A Follow-Up Study of a Genome-wide Association Scan Identifies a Susceptibility Locus for Venous Thrombosis on Chromosome 6p24.1

Pierre-Emmanuel Morange,* Irene Bezemer, Noémie Saut, Lance Bare, Gwenaelle Burgos, Jessy Brocheton, Hervé Durand, Christine Biron-Andreani, Jean-Francois Schved, Gilles Pernod, Pilar Galan, Ludovic Drouet, Diana Zelenika, Marine Germain, Viviane Nicaud, Simon Heath, Ewa Ninio, Aurélien Delluc, Thomas Münzel, Tanja Zeller, Stefan-Martin Brand-Herrmann, Marie-Christine Alessi, Laurence Tiret, Mark Lathrop, François Cambien, Stefan Blankenberg, Joseph Emmerich, David-Alexandre Trégouët, and Frits R. Rosendaal

(The American Journal of Human Genetics *86*, 592–595; April 9, 2010)

In the original online publication of the paper, in the Summary, rs169713C was incorrectly written as rs167913C. This has been corrected in both the online and print versions of the paper, and the authors regret this error.

*Correspondence: pierre.morange@ap-hm.fr
DOI 10.1016/j.ajhg.2010.03.013.

A Single-Nucleotide Deletion in the *POMP* 5′ UTR Causes a Transcriptional Switch and Altered Epidermal Proteasome Distribution in KLICK Genodermatosis

Johanna Dahlqvist, Joakim Klar, Neha Tiwari, Jens Schuster, Hans Törmä, Jitendra Badhai, Ramon Pujol, Maurice A.M. van Steensel, Tjinta Brinkhuizen, Lieke Gijezen, Antonio Chaves, Gianluca Tadini, Anders Vahlquist, and Niklas Dahl*

(The American Journal of Human Genetics *86*, 596–603; April 9, 2010)

In the original online publication of this paper, author Tjinta Brinkhuizen's name was incorrectly spelled as Tjinta Brinkhuijzen. This has been corrected in both the online and print versions of the paper. The authors regret this error.

*Correspondence: niklas.dahl@genpat.uu.se
DOI 10.1016/j.ajhg.2010.03.014.

Ribosomal Protein Genes *RPS10* and *RPS26* Are Commonly Mutated in Diamond-Blackfan Anemia

Leana Doherty, Mee Rie Sheen, Adrianna Vlachos, Valerie Choesmel, Marie-Françoise O'Donohue, Catherine Clinton, Hal E. Schneider, Colin A. Sieff, Peter E. Newburger, Sarah E. Ball, Edyta Niewiadomska, Michal Matysiak, Bertil Glader, Robert J. Arceci, Jason E. Farrar, Eva Atsidaftos, Jeffrey M. Lipton, Pierre-Emmanuel Gleizes, and Hanna T. Gazda*

(The American Journal of Human Genetics *86*, 222–228; February 12, 2010)

An incorrect version of Table 3 was included in the original manuscript and contains some errors in the data for P6, P16, and P17. A correct version of the table is included below.

In addition, there is a typo on page 224, line four from the bottom. The text currently reads RPL26. This should be RPS26.

The authors regret these errors.

Table 3. Sequence Changes in *RPS26* in DBA Patients

Mutation Type	Proband's ID (Gender) Inheritance	Family Members	DNA Mutation	Exon/ Intron	Predicted Amino Acid Change	Age at Diagnosis	Malformation Status	Response at First Steroid Therapy	Present therapy
Missense mutation	*P6* (M) *de novo*	*f, m, s*: normal sequence	c. 1A>T	Ex1	Met1Leu	NA	cleft lip and palate	responsive	steroid therapy
	P7 (M) familial	*f, m*: normal sequence	c. 1A>G	Ex1	Met1Val	2 months	none	responsive	steroid therapy
		d of P7	c. 1A>G	Ex1	Met1Val	6 weeks	none	unresponsive	RBC trx
	P8 (M) *de novo*	*f, m*: normal sequence	c. 1A>G	Ex1	Met1Val	NA	none	unresponsive	RBC trx
	P9 (M) familial		c. 1A>G	Ex1	Met1Val	NA	none	responsive	steroid therapy
		d of P9	c. 1A>G	Ex1	Met1Val	NA	none	responsive	no therapy
	P10 (M) sporadic		c. 1A>G	Ex1	Met1Val	7 weeks	duplicated pelvocalicon on right kidney	unresponsive	RBC trx
	P11 (M) familial	*d*: normal sequence	c. 1T>G	Ex1	Met1Arg	NA	NA	NA	NA
	P12 (M) de novo	*f, m, s*: normal sequence	c. 97G>A	Ex2	Asp33Asn	8 months	inguinal hernia, missing vas deferens (unilateral), slightly abnormal epidymis, pronounced boney prominence of a cervical spinous process	unresponsive	RBC trx
	P13 (M) de novo	*f, m, s:* normal sequence	c.344T>C	Ex4	Met115Thr	8 months	none	responsive	no therapy
Insertion	*P14 (F) sporadic*		c. 31_32 ins.G	Ex2	Frameshift at codon 11; stop at 25	NA	none	unresponsive	deceased
Splice-Site Mutations	*P15 (M) sporadic*	*s*: normal sequence	Donor splice site IVS1 +1g>c	Int1					
	P16 (M)	*f, m, s, b*: normal sequence	Donor splice site IVS1 +1g>a	Int1		NA	none	unresponsive	deceased
	P17 (F) de novo	*f, m*: normal sequence	Donor splice site IVS1 + 1g>t	Int1		NA	NA	NA	NA

Abbreviations are as follows: P, proband; *f*, father; *m*, mother; *d*, daughter; *s*, sister; *b*, brother ins, insertion; Ex, exon; In, intron; mo, month; y, year; NA, not available; RBC trx; red blood cell transfusion.

*Correspondence: hanna.gazda@childrens.harvard.edu
DOI 10.1016/j.ajhg.2010.03.008.