

Sample Annotation File: ITS (Internal Transcribed Spacer)

This line is not required for annotation file.

	Entry	Feature	Location	Qualifier	Value
Common information of all entries	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
First entry		REFERENCE		ab_name	Mishima,H.
				ab_name	Yamada,T.
				ab_name	Kim,K.-H.
				ab_name	Liu,G.Q.
				title	Diversity of fungus DNA
				status	Unpublished
				year	2017
		DATE		hold_date	20191130
	ENT01	source	1..E	isolate	MUMH29s
				organism	Erysiphe trifolii var. trifolii
				variety	trifolii
				mol_type	genomic DNA
				host	Trifolium vulgaris
				tissue_type	conidia
		rRNA	<1..4	product	18S rRNA
	misc_RNA	5..224	note	internal transcribed spacer 1	
	rRNA	225..378	product	5.8S rRNA	
	misc_RNA	379..562	note	internal transcribed spacer 2	
	rRNA	563..>597	product	28S rRNA	
Second entry	ENT02	source	1..E	isolate	MUMH30s
				organism	Erysiphe trifolii var. trifolii
				variety	trifolii
				mol_type	genomic DNA
				host	Trifolium vulgaris
				tissue_type	conidia
		misc_RNA	<1..>597	note	contains 18S rRNA, ITS1, 5.8S rRNA, ITS2 and 28S rRNA

Common information of all entries

First entry

Second entry

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

When the end of the feature is truncated, add "<" or ">".

"contact" is mandatory. Enter the full name.

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

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_format-e.html#annotation-4