Errata

In the August 1997 issue of the *Journal*, in the article "Hereditary Nonpolyposis Colorectal Cancer Families Not Complying with the Amsterdam Criteria Show Extremely Low Frequency of Mismatch-Repair-Gene Mutations," by Wijnen et al. (61:329–335), an error appeared in Table 2. The corrected table follows:

Table 2

MMR-Gene Alterations in HNPCC Families

Gene and Family	Codon(s)	Nucleotide Change[a,b]	Nature of the Mutation[a]
hMSH2:			
NLB-376	Intron 1	aagGAG→aggGAG	a→g substitution at splice acceptor site
NLB-600	76 (exon 2)	CAGAGT→CAGT	AG deletion, frameshift; termination at codon 80
NL-10[c]	288 (exon 5)	CAG→TAG	C→T substitution, Gln→STOP
NL-39	288 (exon 5)	CAG→TAG	C→T substitution, Gln→STOP
NL-38	305 (exon 5)	GCA→ACA	G→A substitution, Ala→Thr
N-HS3	Intron 5	AGgta→AGgtt	a→t substitution at splice donor site
NL-21[c]	339–340 (exon 6)	CAAAGA→CAGA	AA deletion, frameshift; termination at codon 343
NL-7[c]	380–381 (exon 7)	GATTTA→GATTA	T deletion, frameshift; termination at codon 387
NL-23[c]	429 (exon 8)	CAG→TAG	C→T substitution, Gln→STOP
NL-220	429 (exon 8)	CAG→TAG	C→T substitution, Gln→STOP
I-219[c]	481–482 (exon 9)	TTAAG→TTAAAG	A insertion, frameshift; termination at codon 487
NLB-172	Intron 9	aagGC→aggGC	a→g substitution at splice acceptor site
NL-13[c]	532 (exon 10)	AAAGTC→AAAGGTC	G insertion, frameshift; termination at codon 535
NL-221	569 (exon 11)	AACAGAAT→AACAAT	GA deletion, frameshift; termination at codon 570
N-534	596 (exon 12)	CTCAATGAT→CTCGAT	AAT (Asn) in-frame deletion
N-554	596 (exon 12)	CTCAATGAT→CTCGAT	AAT (Asn) in-frame deletion
N-414	680 (exon 13)	CGA→TGA	C→T substitution, Arg→STOP
NL-203[c]	782–783 (exon 14)	ACCCAT→ACCAT	C deletion, frameshift; termination at codon 811
NL-57	834 (exon 15)	GCT→ACT	g→a substitution, Ala→Thr
hMLH1:			
NL-205	6–11 (exon 1)	Deletion of 17 nucleotides	Out-of-frame deletion, termination at codon 29
NLB-1069	6–11 (exon 1)	Deletion of 17 nucleotides	Out-of-frame deletion, termination at codon 29
NLB-100	34 (exon 1)	GAGATG→GATG	GA deletion, frameshift; termination at codon 36
N-498	62 (exon 2)	CAA→AAA	C→A substitution, Gln→Lys
N-2104	64 (exon 2)	AAT→AGT	A→G substitution, Asn→Ser
NL-20[c]	226 (exon 8)	TCGgta→TCAgta	G→A substitution at splice donor site
NL-27[c]	226 (exon 8)	TCGgta→TCGta	G or g deletion at splice donor site
NLB-296	Intron 8	TCGgtatg→TCGgtattg	t insertion at splice donor site
NL-6[c]	Intron 9	tagATC→tacATC	g→c substitution at splice acceptor site
NL-30	269 (exon 10)	TCA→TGA	C→G substitution, Ser→STOP
NL-37	295 (exon 10)	CAGgt→CGGgt	A→G substitution at splice donor site
NL-24[c]	307–308 (exon 11)	TGCACC→TGCGCACC	CG insertion, frameshift; termination at codon 367
CH-1	496–497 (exon 13)	ACCCCCCGG→ACCCCCCCGG	C insertion, frameshift; termination at codon 502
NL-25[c]	Intron 15	TCGgta→TCGata	g→a substitution at splice donor site
D-105[c]	Intron 15	tagGAG→ttgGAG	a→t substitution at splice acceptor site
NL-28[c]	593–594 (exon 16)	CAGAGAGTG→CAGAGTG	AG deletion, frameshift; termination at codon 608
NL-29[c]	616–618 (exon 16)	AAGAAGAAG→AAGAAG	AAG (Lys) in-frame deletion
I-202[c]	616–618 (exon 16)	AAGAAGAAG→AAGAAG	AAG (Lys) in-frame deletion
NL-40	616–618 (exon 16)	AAGAAGAAG→AAGAAG	AAG (Lys) in-frame deletion
NL-59	616–618 (exon 16)	AAGAAGAAG→AAGAAG	AAG (Lys) in-frame deletion
NLB-526	618 (exon 16)	AAGAAGAAG→AAGAAGGCG	AA→GC substitution, Lys→Ala
NL-4[c]	632 (exon 16)	GAGgtg→GAAgtg	G→A substitution at splice donor site
NL-204[c]	632 (exon 16)	GAGgtg→GAAgtg	G→A substitution at splice donor site
NL-56	659 (exon 17)	CGA→CCA	G→C substitution, Arg→Pro
NL-9[c]	Intron 18	AGgta→AGata	g→a substitution at splice donor site
NLB-35	Intron 18	AGgta→AGata	g→a substitution at splice donor site

[a] Uppercase letters represent exonic nucleotides, and lowercase letters represent intronic nucleotides.
[b] Nucleotides involved in the substitution/deletion events are underlined.
[c] Previously described in the studies by Wijnen et al. (1995, 1996).