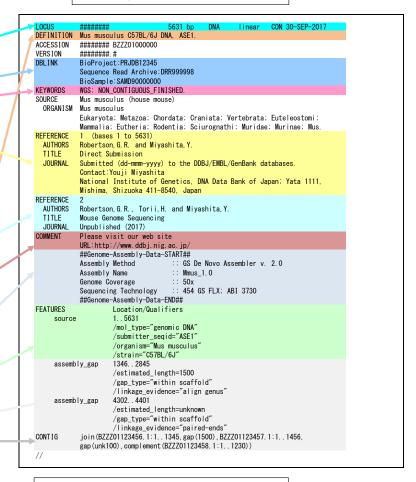
## CON data for WGS scaffold

## Input: Annotation file

Entry	Feature	Location	Qualifier	Value			
COMMON	DATE		hold_date	20191130			
	DATATYPE		type	WGS			
	DIVISION		division	CON			
	DBLINK		project	PRJDB12345 SAMD9000000			
			biosample				
			sequence read archive	DRR999998			
	KEYWORD		keyword	WGS			
			keyword	NON CONTIGUOUS FINISHED			
	SUBMITTER		ab_name	Robertson, G. R.			
			ab_name	Miyashita, Y.			
			contact	Youji Miyashita			
			institute	National Institute of Genetics			
			department	DNA Data Bank of Japan			
			country	Japan			
			state	Shizuoka			
			city	Mishima			
			street	Yata 1111			
			Zip	411-8540			
			ema i l	mishima@ddbj.nig.ac.jp			
				81-55-981-6853			
			phone	81-55-981-6849			
			fax				
	DEEEDENOE		phext	3207			
	REFERENCE		title	Mouse Genome Sequencing			
			ab_name	Robertson, G. R.			
			ab_name	Torii, H.			
			ab_name	Miyashita, Y.			
			year	2017			
			status	Unpub I i shed			
	COMMENT		line	Please visit our web site			
			line	URL:http://www.ddbj.nig.ac.jp/			
	ST_COMMENT		tagset_id	Genome-Assembly-Data			
			Assembly Method	GS De Novo Assembler v. 2.0			
			Assembly Name	Mmus_1.0			
			Genome Coverage	50x			
			Sequencing Technology	454 GS FLX; ABI 3730			
scaffold1	source	1 E	ff_definition	@@[organism]@@ @@[strain]@@ DNA, @@[submitter_seqid]@@			
			organism	Mus musculus			
			mol_type	genomic DNA			
			strain	C57BL/6J			
			submitter_segid	ASE1			
	assembly_gap	1346 2845	estimated_length	known			
			gap_type	within scaffold			
			linkage_evidence	align genus			
	assembly_gap	4302 4401	estimated_length	unknown			
	association y_gap	.502 4401	gap_type	within scaffold			
			linkage evidence	paired-ends			
			I IIINAGE_EVIUETICE	pari cu cius			



Output: DDBJ flat file



Input: AGP file

	object_ beg		part_ number	COMPONENT				orientation or linkage_evidence
scaffold1	1	1345	1	W	BZZZ01123456. 1	1	1345	+
scaffold1	1346	2845	2	N	1500	scaffold	yes	align_genus
scaffold1	2846	4301	3	W	BZZZ01123457. 1	1	1456	+
scaffold1		4401	4	U	100	scaffold	yes	align_trnscpt
scaffold1	4402	5631	5	W	BZZZ01123458. 1	1	1230	_