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ORIGINAL CONTRIBUTIONS

- 1** Whole-proteome analysis of mesonephric-derived cancers describes new potential biomarkers
Evan Gibbard, Dawn R. Cochrane, Jennifer Pors, Gian Luca Negri, Shane Colborne, Angela S. Cheng, Christine Chow, David Farnell, Basile Tessier-Cloutier, Jessica N. McAlpine, Gregg B. Morin, Dietmar Schmidt, Stefan Kommoss, Friedrich Kommoss, Jacqueline Keul, Blake Gilks, David G. Huntsman, and Lynn Hoang
- 12** Programmed death ligand 1 protein expression is positively correlated with the solid predominant subtype, high MIB-1 labeling index, and p53 expression and negatively correlated with epidermal growth factor receptor mutations in lung adenocarcinoma
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- 22** MYC expression is associated with older age, common morphology, increased MYC copy number, and poorer prognosis in patients with ALK+ anaplastic large cell lymphoma
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- 42** Comparison of three FDA-approved diagnostic immunohistochemistry assays of PD-L1 in triple-negative breast carcinoma
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- 51** Breast cancer tumor heterogeneity has only little impact on the estimation of the Oncotype DX® recurrence score using Magee Equations and Magee Decision Algorithm™
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NOTICE TO CONTRIBUTORS

All manuscript submissions to HUMAN PATHOLOGY should be submitted electronically to <https://www.editorialmanager.com/yhupa/default.aspx>. Authors with circumstances that prevent online submission should send inquiries to the following address: David Newcombe, Managing Editor, HumanPathology@stellarmed.com.

Cover legend: Photomicrograph of pericanalicular and intracanalicular growth pattern in a fibroadeonoma from the manuscript "Expression of Lymphoid Enhancer Binding Factor 1 (LEF-1) in Breast Fibroepithelial Lesions" by Dr. Reisenbichler and colleagues.

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