

GENESEQ™ on STN®



GENESEQ™ (file DGENE) covers peptide and nucleic acid sequences from the basic patent publications of 52 authorities of the Derwent World Patents Index (DWPI).

DGENE offers three sequence searching methods

- BLAST for advanced similarity searching based on NCBI BLAST® algorithm
- GETSIM for advanced similarity searching based on FASTA algorithm
- GETSEQ for simple fragment or motif sequence queries

Biosequences in DGENE

- Peptide and nucleic acid sequences from 1981 to date
- From the basic patent publications of 41 authorities of DWPI
- Organism and sequence length
- Feature tables for modifications and other features

DGENE records also provide

- English abstracts written per sequence by Clarivate Analytics experts
- Enhanced patent titles and bibliographic data from DWPI
- Integrated Patent Family and Legal Status display
- Full-text links to Esp@cenet and the USPTO

GENESEQ™ is produced by Clarivate Analytics and is provided on STN® as file DGENE in association with FIZ Karlsruhe.

ACCESSION NUMBER: AAE10698 Protein **1** DGENE

TITLE: **2** An isolated polypeptide (I) possessing beta-(1,3) exoglucanase activity for improvement of plant resistance to fungal phytopathogens and to promote growth

INVENTOR: Frick M M; Huang T Y; Cheng K J; Lu Z; Laroche A J; Huang H C

PATENT ASSIGNEE: (MIAC) CANADA MIN AGRIC & AGRI-FOOD CANADA.

PATENT INFO: CA 2325774 A1 20010610 **3** 86p

APPLICATION INFO: CA 2000-2325774 20001208

PRIORITY INFO: US 1999-170168P **4** 19991210

PAT. SEQ. LOC: Claim 8; fig 2

DATA ENTRY DATE: 10 DEC 2001 (first entry)

DOCUMENT TYPE: Patent

LANGUAGE: English

OTHER SOURCE: 2001-409063 [44] **5**

CROSS REFERENCES: N-PSDB: AAD18016

DESCRIPTION: Coniothyrium minitans beta-(1,3) exoglucanase, cbeg1.

KEYWORD: **6** Beta-(1,3) exoglucanase gene; cbeg1; laminarin; plant resistance; fungal phytopathogen; microbial transgenic strategy; feed digestion; forage feed; industrial application; pulp bleaching; monocot; dicot; antifungal; growth promoter; EC 3.2.1.58.

ORGANISM: Coniothyrium minitans. **7**

ABSTRACT:

The invention relates to nucleotide sequence of a novel beta-(1,3) exoglucanase gene denoted as cbeg1 of the soil borne fungus Coniothyrium minitans. Beta-(1,3) exoglucanase (EC 3.2.1.58) is an enzyme that catalyses the successive hydrolysis of beta-D-glucose units from the non-reducing ends of 1,3-beta-D-glucans, releasing alpha-glucose. cbeg1 is specific for the substrate laminarin. cbeg1 sequences are useful for improvement of plant resistance to fungal phytopathogens or use in ruminant microbial transgenic strategies to improve feed digestion and nutritive carbohydrate availability from forage feed. cbeg1 is also useful for use in high temperature industrial applications such as bleaching of pulp. cbeg1 is useful as an antifungal in dicots and to promote plant growth in monocots and dicots. The present sequence is Coniothyrium minitans cbeg1 protein.

AMINO ACID COUNTS: 73 A; 19 R; 61 N; 39 D; 0 B; 11 C; 33 Q; 18 E; 0 Z; **8** 77 G; 13 H; 52 I; 50 L; 31 K; 14 M; 25 F; 42 P; 68 S; 59 T; 16 W; 29 Y; 55 V; 0 Others

SEQUENCE LENGTH: 785

SEQUENCE

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1 mrlsffscl laagppasal alpspianda tsapleerqa ssywleniqh
51 qgraafnanp agykvfrnvk dygakgdgvt ddsaaainai adgnrcapwv
101 cdsstdtpai vyfpgstyvi gkpiimyymt qlhgnpnnrp vlkaspnlra 9
151 ialidaspyq dgtgkpgwts tnvftrqirn fvidltpipa tsgaqgihwp
201 asqatsiqdv kiqmnvaans vhgifieng sgghltdiet vggllhglnvq
251 nqgftmkniv isnavvginq iwnwgwllwkgt ltisdcstaa fsmkslkdns
301 pdqnvsvii idstitncpi fvdswtrts taagsgqlil enialnnvpv
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FEATURE TABLE: **10**

Key	Location	Qualifier	
Region	1..337	note	"N-terminal region"
Peptide	1..21	label	Signal_peptide
Protein	22..785	note	"Coniothyrium minitans mature cbeg1 protein"
Domain	63..82	label	GAK_box
Region	76..82	note	"Region targetted by Gf1 semidegenerate primer"
Region	338..785	note	"C-terminal region"
Domain	425..434	label	GAX_box

1 Molecule Type (MTY)

2 Clarivate Analytics enhanced patent title describing the over all invention

3 Bibliographic information – Publication, application, assignee & inventor data

4 Patent Sequence Location – claim, example, etc

5 Other Source accession number – link to Derwent World Patents Index® patent family data

6 Clarivate Analytics description, keywords and enhanced abstract describe the context and use of each patent sequence

7 Organism – latin genus & species, from which the sequence is derived

8 Sequence Length and individual amino acid & nucleotide count fields

9 Patent sequence – each DGENE record is based upon a sequence

10 Feature table – includes modifications and features added by the applicant, and by the Clarivate Analytics Analyst