



**Swen Wessendorf**

**Contact**

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## Publications (5)

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.

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.

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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.

## Projects (0)

No results found.

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