Errata

In the February 2002 issue of the *Journal,* in the article "Genomic Screening of Fibroblast Growth-Factor Receptor 2 Reveals a Wide Spectrum of Mutations in Patients with Syndromic Craniosynostosis," by Kan et al. (70:472–486), the following errors occurred: (1) on the 10th line in the left hand column on page 476, "mM" should read "μM," in both cases; and (2) in table 4, the exon 15 polymorphism "1673-47G→A" should be "1673-49G→A," and "1673-10C→T" should be "1673-12C→T." The authors regret these errors.

In the July 2002 issue of the *Journal,* in the article "Chronic Infantile Neurological Cutaneous and Articular Syndrome Is Caused by Mutations in *CIAS1,* a Gene Highly Expressed in Polymorphonuclear Cells and Chondrocytes," by Feldmann et al. (71:198–203), in table 1, the amino acid change in the second row (family 2) should be Q306K. The authors regret this error.

In the October 2002 issue, the *Journal* published a letter entitled "Increased Rate of Twins among Affected Sib Pairs," by Peter Visscher (71:995–996), and a response to this letter by Hodge et al. (71:996). The Hodge et al. response discussed, among other things, an error in a formula in Dr. Visscher's letter. This error was corrected in Dr. Visscher's letter before publication, but, because of an oversight by the *Journal,* Hodge et al. were not given an opportunity to alter their response accordingly. The *Journal* regrets this error.

In the June 2002 issue of the *Journal,* in the article "A Comprehensive Mutation Analysis of *RP2* and *RPGR* in a North American Cohort of Families with X-Linked Retinitis Pigmentosa," by Breuer et al. (70:1545–1554), the authors have identified errors in the numbering of *RPGR-ORF15* mutations, in table 4, that were due to the use of prepublication sequence from Vervoort and colleagues (Vervoort R, Lennon A, Bird AC, Tulloch B, Axton R, Miano MG, Meindl A, Meitinger T, Ciccodicola A, Wright AF [2000] Mutational hot spot within a new RPGR exon in X-linked retinitis pigmentosa. Nat Genet 25:462–466). These errors have now been identified and corrected, and a revised version of table 4 is presented below. (We thank Dr. Dror Sharon for bringing this to our attention.) The authors regret the errors.

Table 4

Mutations in *RPGR-ORF15* in Affected Hemizygotes with XLRP

Mutation[a]	Protein Change[b]	No. of Patients Affected
Deletion:		
432delG	Glu144fsTer230	1
483-4delGA	Glu161fsTer182	9
481-4delGAGA	Arg160fsTer229	1
499-502delAGGA	Lys166fsTer229	1
503-6delGGGA	Gly168fsTer231	1
570-1delAG	Arg190fsTer248	1
631delA	Glu210fsTer229	1
652-3delAG	Glu217fsTer248	7
673-4delAG	Glu224fsTer248	2
689-92delAGAG	Glu230fsTer233	3
1184-5delGG	Glu395fsTer493	1
1244-5delGG	Glu415fsTer493	1
1339-40delAG	Glu446fsTer493	1
Insertion:		
185ins, 5bp	Ala61fsTer111	1
352-460dup		1
Nonsense:		
369G→T	Glu123Ter	1
393G→T	Glu131Ter	1
423G→A	Glu141Ter	3
507G→T	Glu169Ter	3

[a] Nucleotide positions and nomenclature are based on the work of Vervoort et al. (2000).

[b] Frameshift mutations are designated according to the following example: Glu144fsTer230 refers to a frameshift mutation in which Glu144 is the first amino acid altered, with termination of the ORF at residue 230.