*

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GTGGGGGACCTTCGGGCCTCATGCCATCAGATGTGCCCAGATGGGATTAGCTGGTAGGTGGG  
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GACTGACGCTCAGGTGCGAAAGCGTGGGGAGCAAACAGGATTAGATACCCTGGTAGTCCAC  
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>BSNB\_0596\_ *Burkholderia\_plantarii*

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>BSNB\_0597\_ *Burkholderia\_plantarii*

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GGCGAAAGCCGGATTAATACCGCATAACGATCTACGGATGAAAGCGGGGGATCTTCGGACCT  
CGCGCTATAGGGTTGGCCGATGGCTGATTAGCTAGTTGGTAGGGTAAAAGCCTACCAAGGC  
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CAGCCGCGGTAATACGTAGGGTGCGAGCGTTAATCGGAATTACTGGGCGTAAAGCGTGCGC  
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>BSNB\_0598\_ *Klebsiella\_variicola*

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GACTGACGCTCAGGTGCGAAAGCGTGGGGAGCAAACAGGATTAGATACCCTGGTAGTCCAC  
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>BSNB\_0616\_ *Burkholderia\_sp*

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>BSNB\_0617\_ *Burkholderia\_arvi*

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CAGGTAAAGCCCGGGGATTTACATCGGTCTTAACAGACCGCCTGCGCACGCTTTACGCCCA  
GTAATTCCGATTAACGCTCGCACCCCTACGTATTACCGCGGCTGCTGGCACGTAGTTAGCCGG  
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AAAGTGCTTTACAACCCGAAGGCCTTCTTCACACACGCGGCATTGCTGGATCAGGGTTGCCC  
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GCATGTGTAAGGCAT

>BSNB\_0618\_ *Bacillus\_cereus*

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GTCTGACGGAGCAACGCCGCGTGAGTGATGAAGGCTTTCGGGTCGTAAAACTCTGTTGTTAG  
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GAGGGTCATTGGAACTGGGAGACTTGAGTGCAGAAGAGGAAAGTGGAATTCATGTGTAG  
CGGTGAAATGCGTAGAGATATGGAGGAACACCAGTGGCGAAGGCGACTTCTGCTGTGTA  
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>BSNB\_0619\_ *Burkholderia\_sp*

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GTCCAAAATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCGGTGCTCTAGTCCCAGTGTGG  
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>BSNB\_0626\_ *Burkholderia\_tropica*

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GCTTTTATCCGTAGATCGTATGCGGTATTAATCCGGCTTTCGCCGGGCTATCCCCACTACAG  
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>BSNB\_0630\_ *Burkholderia\_tropica*

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>BSNB\_0632\_ Burkholderia\_tropica

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GCTTTTCATCCGTAGATCGTATGCGGTATTAATCCGGCTTTCGCCGGGCTATCCCCACTACAG  
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GCCGTTTCGACTTGCATGTGTAAGGCATGCCGCCAGCGTTCAATCTGAGCCATGAATCA

>BSNB\_0635\_ Paraburkholderia\_guartelaensis

AACCAGTTGACATCGTTTAGGGCGTGGAATACCAGGGTATCTAATCCTGTTTGCTCCCCACG  
CTTTCGTGCATGAGCGTCAGTATTGGCCCAGGGGGCTGCCTTCGCCATCGGTATTCTCCACA  
TCTCTACGCATTTCACTGCTACACGTGGAATTCTACCCCCCTCTGCCATACTCCAGCGATGCA  
GTCACCAATGCAGTTCCCAGGTTAAGCCCGGGGATTTACATCGGTCTTACATCACCGCCTG  
CGCACGCTTTACGCCAGTAATTCCGATTAACGCTTGACCCTACGTATTACCGCGGCTGCTG  
GCACGTAGTTAGCCGGTGCTTATTCTTCCGGTACCGTCATCCCACCCGGATATTAGCCAGGC  
GGTTTTCTTCCGGACAAAAGTGCTTTACAACCCGAAGGCCTTCTTACACACGCGGCATTG  
CTGGATCAGGGTTGCCCCATTGTCCAAAATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGG  
CCGTGTCTCAGTCCCAGTGTGGCTGGTCGTCCTCTCAGACCAGCTACGGATCGTCGCCTTGGT  
GGGCCTTTACCCACCAACTAGCTAATCCGCCATCGG

>BSNB\_0636\_ Burkholderia\_sp

CGGTCACTTACGCGTTAGCTACGTTACCAAGTCAATGAAGACCCGACAACCAGTTGACATC  
GTTTAGGGCGTGGAATACCAGGGTATCTAATCCTGTTTGCTCCCCACGCTTTCGTGCATGAGC  
GTCAGTATTGGCCCAGGGGGCTGCCTTCGCCATCGGTATTCTCCACATCTCTACGCATTTCA  
CTGCTACACGTGGAATTCTACCCCCCTCTGCCATACTCCAGCGATGCAGTCACCAATGCAGT  
TCCCAGGTTAAGCCCGGGGATTTACATCGGTCTTACATCACCGCCTGCGCACGCTTTACGC  
CCAGTAATTCCGATTAACGCTTGACCCTACGTATTACCGCGGCTGCTGGCACGTAGTTAGC  
CGGTGCTTATTCTTCCGGTACCGTCATCCCACCCGGATATTAGCCAGGCGGTTTTCTTCCGG  
ACAAAAGTGCTTTACAACCCGAAGGCCTTCTTACACACGCGGCATTGCTGGATCAGGGTTG  
CCCCATTGTCCAAAATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCC

AGTGTGGCTGGTCGTCCTCTCAGACCAGCTACGGATCGTCGCCTTGGTGGGCCTTTACCCAC  
CAACTAGCTAATCCGCCATCGGCCACCCCAATAGCGCGAGGTCTTTTCGATCCCCGCTTTCC  
TCCACAGAGCGTATGCGGTATTAATCCGGCTTTGCGCGGGCTATCCCCACTACTGGACATG  
TTCCGATGTATTACTCACCCGTTTCGCCACTCGCCACCAGGTGCAAGCACCCGTGCTGCCGTT  
GACTTGCATGTGTAAGGCATGCCGCCAGCGTTCAATCT

>BSNB\_0639\_ *Burkholderia\_tropica*

CCAGGCGGTCACTTCACGCGTTAGCTACGTTACCAAGTCAATGAAGACCCGACAACCAGTT  
GACATCGTTTAGGGCGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCCACGCTTTCGTGC  
ATGAGCGTCAGTATTGACCCAGGGGGCTGCCTTCGCCATCGGTATTCCTCCACATCTCTACG  
CATTTCACTGCTACACGTGGAATTCTACCCCCCTCTGCCATACTCAAGCGATGCAGTCACCA  
ATGCAGTTCCCAGGTTGAGCCCCGGGGATTTCACATCGGTCTTACATCACCGCCTGCGCACGC  
TTTACGCCCAGTAATTCCGATTAACGCTTGACCCCTACGTATTACCGCGGCTGCTGGCACGTA  
GTTAGCCGGTGCTTATTCTTCCGGTACCGTCATCCCCGACCATATTAGGATCAAGGATTTCT  
TTCCGGACAAAAGTGCTTTACAACCCGAAGGCCCTTCTTCACACACGCGGCATTGCTGGATCA  
GGCTTTCGCCCCATTGTCCAAAATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTC  
AGTCCCAGTGTGGCTGGTCGTCCTCTCAGACCAGCTACGGATCGTCGCCTTGGTGAGCCTTTA  
CCTCACCAACTAGCTAATCCGCCATCGGCCGCCCTATAGCGCGAGGTCCGAAGATCCCC  
GCTTTCATCCGTAGATCGTATGCGGTATTAATCCGGCTTTGCGCGGGCTATCCCCACTACAG  
GACACGTTCCGATGTATTACTCACCCGTTTCGCCACTCGCCACCAGGTGCAAGCACCCGTGCT  
GCCGTTGACTTGCATGTGTAAGGCATGCCGCCAGCGTTCAATCTGAGCCATGAATCA

>BSNB\_0640\_ *Burkholderia\_oxypbila*

CGGTCACCTTCACGCGTTAGCTACGTTACCAAGTCAATGAAGACCCGACAACCAGTTGACATC  
GTTTAGGGCGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCCACGCTTTCGTGCATGAGC  
GTCAGTATTGGCCCAGGGGGCTGCCTTCGCCATCGGTATTCCTCCACATCTCTACGCATTTCA  
CTGCTACACGTGGAATTCTACCCCCCTCTGCCATACTCCAGCGATGCAGTCACCAATGCAGT  
TCCCAGGTTAAGCCCCGGGGATTTCACATCGGTCTTACATCACCGCCTGCGCACGCTTTACGC  
CCAGTAATTCCGATTAACGCTTGACCCCTACGTATTACCGCGGCTGCTGGCACGTAGTTAGC  
CGGTGCTTATTCTTCCGGTACCGTCATCCCACCCGGATATTAGCCAGGCGGTTTTCTTTCCGG  
ACAAAAGTGCTTTACAACCCGAAGGCCCTTCTTCACACACGCGGCATTGCTGGATCAGGGTTG  
CCCCATTGTCCAAAATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCC  
AGTGTGGCTGGTCGTCCTCTCAGACCAGCTACGGATCGTCGCCTTGGTGGGCCTTTACCCAC  
CAACTAGCTAATCCGCCATCGGCCACCCCAATAGCGCGAGGTCTTTTCGATCCCCGCTTTCC  
TCCACAGAGCGTATGCGGTATTAATCCGGCTTTGCGCGGGCTATCCCCACTACTGGACATG  
TTCCGATGTATTACTCACCCGTTTCGCCACTCGCCACCAGGTGCAAGCACCCGTGCTGCCGTT  
GACTTGCATGTGTAAGGCATGCCGCCAGCGTTCAATCT

>BSNB\_0643\_ *Paenibacillus\_glucanolyticus*

GCGGATGCTTAATGTGTAACTTCGGCACCAAGGGTATCGAAACCCCTAACACCTAGCATTC  
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AGCGTCAGTTACAGCCCAGAGAGTCGCCTTCGCCACTGGTGTTCTCTCCACATATCTACGCAT  
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GACCTGAAGTTGAGCCCCAGGTTTAAACACCAGACTTAAAGAACCGCCTGCGCGCGCTTTA  
CGCCCAATAATTCCGGACAACGCTTGCCCCCTACGTATTACCGCGGCTGCTGGCACGTAGTT  
AGCCGGGGCTTTCTTCTCAAGTACCGTCACTCTCATAGCAGTTACTCTATGAGACGTTCTTCC  
TTGGCAACAGAGCTTTACGATCCGAAAACCTTCATCACTCACGCGGCGTTGCTCCGTCAGGC  
TTTCGCCCCATTGCGGAAGATTCCCTACTGCTGCCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGT  
CCCAGTGTGGCCGTTACCCCTCTCAGGTTCGGCTACGCATCGTCGCCTTGGTGGGCCGTTACCC

CACCAACTAGCTAATGCGCCGAGGCCCATCCCCAAGTGACAGATTGCTCCGTCTTTCATTA  
TCACATAATGCTATGTAATAAATTATCCGGTATTAGCTACCGTTTCCGGTAGTTATCCCAGTC  
TTGAGGGCAGGTTGCCTACGTGTTACTACCCGTCGCGCGCTAACTCTCAGGAGTGCAAGCA  
CTCCTTCAAGTCCGCTCGACTTGCATGTATTAGGCACGCCGCCAGCGTTCGT

