



**Reiner Siebert**

**Kontakt**

Reiner Siebert

## ■ 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, Wilscher 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 PCBPI 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.

---

Kantonsspital St.Gallen

Rorschacher Strasse 95

CH-9007 St.Gallen

T: +41 71 494 11 11

[support.forschung@kssg.ch](mailto:support.forschung@kssg.ch)