

CANCER DISCOVERY CONTENTS

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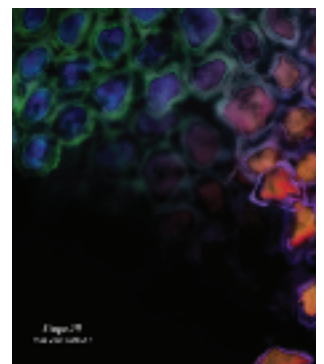
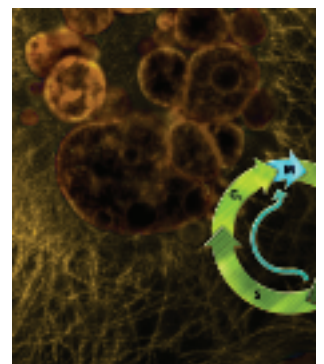
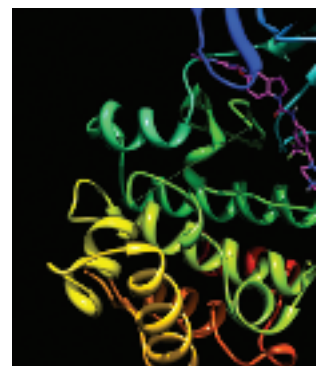
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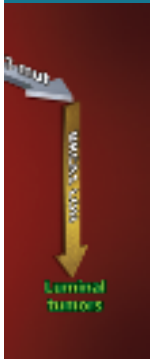
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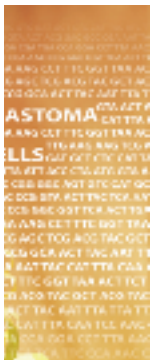
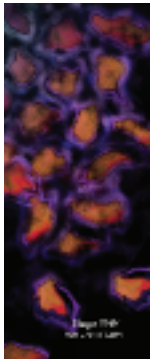
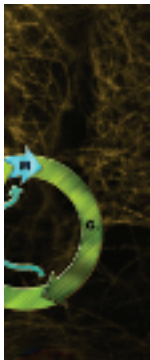
S. Jahid, J. Sun, R.A. Edwards, D. Dizon, N.C. Panarelli, J.W. Milsom, S.S. Sikandar, Z.H. Gümüş, and S.M. Lipkin

Précis: Upregulation of miR-23a in the early stages of colorectal cancer stimulates cell migration and invasion.

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Précis: Murine medulloblastoma cancer stem cells that recapitulate distinct human molecular medulloblastoma subtypes can be valuable preclinical models.

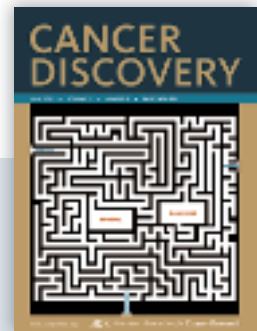


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- Nanoparticles Could Pinpoint Brain Tumors
- Modified T Cells Survive Over Decade
- Gene Expression Signature Predicts Lung Cancer Relapse
- Assay Could Identify Indolent Prostate Cancers

ON THE COVER

Martins and colleagues determined the order of *BRCA1* LOH, *PTEN* loss, and *TP53* mutation in single cells from breast tumors with germline *BRCA1* mutations. Surprisingly, *BRCA1* LOH was rarely the initiating event, and wild-type *BRCA1* expression was not lost in every cell within a tumor. Instead, *PTEN* loss occurred first in the majority of cases, particularly in basal-like tumors, and *TP53* mutation was the initiating event in most luminal tumors. These findings provide insight into the evolution of *BRCA1*-mutant breast cancers and suggest that *BRCA1* loss is not a rate-limiting step in breast tumorigenesis. For details, please see the article by Martins and colleagues on page 503.



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