>BSNB\_0648\_ *Bacillus\_sp*

CGGTCACTTCACGCGTTAGCTACGTTACCAAGTCAATGAAGACCCGACAACCAGTTGACATC  
GTTTAGGGCGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCCACGCTTTCGTGCATGAGC  
GTCAGTATTGGCCAGGGGGCTGCCTTCGCCATCGGTATTCCTCCACATCTCTACGCATTTCA  
CTGCTACACGTGGAATTCTACCCCCCTCTGCCATACTCCAGCGATGCAGTCACCAATGCAGT  
TCCCAGGTTAAGCCCGGGGATTTACATCGGTCTTACATCACCGCCTGCGCACGCTTTACGC  
CCAGTAATTCCGATTAACGCTTGACACCCTACGTATTACCGCGGCTGCTGGCACGTAGTTAGC  
CGGTGCTTATTCTTCCGGTACCGTCATCCCACCCGGATATTAGCCAGGCGGTTTTCTTTCCGG  
ACAAAAGTGCTTTACAACCCGAAGGCCTTCTTACACACGCGGCATTGCTGGATCAGGGTTG  
CCCCATTGTCCAAAATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCGGTGTCTCAGTCCC  
AGTGTGGCTGGTCGTCCTCTCAGACCAGCTACGGATCGTCGCCTTGGTGGGCCTTTACCCAC  
CAACTAGCTAATCCGCCATCGGCCACCCAATAGCGCGAGGTCTTTCGATCCCCCGCTTTC  
TCCACAGAGCGTATGCGGTATTAATCCGGCTTTCGCCGGGCTATCCCCACTACTGGACATG  
TTCCGATGTATTACTACCCGTTTCGCCACTCGCCACCAGGTGCAAGCACCCGTGCTGCCGTT  
GACTTGCATGTGTAAGGCATGCCGCCAGCGTTCAATCT

>BSNB\_0656\_ *Burkholderia\_tropica*

CCAGGCGGTCACTTCACGCGTTAGCTACGTTACCAAGTCAATGAAGACCCGACAACCAGTT  
GACATCGTTTAGGGCGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCCACGCTTTCGTGC  
ATGAGCGTCAGTATTGACCCAGGGGGCTGCCTTCGCCATCGGTATTCCTCCACATCTCTACG  
CATTTCACTGCTACACGTGGAATTCTACCCCCCTCTGCCATACTCAAGCGATGCAGTCACCA  
ATGCAGTTCCCAGGTTGAGCCCGGGGATTTACATCGGTCTTACATCACCGCCTGCGCACGC  
TTTACGCCCAGTAATTCCGATTAACGCTTGACACCCTACGTATTACCGCGGCTGCTGGCACGTA  
GTTAGCCGGTGCTTATTCTTCCGGTACCGTCATCCCCGACCATATTAGGATCAAGGATTTCT  
TTCCGGACAAAAGTGCTTTACAACCCGAAGGCCTTCTTACACACGCGGCATTGCTGGATCA  
GGCTTTCGCCATTGTCCAAAATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCGGTGTCTC  
AGTCCCAGTGTGGCTGGTCGTCCTCTCAGACCAGCTACGGATCGTCGCCTTGGTGAGCCTTTA  
CCTCACCAACTAGCTAATCCGCCATCGGCCGCCCCCTATAGCGCGAGGTCCGAAGATCCCC  
GCTTTCATCCGTAGATCGTATGCGGTATTAATCCGGCTTTCGCCGGGCTATCCCCACTACAG  
GACACGTTCCGATGTATTACTACCCGTTTCGCCACTCGCCACCAGGTGCAAGCACCCGTGCT  
GCCGTTTCGACTTGCATGTGTAAGGCATGCCGCCAGCGTTCAATCTGAGCCATGAATCA

>BSNB\_0658\_ *Burkholderia\_tropica*

CCAGGCGGTCACTTCACGCGTTAGCTACGTTACCAAGTCAATGAAGACCCGACAACCAGTT  
GACATCGTTTAGGGCGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCCACGCTTTCGTGC  
ATGAGCGTCAGTATTGACCCAGGGGGCTGCCTTCGCCATCGGTATTCCTCCACATCTCTACG  
CATTTCACTGCTACACGTGGAATTCTACCCCCCTCTGCCATACTCAAGCGATGCAGTCACCA  
ATGCAGTTCCCAGGTTGAGCCCGGGGATTTACATCGGTCTTACATCACCGCCTGCGCACGC  
TTTACGCCCAGTAATTCCGATTAACGCTTGACACCCTACGTATTACCGCGGCTGCTGGCACGTA  
GTTAGCCGGTGCTTATTCTTCCGGTACCGTCATCCCCGACCATATTAGGATCAAGGATTTCT  
TTCCGGACAAAAGTGCTTTACAACCCGAAGGCCTTCTTACACACGCGGCATTGCTGGATCA  
GGCTTTCGCCATTGTCCAAAATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCGGTGTCTC  
AGTCCCAGTGTGGCTGGTCGTCCTCTCAGACCAGCTACGGATCGTCGCCTTGGTGAGCCTTTA  
CCTCACCAACTAGCTAATCCGCCATCGGCCGCCCCCTATAGCGCGAGGTCCGAAGATCCCC

GCTTTTCATCCGTAGATCGTATGCGGTATTAATCCGGCTTTCGCCGGGCTATCCCCACTACAG  
GACACGTTCCGATGTATTACTACCCGTTTCGCCACTCGCCACCAGGTGCAAGCACCCGTGCT  
GCCGTTGACTTGCATGTGTAAGGCATGCCGCCAGCGTTCAATCTGAGCCATGAATCA

>BSNB\_0659\_ *Bacillus\_subtilis*

GGCTCAGGACGAACGCTGGCGGCGTGCCTAATACATGCAAGTCGAGCGGACAGATGGGAG  
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GGGATAACTCCGGGAAACCGGGGCTAATACCGGATGGTTGTTTGAACCGCATGGTTCAAAC  
ATAAAAGGTGGCTTCGGCTACCACTTACAGATGGACCCGCGGCGCATTAGCTAGTTGGTGA  
GGTAACGGCTCACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGTGATCGGCCACACTGG  
GACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTAGGGAATCTTCCGCAATGGAC  
GAAAGTCTGACGGAGCAACGCCGCGTGAGTGATGAAGGTTTTCCGGATCGTAAAGCTCTGTT  
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CCACGGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGTGGCAAGCGTTGTCCGGAAT  
TATTGGGCGTAAAGGGGCTCGCAGGCGGTTTCTTAAGTCTGATGTGAAAGCCCCCGGCTCAAC  
CGGGGAGGGTCATTGGAAACTGGGGAACCTTGAGTGCAGAAGAGGAGAGTGGAATTCCACG  
TGTAAGCGGTGAAATGCGTAGAGATGTGGAGGAACACCAGTGGCGAAGGCGACTCTCTGGTC  
TGTAAGTACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTC  
CACGCCGTAAACGATGAGTGCTAAGTGTTAGGGGGTTTCCGCCCCCTTAGTGCTGCAGCTAAC  
GCATTAAGCACTCCGCC

>BSNB\_0661\_ *Burkholderia\_tropica*

CCAGGCGGTCACTTCACGCGTTAGCTACGTTACCAAGTCAATGAAGACCCGACAACCAAGTT  
GACATCGTTTTAGGGCGTGGACTIONACCAGGGTATCTAATCCTGTTTGCTCCCCACGCTTTCGTGC  
ATGAGCGTCAGTATTGACCCAGGGGGCTGCCTTCGCCATCGGTATTCCTCCACATCTCTACG  
CATTTCACTGCTACACGTGGAATTCTACCCCCCTCTGCCATACTCAAGCGATGCAGTCACCA  
ATGCAGTTCCCAGGTTGAGCCCCGGGGATTTACATCGGTCTTACATCACCGCCTGCGCACGC  
TTTACGCCCAGTAATTCCGATTAACGCTTGACACCCTACGTATTACCGCGGCTGCTGGCACGTA  
GTTAGCCGGTGCTTATTCTTCCGGTACCGTCATCCCCCGACCATATTAGGATCAAGGATTTCT  
TTCCGGACAAAAGTGCTTTACAACCCGAAGGCCTTCTTACACACGCGGCATTGCTGGATCA  
GGCTTTCGCCCATTTGTCCAAAATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTC  
AGTCCCAGTGTGGCTGGTCCTCTCAGACCAGCTACGGATCGTCGCCTTGGTGAGCCTTTA  
CCTCACCAACTAGCTAATCCGCCATCGGCCGCCCTATAGCGCGAGGTCCGAAGATCCCCC  
GCTTTTCATCCGTAGATCGTATGCGGTATTAATCCGGCTTTCGCCGGGCTATCCCCACTACAG  
GACACGTTCCGATGTATTACTACCCGTTTCGCCACTCGCCACCAGGTGCAAGCACCCGTGCT  
GCCGTTGACTTGCATGTGTAAGGCATGCCGCCAGCGTTCAATCTGAGCCATGAATCA

>BSNB\_0663\_ *Burkholderia\_tropica*

CCAGGCGGTCACTTCACGCGTTAGCTACGTTACCAAGTCAATGAAGACCCGACAACCAAGTT  
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ATGAGCGTCAGTATTGACCCAGGGGGCTGCCTTCGCCATCGGTATTCCTCCACATCTCTACG  
CATTTCACTGCTACACGTGGAATTCTACCCCCCTCTGCCATACTCAAGCGATGCAGTCACCA  
ATGCAGTTCCCAGGTTGAGCCCCGGGGATTTACATCGGTCTTACATCACCGCCTGCGCACGC  
TTTACGCCCAGTAATTCCGATTAACGCTTGACACCCTACGTATTACCGCGGCTGCTGGCACGTA  
GTTAGCCGGTGCTTATTCTTCCGGTACCGTCATCCCCCGACCATATTAGGATCAAGGATTTCT  
TTCCGGACAAAAGTGCTTTACAACCCGAAGGCCTTCTTACACACGCGGCATTGCTGGATCA  
GGCTTTCGCCCATTTGTCCAAAATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTC  
AGTCCCAGTGTGGCTGGTCCTCTCAGACCAGCTACGGATCGTCGCCTTGGTGAGCCTTTA  
CCTCACCAACTAGCTAATCCGCCATCGGCCGCCCTATAGCGCGAGGTCCGAAGATCCCCC  
GCTTTTCATCCGTAGATCGTATGCGGTATTAATCCGGCTTTCGCCGGGCTATCCCCACTACAG



GACACGTTCCGATGTATTACTCACCCGTTCCGCCACTCGCCACCAGGTGCAAGCACCCGTGCT  
GCCGTTGACTTGATGTGTAAGGCATGCCGCCAGCGTTCAATCTGAGCCATGAATCA

