Erratum

Fluorescence-based DNA sequencing with hexamer primers

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The Publishers wish to apologize for the omission of the figure which accompanied this paper. The figure and explanatory text appear below

The figure shows the raw data obtained in an A sequence reaction performed according to the described procedure. The fragment peaks are resolved out to about 400 bases. The intensity and readable sequence length obtained from the other three sequencing reactions are similar (not shown). The signal intensity obtained with the hexamers was comparable to that obtained from a sequencing reaction obtained with a 17mer primer and labeled with F-dATP as described in (11). This is a signal level comparable to that obtained in standard fluorescence-based sequencing protocols.