

happy to report that I had not and that *Science* has not lost its subtle sense of humor.

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## Elephant Appropriation

I saw the News Briefing in the 24 November issue entitled "Bonfire to save the rhino" (p. 1001) and, whilst delighted you were able to cover the news conference, I am concerned about the emphasis. Rather than chiding Senators Bob Kasten (R-WI) and Patrick Leahy (D-VT), I, in fact, congratulated them for their endeavors to get help. What I did say was that we in Kenya were doing our part, and this made it easier for me to urge others to help even further. The \$2-million appropriation for the African Elephant Program is a most worthwhile commitment for which we are all grateful.

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Director, Kenya Wildlife Service,  
Post Office Box 40241, Nairobi, Kenya

## THE RESEARCH INSTITUTE OF SCRIPPS CLINIC

### Ph.D. Program in Macromolecular and Cellular Structure and Chemistry

The Research Institute of Scripps Clinic will admit five to ten highly qualified students for the fall of 1990 to begin studies for a Ph.D. program integrating cellular biology, structure and chemistry.

Candidates for this program must have earned a bachelor's degree and have a strong background in biology and chemistry. Qualified applicants will be asked to visit the Research Institute for an interview. Financial support will be provided to all accepted students.

Individuals interested in applying to the graduate program should send a letter describing their scientific interests and goals as well as a resume of their educational background to:

Dean of Graduate Studies, MB6  
The Research Institute of Scripps Clinic  
10666 North Torrey Pines Road  
La Jolla, California 92037

This graduate program is supported in part by the Rockefeller Foundation and is authorized by the California State Department of Education.



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*Erratum:* In the Research Article "The cholinergic neuronal differentiation factor from heart cells is identical to leukemia inhibitory factor" by T. Yamamori *et al.* (15 Dec., p. 1412), figure 2 (p. 1414) was printed incorrectly. The correct figure appears below.

```
rlif ATGAAGTCTTGGCCGAGGGATTGTGCCCTACTGCTC---ATTCTGCAC 48
mlif ATGAAGTCTTGGCCGAGGGATTGTGCCCT      GCTGG
RLIF                                     LeuLeuLeu---IleLeuHis -7
MLIF MetLysValLeuAlaAlaGlyIleValPro      LeuVal
HLIF                                     Val      ---Val
```

```
          +1
rlif TGGAAACACGGGGCAGGGAGCCCTTCCCATCAGCCCTGTAAATGCCACC 99
mlif                                     T
RLIF TrpLysHisGlyAlaGlySerProLeuProIleThrProValAsnAlaThr 11
MLIF
HLIF
```

```
rlif TGGCCATACGCCACCCGTGTACGGCAACCTCATGAACCAGATCAAGAGT 150
mlif T      A C      A
RLIF CysAlaIleArgHisProCysHisGlyAsnLeuMetAsnGlnIleLysSer 28
MLIF                                     Asn
HLIF                                     Asn      ArgSer
```

```
rlif CAACTGGCTCAACTCAACGGCAGTGCCTTCTTTATTCTCTATTAC 201
mlif A G T C T C
RLIF GlnLeuAlaGlnLeuAsnGlySerAlaAsnAlaLeuPheIleSerTyrTyr 45
MLIF
HLIF Leu
```

```
rlif ACAGCTCAAGGGAACCATTTCCCAACAACGTGGATAAGCTATGTGCCCA 252
mlif A G G      A T
RLIF ThrAlaGlnGlyGluProPheProAsnAsnValAspLysLeuCysAlaPro 62
MLIF                                     Glu
HLIF Leu Gly
```

```
rlif AACATGACGGATTTCACCTTCCATGCCAATGGGACAGAGAAGACCAAG 303
mlif A C T G C
RLIF AsnMetThrAspPheProProPheHisAlaAsnGlyThrGluLysThrLys 79
MLIF Ser Gly
HLIF Val Ala
```

```
rlif TTGGTCGAGCTGTATCGGATGGTGGGTACCTGGGAGCCTCCCTGACCAAC 354
mlif G      A A C T
RLIF LeuValGluLeuTyrArgMetValAlaTyrLeuGlyAlaSerLeuThrAsn 96
MLIF Ser
HLIF Ile Val Thr Gly
```

```
rlif ATCACCTGGGATCAGAAAAACCTCAACCCCACTGCCGTGAGCTCCAGATC 405
mlif C C GGT G G
RLIF IleThrTrpAspGlnLysAsnLeuAsnProThrAlaValSerLeuGlnIle 113
MLIF Arg Val Val
HLIF Arg Ile Ser Leu HisSer
```

```
rlif AAACCTCAATGCGACTACAGACGTCATGAGGGGGCTCTTAGCAACGTGCTT 456
mlif G T T C C T
RLIF LysLeuAsnAlaThrThrAspValMetArgGlyLeuLeuSerAsnValLeu 130
MLIF Ile
HLIF Ala IleLeu
```

```
rlif TGCCGTCTGTGCAACAAGTACCATGTGGGCCATGTGGATGTGCCCTGTGTC 507
mlif G C ACC
RLIF CysArgLeuCysAsnLysTyrHisValGlyHisValAspValProCysVal 147
MLIF Arg Pro
HLIF Ser ThrTyrGly
```

```
rlif CCCGACAACCTAGCAAAGAAGCCTTCAAAGGAAGAAGTGGGCTGCCAG 558
mlif C GA A T
RLIF ProAspAsnSerSerLysGluAlaPheGlnArgLysLysLeuGlyCysGln 164
MLIF His Asp
HLIF Thr Gly AspVal Lys
```

```
rlif CTCCTGGGGACATACAAGCAAGTCATAAGTGTGTGGCCAGGCCTTCTAG 609
mlif T      GTCATAAGTGTGGTGGTCCAGGCCTTCTAG
RLIF LeuLeuGlyThrTyrLysGln      180
MLIF ValIleSerValValValGlnAlaPheTER
HLIF Lys Ile Ala LeuAla TER
```