

What then is the function of nuclear speckles? They could be sites for storage of splicing factors, or more simply sites where excess splicing factors accumulate, as indicated in a study of Drosophila polytene nuclei<sup>28</sup>. It has been shown in yeast that splicing factors are present in large functional excess<sup>2</sup>

The approach described here can be extended to other genes in order to provide high spatial and temporal resolution of the transcription and processing of its pre-mRNA. The precise targeting of fluorochrome-labelled oligonucleotide probes to different regions of a large transcript, and the exact superimposition of these images by digital imaging microscopy, will allow a correlation of biochemical and structural events in the early life of an mRNA. 

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FIG. 4 Localizing sites of endogenous  $\beta$ -actin pre-mRNA splicing. A, Spatial correlation of unspliced and spliced  $\beta$ -actin RNAs. In situ hybridization signals detected using intron (a) and splice-junction (b) probes. (c), Overlay. B, Distribution of  $\beta$ -actin RNA relative to intranuclear speckles. Distributions of actively transcribed  $\beta$ -actin RNA (actin) relative to intranuclear speckles (SC35) in five representative cells. Nuclear  $\beta$ -actin RNA signals are indicated by arrows.

METHODS. Oligonucleotide probes contained 24 nucleotides complementary to actin sequence and 18 nucleotides of nonspecific sequence added to increase the fluorochromes. Intron-specific probes: IN1, 5'-GTCCCTGTGCAGAGAAAGCGCCCT-3'; IN2, 5'-CACGGCTAAGTGTGCT GGGGTCTT-3'; IN3, 5'-ATGAGGGCAGGACTTAGCTTCCAC-3'; 5'-CTGACCTGCCCAGGTCAGCTCAGG-3' complementary to the regions of 1,088-1,111, 1,744-1,768, 2,354-2,377 and 2,587-2,610. Splicejunction probes: SJ1, 5'-CACCATCACGCC/CTGGTGCCTGGG-3'; SJ2, 5'-CTCAAACATGAT/CTGGGTCATCTT-3'; SJ3, 5'-GGACTCCATGCC/CAGG-AAGGAAGG-3'; and SJ4, 5'-AGGAGCAATGAT/CTTGATCTTCAT-3' complementary to 1,028-1,039/1,074-1,085, 1,401-1,412/1,854-1,865, 2,281-2,292/2,388-2,399 and 2,559-2,570/2,683-2,694. Nonspecific sequences were: 5'-TTGCTTGCTTGCTTGCTT-3'. Morphometric analysis of nuclei was done on photographic images of optical sections from ten representative cells using a combined Image-Lab and Image-Measure program (Micro-Science, Inc.). The speckles occupied  $31 \pm 7\%$ of the nucleoplasmic area (excluding nucleoli). Of ten  $\beta$ -actin RNA signals, three coincided with nuclear speckes and one was in contact.

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## **ERRATUM**

# Signal transduction and regulation in smooth muscle

#### Andrew P. Somlyo & Avril V. Somlyo

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