

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 legend and the text are correct. Family H59 provides preliminary evidence for locus heterogeneity in Hallervorden-Spatz syndrome. These completed data will be presented in a forthcoming manuscript.

erratum

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.

SAS2	LACILIFPPYQRRGLGLLLIEFSY
SAS3	LSCILTLPIYQRKGYGQFLMEFSY
Spsas+	VSCILTLPIYQRKGYGVFLIDFSY
TIP60	VACILTLPPYQRRGYRKLIEFSY
MOZ	VSCIMILPQYQRKGYGRFLIDFSY
HAT1	ISQFLIFPPYQNKGHGSCLYEAI
rimI	LFNIAVDPDYQRQGLGRALLEELI
HEAH1	LFNIAILPTYQCGFGKLLGKLI
TtHATA1	VAF LAVTANEQVRGYGTRLMNFK
GCN5	IVFCAISSTEQVRGYGAHLMNHLK
hGCN5	IVFCAVTSNEQVKGYGTHLMNHLK
ARD1	ITSLSVMRTYRRMGIAENLMRQAL
MAK3	IGMLAVESTYRGHGIAKKLVEIAI
NAT2	EEKIYLNRRGKOLIGMGEPEDESKVI