>BSNB\_0669\_ *Enterobacter\_amnigenus*

GGCGGTCGACTTAACGCGTTAGCTCCGGAAGCCACGCCTCAAGGGCACAACCTCCAAGTCG  
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TGAGCGTCAGTCTTTGTCCAGGGGGCCGCCTTCGCCACCGGTATTCTCCAGATCTCTACGCA  
TTTACCCGCTACACCTGGAATTCTACCCCCCTCTACAAGACTCTAGCCTGCCAGTTTCGAATG  
CAGTTCCCAGGTTGAGCCCGGGGATTTACATCCGACTTGACAGACCGCCTGCGTGCGCTTT  
ACGCCCAGTAATTCCGATTAACGCTTGACCCCTCCGTATTACCGCGGCTGCTGGCACGGAGT  
TAGCCGGTGCTTCTTCTGCGGGTAACGTCAATTGCTGAGGTTATTAACCTCAACACCTTCCTC  
CCCGCTGAAAGTACTTTACAACCCGAAGGCCTTCTTCATACACGCGGCATGGCTGCATCAGG  
CTTGCGCCCATTTGTGCAATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGACCGTGTCTCAG  
TTCCAGTGTGGCTGGTCATCCTCTCAGACCAGCTAGGGATCGTCGCCTAGGTGAGCCGTTAC  
CCCACCTACTAGCTAATCCCATCTGGGCACATCTGATGGCAAGAGGCCCGAAGGTCCCCCTC  
TTTGGTCTTGCGACGTTATGCGGTATTAGCTACCGTTTCCAGTAGTTATCCCCCTCCATCAGG  
CAGTTTCCCAGACATTACTCACCCGTCCGCCGCTCGCCGGCAAAGTAGCAAGCTACTTTCCG  
CTGCCGCTCGACTTGATGTGTTAGGCCTGCCGCCAGCG

>BSNB\_0671\_ *Pantoea\_stewartii*

CTTAACGCGTTAGCTCCGGAAGCCACTCCTCAAGGGAACAACCTCCAAGTCGACATCGTTTA  
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GTCTTCGTCCAGGGGGCCGCCTTCGCCACCGGTATTCTCCAGATCTCTACGCATTTACCCGC  
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AGGTTAAGCCCGGGGATTTACATCTGACTTAACAGACCGCCTGCGTGCGCTTTACGCCCAG  
TAATTCCGATTAACGCTTGACCCCTCCGTATTACCGCGGCTGCTGGCACGGAGTTAGCCGGT  
GCTTCTTCTGCGGGTAACGTCAATTGCTGAGGTTATTAACCTCAGCACCTTCCTCCCCGCTGA  
AAGTACTTTACAACCCGAAGGCCTTCTTCATACACGCGGCATGGCTGCATCAGGCTTGCGCC  
CATTGTGCAATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGACCGTGTCTCAGTTCCAGTG  
TGGCTGGTCATCCTCTCAGACCAGCTAGGGATCGTCGCCTAGGTGGGCCGTTACCCCGCCTA  
CCAGCTAATCCCATCTGGGCACATCCGATGGTGTGAGGCCCGAAGGTCCCCCACTTTGGTCT  
TGCGACGTTATGCGGTATTAGCTACCGTTTCCAGT

>BSNB\_0672\_ *Pantoea\_stewartii*

CTTAACGCGTTAGCTCCGGAAGCCACTCCTCAAGGGAACAACCTCCAAGTCGACATCGTTTA  
CGGCGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCCACGCTTTCGCACCTGAGCGTCA  
GTCTTCGTCCAGGGGGCCGCCTTCGCCACCGGTATTCTCCAGATCTCTACGCATTTACCCGC  
TACACCTGGAATTCTACCCCCCTCTACGAGACTCAAGCCTGCCAGTTTCAAATGCAGTTCCC  
AGGTTAAGCCCGGGGATTTACATCTGACTTAACAGACCGCCTGCGTGCGCTTTACGCCCAG  
TAATTCCGATTAACGCTTGACCCCTCCGTATTACCGCGGCTGCTGGCACGGAGTTAGCCGGT  
GCTTCTTCTGCGGGTAACGTCAATTGCTGAGGTTATTAACCTCAGCACCTTCCTCCCCGCTGA  
AAGTACTTTACAACCCGAAGGCCTTCTTCATACACGCGGCATGGCTGCATCAGGCTTGCGCC  
CATTGTGCAATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGACCGTGTCTCAGTTCCAGTG  
TGGCTGGTCATCCTCTCAGACCAGCTAGGGATCGTCGCCTAGGTGGGCCGTTACCCCGCCTA  
CCAGCTAATCCCATCTGGGCACATCCGATGGTGTGAGGCCCGAAGGTCCCCCACTTTGGTCT  
TGCGACGTTATGCGGTATTAGCTACCGTTTCCAGT

>BSNB\_0675\_ *Enterobacter\_mori*

GGCGGTCGACTTAACGCGTTAGCTCCGGAAGCCACGCCTCAAGGGCACAACCTCCAAGTCG  
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TGAGCGTCAGTCTTTGTCCAGGGGGCCGCTTCGCCACCGGTATTCTCCAGATCTCTACGCA  
TTTACCGCTACACCTGGAATTCTACCCCCCTCTACAAGACTCTAGCCTGCCAGTTTCGAATG  
CAGTTCCCAGGTTGAGCCCGGGGATTTACATCCGACTTGACAGACCGCCTGCGTGCGCTTT  
ACGCCCAGTAATTCCGATTAACGCTTGACCCCTCCGTATTACCGCGGCTGCTGGCACGGAGT  
TAGCCGGTGCTTCTTCTGCGGGTAACGTCAATTGCTGAGGTTATTAACCTCAACACCTTCCTC  
CCCGCTGAAAGTACTTTACAACCCGAAGGCCTTCTTCATACACGCGGCATGGCTGCATCAGG  
CTTGCGCCCATTTGTGCAATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGACCGTGTCTCAG  
TTCCAGTGTGGCTGGTCATCCTCTCAGACCAGCTAGGGATCGTCGCCTAGGTGAGCCGTTAC  
CCCACCTACTAGCTAATCCCATCTGGGCACATCTGATGGCAAGAGGCCCGAAGGTCCCCCTC  
TTTGGTCTTGCGACGTTATGCGGTATTAGCTACCGTTTCCAGTAGTTATCCCCCTCCATCAGG  
CAGTTTCCCAGACATTACTACCCGTCCGCGCTCGCCGCAAAGTAGCAAGCTACTTTCCG  
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>BSNB\_0683\_ *Bacillus\_subtilis*

CGGAGTGCTTAATGCGTTAGCTGCAGCACTAAGGGGCGGAAACCCCCTAACACTTAGCACT  
CATCGTTTACGGCGTGGACTACCAGGGTATCTAATCCTGTTGCTCCCCACGCTTTCGCTCCT  
CAGCGTCAGTTACAGACCAGAGAGTCGCCTTCGCCACTGGTGTTCTCCACATCTCTACGCA  
TTTACCGCTACACGTGGAATTCCACTCTCCTCTTCTGCACTCAAGTTCCCCAGTTTCCAATG  
ACCCTCCCCGGTTGAGCCGGGGGCTTTCACATCAGACTTAAGAAACCGCCTGCGAGCCCTTT  
ACGCCCAATAATTCCGGACAACGCTTGCCACCTACGTATTACCGCGGCTGCTGGCACGTAGT  
TAGCCGTGGCTTTCTGGTTAGGTACCGTCAAGGTACCGCCCTATTCTGAACGGTACTTGTTCTT  
CCCTAACAACAGAGCTTTACGATCCGAAAACCTTCATCACTCACGCGGCGTTGCTCCGTCAG  
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GTCCCAGTGTGGCCGATCACCTCTCAGGTGGCTACGCATCGTTGCCTTGGTGAGCCGTTAC  
CTACCAACTAGCTAATGCGCCGCGGGTCCATCTGTAAGTGGTAGCCGAAGCCACCTTTTAT  
GTTTGAACCATGCGGTTCAAACAACCATCCGGTATTAGCCCCGGTTTCCCGGAGTTATCCCA  
GTCTTACAGGCAGGTTACCCACGTGTTACTACCCGTCCGCCGCTAACATCAGGGAGCAAGC  
TCCCATCTGTCCGCTCGACTTGCATGTATTAGGCACGCCGCCAGCGTTCGTC

>BSNB\_0685\_ *Bacillus\_subtilis*

CGGAGTGCTTAATGCGTTAGCTGCAGCACTAAGGGGCGGAAACCCCCTAACACTTAGCACT  
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CAGCGTCAGTTACAGACCAGAGAGTCGCCTTCGCCACTGGTGTTCTCCACATCTCTACGCA  
TTTACCGCTACACGTGGAATTCCACTCTCCTCTTCTGCACTCAAGTTCCCCAGTTTCCAATG  
ACCCTCCCCGGTTGAGCCGGGGGCTTTCACATCAGACTTAAGAAACCGCCTGCGAGCCCTTT  
ACGCCCAATAATTCCGGACAACGCTTGCCACCTACGTATTACCGCGGCTGCTGGCACGTAGT  
TAGCCGTGGCTTTCTGGTTAGGTACCGTCAAGGTACCGCCCTATTCTGAACGGTACTTGTTCTT  
CCCTAACAACAGAGCTTTACGATCCGAAAACCTTCATCACTCACGCGGCGTTGCTCCGTCAG  
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GTCCCAGTGTGGCCGATCACCTCTCAGGTGGCTACGCATCGTTGCCTTGGTGAGCCGTTAC  
CTACCAACTAGCTAATGCGCCGCGGGTCCATCTGTAAGTGGTAGCCGAAGCCACCTTTTAT  
GTTTGAACCATGCGGTTCAAACAACCATCCGGTATTAGCCCCGGTTTCCCGGAGTTATCCCA  
GTCTTACAGGCAGGTTACCCACGTGTTACTACCCGTCCGCCGCTAACATCAGGGAGCAAGC  
TCCCATCTGTCCGCTCGACTTGCATGTATTAGGCACGCCGCCAGCGTTCGTC

>BSNB\_0704\_ *Stenotrophomonas\_maltophilia*

CAGGCGGCGAACTTAACGCGTTAGCTTCGATACTGCGTGCCAAATTGCACCCAACATCCATG

TTCGCATCGTTT TAGGGCGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCCACGCTTTCGT  
GCCTCAGTGTCAATGTTGGTCCAGGTAGCTGCCTTCGCCATGGATGTTCTCCTGATCTCTAC  
GCATTTCACTGCTACACCAGGAATTCGCTACCCTCTACCACATTCTAGTCATCCAGTATCCA  
CTGCAGTTCCCAGGTTGAGCCCAGGGCTTTCACAACGGACTTAAATAAACCACCTACGCACGC  
TTTACGCCCAGTAATTCCGAGTAACGCTTGACCCCTTCGTATTACCGCGGCTGCTGGCACGA  
AGTTAGCCGGTGCTTATTCTTTGGGTACCGTCATCCCAACCAGGTATTAACCGGCTGGATTTC  
TTCCCAACAAAAGGGCTTTACAACCCGAAGGCCTTCTTCACCCACGCGGTATGGCTGGATC  
AGGCTTGCGCCCATTTGTCCAATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGACCGTGTCT  
CAGTTCAGTGTGGCTGATCATCCTCTCAAACCAGCTACGGATCGTCGCCTTGGTGGGCCTTT  
ACCCCGCCAACTAGCTAATCCGACATCGGCTCATTCAATCGCGCAAGGTCCGAAGATCCCCT  
GCTTTCACCCGTAGGTCGTATGCG

