Sample Annotation File: GSS (Genome Survey Sequence)

This line is not required for annotation file.

Entry	Feature	Location	Qualifier	Value
COMMON	SUBMITTER		contact	Hanako Mishima
			ab_name	Mishima,H.
			ab_name	Yamada,T.
			ab_name	Park,C.S.
			ab_name	Liu,G.Q.
			email	mishima@ddbj.nig.ac.jp
			phone	81-55-981-6853
			fax	81-55-981-6849
			institute	National Institute of Genetics
			department	DNA Data Bank of Japan
			country	Japan
			state	Shizuoka
			city	Mishima
			street	Yata 1111
			zip	411-8540
	REFERENCE		ab_name	Mishima,H.
			ab_name	Yamada,T.
			ab_name	Park,C.S.
			ab_name	Liu,G.Q.
			title	Arabidopsis thaliana DNA
			status	Unpublished
			year	2017
	DATE		hold_date	20191130
	DIVISION		division	GSS
	KEYWORD		keyword	GSS
ENT001	source	1..E	clone	CIC5D1
			clone_lib	AT01 YAC
			organism	Arabidopsis thaliana
			mol_type	genomic DNA
			chromosome	1
			ecotype	columbia
			ff_definition	@@[organism]@@ DNA, clone: @@[clone]@@, left end, chromosome @@[chromosome]@@, between mi103 and mi169
			map	between mi103 and mi169
ENT002	source	1..E	clone	CIC6D5
			clone_lib	AT01 YAC
			organism	Arabidopsis thaliana
			mol_type	genomic DNA
			chromosome	3
			ecotype	columbia
			ff_definition	@@[organism]@@ DNA, clone: @@[clone]@@, left end, chromosome @@[chromosome]@@, between mi303 and mi259
			map	between mi303 and mi259

Common information of all entries

First entry

Second entry

"contact" is mandatory. Enter the full name.

Enter the date to withhold your data. If the data should be publicized immediately, delete the

Entry name should be unique to all entries, i.e. clone name, isolate name, etc. and identical to the one in the sequence file.

"clone" is mandatory for GSS entry.

Metadecription can be used to represent value for some qualifiers, such as organism, clone, in ff_definition. Please visit the following site for deatil.
http://www.ddbj.nig.ac.jp/sub/mss/file_format-e.html#annotation-15

DDBJ
DNA Data Bank of Japan