Homozygosity mapping of the gene for Hallervoden-Spatz syndrome to chromosome 20p12.3-p13

Todd Taylor, Michael Litt, Patricia Kramer, Massimo Pandolfo, Lucia Angelini, Nardo Nardocci, M.Pineda, Suzanne Davies, Haruo Hattori, Peter J. Flett, M. Roberta Cilio, Enrico Bertini & Susan J. Hayflick

Nature Genetics **14**, 479–481 (1996).

In Fig. 1 the Hs9 pedigree was inadvertently included. The other pedigrees, the figure legened and the text are correct. Family HS9 provides preliminary evidence for locus heterogeneity in Hallervorden-Spatz syndrome. These completed data will be presented in a forthcoming manuscript.

Yeast *SAS* silencing genes and human genes associated with AML and HIV-1 Tat interactions are homologous with acetyltransferases

Cherly Reifnyder, Joanna Lowell, Astrid Clarke & Lorraine Pillus

Nature Genetics **14**, 42–49 (1996).

Owing to a printing error, the lightly shaded residues in Figure 3 appeared a little *too* light. The correct version of the figure is shown below.