>BSNB\_0706\_ Enterobacter\_mori

GGCGGTCGACTTAACGCGTTAGCTCCGGAAGCCACGCCTCAAGGGCACAACCTCCAAGTCG  
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TGAGCGTCAGTCTTTGTCCAGGGGGGCCCTTCGCCACCGGTATTCTCCAGATCTCTACGCA  
TTTCACCGCTACACCTGGAATTCTACCCCCCTCTACAAGACTCTAGCCTGCCAGTTTCGAATG  
CAGTTCACAGGTTGAGCCCGGGGATTTACATCCGACTTGACAGACCGCCTGCGTGCGCTTT  
ACGCCCAGTAATTCCGATTAACGCTTGACCCCTCCGTATTACCGCGGCTGCTGGCACGGAGT  
TAGCCGGTGCTTCTTCTGCGGGTAACGTCAATTGCTGAGGTTATTAACCTCAACACCTTCCTC  
CCCGCTGAAAGTACTTTACAACCCGAAGGCCTTCTTCATACACGCGGCATGGCTGCATCAGG  
CTTGCGCCCATTTGTGCAATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGACCGTGTCTCAG  
TTCCAGTGTGGCTGGTCATCCTCTCAGACCAGCTAGGGATCGTCGCCTAGGTGAGCCGTTAC  
CCCACCTACTAGCTAATCCCATCTGGGCACATCTGATGGCAAGAGGCCCGAAGGTCCCCCTC  
TTTGGTCTTGCGACGTTATGCGGTATTAGCTACCGTTTCCAGTAGTTATCCCCCTCCATCAGG  
CAGTTTCCAGACATTACTACCCGTCCGCGGCTCGCCGGCAAAGTAGCAAGCTACTTTCCG  
CTGCCGCTCGACTTGATGTGTTAGGCCTGCCGCCAGCG

>BSNB\_0708\_ Enterobacter\_mori

GGCGGTCGACTTAACGCGTTAGCTCCGGAAGCCACGCCTCAAGGGCACAACCTCCAAGTCG  
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TTTCACCGCTACACCTGGAATTCTACCCCCCTCTACAAGACTCTAGCCTGCCAGTTTCGAATG  
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>BSNB\_0709\_ Enterobacter\_mori

GGCGGTCGACTTAACGCGTTAGCTCCGGAAGCCACGCCTCAAGGGCACAACCTCCAAGTCG  
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>BSNB\_0712\_ *Bacillus\_subtilis*

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>BSNB\_0716\_ *Bacillus\_subtilis*

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>BSNB\_0719\_ *Luteibacter\_yeojuensis*

GGCGAACTTAACGCGTTAGCTTCGACACTGATCTCCGAGTTGAGACCAACATCCAGTTGCA  
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CACCGCTACACCGGGAATTCCACCATCCTCTATCACACTCTAGCTCGCCAGTATCCACTGCC  
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GCCCAGTAATTCCGATTAACGCTTGCACCCTCCGTATTACCGCGGCTGCTGGCACGGAGTTA  
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TGCGCCCATTTGTCCAATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTC  
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>BSNB\_0721\_ *Luteibacter\_yejuensis*

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GCGTCAGTGTGATCCAGATGGCCGCCTTCGCCACTGATGTTCTCCCGATCTCTACGCATTT  
CACCGCTACACCGGGAATTCACCATCCTCTATCACACTCTAGCTCGCCAGTATCCACTGCC  
ATTCCCAGGTTGAGCCCCGGGGCTTTCACAGCAGACTTAACGAACCGCCTACGCACGCTTAC  
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TGCGCCCATTTGTCCAATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTC  
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>BSNB\_0722\_ *Bacillus\_subtilis*

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GACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTAGGGAATCTTCCGCAATGGAC  
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>BSNB\_0723\_ *Enterobacter\_mori*

GGCGGTCGACTTAACGCGTTAGCTCCGGAAGCCACGCCTCAAGGGCACAACCTCCAAGTCG  
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>BSNB\_0725\_ Enterobacter\_mori

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>BSNB\_0726\_ Enterobacter\_mori

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>BSNB\_0727\_ Enterobacter\_mori

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>BSNB\_0730\_ *Bacillus\_subtilis*

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CTACCAACTAGCTAATGCGCCGCGGGTCCATCTGTAAGTGGTAGCCGAAGCCACCTTTTAT  
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>BSNB\_0762\_ *Bacillus\_subtilis*

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>BSNB\_0803\_ *Bacillus\_subtilis*

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GACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTAGGGAATCTTCCGCAATGGAC  
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>BSNB\_0804\_ *Bacillus\_subtilis*

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>BSNB\_0805\_ *Methylobacterium\_mesophilicum*

CCGGATACGCCCTTTTGGGGAAAGGTTTACTGCCGGAAGATCGGCCCGCGTCTGATTAGCTA  
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>BSNB\_0816\_ *Bacillus\_subtilis*

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>BSNB\_0817\_ Bacillus\_subtilis

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>BSNB\_0955\_ Bacillus\_subtilis

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ATAAAAGGTGGCTTCGGCTACCACTTACAGATGGACCCGCGGCGCATTAGCTAGTTGGTGA  
GGTAACGGCTCACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGTGATCGGCCACACTGG  
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GAAAGTCTGACGGAGCAACGCCGCGTGAGTGATGAAGGTTTTCCGATCGTAAAGCTCTGTT  
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CGGGGAGGGTCATTGGAAACTGGGGAACCTTGAGTGCAAGAGAGGAGAGTGGAATTCCACG  
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>BSNB\_0956\_ Bacillus\_subtilis

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GACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTAGGGAATCTTCCGCAATGGAC  
GAAAGTCTGACGGAGCAACGCCGCGTGAGTGATGAAGGTTTTCCGATCGTAAAGCTCTGTT  
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CGGGGAGGGTCATTGGAAACTGGGGAACCTTGAGTGCAAGAGAGGAGAGTGGAATTCCACG  
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TGTAAGTACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTC  
CACGCCGTAAACGATGAGTGCTAAGTGTTAGGGGGTTTCCGCCCCCTTAGTGCTGCAGCTAAC  
GCATTAAGCACTCCGCC  
>BSNB\_0959\_ *Bacillus\_subtilis*

GGCTCAGGACGAACGCTGGCGGCGTGCCTAATACATGCAAGTCGAGCGGACAGATGGGAG  
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GACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTAGGGAATCTTCCGCAATGGAC  
GAAAGTCTGACGGAGCAACGCCGCGTGAGTGATGAAGGTTTTCCGATCGTAAAGCTCTGTT  
GTTAGGGAAGAACAAGTACCGTTTCAATAGGGCGGTACCTTGACGGTACCTAACCAGAAAAG  
CCACGGCTAACTACGTGCCAGCAGCCGCGTAATACGTAGGTGGCAAGCGTTGTCCGGAAT  
TATTGGGCGTAAAGGGCTCGCAGGCGGTTTCTTAAGTCTGATGTGAAAGCCCCCGGCTCAAC  
CGGGGAGGGTCATTGGAACTGGGGAACTTGAGTGCAAGAGGAGAGTGGAATTCCACG  
TGAGCGGTGAAATGCGTAGAGATGTGGAGGAACACCAGTGGCGAAGGCGACTCTCTGGTC  
TGTAAGTACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTC  
CACGCCGTAAACGATGAGTGCTAAGTGTTAGGGGGTTTCCGCCCCCTTAGTGCTGCAGCTAAC  
GCATTAAGCACTCCGCC  
>BSNB\_0978\_ *Acidisoma\_sp*

AGAATAAGCCCCGGATAACTTTGTGCCAGCAGCCGCGTAATACGAAGGGGGCTAGCGTTG  
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GGGCTCAACCTGGGGGCTGCGCTTGATACTTGTTGCTTGAGTGGGGAAGAGGGTCGTGGA  
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ACCTGGTCCTTGACTGACGCTGAGGCGCGAAAGCGTGGGGAGCAAACAGGATTAGATACCC  
TGGTAGTCCACGCTGTAAACGATGTGTGCTGGATGTTGGGTGACTTAGTCACTCAGTGTGCTA  
GCTAACGCGATAAGCATCA

>BSNB\_1020\_ *Burkholderia\_tropica*

CCAGGCGGTCACTTCACGCGTTAGCTACGTTACCAAGTCAATGAAGACCCGACAACCAGTT  
GACATCGTTTAGGGCGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCCACGCTTTCGTGC  
ATGAGCGTCAGTATTGACCCAGGGGGCTGCCTTCGCCATCGGTATTCCTCCACATCTCTACG  
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ATGCAGTTCCCAGGTTGAGCCCCGGGGATTTCACATCGGTCTTACATCACCGCCTGCGCACGC  
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TTCCGGACAAAAGTGCTTTACAACCCGAAGGCCTTCTTCACACACGCGGCATTGCTGGATCA  
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CCTCACCAACTAGCTAATCCGCCATCGGCCGCCCTATAGCGCGAGGTCCGAAGATCCCCC  
GCTTTTATCCGTAGATCGTATGCGGTATTAATCCGGCTTTCGCCGGGCTATCCCCACTACAG  
GACACGTTCCGATGTATTACTACCCGTTTCGCCACTCGCCACCAGGTGCAAGCACCCGTGCT  
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>BSNB\_1022\_ *Methylobacterium\_populi*

CCCCTATCTTCCGGTMCGTCAATTATCGTCCCGGACAAAAGAGCTTTACAACCCTAAGGCCTT  
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CCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCTGATCATCCTCTCAGACCAG  
CTACTGATCGTCGCCTTGGTAGGCCGTTACCCACCAACAAGCTAATCAGACGCGGGCCGAT  
CCTTCGGCAGTAAACCTTTCCCCATAAGGGCGTATCCGGTATTAGCTCAAGTTTCCCTGAGTT  
ATTCCGAACCGAAGGGCACGTTCCACGTGTTACTACCCGTCTGCCACTGACTTCCGAAGA  
AGCCCGTTCGACTTGATGTGTTAAGCCTGCCGCCAGCGTTCGCTCTGAGCCAGGATCAAAC  
TCTA

>BSNB\_1023\_ *Curtobacterium\_sp*

CTTCTTTCTGCAGGTACGTCACTTTCGCTTCTTCCCTACTAAAAGAGGTTTACAACCCGAAGG  
CCGTCATCCCTCACGCGGCGTTGCTGCATCAGGCTTTCGCCCATTTGTGCAATATCCCCACTG  
CTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCCGGTACCCCTCTCAGGC  
CGGCTACCCGTCGTCGCCTTGGTGAGCCATTACCTACCAACAAGCTGATAGGCCGCGAGTC  
CATCCCCAACCAAAAAATCTTTCCACCACCAGACCATGCGGCCGGCAGTCCTATCCAGTATT  
AGACGTTCGTTTCCAACGCTTATCCCAGAGTCAAGGGCAGGTTACTCACGTGTTACTACCCG  
TTCGCCACTAATCCACCCAGCAAGCTGGGCATCATCGTTTCGACTTGATGTGTTAAGCACGC  
CGCCAGCGTTCGTCTGAGCCAGGTYAAAACTCTA

>BSNB\_1024\_ *Pantoea\_dispersa*

GGTCGGTCTCTCTGCGGGTAACGTCAATCGGCGAGGTTATTAACCTCACCGCCTTCTCCCCG  
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CGCCCATTTGTGCAATATCCCCACTGCTGCCTCCCGTAGGAGTCTGGACCGTGTCTCAGTTCC  
AGTGTGGCTGGTCATCCTCTCAGACCAGCTAGGGATCGTCGCCTAGGTGAGCCATTACCCCA  
CCTACTAGCTAATCCCATCTGGGCACATCCGATGGTGTGAGGCCCCGAAGGTCCCCCACTTTG  
GTCTTGCGACGTTATGCGGTATTAGCTACCGTTTCCAGTAGTTATCCCCCTCCATCGGGCAGT  
TTCCCAGACATTACTACCCGTCCGCCACTCGCCACCCAAAGAGCAAGCTCTTCTGTGCTGC  
CGTTTCGACTTGATGTGTTAGGCCTGCCGCCAGCGTTCAATATGAGCCAGGTYAAAACTC  
TA

