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Erratum

DNA databank accession numbers were omitted from two articles published in Volume 18 issue number 3. The complete titles are published below.

Nucleotide sequence of the *aroP* gene encoding the general aromatic amino acid transport protein of *Escherichia coli* K-12: homology with yeast transport proteins
 by Nadine Honoré and Stewart T.Cole

Nucleic Acids Research, **18**, p. 653 (1990) EMBL accession no. X17333

Structural homology between the human *fur* gene product and the subtilisin-like protease encoded by yeast *KEX2*
 by Ans M.W.van den Ouweland, Hans L.P.van Duijnhoven, Gerrit D.Keizer, Lambert C.J.Dorssers and Wim J.M.Van de Ven

Nucleic Acids Research, **18**, p. 664 (1990) EMBL accession no. X17094

Corrigendum

Nucleotide sequence of the coat protein gene and flanking regions of cucumber virus (CMV) strain I17F
 by Marianne J.T.Noel and Sofia Ben Tahar

Nucleic Acids Research, **17**, 10492 (1989)

An incorrect sequence was published with the above article. The correct sequence is as follows:

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AGAGAGTGTGTGTGTGTGTCTTTCTCTTTGTGTGTGTAGTAATTGAGTCGAGTCATGGACAAATCTGAATCAACCA
                                     M D K S E S T
GTGCTGGTGTGTAACCGTCGACGTCGTCGCGCTCGTGGTTCGCGCTCCGCCCCCTCCTCCGCGGATGCTAACTTTA
S A G R N R R R R R P R R G S R S A P S S A D A N F
GAGTCTTGTGCGCAGCAGCTTTCGCGACTTAATAAGCGTTAGCAGCTGGTGTGCTCAACTATTAACCCCAACCT
R V L S Q Q L S R L N K T L A A G R P T I N H P T
TTGTAGGGAGTGAACGCTGTAGACCTGGGTACACGTTACATCTATTACCCCTAAAGCCACCAAAATAGACCGTG
F V G S E R C R P G Y T F T S I T L K P P K I D R
GGTCTTATTACGGTAAAGGTTGTACTACCTGATTGAGTCAGTCACGGAATATGATAAGAAGCTTGTTCGCGCATTC
G S Y Y G K R L L L L P D S V T E Y D K K L V S R I
AAATTCGAGTTAATCTTTGCCGAAATTTGATTCTACCGTGTGGGTGACAGTCCGTAAAGTTCTCGCTCCTCGG
Q I R V N P L P K F D S T V W V T V R K V P A S S
ACTTATCCGTTGCGGCATCTCTGCTATGTTTCGCGGACGGAGCCTACCGGTACTGGTTATCAGTATGCGCGAT
D L S V A A I S A M F A D G A S P V L V Y Q Y A A
CTGAGTCCAAGCAACAACTGTTGTATGATCTTTCGCGCATGCGCGCTGATATAGGTGACATGAGAAAGT
S G V Q A N N K L L Y D L S A M R A D I G D M R K
ACGCGTCTCTGTATTCAAAAGACGATGCGCTAGAGACGGACGAGTAGTACTCATGTTGACATCGAGCACC
Y A V L V Y S K D D A L E T D E L V L H V D I E H
AACGATTCCTCCAGCTGCGAGTCTCCAGTCTGATTCTGTGTTCCAGAACTCTCCCTCCGATCTCTGTGGCGGG
Q R I P T S G V L P V
AGCTGAGTGGCAGTTCTGCTATAAACTGTCTGAAGTCACTAAAGTTTTTACGGTGAACGGGTGTCATCCAG
CTTACGGCTAAAATGGTCAGTCGTGGAGAAATCCACCGCAGTAGATTACAAATCTCTGAGGCGCCTTTGAAACC
ATCTCTAGGTTTCTTCGGAAGGACTTCGGTCCGTGACCTTAGCACAACGTGCTAGTTTCAGGGTACGGGTGC
CCCCCACTTTCGTGGGGGCTCCAAAAGGAG

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