

LUBAC activity is required for ABC-DLBCL pathogenesis

D. Bonasera^{1,2}, J. Saggau^{1,2}, J. Valiulis³, J. Löber⁴, M. Peifer³, C. Rheinhardt⁵, B. Chapuy⁴, A. Montinaro⁷, G. Liccardi^{2,6}, H. Walczak^{1,7}

¹ University of Cologne, Cell death, inflammation and immunity laboratory, Institute of Biochemistry I, Centre for Biochemistry, Faculty of Medicine, Cologne, North Rhine-Westphalia, Germany

² University of Cologne, Genome instability, inflammation and cell death laboratory, Institute of Biochemistry I, Centre for Biochemistry, Faculty of Medicine, Cologne, North Rhine-Westphalia, Germany

³ University of Cologne, Department of Translational Genomics, Cologne, North Rhine-Westphalia, Germany

⁴ Charité, Department of Hematology, Oncology and Tumorimmunology, Berlin, Brandenburg, Germany

⁵ Univeristy Hospital Essen, Department of Hematology and Stem Cell Transplantation, Essen, North Rhine-Westphalia, Germany

⁶ University of Cologne, Center for Molecular Medicine Cologne (CMMC), Cologne, North Rhine-Westphalia, Germany

⁷ University College London, Centre for Cell Death, Cancer and Inflammation, UCL Cancer Institute, London, United Kingdom

Activated B-cell diffuse large B-cell lymphoma (ABC-DLBCL) is a subclass of DLBCL associated with a high relapse rate after treatment, and a poor prognosis. At the molecular level, the pathogenesis of ABC-DLBCL is distinguished by mutations in the B-cell receptor signaling pathway resulting in chronic activation of NF- κ B which is known to drive tumour proliferation and migration, significantly contributing to the poor clinical outcome. The Linear Ubiquitin Chain Assembly Complex (LUBAC) consisting of HOIL-1, SHARPIN and HOIP is the only currently known E3 ligase capable of the *de novo* synthesis of linear (M1) ubiquitin chains known to act downstream of many immune receptor signaling, including the B-cell receptor, and to regulate NF- κ B signaling, cell death and inflammation. While a role for LUBAC in lymphomagenesis has been described, currently, it is unclear how LUBAC activity affects the etiology of ABC-DLBCL and importantly the molecular mechanisms that require linear ubiquitination to promote the disease. In this study we employed the MBC (*Myd88*^{c-p.L252P/WT}; *Rosa26*^{LSL.BCL2.IRES.GFP/WT}; *Cd19*^{Cre/WT}) mouse model and investigated the role of linear ubiquitin by modulation the catalytic activity of either HOIP or HOIL-1. We therefore generated two additional MBC models one in which HOIL1 was catalytically inactive (MBC HOIL^{ci}) and one in which malignant B cell were devoid of HOIP catalytical activity (MBC HOIP^{cond.ci}). Our work shows the importance of LUBAC activity in the pathogenesis of ABC-DLBCL lymphomas *in vivo* since HOIP activity depletion completely prevented lymphomagenesis while loss of HOIL-1 activity significantly affected the onset and the severity of the disease. Furthermore, our data show the efficacy of LUBAC targeted inhibition for the treatment of this disease as a first line - single agent and also a synergistic affect in combination with standard of care.

With our findings we are able to attribute an indispensable role for LUBAC activity downstream of the My-T-BCR-receptor aberrant signaling rendering linear ubiquitin a crucial regulator of malignant B-cell transformation and signaling activation and an important target to warrant therapeutic benefit.