>BSNB\_1025\_ *Bacillus\_megaterium*

ATCGTGCTTCTGGTAGGTACCGTCAAGGTACGAGCAGTTACTCTCGTACTTGTTCTTCCCTAA  
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GTCCATTGCGGAAGATTCCCTACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCC  
AGTGTGGCCGATCACCTCTCAGGTCGGCTATGCATCGTTGCCTTGGTGAGCCGTTACCTCAC  
CAACTAGCTAATGCACCGCGGGCCCATCTGTAAGTGATAGCCGAAACCATCTTTCAATCATC  
TCCCATGAAGGAGAAGATCCTATCCGGTATTAGCTTCGGTTTCCCGAAGTTATCCAGTCTTA  
CAGGCAGGTTGCCACGTGTTACTACCCGTCCGCCGCTAACGTCATAGAAGCAAGCTTCTA  
ATCAGTTCGCTCGACTTGATGTATTAGGCACGCCGCCAGCGTTCATCCTGAGCCAGGTYMA  
AAACTCTA

>BSNB\_1026\_ *Methylobacterium\_sp*

CTCTCCTATCTCGGTACGTCAATTATCGTCCCGGATAAAAGAGCTTTACAACCCTAAGGCCTTC  
ATCACTACGCGGCATGGCTGGATCAGGCTTGCGCCCATTTGTCCAATATTCCCCACTGCTGC  
CTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCTGATCATCCTCTCAGACCAGC  
TACTGATCGTCGCCTTGGTAGGCCGTTACCCACCAACTAGCTAATCAGACGCGGGGCCGATC  
TTCCGGCAGTAAACCTTTCCCCATAAGGGCGTATCCGGTATTAGCCCTAGTTTCCCAGGGTTA  
TTCCGAACCGGAAGGCACGTTCCACGCGTACTACCCGTCCGCCGCTGACCCCGAAGGG  
CCCGCTCGACTTGATGTGTTAAGCCTGCCGCCAGCGTTCGCTCTGAGCCAGGATCAAACCTC  
TAA

>BSNB\_1027\_ *Asaia\_sp*

GTCTGGCTCTTCTACGGGTACCGTCATCATCGTCCCCGTCGAAAGTGCTTTACAATCCGAAG  
ACCTTCTTCACACACGCGGCATTGCTGGATCAGGCTTTCGCCCATTTGTCCAATATTCCCCACT  
GCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCTGATCATCCTCTCAGAC  
CAGCTATCGATCATCGCCTTGGTAGGCCTTTACCCACCAACAAGCTAATCGAACGCAGGCT  
CCTCCACAGGCGACTTGCGCCTTTGACCCTCAGGTATCATGCGGTATTAGCACCAGTTTCCCG  
ATGTTATCCCCACCCGTGGATAGATCCCTACGCGATACTCACCCGTCCGCCACTCACCCCG  
AAGGGTCCGTGCGACTTGCAATGTGTTAAGCATGCCGCCAGCGTTTCGCTCTGAGCCAGATCAA  
AACTCTA

>BSNB\_1030\_ *Pseudomonas\_sp*

CTTATTCTGTCGGTACGTCAAAACTCACAGGTATTTCGCTGTAAGCCCTTCCTCCCAACTTAAA  
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TTGTCCAATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGACCGTGTCTCAGTTCAGTGTG  
ACTGATCATCCTCTCAGACCAGTTACGGATCGTCGCCTTGGTAGGCCTTTACCTACCAACTA  
GCTAATCCGACCTAGGCTCATCTAATAGCGTGAGGTCCGAAGATCCCCCACTTTCTCCCGTA  
GGACGTATGCGGTATTAGCGTTCCTTTCGAAACGTTGTCCCCCACTACTAGGCAGATTCCCTAG  
GCATTACTCACCCGTCCGCCGCTGAATCGARGAGCAAGCTCCTCTCATCCGCTCGACTTGCA  
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>BSNB\_1032\_ *Arthrobacter\_sp*

CCGGCCTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTACTGAAAGAGGTTTACAACCCG  
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CACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCCGGTCACCCTCTC  
AGGCCGGCTACCCGTGCTCGCCTTGGTAGGCCATTACCCACCAACAAGCTGATAGGCCGC  
GAGTCCATCCAAAACCACAAAAGCTTTCACCACCATGACATGCGCCAGATGGTCGTATCC  
GGTATTAGACCCAGTTTCCCAGGCTTATCCAGAGTCAAGGGCAGGTTACTCACGTGTTACT  
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ACGCCGCCAGCGTTCATCCTGAGCCAGGATCAAACTCTA

>BSNB\_1034\_ *Pantoea\_agglomerans*

GGGTGCGGGGCTCTTCTGCGGGTACGTATCGACASGGTTATTAACCCCRTCGCCTTCCTCCC  
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CCAGTGTGGCTGGTCATCCTCTCAGACCAGCTAGGGATCGTCGCCTAGGTGGGCCATTACCC  
CGCCTACTAGCTAATCCCATCTGGGTTTCATCCGATAGTGAGAGGCCCGAAGGTCCCCCTCTT  
TGGTCTTGCGACGTTATGCGGTATTAGCCACCGTTTCCAGTGGTTATCCCCCTCTATCGGGCA  
GATCCCCAGACATTACTCACCCGTCCGCCACTCGTCACCCAAGRRCMAGCTYTCTSTGCKAC  
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TSAAAA

>BSNB\_1116\_ *Methylobacterium\_mesophilicum*

CCGGATACGCCCTTTTGGGGAAAGGTTTACTGCCGGAAGATCGGCCCGCGTCTGATTAGCTA  
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CACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGGA  
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GCTCTTTTATCCGGGACGATAATGACGGTACCGGAGGAATAAGCCCCGGCTAACTTCGTGCC  
AGCAGCCGCGGTAATACGAAGGGGGTACGCTTGCTCGGAATCACTGGGCGTAAAGGGCGC  
GTAGGCGGCGTTTTAAGTCGGGGGTGAAAGCCTGTGGCTCAACCACAGAATGGCCTTCGAT  
ACTGGGACGCTTGAGTATGGTAGAGGTTGGTGGAACTGCGAGTGTAGAGGTGAAATTCGTA  
GATATTCGCAAGAACACCGGTGGCGAAGGCGGCCAACTGGACCATTACTGACGCTGAGGCG

CGAAAGCGTGGGGAGCAAACAGGATTAGATACCCTGGTAGTCCACGCCGTAAACGATGAAT  
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>BSNB\_1123\_ *Methylobacterium\_mesophilicum*

CCGGATACGCCCTTTTGGGGAAAGGTTTACTGCCGGAAGATCGGCCCCGCGTCTGATTAGCTA  
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CACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGGA  
CAATGGGCGCAAGCCTGATCCAGCCATGCCGCGTGAGTGATGAAGGCCTTAGGGTTGTAAA  
GCTCTTTTATCCGGGACGATAATGACGGTACCGGAGGAATAAGCCCCGGCTAACTTCGTGCC  
AGCAGCCGCGGTAATACGAAGGGGGCTAGCGTTGCTCGGAATCACTGGGCGTAAAGGGGCG  
GTAGGCGGCGTTTTAAGTCGGGGGTGAAAGCCTGTGGCTCAACCACAGAATGGCCTTCGAT  
ACTGGGACGCTTGAGTATGGTAGAGGTTGGTGGAACTGCGAGTGTAGAGGTGAAATTCGTA  
GATATTCGCAAGAACACCGGTGGCGAAGGCGGCCAACTGGACCATTACTGACGCTGAGGCG  
CGAAAGCGTGGGGAGCAAACAGGATTAGATACCCTGGTAGTCCACGCCGTAAACGATGAAT  
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>BSNB\_1136\_ *Methylobacterium\_mesophilicum*

CCGGATACGCCCTTTTGGGGAAAGGTTTACTGCCGGAAGATCGGCCCCGCGTCTGATTAGCTA  
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CACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGGA  
CAATGGGCGCAAGCCTGATCCAGCCATGCCGCGTGAGTGATGAAGGCCTTAGGGTTGTAAA  
GCTCTTTTATCCGGGACGATAATGACGGTACCGGAGGAATAAGCCCCGGCTAACTTCGTGCC  
AGCAGCCGCGGTAATACGAAGGGGGCTAGCGTTGCTCGGAATCACTGGGCGTAAAGGGGCG  
GTAGGCGGCGTTTTAAGTCGGGGGTGAAAGCCTGTGGCTCAACCACAGAATGGCCTTCGAT  
ACTGGGACGCTTGAGTATGGTAGAGGTTGGTGGAACTGCGAGTGTAGAGGTGAAATTCGTA  
GATATTCGCAAGAACACCGGTGGCGAAGGCGGCCAACTGGACCATTACTGACGCTGAGGCG  
CGAAAGCGTGGGGAGCAAACAGGATTAGATACCCTGGTAGTCCACGCCGTAAACGATGAAT  
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>BSNB\_1137\_ *Methylobacterium\_mesophilicum*

CCGGATACGCCCTTTTGGGGAAAGGTTTACTGCCGGAAGATCGGCCCCGCGTCTGATTAGCTA  
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CACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGGA  
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GCTCTTTTATCCGGGACGATAATGACGGTACCGGAGGAATAAGCCCCGGCTAACTTCGTGCC  
AGCAGCCGCGGTAATACGAAGGGGGCTAGCGTTGCTCGGAATCACTGGGCGTAAAGGGGCG  
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ACTGGGACGCTTGAGTATGGTAGAGGTTGGTGGAACTGCGAGTGTAGAGGTGAAATTCGTA  
GATATTCGCAAGAACACCGGTGGCGAAGGCGGCCAACTGGACCATTACTGACGCTGAGGCG  
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>BSNB\_1141\_ *Methylobacterium\_mesophilicum*

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CACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGGA  
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GCTCTTTTATCCGGGACGATAATGACGGTACCGGAGGAATAAGCCCCGGCTAACTTCGTGCC  
AGCAGCCGCGGTAATACGAAGGGGGCTAGCGTTGCTCGGAATCACTGGGCGTAAAGGGGCG  
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ACTGGGACGCTTGAGTATGGTAGAGGTTGGTGGAACTGCGAGTGTAGAGGTGAAATTCGTA

GATATTCGCAAGAACACCGGTGGCGAAGGCGGCCAACTGGACCATTACTGACGCTGAGGCG  
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>BSNB\_1179\_ *Rhodococcus\_kroppenstedtii*

CCGCTTAATGCGTTAGCTACGGCACGGATCCCGTGGAAGGAAACCCACACCTAGCGCCCAC  
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ACCGCTACACCAGGAATTCCAGTCTCCCCTGAAATACTCAAGTCTGCCCGTATCGCCTGCAA  
GCCAACAGTTGAGCTGCTGGTTTTACAGACGACGCGACAAACCGCCTACGAGCTCTTTACG  
CCCAGTAATTCCGGACAACGCTTGACCCCTACGTATTACCGCGGCTGCTGGCACGTAGTTGG  
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CCCCTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCCGGTCGCCCT  
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GCGGGCCCATCCTGCACCGATAAATCTTTCCACCACACGGCATGCACCGCGCAGTCCTATCC  
GGTATTAGACCCAGTTTCCCGGGCTTATCCCGAAGTGCAGGGCAGATCACCCACGTGTTACT  
CACCCGTTCCGCACTCGTGTACCCCGAAGGGCCTTACCGTTCGACTTGCATGTGTTAAAGCA  
CGCCGCCAGCGTTCGTCCTGA  
>BSNB\_1190\_ *Rhodococcus\_kroppenstedtii*

