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Retrospective Study

Evaluation of a five gene signature associated with stromal infiltration for diffuse large B-cell lymphoma

Nan YY *et al.* A gene signature for DLBCL

Ying-Yu Nan, Wen-Jun Zhang, De-Hong Huang, Qi-Ying Li, Yang Shi, Tao Yang, Xi-Ping Liang, Chun-Yan Xiao, Bing-Ling Guo, Ying Xiang

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[Stromal Gene Signatures in Large-B-Cell Lymphomas | NEJM](https://www.nejm.org/doi/full/10.1056/NEJMoa0802885)

<https://www.nejm.org/doi/full/10.1056/NEJMoa0802885>

A different analytic approach identified four **gene**-expression signatures that reflected distinct biologic attributes of **diffuse large-B-cell lymphoma** tumors that were **associated** with survival ...

Cited by: 1546 **Author:** G Lenz, G Wright, S S Dave, W Xiao, J Powe...

Publish Year: 2008

[A Unified 35-Gene Signature for both Subtype ...](https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2938343)

<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2938343>

LMO2, MYBL1, BCL6, LRMP, and CCND2 in our 35 **signature genes** were also reported in **Lossos's 36 genes**, which predicted survival in **diffuse large-B-cell lymphoma** [9]. LMO2, ranking second in our...

Cited by: 27 **Author:** Yu-Dong Cai, Yu-Dong Cai, Tao Huang, Kai...

Publish Year: 2010

[A Gene Signature Predictive for Outcome in Advanced ...](https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3008560)

<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3008560>

Dec 08, 2009 · Independent **evaluation** confirmed the association of a prognostic **gene** ... we used the data to characterize the top survival-associated **signature gene** ... Ma C, Lossos IS, Rosenwald A, Boldri...

Cited by: 205 **Author:** Samuel C. Mok, Tomas Bonome, Vinod Vat...

Publish Year: 2009

[A prognostic gene model of immune cell infiltration in ...](https://europepmc.org/article/MED/32844062)

<https://europepmc.org/article/MED/32844062> ▾

A prognostic **gene** model of immune cell **infiltration** in **diffuse large B-cell lymphoma**. Coronavirus: Find the latest articles and preprints Sign in or create an account. <https://orcid.org>. Europe PMC. Menu. Abou...

[A prognostic gene model of immune cell infiltration in ...](https://peerj.com/articles/9658)

<https://peerj.com/articles/9658> ▾

Aug 05, 2020 · Background Immune cells in the tumor microenvironment are an important prognostic indicator in **diffuse large B-cell lymphoma** (DLBCL). However, information on the heterogeneity and risk...

Author: Hao Zhou, Chang Zheng, De-Sheng Hua... **Publish Year:** 2020

[Tumor Microenvironment in Diffuse Large B-Cell Lymphoma](#)

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(PDF) Stromal Gene Signatures in Large-B-Cell Lymphomas

<https://www.researchgate.net/publication/23502648...>

Stromal Gene Signatures in Large-B-Cell Lymphomas. ... Survival after treatment of **diffuse large-B-cell** lymphoma is influenced by differences in immune cells, fibrosis, and angiogenesis in ...

Gene expression profiling-based risk prediction and ...

<https://www.nature.com/articles/s41408-020-00404-0>

Jan 07, 2021 · **Diffuse large B-cell** lymphoma (DLBCL) is the most common lymphoma in the United States, accounting for about one-third of adult cases of non-Hodgkin's lymphoma
1.Despite the high cure rates for ...

A prognostic gene signature for predicting survival ...

<https://europepmc.org/article/MED/33486462> ▾

Jan 13, 2021 · A prognostic **gene signature** for predicting survival outcome in **diffuse large B-cell** lymphoma. Santosh Khanal Center for Pediatric Genomic Medicine, Children's Mercy Kansas City, Kansas City, MO 64110, United States.

A prognostic gene signature for predicting survival ...

<https://www.sciencedirect.com/science/article/pii/S2210776221000016>

Introduction. **Diffuse large B-cell** lymphoma (DLBCL) is the most common type of non-Hodgkin lymphoma and can have variable response to therapy and long-term clinical outcomes [1,2].DLBCL is of **B-cell** origin and was typically treated with a regimen of cyclophosphamide, hydroxydaunorubicin, oncovin and prednisone (CHOP) but the addition of the anti-CD20 ...

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Stromal Gene Signatures in Large-B-Cell Lymphomas | NEJM

<https://www.nejm.org/doi/full/10.1056/NEJMoa0802885>

The survival model includes two gene-expression signatures, **stromal-1** and **stromal-2**, that reflect the character of the nonmalignant cells in **diffuse large-B-cell lymphoma**. The **stromal-1 signature** ...

Cited by: 1560

Author: G Lenz, G Wright, S S Dave, W Xiao, J P...

Publish Year: 2008

Gene expression profiling-based risk prediction and ...

<https://www.nature.com/articles/s41408-020-00404-0>

Jan 07, 2021 · **Diffuse large B-cell lymphoma** (DLBCL) is the most common **lymphoma** in the United States, accounting for about one-third of adult cases of non-Hodgkin's **lymphoma** 1.Despite the high cure rates for ...

Significance of stromal-1 and stromal-2 signatures and ...

<https://europepmc.org/article/MED/28607806> ▾

Introduction. **Diffuse large B-cell lymphoma** (DLBCL) accounts for 30% of non-Hodgkin's **lymphoma** (NHL) diagnosed in adults 1.In Egypt, DLBCL is the most common subtype of NHL and represents roughly 54.4% of NHL cases at the National Cancer Institute, Cairo University 2.Based on the pathology-based registry of Ain Shams, Faculty of Medicine (2001–2010), NHL cases ...

Diffuse large B-cell lymphoma - ScienceDirect

<https://www.sciencedirect.com/science/article/pii/S0031302517304713>

Jan 01, 2018 · **Diffuse large B cell lymphoma** (DLBCL) is the most common type of non-Hodgkin **lymphoma** worldwide, representing approximately 30–40% of all cases in different geographic regions. Patients most often present with a rapidly growing tumour mass in single or multiple, nodal or extranodal sites.

Cited by: 156

Author: Shaoying Li, Ken H. Young, L. Jeffrey M...

Publish Year: 2018

[PDF] A prognostic gene model of immune cell infiltration in ...

<https://peeri.com/articles/9658.pdf>