

# paten\_MKAFGlu1.txt

*by*

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**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JOURNAL Unpublished

<sup>2</sup> REFERENCE 2 (bases 1 to 717)

<sup>1</sup> 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

1 atgcgacgtt ctttcaaaa tgcgttttg tactgtcaa ggaagttgt agctccaac  
61 tgccagtcca tggccaagg acttctagg tggcctcctc gctcttctt tgggctgcac  
121 tgcgctgcg attgttgc tgcctttgc tattactgtg gctttgggtc acattctgt  
181 gcgggcagtc ggcattgaga tggcttctc catgatgagc ggagtggatt cattccatta  
241 tctaagcagg ggctcatctc actcacacac tcgaccagc ccatatccat ttgggtggc  
301 tcattcattc attcattcat tcattcatta attgtttca ctcgctcgc tctcattcg  
361 ctgccagctt atctatatat ttgtcgaagt gggcccaagt acttgagtgt actcgtagta  
421 tgcccagat gccagatac tcggaatgca ttctatgcc tgtgctctgg cttatctgat  
481 gaagtggaac tctggctggt ggtcctcctt tccaagccat ttccagctg tacgtttcc

541 atgctgactt ctggggttt tagttatt ttctagtc aaaatccact tcaattgtt

601 gatttctgc tccaaaatc gatgtgctt aaaactcac tgcaaatgca tgttttcct

661 gccacgaagt cgggggtccc aaaaaaaaaac agactaacca ggaaaccctc gccatcc

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