CCGCTTAATGCGTTAGCTACGGCACGGATCCCGTGGAAGGAAACCCACACCTAGCGCCCAC  
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CGTCAGTTATTTCCCAGAGACCCGCCTTCGCCACCGGTGTTCCCTCCTGATATCTGCGCATTTTC  
ACCGCTACACCAGGAATTCCAGTCTCCCCTGAAATACTCAAGTCTGCCCGTATCGCCTGCAA  
GCCAACAGTTGAGCTGCTGGTTTTACAGACGACGCGACAAACCGCCTACGAGCTCTTTACG  
CCCAGTAATTCCGGACAACGCTTGACCCCTACGTATTACCGCGGCTGCTGGCACGTAGTTGG  
CCGGTGCTTCTTCTGCAGGTACCGTCACTCACGCTTCGTCCCTGCTGAAAGAGGTTTACAACC  
CGAAGGCCGTCATCCCTCACGCGGCGTCGCTGCATCAGGCTTGCGCCCATTGTGCAATATTC  
CCCCTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCCGGTCGCCCT  
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GCGGGCCCATCCTGCACCGATAAATCTTTCCACCACACGGCATGCACCGCGCAGTCCTATCC  
GGTATTAGACCCAGTTTCCCGGGCTTATCCCGAAGTGCAGGGCAGATCACCCACGTGTTACT  
CACCCGTTCCGCACTCGTGTACCCCGAAGGGCCTTACCGTTCGACTTGCATGTGTTAAAGCA  
CGCCGCCAGCGTTCGTCCTGA  
>BSNB\_1191\_ *Microbacterium\_testaceum*

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GGGAGCAAACAGGCTTAGATACCTGGTAGTCCACCCCGTAAACGTTGGGAAGTAGTTGTG  
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>BSNB\_1192\_ *Microbacterium\_testaceum*

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CCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGGAAGCCTGATGCA  
GCAACGCCGCGTGAGGGATGACGGCCTTCGGGTTGTAAACCTCTTTTAGCAGGGAAGAAGC  
GAAAGTGACGGTACCTGCAGAAAAAGCGCCGGCTAACTACGTGCCAGCAGCCGCGGTAAT  
ACGTAGGGCGCAAGCGTTATCCGGAATTATTGGGCGTAAAGAGCTCGTAGGCGGTTTGTTCG  
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GCGGTAGGGGAGATTGGAATTCCTGGTGTAGCGGTGGAATGCGCAGATATCAGGAGGAACA  
CCGATGGCGAAGGCAGATCTCTGGGCCGTAACCTGACGCTGAGGAGCGAAAGGGTGGGGAG  
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C

>BSNB\_1207\_ *Bacillus\_subtilis*

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GACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTAGGGAATCTTCCGCAATGGAC  
GAAAGTCTGACGGAGCAACGCCGCGTGAGTGATGAAGGTTTTTCGGATCGTAAAGCTCTGTT  
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CCACGGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGTGGCAAGCGTTGTCCGGAAT  
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TGTAAGCGGTGAAATGCGTAGAGATGTGGAGGAACACCAGTGGCGAAGGCGACTCTCTGGTC  
TGTAAGTACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTC  
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GCATTAAGCACTCCGCC

>BSNB\_1208\_ *Bacillus\_subtilis*

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>BSNB\_1209\_ *Bacillus\_subtilis*

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>BSNB\_1210\_ *Bacillus\_subtilis*

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>BSNB\_1220\_ *Rhodococcus\_cerastii*

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>BSNB\_1238\_ *Bacillus\_subtilis*

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>BSNB\_1246\_ *Rhodococcus\_cerastii*

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>BSNB\_A1F2A1\_ *Klebsiella\_michiganensis*

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>BSNB\_A4F1\_12\_ *Klebsiella\_michiganensis*

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>BSNB\_LRFP\_069\_ *Methylobacterium\_mesophilicum*

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TABLE S5. ITS OF STUDIED FUNGAL STRAINS (BSNB)

>BSNB\_0001\_ *Epicoccum\_nigrum*

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>BSNB\_0002\_ *Colletotrichum\_gloeosporioides*

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>BSNB\_0006\_ *Pestalotiopsis\_sp*

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>BSNB\_0008\_ *Pestalotiopsis\_sp*

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>BSNB\_0018\_ *Chaetomium\_globosum*

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>BSNB\_0035\_Colletotrichum\_boninense

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>BSNB\_0196\_Cladosporium\_sp

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>BSNB\_0197\_Penicillium\_sanguifluum

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>BSNB\_0201\_Exophiala\_xenobiotica

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>BSNB\_0212\_Pseudallescheria\_boydii

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>BSNB\_0218\_Penicillium\_citrinum

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>BSNB\_0220\_Paecilomyces\_formosus

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>BSNB\_0227\_*Aspergillus\_aculeatus*

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>BSNB\_0266\_*Rhizomucor\_variabilis*

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>BSNB\_0290\_Colletotrichium\_gloeosporioides

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>BSNB\_0549\_Colletotrichum\_sp

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>BSNB\_0642\_Hypoxylon\_investiens

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>BSNB\_1040\_Trichoderma\_sp

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GCAGCCCCGGAACCAGGCGCCCGCCGGAGGGACCAACCAAACCTCTTCTGTAGTCCCCTCG  
CGGACGTTATTTCTTACAGCTCTGAGCAAAAATTCAAAATGAATCAAAACTTTCAACAACGG  
ATCTCTTGTTCTGGCATCGATGAAGAACGCAGCGAAATGCGATAAGTAATGTGAATTGCAG  
AATTCAGTGAATCATCGAATCTTTGAACGCACATTGCGCCCGCCAGTATTCTGGCGGGCATG  
CCTGTCCGAGCGTCATTTCAACCCTCGAACCCTCCGGGGGGTTCGGCGTTGGGGATCGGGAA  
CCCCTAAGACGGGATCCCGGCCCGGAAATACAGTGGCGGTCTCGCCGAGCCTCTCCTGCG  
CAGTAGTTTGCACAACTCGCACCGGGAGCGCGGCGCTCCACGTCCGTAAAACACCCAACCT  
TCTGAAATGTTGACCTCGGATCAGGTAGGAATACCCGCTGAACTTAAGCATATCAA

>BSNB\_1041\_Beuveria\_bassiana

CCCCTAAATCCCTTCTGTGACCTACCTATCGTTGCTTCGGCGGACTCGCCCCAGCCCGGACGC  
GACTGGACCAGCGGCCCGCCGGGGACCTCAAACCTCTTGTTATTCAGCATCTTCTGAATACG  
CCGCAAGGCAAAACAAATGAATCAAAACTTTCAACAACGGATCTCTTGCTCTGGCATCGA

TGAAGAACGCAGCGAAATGCGATAAGTAATGTGAATTGCAGAATCCAGTGAATCATCGAAT  
CTTTGAACGCACATTGCGCCCGCCAGCATTCTGGCGGGCATGCCTGTTTCGAGCGTCATTTCA  
ACCCTCGACCTCCCTTGGGGAGGTGCGCGTTGGGGACCGGCAGCACACCGCCGGCCCTGA  
AATGGAGTGGCGGCCCCGTCCGCGGCGACCTCTGCGTAGTAATACAGCTCGCACCGGAACCC  
CGACGCGGCCACGCCGTAAAACACCCAACCTTCTGAACGTTGACCTCGAATCAGGTAGGACT  
ACCCGCTGAACTTAAGCATATCAATAAGCGGAGGAA

>BSNB\_1042\_Beauveria\_bassiana

TCTGTGAGCCTACCTATCGTTGCTTCGGCGGACTCGCCCCAGCCCGGACGCGGACTGGACCA  
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AACAAATGAATCAAAACTTTCAACAACGGATCTCTTGGCTCTGGCATCGATGAAGAACGCA  
GCGAAATGCGATAAGTAATGTGAATTGCAGAATCCAGTGAATCATCGAATCTTTGAACGCA  
CATTGCGCCCCGCGAGCATTCTGGCGGGCATGCCTGTTTCGAGCGTCATTTCAACCCTCGACCTC  
CCCTTGGGGAGGTGCGCGTTGGGGACCGGCAGCACACCGCCGGCCCTGAAA  
TGGAGTGGCGGCCCCGTCCGCGGCGACCTCTGCGTAGTAATACAGCTCGCACCGGAACCCCG  
ACGCGGCCACGCCGTAAAACACCCAACCTTCTGAACGTTGACCTCGAATCAGGTAGGACTAC  
CCGCTGAACTTAAGCATATCAATAAGCGGAGGAA

>BSNB\_1043\_Mucor\_sp

CCGTATTTCTTATTTACTGTGAACTGTTTTATTGTTTGRCGCTTGAGGAATGTTGTTTAGCCAT  
AGGGATAGGCTAGGCAAATGTTAACCGAGTCAAAGTCAGGCTTAGGCCTGGTATCCTATTC  
ATTATTTACCAAAAGAATTCAGAATTATTATTGTAACATAAGCGTAAAAAACTTATAAAACA  
ACTTTTAACAACGGATCTCTTGGTTCTCGCATCGATGAAGAACGTAGCAAAGTGCGATAACT  
AGTGTTGAATTGCATATTCAGTGAATCATCGAGTCTTTGAACGCATCTTGCGCTCAATGGTATT  
CCATTGAGCACGCCTGTTTCAGTATCAAAAACACCCACATTACAACTTGTGTTGTTGAATG  
GAATTGAGAGTTTCGGCTTAACGCTGATCTCTTTAAAWTATTAGGCCTGAACTTTTGTCTT  
TYYGCTGAACATTTTTTTAATATAAAGGAATGCTCTAGTAAAAAGACTCTTTCTGGGGCCTC  
CCAAATAAATCATTCTTAACTTGATCTGAAATCAGGTGGGATTACCCGCTGAACTTAAGCA  
TATCATAAGCCGGAGGA

>BSNB\_1044\_Penicillium\_sp

AAAGCTGTACGGCGAGTGAGCGGCAAGAGCTCAAATTTGARRGCTGGCTCCTTCGGGGTCC  
GCATTGTAATTTGCAGAGGATGCTTCGGGAGTGGCCCCCATCTAAGTGCCCTGGAACGGGGCC  
GTCATAGAGGGTGAGAATCCCGTCTGGGATGGGGTGTCACGCCCCGTGTGAAGCTCCTTCGA  
CGAGTCGAGTTGTTTGGGAATGCAGCTCTAAATGGGTGGTAAATTTTCATCTAAAGCTAAATA  
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GAGAGTTAAACAGCACGTGAAATTGTTGAAAGGGAAGCGCTTGCGACCAGACTCGCCACG  
GGGTCAGCCGGCATTTCGTGCCGGTGTACTTCCCCGCGGGCGGGCCAGCGTCGGTTTGGGCG  
GCCGGTCAAAGGCCCTCGGAATGTAACGCCCCCGGGGCGTCTTATAGCCGAGGGTGCCAT  
GCGGCCAGCCCGGACCGAGGAACGCGCTTCGGCTCGGACGCTGGCATAATGGTCGTAAGCG  
ACCCGTCTTGAAACACGGACCAAA

