

## This week in techniques

| Approach   | Summary   | Licensing status   | Publication and contact information  |
|--|---|--|--|
| <b>Assays &amp; screens</b>  |   |  |  |
| Assay to evaluate gene-drug interactions and predict chemotherapeutic resistance | <p>An assay to evaluate gene-drug interactions could help guide treatment choices. A library of 87 small molecules was screened against 89 human breast epithelial cell lines that expressed or lacked a cancer-associated gene. As proof of principle, the screen identified cell lines expressing an activated form of notch 1 (NOTCH1) that were sensitive to two aurora kinase inhibitors but resistant to a dual phosphoinositide 3-kinase (PI3K) and mammalian target of rapamycin (mTOR; FRAP; RAFT1) inhibitor. Ongoing work includes working with clinical oncologists to obtain samples from patients treated with PI3K and mTOR inhibitors to determine whether the proposed mechanism of resistance is valid in patients.</p> <p><b>SciBX 4(40); doi:10.1038/scibx.2011.1125</b><br/>Published online Oct. 13, 2011</p> | <p>Patent application filed for cancer; available for licensing through Austria Wirtschaftsservice<br/><b>Contact:</b> Angela Siegling, Austria Wirtschaftsservice, Vienna, Austria<br/>e-mail: <a href="mailto:A.Siegling@awsg.at">A.Siegling@awsg.at</a></p> | <p>Muellner, M.K. <i>et al. Nat. Chem. Biol.</i>; published online Sept. 25, 2011;<br/>doi:10.1038/nchembio.695<br/><b>Contact:</b> Sebastian M.B. Nijman, Austrian Academy of Science, Vienna, Austria<br/>e-mail: <a href="mailto:snijman@cemm.oeaw.ac.at">snijman@cemm.oeaw.ac.at</a></p> |