Sample Annotation file: Environmental Sample

This line is not required for annotation file. **Entry** Location Qualifier Value **Feature** COMMON SUBMITTER Hanako Mishima contact ab_name Mishima,H. "contact" is mandatory. ab_name Yamada,T. Kim,K.-H. Enter the full name. ab_name Park, C.S. ab name ab_name Liu, G.Q. mishima@ddbj.nig.ac.jp email Common information of all entries 81-55-981-6853 phone 81-55-981-6849 fax National Institute of Genetics institute department DNA Data Bank of Japan Entry name should be unique to Japan country all entries, i.e. clone name, isolate Shizuoka state name, etc. and identical to the Mishima city Yata 1111 one in the sequence file. street 411-8540 zip Enter the date to withhold your REFERENCE ab_name Mishima,H. data. If the data should be Yamada,T ab_name publicized immediately, delete Kim,K.-H. ab_name the DATE line. Park, C.S. ab_name These lines are mandatory, ab_name Liu, G.Q. when the sequences are Phylogenetic diversity of 16S ANA from title derived from environmental uncultured bacteria sample. Unpublished status 2017 year DATE 20191130 hold_date The identified qualifier, ie. DIVISION ▼ division ENV clone or isolate name, is **KEYWORD** keyword ENV necessary for uncultured CLN01 1..E A04 source clone organism entry. entry uncultured bacterium organism genomic DNA mol_type First environmental_sample solid waste compost isolation_source Japan:Okinawa country The identified qualifier, ie. rRNA <1..>E product 16S rRNA clone or isolate name, is CLN02 source clone B01 necessary for uncultured Second entry uncultured bacterium organism organism entry. When the end of the mol_type genomic DNA environmental_sample feature is truncated, add "<" or ">". solid waste compost isolation_source Taiwan:Tainan country <1..>E rRNA product 16S rRNA

Meta-base position, "E", can be used to describe location of a feature instead of numerical number of sequence end point, if a given feature, including source, spans full length of a submitted sequence. Please visit a following site for detail,

http://www.ddbj.nig.ac.jp/sub/mss/file_formate.html#annotation=4

When the organism is uncultured and the sequence is derived by direct molecular isolation (PCR, etc.), "environmental_sample" and "isolation_source" are mandatory.

The qualifier "environmental_sample" is used without value.

