Plant Molecular Biology **21**: 581–582, 1993.

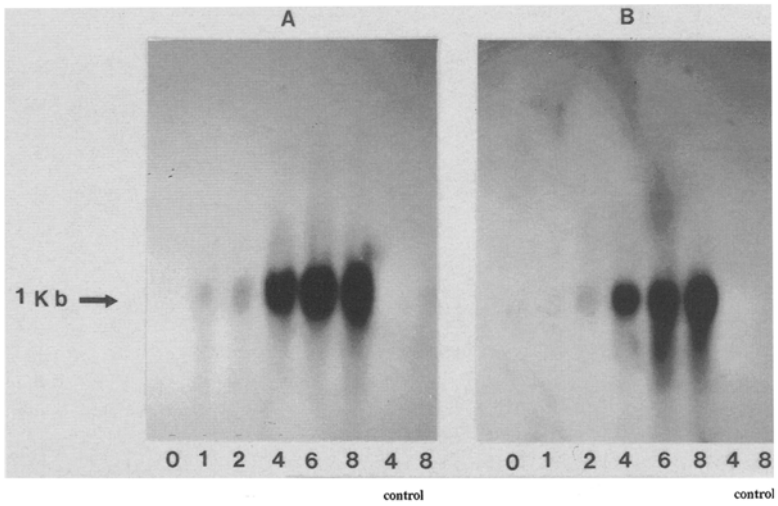
Erratum

Cloning and characterization of a pathogen-induced chitinase in *Brassica napus*

Ulla Rasmussen [1,3], Kirsten Bojsen [2] and David B. Collinge [1]
[1] *Plant Biology Section, Environmental Science and Technology Department, Risø National Laboratory, P.O. Box 49, DK-4000 Roskilde, Denmark;* [2] *Maribo Seed Biotechnology Copenhagen, Danisco A/S, Langebrogade 1, DK-1001 Copenhagen K, Denmark;* [3] *Present address: Botaniska Institutionen, Stockholms Universitet, S-1069 Stockholm, Sweden*

Plant Molecular Biology **20**: 277–287, 1992.

In Fig. 5 of this paper, the lanes were incorrectly marked. The correct figure is as follows:



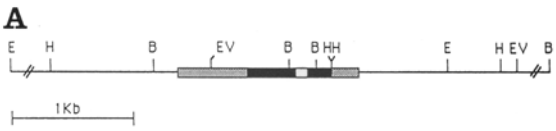
Erratum

The expression of a *rab*-related gene, *rab 18*, is induced by abscisic acid during the cold acclimation process of *Arabidopsis thaliana* (L.) Heynh.

Viola Lång and E. Tapio Palva
Department of Molecular Genetics, Uppsala Genetic Center, Swedish University of Agricultural Sciences, Box 7003, S-750 07 Uppsala, Sweden

Plant Molecular Biology **20**: 951–962, 1992.

When issue 20:5 was being printed, Fig. 2c of the above-mentioned paper was erroneously deleted from the page. The printers apologize for this error and herewith place it as it should have been:



B

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-613                                                    GATCTAAACGCGG
-600  CGTTTGGTAAAAGTTGAGTAAATTTTGTTAGGGCTTAGTTTTAGTCCATG
-550  GGCTAATTAGTAAGTGATTTACGGCCCACACATGAGCCCAAATGTTTCAG
-500  ACCCAGCCAAGTTTCTTCAAATTCACCCAATCAACGACGATGTACGTGTG
-450  TATGAAAATCATTAACACGACGCATCGCTTTCGAGGAGGAGCATTACGTG
-400  TCCTGTTAGCTACGATAATGTTAGTACCGCCACAAAGAAAAGGATAGATA
-350  TTTTGCTTTCCAGCACCCTGTCATGGGATTGATATGAACACGTACTTGGT
-300  ATCGACATGAAAGCTCAAAAATAAATTCAATCCGATTCCTTTAGTGATAT
-250  CAGAAGTTCATTTTAAATACGAACACGTATGGCGAAACACCACGCCGACA
-200  TTTTCTGCTGCTGCCACGCGTCACTTTCCAAATATTGATTCATTAAACTA
-150  ATAGTTGATCCATATCCGAAACCGGACTATAAAACTATCTTCAATGCGTT
-100  AACGAATCTTCATCGATCAAACTCATCAAAGTCTAATATCACAAAGAAAG
 -50  AGTTTTTTTAACTAGCTTAGCTCAAAGTGTTTGCTTAAGACAAGAAGAAC

