## HTG data

## Input: Annotation file

Entry	Feature	Location	Qualifier	Value
COMMON	DATE		hold_date	20191130
	SUBMITTER		contact	Hanako Mishima
			ab_name	Mishima, H.
			ab_name	Yamada, T.
			ab_name	Park, C. S.
			ab_name	Liu, G. Q.
			ema i I	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.
	KLI LKLIIOL		ab_name	Yamada, T.
			ab_name	Park, C. S.
			ab_name	Liu, G. Q.
			title	Arabidopsis thaliana DNA
				2017
			year	
			status email	Unpublished
			phone	mishima@ddbj.nig.ac.jp 81-55-981-6853
			fax	81-55-981-6849
				3207
	DIVICION		phext	
	DIVISION		division	HTG
	KEYWORD		keyword	HTG
	COMMENT		keyword	HTGS_PHASE1
	COMMENT		line	Please visit our web site
		_	line	URL; http://ddbj.nig.ac.jp
ENTO01	source	1 E	clone	CIC5D1
			clone_lib	ATO1 BAC
			organism	Arabidopsis thaliana
			mol_type	genomic DNA
			chromosome	1
			ecotype	columbia
			ff_definition	@@[organism]@@ DNA, BAC clone: @@[clone]@@, chromosome @@[chromosome]@@, *** SEQUENCING IN PROGRESS ***
	assembly_gap	50015100	estimated_length	unknown
			gap_type	within scaffold
			linkage_evidence	within clone
	assembly_gap	2400124100	estimated_length	unknown
			gap_type	within scaffold
			linkage_evidence	within clone
	assembly_gap	6200162100	estimated_length	unknown
			gap_type	within scaffold
			gap_type linkage evidence	within scaffold within clone
		105001105100	linkage_evidence	within clone
	assembly_gap	105001105100		



## Output in DDBJ flat file

```
123456 bp DNA linear HTG 30-SEP-2017
DEFINITION Arabidopsis thaliana DNA, BAC clone: CIC5D1, chromosome 1, ***
           SEQUENCING IN PROGRESS ***
ACCESSION
          #############
VERSION
           ##########. 1
DBLINK
KEYWORDS
           HTG: HTGS PHASE1.
           Arabidopsis thaliana (thale cress)
  ORGANISM Arabidopsis thaliana
           Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
           Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons;
           rosids; malvids; Brassicales; Brassicaceae; Camelineae;
           Arabidopsis.
REFERENCE 1 (bases 1 to 123456)
  AUTHORS Mishima, H., ada, T., Park, C.S. and Liu, G.Q.
  TITLE
           Direct Submission
  JOURNAL Submitted (dd-mmm-yyyy) to the DDBJ/EMBL/GenBank databases.
           Contact:Hanako Mishima
           National Institute of Genetics, DNA Data Bank of Japan; Yata 1111,
           Mishima, Shizuoka 411-8540, Japan
REFERENCE 2
  AUTHORS Mishima, H., ada, T., Park, C.S. and Liu, G.Q.
  TITLE
           Arabidopsis thaliana Sequencing
  JOURNAL Unpublished (2017)
           Please visit our web site
           URL:http://www.ddbj.nig.ac.jp/
FEATURES
                    Location/Qualifiers
                    1...123456
    source
                    /chromosome="1"
                    /clone="CIC6D5"
                    /clone_lib="AT01 BAC"
                    /ecotype="columbia"
                    /mol_type="genomic DNA"
                    /organism="Arabidopsis thaliana"
                    5001..5100
    assembly_gap
                    /estimated length="unknown"
                    /gap_type="within scaffold"
                    /linkage_evidence="within clone"
    assembly_gap
                    24001...24100
                    /estimated_length="unknown"
                    /gap_type="within scaffold"
                    /linkage_evidence="within clone"
    assembly_gap
                    62001..62100
                    /estimated_length="unknown"
                    /gap_type="within scaffold"
                    /linkage_evidence="within clone"
     assembly_gap
                   105001...105100
                    /estimated_length="unknown"
                    /gap_type="within scaffold"
                    /linkage_evidence="within clone"
(The lines for BASE COUNT, ORIGIN, and nucleotide sequence are omitted)
```