

Screpanti, I. 1153  
 Seidah, N.G. 1298  
 Seifart, K.H. 1225  
 Sesboué, R. 1319  
 Sick, A. 1305  
 Simon, M.I. 1255  
 Singer-Sam, J. 1255  
 Sirugo, G. 1319  
 Skotnicki, M. 1181  
 Soller, M. 1129  
 Solymosy, F. 1295  
 Srifah, P. 1181  
 Stabel, L.F.E.M. 1294  
 Steinmetz, A. 1297  
 Stinissen, P. 1319

Strong, L.C. 1317  
 Suzuki, T. 1302  
 Szkukálek, A. 1295  
 Thole, J.E.R. 1294  
 Tietze, E. 1283  
 Tikochinsky, Y. 1129  
 Torkelson, S. 1099  
 Torronen, M. 1181  
 Ueda, T. 1217  
 Ueyama, H. 1318  
 Uhlenbeck, O.C. 1103  
 Vacca, A. 1153  
 Van Broeckhoven, C. 1319  
 van der Logt, P. 1189  
 Vandenbergh, A. 1319

Vassart, G. 1301  
 Vignault, J.C. 1293  
 Walker, M.D. 1159  
 Watson, C.J. 1167  
 Weigel, C. 1285  
 Weil, J.-H. 1115  
 Wetzstein, M. 1289  
 Wilson, G.G. 1145  
 Wilson, M.J. 1308  
 Winship, P.R. 1310  
 Woloszyn, N. 1303  
 Wong, A. 1305  
 Wu, J. 1316, 1316  
 Yamagishi, A. 1133  
 Zarling, D.A. 1099  
 Zhang, Z.H. 1284

**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:

```

AGAGAGTGTGTGCTGTGTTTTCTCTTTTGTGTGCTAGAATTGAGTCGAGTCATGGACAAATCTGAATCAACCA
                                     M D K S E S T
GTGCTGGTTCGTAACCGTCGACGCTCGTCCGGCTCGTGGTCCCGCTCCGCCCTCCTCCGGGATGCTAACTTTA
S A G R N R R R R P R R G S R S A P S S A D A N F
GAGCTTGTCCGACGACTTTCGCGACTTAATAAGCGTTAGCAGCTGGTCCCACTATTAACCCAACTT
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
TTGTAGGGAGTGAACGCTGTAGACCTGGGTACACGTTACATCTATTACCCCTAAAGCCACCAAAAATAGACCGTG
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
GGTCTTATTACGGTAAAGGTTGTTACTACCTGATTGATTCAGTCACGGAATATGATAAGAAGCTTGTTCGGCATT
G S Y Y G K R L L L P D S V T E Y D K K L V S R I
AAATTCGAGTTAATCCTTTGCCGAAATTTGATTCACCGTGGGTGACAGTCCGTAAGTTCCTGCCTCCTCGG
O I R V N P L P K F D S T V W V T V R K V P A S S
ACTTATCCGTTCCGCCATCTCTGCTATGTTCCGGACGGAGCCTCACCGTACTGGTTATCAGTATGCCGAT
D L S V A A I S A M F A D G A S P V L V Y O Y A A
CTGAGTCCAAGCAACAACAACTGTTGATGATCTTTCGGCGATGGCGCTGATATAGGTGACATGAGAAAGT
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
ACGCCCTCCTCGTGTATTCAAAGACGATGCCCTAGAGACGGACGAGCTAGTACTTATGTTGACATCGAGCACC
Y A V L V Y S R D D A L E T D E L V L H V D I E H
AACGCATCCCACTGCTGGAGTCTCCGAGTCTGATTCGTTCCAGAAATCCTCCCTCCGATCTGTGGGGGG
O R I P T S G V L P V
AGCTGAGTTGGCAGTTCTGCTATAAATGTCTGAAGTCACTAAAGCTTTTACGGTGAACGGGTTGCCATCCAG
CTTACGGCTAAAATGGTCACTGTTGGAGAAATCCACGCCAGTAGATTTACAAATCTCTGAGGCGCTTTGAAACC
ATCTCCTAGGTTTCTTCGGAAGGACTTCGGTCCGTGACCTTAGCACAACTGCTAGTTTCAGGTAACGGGTGC
CCCCCACTTTCGTGGGGCTCCAAAAGGAG
    
```