



Markus Löffler

Kontakt

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Publikationen (4)

Staiger A, Hartmann S, Möller P, Cogliatti S, Lenz G, Trümper L, Löffler M, Schmitz N, Pfreundschuh M, Rosenwald A, Ott G, Hansmann M, Lenze D, Ziepert M, Horn H, Scott D, Barth T, Bernd H, Feller A, Klapper W, Szczepanowski M, Hummel M, Stein H, German High-Grade Lymphoma Study Group. Clinical Impact of the Cell-of-Origin Classification and the MYC/ BCL2 Dual Expresser Status in Diffuse Large B-Cell Lymphoma Treated Within Prospective Clinical Trials of the German High-Grade Non-Hodgkin's Lymphoma Study Group. *J Clin Oncol* 2017; 35:2515-2526.

Salaverria I, Oschlies I, Rosolowski M, Russell R, Rymkiewicz G, Schindler D, Schlesner M, Scholtysik R, Schwaenen C, Spang R, Szczepanowski M, Trümper L, Vater I, Wessendorf S, Klapper W, Siebert R, Molecular Mechanisms in Malignant Lymphoma Network Project, Nagel I, Macleod R, Löffler M, Martin-Guerrero I, Wagener R, Kreuz M, Kohler C, Richter J, Pienkowska-Grela B, Adam P, Burkhardt B, Claviez A, Damm-Welk C, Drexler H, Hummel M, Jaffe E, Küppers R, Lefebvre C, Lisfeld J, Berlin-Frankfurt-Münster Non-Hodgkin Lymphoma Group. A recurrent 11q aberration pattern characterizes a subset of MYC-negative high-grade B-cell lymphomas resembling Burkitt lymphoma. *Blood* 2014; 123:1187-98.

Bernd H, Möller P, Cogliatti S, Pfreundschuh M, Schmitz N, Trümper L, Höller S, Löffler M, Feller A, Barth T, Müller-Hermelink H, Rosenwald A, Ziepert M, Thorns C, Klapper W, Wacker H, Hummel M, Stein H, Hansmann M, Ott G, German High Grade Non-Hodgkin's Lymphoma Study Group (DSHNHL). Loss of HLA-DR expression and immunoblastic morphology predict adverse outcome in diffuse large B-cell lymphoma - analyses of cases from two prospective randomized clinical trials. *Haematologica* 2009; 94:1569-80.

Dierlamm J, Cogliatti S, Möller P, Schwaenen C, Stein H, Löffler M, Spang R, Trümper L, Siebert R, Ott G, Haralambieva E, Murga Penas E, Bentink S, Wessendorf S, Berger H, Hummel M, Klapper W, Lenze D, Rosenwald A, Deutsche Krebshilfe Network Project "Molecular Mechanisms in Malignant Lymphomas". Gain of chromosome region 18q21 including the MALT1 gene is associated with the activated B-cell-like gene expression subtype and increased BCL2 gene dosage and protein expression in diffuse large B-cell lymphoma. *Haematologica* 2008; 93:688-96.

Projekte (0)

Keine Resultate gefunden.

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