nation codon, resulting in a predicted NPM-ALK protein of 525 amino acids. The correct *NPM-ALK* nucleotide sequence, now entered in GenBank, encodes a protein of 679 amino acids with a predicted molecular weight of 75.3 kilodaltons. A polypeptide with this relative mobility, generated by in vitro transcription/translation of the full-length *NPM-ALK* complementary DNA, co-migrates with the NPM-ALK protein immunoprecipitated from the t(2;5)-containing lymphoma cell lines SU-DHL-1, SUP-M2, and UCONN-L2.

To our knowledge, all other data reported in our paper are correct. We regret any inconvenience or confusion that this sequencing error may have caused our scientific colleagues.

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### References

1. S. W. Morris *et al.*, *Science* **263**, 1281 (1994).

### Corrections and Clarifications

In the article "Molecular basis of mammalian sexual determination: Activation of Müllerian inhibiting substance gene expression by SRY" by C. M. Haqq *et al.* (2 Dec., p. 1494), parts B and C of figure 1 were inadvertently interchanged. In figure 3A, the third, fourth, and fifth lanes should have been labeled "$A_7 \rightarrow$ T," "$A_7 \rightarrow$ C," and "$A_7 \rightarrow$ G," respectively. In figure 5, the factor labeled "SRIF's" should have been labeled "SRYIF's."