

This week in techniques

Approach	Summary	Licensing status	Publication and contact information
Markers			
Integrated genomic analysis of ovarian carcinoma	<p>An analysis of patient ovarian carcinoma samples could aid the development of new treatments and diagnostics. In 316 high-grade ovarian adenocarcinoma samples, 96% had <i>p53</i> mutations and 22% had <i>breast cancer 1 early onset (BRCA1)</i> or <i>BRCA2</i> mutations. Recurrent mutations in 7 other genes were also identified in 2%–6% of the analyzed sample cases. Next steps include analyzing additional tumor types.</p> <p><i>SciBX</i> 4(27); doi:10.1038/scibx.2011.782 Published online July 14, 2011</p>	Work unpatented; licensing status not applicable	<p>The Cancer Genome Atlas Research Network. <i>Nature</i>; published online June 29, 2011; doi:10.1038/nature10166 Contact: P.T. Spellman, Lawrence Berkeley National Laboratory, Berkeley, Calif. e-mail: spellmap@ohsu.edu</p>