Kantonsspital St.Gallen

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Publikationen (9)

Loeffler-Wirth H, Pott C, Schwaenen C, Trautmann H, Wessendorf S, Stein H, Szczepanowski M, Trümper L, Hummel M, Klapper W, Siebert R, Loeffler M, Binder H, Rosenwald A, Ott G, Kreuz M, Hopp L, Arakelyan A, Haake A, Cogliatti S, Feller A, Hansmann M, Lenze D, Möller P, Müller-Hermelink H, Fortenbacher E, Willscher E, German Cancer Aid consortium Molecular Mechanisms for Malignant Lymphoma. A modular transcriptome map of mature B cell lymphomas. Genome Med 2019; 11:27.

Paul U, Stein H, Hansmann M, Möller P, Szczepanowski M, Burkhardt B, Pfreundschuh M, Schmitz N, Loeffler M, Trümper L, Siebert R, Cogliatti S, Feller A, Rosenwald A, Richter J, Stuhlmann-Laiesz C, Kreuz M, Nagel I, Horn H, Staiger A, Aukema S, Hummel M, Ott G, Spang R, Klapper W. Advanced patient age at diagnosis of diffuse large B-cell lymphoma is associated with molecular characteristics including ABC-subtype and high expression of MYC. Leuk Lymphoma 2017:1-9.

Wagener R, Betts M, Russell R, Bernhart S, Hoffmann S, Rosenstiel P, Schilhabel M, Szczepanowski M, Trümper L, Klapper W, Siebert R, ICGC MMML-Seq-Project, Rohde M, Richter J, Möller P, Aukema S, Schlesner M, Haake A, Burkhardt B, Claviez A, Drexler H, Hummel M, Kreuz M, Loeffler M, Rosolowski M, López C, "Molecular Mechanisms in Malignant Lymphomas" Network Project of the Deutsche Krebshilfe. The PCBP1 gene encoding poly(rC) binding protein I is recurrently mutated in Burkitt lymphoma. Genes, chromosomes & cancer 2015; 54:555-64.

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.

Aukema S, Stein H, Trautmann H, Wessendorf S, Trümper L, Loeffler M, Spang R, Kluin P, Klapper W, Siebert R, Molecular Mechanisms in Malignant Lymphomas Network Project, Schwaenen C, Szczepanowski M, Rosenwald A, Kreuz M, Kohler C, Rosolowski M, Hasenclever D, Hummel M, Küppers R, Lenze D, Ott G, Pott C, Richter J, Cogliatti S. Biological characterization of adult MYC-translocation-positive mature B-cell lymphomas other than molecular Burkitt lymphoma. Haematologica 2013; 99:726-35.

Horn H, Cogliatti S, Pfreundschuh M, Schmitz N, Trümper L, Siebert R, Loeffler M, Rosenwald A, Ott G, Möller P, Schmelter C, Hansmann M, Ziepert M, Becher C, Barth T, Bernd H, Feller A, Klapper W, Hummel M, Stein H, German High-Grade Non-Hodgkin Lymphoma Study Group. MYC status in concert with BCL2 and BCL6 expression predicts outcome in diffuse large B-cell lymphoma. Blood 2013; 121:2253-63.

Hartmann S, Siebert R, Küppers R, Hoefler G, Pileri S, Piccaluga P, Porwit A, Kwiecinska A, Merlio J, Parrens M, Cogliatti S, Döring C, Vater I, Bug S, Kreuz M, Scholtysik R, Gesk S, Hansmann M. High resolution SNP array genomic profiling of peripheral T cell lymphomas, not otherwise specified, identifies a subgroup with chromosomal aberrations affecting the REL locus. British journal of haematology 2009

Molecular Mechanisms in Malignant Lymphomas Network Project of the Deutsche Krebshilfe, Seifert M, Schübeler D, Schwaenen C, Rosenwald A, Prosper F, Pott C, Ott G, Möller P, Macleod R, Stürzenhofecker B, Weber M, Siebert R, Hasenclever D, Barker D, Esteller M, Spang R, Stein H, Trümper L, Loeffler M, Wessendorf S, Küppers R, Korn B, Ballestar E, Lopez-Serra L, Richter J, Rosolowski M, Wickham-Garcia E, Ammerpohl O, Bentink S, Bibikova M, Kreuz M, Berger H, Agirre X, Klapper W, Hummel M, Hansmann M, Fraga M, Fan J, Drexler H, Cogliatti S, Calvanese V, Bernd H, Martín-Subero J. New insights into the biology and origin of mature aggressive B-cell lymphomas by combined epigenomic, genomic, and transcriptional profiling. Blood 2009; 113:2488-97.

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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