

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
			biosample	SAMD900000000
			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
			email	mishima@ddbj.nig.ac.jp
			phone	81-55-981-6853
			fax	81-55-981-6849
			phext	3207
	REFERENCE		title	Mouse Genome Sequencing
			ab_name	Robertson, G. R.
			ab_name	Torii, H.
			ab_name	Miyashita, Y.
			year	2017
			status	Unpublished
	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_seqid	ASE1
	assembly_gap	1346..2845	estimated_length	known
			gap_type	within scaffold
			linkage_evidence	align genus
	assembly_gap	4302..4401	estimated_length	unknown
			gap_type	within scaffold
			linkage_evidence	paired-ends

Output: DDBJ flat file

```

LOCUS      #####          5631 bp   DNA   linear   CON 30-SEP-2017
DEFINITION Mus musculus C57BL/6J DNA, ASE1.
ACCESSION  ##### BZZZ01000000
VERSION    #####. #
DBLINK     BioProject:PRJDB12345
           Sequence Read Archive:DRR999998
           BioSample:SAMD900000000
KEYWORDS   WGS: NON_CONTIGUOUS_FINISHED.
SOURCE     Mus musculus (house mouse)
ORGANISM   Mus musculus
           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
           Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
REFERENCE  1 (bases 1 to 5631)
AUTHORS    Robertson, G. R. and Miyashita, Y.
TITLE      Direct Submission
JOURNAL    Submitted (dd-mmm-yyyy) to the DDBJ/EMBL/GenBank databases.
           Contact: Youji Miyashita
           National Institute of Genetics, DNA Data Bank of Japan: Yata 1111,
           Mishima, Shizuoka 411-8540, Japan
REFERENCE  2
AUTHORS    Robertson, G. R., Torii, H. and Miyashita, Y.
TITLE      Mouse Genome Sequencing
JOURNAL    Unpublished (2017)
COMMENT     Please visit our web site
           URL: http://www.ddbj.nig.ac.jp/
           ##Genome-Assembly-Data-START##
           Assembly Method      : GS De Novo Assembler v. 2.0
           Assembly Name        : Mmus_1.0
           Genome Coverage      : 50x
           Sequencing Technology : 454 GS FLX: ABI 3730
           ##Genome-Assembly-Data-END##
FEATURES   Location/Qualifiers
           source                1..5631
                                   /mol_type="genomic DNA"
                                   /submitter_seqid="ASE1"
                                   /organism="Mus musculus"
                                   /strain="C57BL/6J"
           assembly_gap          1346..2845
                                   /estimated_length=1500
                                   /gap_type="within scaffold"
                                   /linkage_evidence="align genus"
           assembly_gap          4302..4401
                                   /estimated_length=unknown
                                   /gap_type="within scaffold"
                                   /linkage_evidence="paired-ends"
CONTIG     join(BZZZ01123456.1:1..1345, gap(1500), BZZZ01123457.1:1..1456,
           gap(unk100), complement(BZZZ01123458.1:1..1230))
           //
  
```



Input: AGP file

#object	object_	object_	part_	component	component_	component_beg	component_end	orientation
#	beg	end	number	type	id	or	or	or
					or gap length	gap_type	linkage	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	4302	4401	4	U	100	scaffold	yes	align trnsct
scaffold1	4402	5631	5	W	BZZZ01123458.1	1	1230	-