>BSNB\_1045\_Penicillium\_sp

CCACCTCCACCCCTTGTCTCCACACCTGTTGCTTCGGCGGGGCCACCGGGGCCACCCGGTCG  
CCGGGGGACATCCGTCCCCGGGCCCCGCGCCCCGCCGAGGCGCTCTGTGAACCCTGATGAAGA  
TGGGCTGTCTGAGTGATATGAAAATTGTCAAACTTTCAACAATGGATCTCTKGGTTCCGGC  
ATCGATSAAGAACGCAGCGAAWTGCGATAAGTAATGTGAATTGCAGAATTCGGTGAATCAT  
CGAATCTTTGAACGCACATTGCGCCCCCTGGCATTCCGGGGGGCATGCCTGTCCGAGCGTCA  
TTTCTGCCCTCAAGCACGGCTTGTGTGTTGGGTGTGGTCCCCCTGGGGACCTGCCCCGAAAGG  
CAGCGGCYACGTCCGTCTGGTCCTCGAGCGTATGGGGCTCTGTCACTCGCTCGGGAAGGACC

TGCGGGGGTTGGTCACCACCACATCTGKTACAAGGTTGACCTCGGATCAGGTAGGAGTTAC  
CCGCTTA

>BSNB\_1046\_Penicillium\_simplicissimum

CGGGCCCTCACTCGGCAATGATCCTTCCGTAGGTGAACTTGCGGAAGGATCATTACCGAGTG  
AGGGCCCTCTGGGTCCAACCTCCCACCCGTGTTTATCGTACCTTGTTGCTTCGGCGGGCCCGC  
CTCACGGCCGCGGGGGGGCATCCGCTCCCGGGCCCGCGCCCGCGAAGACACCAATGAACTC  
TGTCTGAAGATTGCAGTCTGAGCAGATTAGCTAAATCAGTTAAAACCTTTCAACAACGGATCT  
CTTGGTTCCGGCATCGATGAAGAACGCAGCGAAATGCGATACGTAATGTGAATTGCAGAAT  
TCAGTGAATCATCGAGTCTTTGAACGCACATTGCGCCCCCTGGTATTCCGGGGGGGCATGCCT  
GTCCGAGCGTCATTGCTGCCCTCAAGCACGGCTTGTGTGTTGGGCTCCGCCCCCGGCTCCC  
GGGGGGCGGGCCCGAAAGGCAGCGGCGGCACCGCGTCCGGTCCTCGAGCGTATGGGGCTTG  
TACCCGCTCTGTAGGCCCGGCCGCGCCCGCGCGACCCCAATCAATCTCCAGGTTACCTC  
GGAATCAGGT

>BSNB\_1048\_Neosartorya\_sp

CACTCGGTAATGATCCTTCCGTAGGTGAACCTGCGGAAGGATCATTACCGAGTGAGGGCCCT  
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CGGCCCGCGGGGAGGCCTCGCGCCCCGGGCCCGCGCCCGCCGAAGACCCCAACTGAACG  
CTGTTCTGAAAGTATGCAGTCTGAGTTGATTATCATAATCAGTTAAAACCTTTCAACAACGGA  
TCTCTTGGTTCCGGCATCGATGAAGAACGCAGCGAAATGCGATAAGTAATGTGAATTGCAG  
AATTCAAGTGAATCATCGAGTCTTTGAACGCACATTGCGCCCCCTGGTATTCCGGGGGGCATG  
CCTGTCCGAGCGTCATTGCTGCCCTCAAGCACGGCTTGTGTGTTGGGCCCGCGTCCCCGGTTT  
CCCCCGGGGACGGGCCCGAAAGGCAGCGGCGGCACCGCGTCCGGTCCTCGAGCGTATGGG  
GCTTTGTACCCGCTCTGTAGGCCTGGCCGCGGCCAGCCGACCCCAACTTTATTTCAGGTG  
CACCTCGATTCAATC

>BSNB\_1049\_Penicillium\_oxalicum

CGGTATGATCCTTCCGTAGGTGAACCTGCGGAAGGATCATACCGAGTGAGGGCCCTCTGGGT  
CCAACCTCCCACCCGTGTTTATCGTACCTTGTTGCTTCGGCGAGCCCGCCTCACGGCCGCGG  
GGGGCATCTGCCCCCGGGCCTGCGCTCGCCGAAGACACACAAACGAACTCTTGTCTGAAGA  
TTGCAGTCTGAGTACTTGACTAAATCAGTTAAAACCTTTCAACAACGGATCTCTTGGTTCCGGC  
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AGTCTTTGAACGCACATTGCGCCCCCTGGTATTCCGGGGGGCATGCCTGTCCGAGCGTCATT  
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CGAAAGGCAGCGGCGGCACCGTGTCCGGTCCTCGAGCGTATGGGGTTTGTACCCGCTCTGT  
AGGCCCGGCGGGCCCGCGCGGAACCAATCAACTTACCCATCTCACCTATCAAACCCCCC  
G

>BSNB\_1053\_Beuveria\_bassiana

GGCACCGCCAACAGGCGGCAAGGCTGGAGTATTTTATTACCCTTGTCTTTTGCGCACTTGTTG  
TTTCCTGGGCGGGTTCGCCCCGCTCCAGGACCACATGATAAACCTTTTTTATGCAGTTGCAAT  
CAGCGTCAGTACAACAAATGTAAATCATTTACAACCTTTCAACAACGGATCTCTTGGTTCTGG  
CATCGATGAAGAACGCAGCGAAATGCGATACGTAAGTGAATTGCAGAATTCAGTGAATCA  
TCGAATCTTTGAACGCACATTGCGCCCTTTGGTATTCCAAAGGGCATGCCTGTTTCGAGCGTCA  
TTTGTACCCTCAAGCTTTGCTTGGTGTTGGGCGTTTTTGTCTTTGGTTTTGCCCAAAGACTCGC  
CTTAAAACGATTGGCAGCCGGCCTACTGGTTTCGAGCGCAGCACATTTTTCGCTTGCAAT  
CAGCAAAAGAGGACGGCACTCCATCAAGACTCTTTATCACTTTTGACCTCGGATCAGGTAGG  
GATACCCGCTGAACTTAAGCATATCAATAAGCCGGAGGAA

>BSNB\_1054\_Pestalotiopsis\_theae

AAGCTCCAACCCATGTGACTTACCTTTTTGTTGCCTCGGCAGAGGTTACCTGGTACCTGGAGA  
CAGGTTACCCCTGTAGCAACTGCCGGTGGACTACTAAACTCTTGTTATTTTATGTAATCTGAGC  
GTCTTATTTTAATAAGTCAAAACTTTCAACAACGGATCTCTGGGTTCTGGCATCGATGAAGA  
ACGCAGCGAAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGA  
ACGCACATTGCGCCCATTAGTATTCTAGTGGGCATGCCTGTTTCGAGCGTCATTTCAACCCTTA  
AGCCTAGCTTAGTGTTGGGAATTTACAGTTATGTAATTCCTGAAATACAACGGCGGATCTGT  
GGTATCCTCTGAGCGTAGTAAATTATTTCTCGCTTTTGTTAGGTGCTGCAGCTCCCAGCCGCT  
AAACCCCCAATTTTTTGTGGTTGACCTCGGATCAGGTAGGAATACCCGCTGAACTTAAGCAT  
ATCAATAAGCGGAGGA

>BSNB\_1055\_Fusarium\_sp

CTCCAAACCCCTGTGAACATACCTTATGTTGCCTCGGCGGATCAGCCCGCGCCCCGTAAAAA  
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CAAATAAATCAAAACTTTCAACAACGGATCTCTTGGTTCTGGCATCGATGAAGAACGCAGC  
AAAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCACAT  
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CTTGGTGTTGGGAGCTGCAGTCCTGCTGCACTCCCCAAATACATTGGCGGTCACGTCGAGCT  
TCCATAGCGTAGTAATTTACACATCGTTACTGGTAATCGTCGCGGCCACGCCGTTAAACCCC  
AACTTCTGAATGTTGACCTCGGATCAGGTAGGAATACCCGCTGAACTTAAGCATATCATAAA  
GCGGAGGA

>BSNB\_1056\_Lecanicillium\_sp

ACCAAAACCCCTTATGTGAACATACCACGATGTTGCTTCGGCGGACTCGCCCCGGCGTCCGGA  
CGGCCTAGCGCCGCCCGCGGCCCGGATCCAGGCGGCCGCCGGAGACCACCAAAACTATTTT  
GTATCAGCAGTTTTTTCTGAATCCGCCGCAAGGCAAAACAAATGAATCAAAACTTTCAACAA  
CGGATCTCTTGGTTCTGGCATCGATGAAGAACGCAGCGAAATGCGATAAGTAATGTGAATTG  
CAGAATTCAGTGAATCATCGAATCTTTGAACGCACATTGCGCCCGCCAGCATTCTGGCGGGC  
ATGCCTGTTTCGAGCGTCATTTCAACCCTCGACTTCCCTTTGGGGAAATCGGCGTTGGGGACTG  
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ATCCAACCTCGCACCGGAACCCCGACGTGGCCACGCCGTAAACACCCCCACTTTCTGAACG  
TTGACCTCGGATCAGGTAGGAATACCCGCTGAACTTAAGCATATCAATAAGGCRGAGGA

>BSNB\_1057\_Fusarium\_sp

TAGAACGTACGTAGAAAAAAGTAGTAGCTCYTYGCTCCYACCCCTCGACGTCCTT  
ATGTTGCCTCGGCGGACACCCSCGCCCGKAAAACGGGACGGCCCGCCAGAAAACCGAARC  
TCTAATGTTTCTTATTGTAACCTTCTGAGTWWWACAAACAAATAAATCAAAACTTTCAACAA  
CGGATCTCTTGGTTCTGGCATCGATGAASAACGCAGCAAAATGCGATAAGTAATGTGAATTG  
CAGAATTCWGTGAATCATCGAATCTTTGAACGCACATTGCGCCCGCTGGTATTCCGGCGGGC  
ATGCCTGTTTCGAGCGTCATTTCAACCCTCAAGCCCCCGGGTTGGTGTTGGGGATCGGCTCTG  
CCCTTCTGGGCGGTGCCGCCCCGAAMTACATTGGCGGTCTCGCTGCASCCTCCATTGCGTA  
GTAGCTAACACCTCGCAACTGGAACGCGGCGGGCCATGCCGTAAAACCCCAACTTCTGAA  
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>BSNB\_1058\_Fusarium\_sp

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GGACGGCCCCGCCAGAGGACCCAAACTCTAATGTTTCTTATTGTAACCTTCTGAGTAAAACAAA  
CAAATAAATCAAAACTTTCAACAACGGATCTCTTGGTTCTGGCATCGATGAAGAACGCAGC  
AAAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCACAT  
TGCGCCCGCTGGTATTCCGGCGGGCATGCCTGTTTCGAGCGTCATTTCAACCCTCAAGCCCC

GGGTTTGGTGTGTTGGGGATCGGCTCTGCCCTTCTGGGCGGTGCCGCCCCGAAATACATTGGC  
GGTCTCGCTGCAGCCTCCATTGCGTAGTAGCTAACACCTCGCAACTGGAACGCGGCGCGGCC  
ATGCCGTAAAACCCCAACTTCTGAATGTTGACCTCGGATCAGGTAGGAATACCCGCTGAACT  
TAAGCATATCAATAAGCCGGAGGAA

