Biochemistry. In the article "A family of proteins structurally and functionally related to the E6-AP ubiquitin–protein ligase" by Jon M. Huibregtse, Martin Scheffner, Sylvie Beaudenon, and Peter M. Howley, which appeared in number 7, March 28, 1995, of *Proc. Natl. Acad. Sci. USA* (**92,** 2563–2567), the authors wish that the following be noted. During preparation of Fig. 1*B*, the first 10–13 amino acids of five of the aligned protein sequences were inadvertently interchanged so that the first 13 amino acids of the rat p100 sequence are found in the NEDD-4 sequence, the corresponding region of NEDD-4 is found in YKL162, and so on. The error does not in any way affect the interpretation of the results or the conclusions drawn in the paper. The corrected Fig. 1*B* is shown here.

B

```
                -100
                 •
E6-AP:          YDGGYTRDSVL----IREFWEIVHSFTDEQKRLFLQFTTGTDRAPVGGLGKL----
rat p100:       FNDESGENAEKLLQFKRWFWSIVERMSMTERQDLVYFWTSSPSLPASEEGFQPM--
RSP5:           YR-GYQESDEV----IQWFWKCVSEWDNEQRARLLQFTTGTSRIPVNGFKDLQGSD
D13635:         YSGGYSADHPV----IKVFWRVVEGFTDEEKRKLLKFVTSCSRPPLLGFKELYP--
D25215:         YKGDYSATHPT----VKLFWETFHEFPLEKKKKFLLFLTGSDRIPIYGMASL----
D28476:         PDHGYTHDSRAV---KFLF-EILSSFDNEQQRLFLQFVTGSPRLPVGGFRSLNPPL
NEDD-4:         YKNGYSMNHQV----IHWFWKAVWMMDSEKRIRLLQFVTGTSRVPMNGFAELYGSN
Dros.HYD:       FNDESSEGPDKLXKFKKWFWSIVEKMNIMERQHLVYFWTGSPALPASEEGFQPL--
YKL162:         AEHGYTMDSSI----IHDFISIISAFGKHERRLFLQFLTGSPKLPIGGFKSLNPKF
ScORF:          YVGGFSDDSRA----VCWFWEIIESWDYPLQKKLLQFVTASDRIPATGISTI----
UreB1:          Y-HKYQSNSIQ----IQWFWRALRSFDQADRAKFLQFVTGTSKVPLQGFAALEGMN

                     -50
                      •
                                      ▼
E6-AP:          ---KMIIAKNGPDTERLPTSHTCFNVLLLPEYSSKEKLKERLLKAIT-YAKGFGML
rat p100:       ----PSITIRPPDDQHLPTANTCISRLYVPLYSSKQILKQKLLLAIK--TKNFGFV   42
RSP5:           GPRRFTIE-KAGEVQQLPKSHTCFNRVDLPQYVDYDSMKQKLTLAV-EETIGFGQE   48
D13635:         ---AFCIHNGGSDLERLPTASTCMNLLKLPEFYDETLLRSKLLYAI-ECAAGFELS   54
D25215:         ---QIVIQSTASGEEYLPVAHTCYNLLDLPKYSSKEILSARLTQAL-DNYEGFSLA   54
D28476:         TIVRKTFESTENPDDFLPSVMTCVNYLKLPDYSSIEIMREKLLIAAREGQQSFHLS   55
NEDD-4:         GPQSFTVE-QWGTPDKLPRAHTCFNRLDLPPYESFDELWDKLQMAI-ENTQGFDHVD   50
Dros.HYD:       --PSVTI--RPADDSHLPTANTCISRLYIPLYSSKSILRSKMLMAIK--SKNFGFV·   40
YKL162:         TVVLKHAEDGLTADEYLPSVMTCANYLKLPKYTSKDIMRSRLCQAIEEGAGAFLLS   54
ScORF:          --PFKISLLGSHDSDDLPLAHTCFNEICLWNYSSKKKLELKLLWAI-NESEGYGFR   55
UreB1:          GIQKFQIHRDDRSTDRLPSAHTCFNQLDLPAYESFEKLRHMLLLAIQECSEGFGLANK 57

                                                  % similarity to E6-AP
```

Biochemistry. In regard to the article "Rapid gene-specific repair of cisplatin lesions at the human *DUG/DHFR* locus comprising the divergent upstream gene and dihydrofolate reductase gene during early G_1 phase of the cell cycle assayed by using the exonucleolytic activity of T4 DNA polymerase" by Nicholas J. Rampino and Vilhelm Bohr, which appeared in number 23, November 8, 1994, of *Proc. Natl. Acad. Sci. USA* (**91,** 10977–10981), the authors make the following modifying statement.

"We have become aware of a discrepancy between the percentage of hybridization measured in Fig. 3 and Fig. 4 in our paper. We think the discrepancy may be explained by an age-associated decline in the potency of our cisplatin solution (the data in Fig. 3 were obtained much later than those in Fig. 4). We have not measured the cisplatin concentration by other separate techniques, and there is a possibility that the assay is less sensitive to cisplatin than what we report. The main observations and implications regarding gene specific and preferential DNA repair are not affected by this modifying statement."