

Fusarium graminearum chromosome 4, complete genome

GenBank: HG970335.2

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LOCUS HG970335 448 bp DNA linear PLN 07-JUL-2017

DEFINITION Fusarium graminearum chromosome 4, complete genome.

ACCESSION [HG970335](#) REGION: 6206552..6206999

VERSION HG970335.2

DBLINK BioProject: [PRJEB5475](#)
BioSample: [SAMEA3283145](#)

KEYWORDS complete genome.

SOURCE Fusarium graminearum (Gibberella zeae)

ORGANISM [Fusarium graminearum](#)
Eukaryota; Fungi; Dikarya; Ascomycota; Pezizomycotina;
Sordariomycetes; Hypocreomycetidae; Hypocreales; Nectriaceae;
Fusarium.

REFERENCE 1

AUTHORS King,R., Urban,M., Hammond-Kosack,M.C., Hassani-Pak,K. and
Hammond-Kosack,K.E.

TITLE The completed genome sequence of the pathogenic ascomycete fungus
Fusarium graminearum

JOURNAL BMC Genomics 16, 544 (2015)

PUBMED [26198851](#)

REMARK Publication Status: Online-Only

REFERENCE 2 (bases 1 to 448)

AUTHORS King,R.

TITLE Direct Submission

JOURNAL Submitted (19-FEB-2014) Applied Bioinformatics, ROTHAMSTED
RESEARCH, W Common, Harpenden, AL5 2JQ, UNITED KINGDOM

COMMENT On Aug 3, 2016 this sequence version replaced [HG970335.1](#).

FEATURES Location/Qualifiers

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mRNA complement(<1..>448)

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ORIGIN

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