>BSNB\_1059\_Arthrinium\_sp

CAACTCCATACCATCTGTAACTACCCAGTTATGCCTCGGCGTAAGCTCGGTTGGAGGCAC  
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AACTCTGTTTTATTTTATATTCTGAGCGTCTTATTTTAATAAGTTAAAACTTTCAACAACGGAT  
CTCTTGGTTCTGGCATCGATGAAGAACGCAGCGAAATGCGATAAGTAATGTGAATTGCAGA  
ATTCAGTGAATCATCGAATCTTTGAACGCACATTGCGCCCATCAGTATTCTGGTGGGCATGC  
CTGTTGAGCGTCATTTCAACCCTTAAGCCTAGCTTAGTGTTGGGAATCTGCTGTACTGCAGT  
TCCTTAAAGACAGTGGCGGAGCGGCGGTAGTCTCTGAGCGTAGTAATTTATTTCTCGCTTTT  
GTCAGGCTCTGTCTCCCGCCATAAAACCCCAATTTTTTAGTGTTGACCTCGGATCAGGTA  
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>BSNB\_1061\_Penicillium\_sp

GAACCCGCACTCGGTAATGATCCTTCCGTAGGTGAACCTGCGGAAGGATCATTACCGAGTG  
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CACTGGGGCCTGGCCCCGGTCGCCGGGGGGCTTCTGCCCCCGGGCCCGCGCCCGCCGAAGC  
ACCCTAGAACCCTGTCTGAACAGTGAGTCTGAGTCGGATATTGAATCATTAAAACTTTCAAC  
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TTGCAGAATTCCGTGAATCATCGAATCTTTGAACGCACATTGCGCCCCCTGGCATTCCGGGG  
GGCATGCCTGTCCGAGCGTCATTTCTGCCCTCCAGCACGGCTGGGTGTTGGGCGCTGTCCCCC  
CGGGGACACGCCCCAAAAGCAGTGCGGCGCCGGTCGGGTCTCGAGCGTATGGGGCTTTGT  
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CCGGTCCCCCTCGGAATTCCAAGG

>BSNB\_1063\_Mucor\_hiemalis

CTTTGCTAGTTTTCTAGCGAATGGTTCATTCTTTTTTACTGTGAACTGTTTTAATTTTTCAGCGT  
CTGAGGAATGTCTTTTAGCCATAGGGATAGGCTACTAGAATGTTAACCAGAGCTGAAAGTCAG  
GCTTAGGCCTGGTATCCTATTAATTATTTACCAAAAAGAATTCAGTATTATAATTGTAACATAA  
GCGTAAAAAACTTATAAAACAACTTTTAACAACGGATCTCTTGGTTCTCGCATCGATGAAGA  
ACGTAGCAAAGTGCAGATAACTAGTGTGAATTGCATATTCAGTGAATCATCGAGTCTTTGAAC  
GCAACTTGCGCTCAATGGTATTCCATTGAGCACGCCTGTTTCAGTATCAAAAACACCCCA  
TTCATAATTTTGTGTGAATGGAAATGAGAGTTTCGGCTTTATTGCTGAATTCTTTAAATTTAT  
TAGGCCTGAACTATTGTTCTTTCTGCCTGAACATTTTTTTAATATAAAGGAATGCTCTAGTAA  
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>BSNB\_1065\_Fusarium\_sp

AACTGCCAACCCTGTGACATACCTTRATGTTGCCTCGGCGGATCAGCCCGCGCCCCGTAAA  
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AAACAAATAAATCAAAACTTTCAACAACGGATCTCTTGGTTCTGGCATCGATGAAGAACGC  
AGCAAAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGC  
ACATTGCGCCCGCTGGTATTCCGGCGGGCATGCCTGTTGAGCGTCATTTCAACCCTCAAGC  
CCCCGGGTTTGGTGTGGGGATCGGCTCTGCCCTTCTGGGCGGTGCCGCCCCGAAATACAT  
TGGCGGTCTCGCTGCAGCCTCCATTGCGTAGTAGCTAACACCTCGCAACTGGAACGCGGCGC  
GGCCATGCCGTAAAACCCCAACTTCTGAATGTTGACCTCGGATCAGGTAGGAATACCCGCTG  
AACTTAAGCATATCAATAAGCGGAGGAA

>BSNB\_1067\_Penicillium\_sp

CCTTCCGTAGGTGAACCTGCGGAAGGATCATTACCGAGTGCGGGGCCCTCGCGGCCCAACCTC  
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GGACGCTCGTCCCCGGGCCCCGCGCCCGCCGAAGCGCCCTGTGAACCCTGATGAAGATGGAC  
TGTCTGAGTACCATGAAAATTGTCAAACTTTCAACAATGGATCTCTTGGTTCCGGCATCGAT  
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CCTCAAGCACGGCTTGTGTGTTGGGTGCGGTCCCCCGGGGACCTGCCCCGAAAGGCAGCGG  
CGACGTCCGTCCGGTCTCGAGCGTATGGGGCTTTGTCACTCGCTCGGGAAGGACCTGCGGG  
GCGTTGGTCACCACCATATTTGACCTACGGCTGATTACCC

>BSNB\_1069\_Beuveria\_sp

CTACCAACCCTTATGTGACCTACCTATCGTTGCTTCGGCGGACTCGCCCCAGCCGGACGCGG  
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CGCAWGGCAAAACAAATAAATTAAACTTTCAACAACGGATCTCTTGGCTCTGGCATCGAT  
GAAGAACGCAGCGAAATGCGATAAGTAATGTGAATTGCAGAATCCAGTGAATCATCGAATC  
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GACGTGGCCACGCCGTAAACACCCAACTTCTGAACGTTGACCTCGAATCAGGTAGGACTA  
CCCCTGAACCTAAGCATATCAATAAGCCGGAGGA

>BSNB\_1250\_Isaria\_farinos

GCGGAGGGATCATTACCGAGTTTTCAACTCCCAAACCCCTTTGTGAACATACCTATCGTTGCT  
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CCGCCGGAGACCCCCAACTCTGTATTCTCAGTATCTTCTGAATCCGCCGCAAGGCAAAACA  
AATGAATCAAACTTTCAACAACGGATCTCTTGGTTCTGGCATCGATGAAGAACGCAGCGA  
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CGCCCCGCCAGCATTCTGGCGGGCATGCCTGTTGAGCGTCATTTCAACCCTCGACTTCCCTTT  
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GTCCGCGGCGACCTCTGCGTAGTAATCCAACTCGCACCGGAACCCCCGACGTGGCCACGCCG  
TAAACCCCCGACTTCTGAACGTTGACCTCGGATCAGGTAGGAATACCCGCTGAACCTAAGC  
ATATCA

>BSNB\_1252\_Cordyceps\_militaris

GCGGAGGGATCATTAACGAGTTTCCCAAACCTCCCAAACCCCTTTGTGAACATACCTATCGTTGCTT  
CGGCGGACTCGCCAGCGCCTGGACGCGGGCCTGGGCGGCGGCCGTGCGGGGCCCAAC  
ACTGTATCTACCAGTTTTTCTGAATCCGCCGCAAGGCAAAACAAATGAATCAAACTTTCAA  
CAACGGATCTCTTGGCTCTGGCATCGATGAAGAACGCAGCGAAATGCGATAAGTAATGTGA  
ATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCACATTGCGCCCCGCCAGCATTCTGGC  
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GACCGGCAGCACACCGCCGCCCGCGAAATGAAGTGGCGGCCCGTCCGCGGCGACCTCTGCG  
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>BSNB\_1253\_Ophiocordyceps\_forquignonii

GCGGAGGGATCATTACTGAGTGTTAAAACTCTCTAACCCCCTATGTGATTACTACCATTTTTTA  
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TCTATTGAGTCTTCTGAGTTTATAATAACAAAACGCATAAAAACTTTCAGCAACGGATCTCTT  
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CCGAGCGTCATTATCAACCCCTCTCAGGCTTTCTTATAGGGTTTTAATAGGACCCTATATCGA  
TTGACTGGCACCTGGAGATTGGCCGTGCCCCTCTTGTTTTTTCTTAAAACCAAGGGGGAAGG  
CCAGCTCTGAAATACAGTGGCGACTCGAGGTCTTACTTCTTCCCCTGGCAGTAAAACCTCTTG  
CTCGCAACGGGACTAGTGACCTTGTGTTAGCCATCTTAACACCCTAT



FIGURE S1. R SCRIPT USED: GENERATION OF A .MGF AND .CSV FILES

```
```{r setup, include=FALSE}

library(MALDIquant)
library(MALDIquantForeign)

path <- "C:/Users/Desktop/..." #enter directory path
setwd(path)

spectra <- import(path, verbose=FALSE, excludePattern = NULL)

liste_fichiers <- list.files(path)
NameListe = as.list(liste_fichiers)
condition = c(15,      ## HwS
              30,      ## baseline
              6,       ## SnR
              2000, 20000) ## Trim
spectra2 = transformIntensity(spectra, method="sqrt")
spectra2 = trim(spectra2, range = c(condition[4], condition[5] ))
spectra2 = smoothIntensity(spectra2, method="SavitzkyGolay",
                           halfWindowSize=condition[1])
spectra2 <- removeBaseline(spectra2, method="SNIP",
                           iterations=condition[2])
spectra2 <- calibrateIntensity(spectra2, method="TIC")
peaks <- detectPeaks(spectra2, SNR=condition[3],
halfWindowSize=condition[1])
a = plot(peaks[[1]])

peaks <- binPeaks(peaks)

liste = list()
```

```

mgf <- paste(path, ".mgf", sep = "")

for (i in 1:length(peaks)) {

  test = matrix(data = NA, nrow = length(peaks[[i]]@mass), ncol= 2)

  colnames(test) = c("m/z", "intensity")

  test[,1] = peaks[[i]]@mass

  test[,2] = peaks[[i]]@intensity


  write.table("BEGIN IONS", file = mgf, append = TRUE, quote = FALSE, sep
= "\t ", row.names = FALSE, col.names = FALSE)

  write.table("PEPMASS=0", file = mgf, append = TRUE, quote = FALSE, sep
= "\t ", row.names = FALSE, col.names = FALSE)


  write.table(list(test), file = mgf, append = TRUE, quote = FALSE, sep
= "\t ", row.names = FALSE, col.names = FALSE)

  write.table("END IONS", file = mgf, append = TRUE, quote = FALSE, sep
= "\t ", row.names = FALSE, col.names = FALSE)

  write.table("", file = mgf, append = TRUE, quote = FALSE, sep = "\t ",
row.names = FALSE, col.names = FALSE)
}

csv <- paste(path, ".txt", sep = "")

for (i in 1:length(liste_fichiers)) {

  write.table(liste_fichiers[i], file = csv, append = TRUE, quote =
FALSE, sep = "\t ", eol = "\n", na = "NA", dec = ".", row.names = FALSE,
col.names = FALSE, qmethod = c("escape", "double"))
}

...

```

TABLE S6. PARAMETERS USED FOR CONSTRUCTING CHEMOTAXONOMIC NETWORKS ON METGEM

Figures	Treat as MS <sup>1</sup> Data	<i>m/z</i> Tolerance	Minimum Matched Peaks	Keep peaks above	Cosine score(s) above	Number of iterations	Learning Rate	Perplexity
2	Used	2	2	1%	0.65	10 000	10	11
3	Used	2	2	1%	0.4	10 000	10	11
4	Used	0.1	4	1%	0.65	10 000	10	11
5	Used	4	2	1%	0.35	10 000	10	16
6	Used	0.1	4	1%	0.45	10 000	10	11