



Bacteroides uniformis extra-ERR982822.32\_1677794517 (GCA\_949521815.1)  
Bacteroides uniformis MGBC101181 (GCA\_948557725.1)  
Bacteroides uniformis 34\_110531.37\_1677753728 (GCA\_949387155.1)  
Bacteroides uniformis MGBC108267 (GCA\_948504755.1)  
Bacteroides uniformis dnLKV2 (GCA\_000403175.2)  
Bacteroides uniformis MGBC165632 (GCA\_948810825.1)  
Bacteroides uniformis extra-ERR982846.41\_1677794780 (GCA\_949452615.1)  
Bacteroides uniformis SRR15730556\_bin.70\_metawrap\_v1.3\_MAG (GCA\_947642775.1)  
Bacteroides uniformis ERR8099105\_bin.5\_metawrap\_v1.3\_MAG (GCA\_947036735.1)  
Bacteroides uniformis MGBC165663 (GCA\_948878225.1)  
Bacteroides uniformis MGBC101827 (GCA\_948940695.1)  
Bacteroides uniformis D-Fe5\_120416.25\_1677753879 (GCA\_949383745.1)  
Bacteroides uniformis MGBC108160 (GCA\_949011935.1)  
Bacteroides uniformis extra-ERR982795.22\_1677760209 (GCA\_949469305.1)  
Bacteroides uniformis 10\_110627.34\_1677752997 (GCA\_949386435.1)  
Bacteroides uniformis MGBC108142 (GCA\_949085795.1)  
Bacteroides uniformis MGBC108246 (GCA\_948575575.1)  
Bacteroides uniformis D-Fe7\_120416.58\_1677753913 (GCA\_949404485.1)  
**Bacteroides uniformis MGBC106141 (GCA\_949011115.1)**  
Bacteroides uniformis MGBC106314 (GCA\_948586835.1)