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Strong expression of FOXP1 identifies a distinct subset of diffuse large B-cell lymphoma (DLBCL) patients with poor outcome

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#	Paper	IF	Citations
189	Organisation of neoplastic haematopathology services: a UK perspective. <i>Pathology</i> , <b>2005</b> , 37, 479-92	1.6	7
188	Diagnostic pathology of lymphoproliferative disorders. <i>Pathology</i> , <b>2005</b> , 37, 434-56	1.6	8
187	Primary cutaneous large B-cell lymphomas: clinicopathologic features, classification, and prognostic factors in a large series of patients. <i>Blood</i> , <b>2005</b> , 106, 2491-7	2.2	186
186	T(3;14)(p14.1;q32) involving IGH and FOXP1 is a novel recurrent chromosomal aberration in MALT lymphoma. <i>Leukemia</i> , <b>2005</b> , 19, 652-8	10.7	309
185	FOXP1, a gene highly expressed in a subset of diffuse large B-cell lymphoma, is recurrently targeted by genomic aberrations. <i>Leukemia</i> , <b>2005</b> , 19, 1299-305	10.7	125
184	Expression of PKC-beta or cyclin D2 predicts for inferior survival in diffuse large B-cell lymphoma. <b>2005</b> , 18, 1377-84		106
183	Foxp3 interacts with nuclear factor of activated T cells and NF-kappa B to repress cytokine gene expression and effector functions of T helper cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2005</b> , 102, 5138-43	11.5	431
182	Recent advances in the molecular diagnosis of diffuse large B-cell lymphoma. <i>Expert Review of Molecular Diagnostics</i> , <b>2005</b> , 5, 397-408	3.8	1
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