

redlosses_phyloseq_custom.R

Stephane CHAILLOU, Simon POIRIER

Abstract

R script used for the phyloseq analysis of the Poirier et al., food microbiota dataset and comparative analysis between 16S and gyrB amplicon sequencing.

Citation: Stephane CHAILLOU, Simon POIRIER redlosses_phyloseq_custom.R. protocols.io

dx.doi.org/10.17504/protocols.io.rhsd36e

Published: 04 Jul 2018

Protocol