   1  ATG GCG TCT TAC CAG AAC CGT CCA GGA GGT CAG GCC ACT
      M  A  S  Y  Q  N  R  P  G  G  Q  A  T

  40  GAC GAG TAC GGA AAC CCG ATC CAG CAG CAG TAT GAC GAG
      D  E  Y  G  N  P  I  Q  Q  Q  Y  D  E

  79  TAC GGA AAT CCG ATG GGA GGA GGA GGA TAC GGA ACT GGT
      Y  G  N  P  M  G  G  G  G  Y  G  T  G

 118  GGT GGT GGA GGA GCT ACA GGT GGC CAA GGA TAC GGA ACA
      G  G  G  G  A  T  G  G  Q  G  Y  G  T

 157  GGT GGC CAA GGG TAC GGA TCA GGT GGC CAA GGG TAC GGA
      G  G  Q  G  Y  G  S  G  G  Q  G  Y  G

 196  ACC GGT GGC CAA GGA TAC GGA ACC GGG ACC GGG ACT GAA
      T  G  G  Q  G  Y  G  T  G  T  G  T  E

 235  GGC TTT GGA ACT GGC GGA GGA GCT AGG CAC CAC GGC CAA
      G  F  G  T  G  G  G  A  R  H  H  G  Q

 274  GAG CAA CTC CAC AAG GAA AGT GGT GGT GGC TTG GGA GGA
      E  Q  L  H  K  E  S  G  G  G  L  G  G

 313  ATG CTT CAC CGC TCC GGA TCT GGA TCC AGC TCT AGC TCG
      M  L  H  R  S  G  S  G  S  S  S  S  S

 352  GTACGTATCAAGTGTATAAAGAAGATTTATGATGTTTTAGTTTAGTATGTG
 403  AAGCGTAATTCGAATGTTATATGTATAACAG GAG GAT GAT GGA CAA
                                      E  D  D  G  Q

 449  GGA GGG AGG AGG AAG AAG GGA ATA ACA CAA AAG ATC AAG
      G  G  R  R  K  K  G  I  T  Q  K  I  K

 488  GAG AAG TTG CCA GGT CAT CAT GAT CAG TCT GGT CAA GCT
      E  K  L  P  G  H  H  D  Q  S  G  Q  A

 527  CAA GCG ATG GGC GGC ATG GGA TCC GGA TAT GAT GCT GGT
      Q  A  M  G  G  M  G  S  G  Y  D  A  G

 566  GGC TAC GGT GGT GAG CAC CAC GAG AAG AAG GGG ATG ATG
      G  Y  G  G  E  H  H  E  K  K  G  M  M

 605  GAC AAG ATC AAG GAA AAG CTT CCC GGT GGT GGC CGT TAA
      D  K  I  K  E  K  L  P  G  G  G  R  -

 644  GCTTCGAACAATCGTGTATACATATTAAATAAAAATAATGAGGGTTTGTA
 694  ACGCAGTCGCATTCGGTCGTTGTATTGTGCTTTTTATGTATGTACGTCCT
 744  GCGATGTGTGTTGTTTACTTACATGAGTGTGTAATGAGCATCTGGCTCTT
 794  TTTATGTTCTGAGATGTTTGTGTTATGTAATTTCACATCTATATAAATCT
 844  ACTTTCTTCTGCTTAGTTTGTGTTACACCTCTCGTCACTGTTGAACTCCG
 894  TTAACTGACACTACAATTAGTG
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