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DNA vaccination affords significant protection against feline immunodeficiency virus infection without inducing detectable antiviral antibodies (vol 72, pg 7310, 1998)

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ERRATA

Moloney Murine Leukemia Virus Envelope Protein Subunits, gp70 and Pr15E, Form a Stable Disulfide-Linked Complex

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Volume 72, no. 8, p. 6537–6545, 1998. Page 6540, Fig. 3C should appear as shown below.

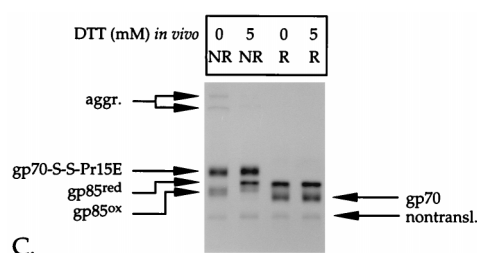


FIG. 3C.

The Second Extracellular Loop of CXCR4 Determines Its Function as a Receptor for Feline Immunodeficiency Virus

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Volume 72, no. 8, p. 6475–6481, 1998. Page 6475, column 2, line 16: “C” should read “R;” line 17, “C5” should read “R5.”

DNA Vaccination Affords Significant Protection against Feline Immunodeficiency Virus Infection without Inducing Detectable Antiviral Antibodies

MARGARET J. HOSIE, J. NORMAN FLYNN, MARK A. RIGBY, CELIA CANNON, THOMAS DUNSFORD, NANCY A. MACKAY, DAVID ARGYLE, BRIAN J. WILLETT, TAKAYUKI MIYAZAWA, DAVID E. ONIONS, OSWALD JARRETT, AND JAMES C. NEIL

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Volume 72, no. 9, p. 7310–7319, 1998. Page 7316, Table 1, line 1 of data: Trial 1, FIVΔRT, 12-week postchallenge response, “–, 0, blank” should read “+, –, 0.”

Functional Interaction of Human Immunodeficiency Virus Type 1 Vpu and Gag with a Novel Member of the Tetratricopeptide Repeat Protein Family

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Volume 72, no. 6, p. 5189–5197, 1998. Page 5192: We found that nucleotide 932 (G) was omitted from the originally published *ubp* sequence. This resulted in changes in all the encoded amino acids following this position. The *ubp* cDNA is 2,222 bp long, with a 942-bp open reading frame and a 1,242-bp 3' untranslated region. The *ubp* open reading frame is predicted to encode a 313-residue protein with a molecular mass of 34.1 kDa, which is consistent with the mobility of UBP on polyacrylamide gels. These changes do not alter the conclusions made in the original publication. Figure 1 should appear as shown below.

