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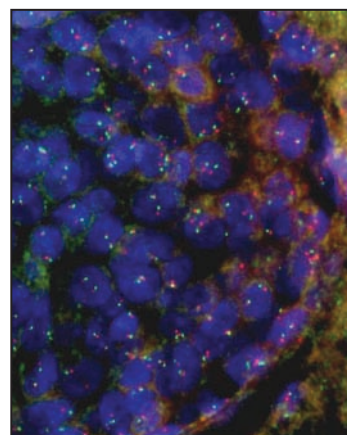
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Three potential driver genes of the 20q13 amplification were studied: *MYBL2*, *ZNF217*, and *STK6*, using fluorescence *in situ* hybridization (FISH) of locus-specific BAC probes on sections of 466 breast tumors arrayed in tissue microarrays. The authors observed two types of amplification. In the first type, the *ZNF217* locus was amplified, but amplification was not detected at the two other loci. In the second type, the amplified region seemed larger and involved two or three loci. The two types showed different gene expression profiles and were associated with different histoclinical features, including lymph node status and survival. For further details, please see Ginestier *et al.* on page 4533 in this issue.



# Clinical Cancer Research

**12 (15)**

*Clin Cancer Res* 2006;12:4459-4785.

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