paten_MKAFGlu1.txt

Submission date: 01-Apr-2020 11:36AM (UTC+0700)

Submission ID: 1286766228

File name: paten_MKAFGlu1.txt (2.62K)

Word count: 242

Character count: 1941

LOCUS MKAFGlu1 717 bp mRNA linear INV 15-APR-2018

DEFINITION UNVERIFIED: [Achatina fulica] 1,3-beta-Glucanase, complete CDS.

ACCESSION MKAFGlu1

VERSION

KEYWORDS UNVERIFIED.

SOURCE mitochondrion Achatina fulica (giant African snail)

ORGANISM Achatina fulica

Eukaryota; Metazoa; Lophotrochozoa; Mollusca; Gastropoda;

Heterobranchia; Euthyneura; Panpulmonata; Eupulmonata;

Stylommatophora; Sigmurethra; Achatinoidea; Achatinidae; Achatina.

REFERENCE 1 (bases 1 to 717)

AUTHORS Kurniawati, M. and Baktir, A.

TITLE A novel 1,3-beta-glucanase gene from the metagenomic expression library of Achatina fulica's digestive gland

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 717)

AUTHORS Kurniawati, M. and Baktir, A.

TITLE Direct Submission

JOURNAL Submitted (15-APR-2018) Science and Technology, Kanjuruhan University, S. Supriyadi 48, Malang, East Java 65148, Indonesia

COMMENT GenBank staff is unable to verify sequence and/or annotation provided by the submitter.

Bankit Comment: ALT EMAIL:mrskurniawati@gmail.com.

Bankit Comment: TOTAL # OF SEQS:1.

##Assembly-Data-START##

Assembly Method :: Alignment v. Clustal W

Sequencing Technology :: HiSeq2000/2500

##Assembly-Data-END##

FEATURES Location/Qualifiers

source 1..717

/organism="Achatina fulica"

/organelle="mitochondrion"

/mol_type="mRNA"

/db_xref="taxon:6530"

BASE COUNT 144 a 183 c 146 g 244 t

ORIGIN

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61 tgccagtcca tggtccaagg acttcctagg tggcctcctc gctcttctct tgggctgcac

121 tgctgctgcg attgttgctt atgcttttgc tattactgtg gctttgggtc acattcttgt

181 gcgggcagtc ggcatggaga tggtcttctc catgatgagc ggagtggatt cattccatta

241 totaagcagg ggotcatotc actoacacac togcaccage coatatocat ttgggotggc

301 teatteatte atteatteat teatteatta attgttttea ettegetege teteattteg

361 ctgccagctt atctatatat ttgtcgaagt gggcccaagt acttgagtgt actcgtagta

421 ttgcccagat gcccagatac tcggaatgca ttctcatgcc tgtgctctgg cttatctgat

481 gaagtggaac totggotggt ggtoctoott tocaagcoat tttocagctg tacgttttoo

	541 atgctgactt cttgggtttt tagtttattt ttcttagttc aaaatccact tcaattgttt
	601 gatttetege tecaaaatte gatgttgett aaaaetteae tgeaaatgea tgttttteet
	661 gccacgaagt cggggtcccc aaaaaaaaac agactaacca ggaaaccctc gccatcc
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