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1   TCGGTCGCCTGAGAGGTATCACCTCTTCTGGGCTCAAGATGGACAACAAGAAGCGCCTGGCCTACGCCATCATCCAGTTCCTGCATGACCAGCTCCGGCA
      M D N K K R L A Y A I I Q F L H D Q L R H>
101  CGGGGGCCTCTCGTCCGATGCTCAGGAGAGCTTGAAGTGCCTCCAGTGCCTGGAGACTGCGTTTGGGGTGACCGTAGAAGACAGTACCTTGGCGTC
      G G L S S D A O E S L E V A I O C L E T A F G V T V E D S D L A L >
201  CCTCAGACTCTGCCGGAGATATTTGAAGCGGCTGCCACGGCAAGGAGATGCCCGCAGGACCTGAGGAGCCCAGCGGAACCCCGCTTCCGAGGAGGACT
      P Q T L P E I F E A A A T G K E M P Q D L R S P A R T P P S E E D S >
301  CAGCAGAGGCAGAGCGCTCAAAAACCGAAGAAACGAGCAGATGAAAGTGGAAAACCTTTGAAGCTGCCGTGCATTTCTACGGAAGCCATCGAGCTCAA
      A E A E R L K T E G N E O M K V E N F E A A V H F Y G K A I E L N >
401  CCCAGCCAACCGCTCTATTTCTGCAACAGAGCCGACGCTACAGCAAACCTCGGCAACTACGCAGGCGGGTGACGACTGTGAGCGGGCCATCTGCATT
      P A N A V Y F C N R A A A Y S K L G N Y A G A V O D C E R A I C I >
501  GACCCGGCTTACAGCAAGGCTACGGCAGGATGGGCTGGCGCTCTCCAGCCTCAACAAGCACGTGGAGGCGGTGGCTTACTACAAGAAGCGCTGGAGC
      D P A Y S K A Y G R M G L A L S S L N K H V E A V A Y Y K K A L E L >
601  TGGACCCGACAACGAGACATACAAGTCAACCTCAAGATAGCGGAGCTGAAGCTGCGGGAGGCCCCAGCCCCAGGGAGGCGTGGGCGAGCTTCGACAT
      D P D N E T Y K S N L K I A E L K L R E A P S P T G G V G S F D I >
701  CGCCGGCTGCTGAACAACCTTGGCTTCATGAGCATGGCTTCGAACCTAATGAACAATCCCCAGATTACAGCAGTTCGCGCATGATTCGGGCATGATTCGGGTGGC
      A G L L N N P G F M S M A S N L M N N P Q I Q Q L C M S G M I S G G >
801  AACAAACCCCTGGGAACCTCCGGCACCAGCCCTCGCAGAACGACCTGGCCAGCCTCATCCAGGCGGGCCAGCAGTTTCCAGCAGATGCAGCAGCAGA
      N N P L G T P G T S P S Q N D L A S L I Q A G Q Q F A Q Q M Q Q Q N >
901  ACCCAGAGTTGATAGAGCAGCTCAGGAGCCAGATCCGGAGTCGGACGCCAGCCAGCCAGCAACGACGACCAGCAGGAGTACGCTGCCTGCCCTCCCGTGTG
      P E L I E Q L R S Q I R S R T P S A S N D D Q Q E *
1001 ACCGCGTCTTCCCTGGCCGACCCGAAGGAGCCTTCTGGTGTCTGCCACTTCCTCCTGTTGGACTGCCTGAGAGAGGGGAAGAGAGACCTCGGACC
1101 TGCATGTCAAGATGGATTTCCCTTTTATCTCTGCCCTCCTCCACTCCCTTTTGTAACTCCCTTACAGCCCCAGACCCCTTCTGAAACGAGAGCCAG
1201 CAAGCTGAGCACAGACCAGCAGCGACCTCCCTTCCAGCCCCAGAAAGCTCGGTCACTTGAGTGTTTCTAGAACTCTGGGGTGCCTCCGGCCGCTCTC
1301 AGAGAAATGGCAGGTTTTCACGTTACAGCGGTGGCGGATCGTGTGGCTTCCAAAGCCCTTTTACAGCCCCGCCCCCATCCCGTGGTCTGTCTGCAGGAA
1401 CTCCTCCGCTCTGTGAGAAGCCTTTTCCGAGTCGACCTCCCGGCCACCCCGGCCCTGTGCCTGCTCGGAAGAGCTCACTGCCAGCTGCGGCCCTGGGCACC
1501 GCGGGCCATGTGTGTTGATGAGGAACCTTTAGTGGCAGACACCTAAGAGACGGCTGCGGTACCCACGCTCCGTGGTTCAGGAGCCGCTCTGGGT
1601 GCATAGGACAGTTTCTGTGACTTTTCTCCAGTTGGGCATGTTGACAGACATGTTCCCTCCCTCCACCTCATTTTCTGTCTCCTCGCAGCTGAGAGCC
1701 AGGGCCGACATCATGACCTTCTGTCCCGCCGCTTAGCCCGGGAACAGGGAAGGAGCTGGGCCGTTTCTGTCTGTGTCCCATCTGCTGTCTCTCTG
1801 TCCTGGATGTTTCATGGGCCCCGGGCCCCAGGGAAGCTTACCCCTCCTGTGCTGGGTGGAGGCCACGGACACCTCAGGTGCCACCCACTTGGCCCT
1901 AAAACAGCCACCAGGAAAGCAGCCGGAGAGCCGACAGCGGGCAGCCTGTCTGGGTCTCTGAGGCTGGGGGTGGCAGACGAACCCACGGCCCGTGGTC
2001 CCAGCAGCAGGTTTGTAGTCGAGCATCTTGGGCTCCCTGGCTCCTGGCGTCTGTGAGGTAGGNGCAGTACCGTGTATCGTAGGATAGCAGTAGGAAC
2101 GGGGCCACCGCGCCCTGCAGCCGTTTCATGGCGGTGAGTGTGTGCCAAGCCACCCGGGGTGCAGGGCGTGACGTGTGGGAATAAATAGGCGTTGTG
2201 ACCTCAAAAAAAAAAAAAAAAAA
    
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FIG. 1.