

Are MYD88 mutations by themselves sufficient to cause WM?

Maria Luisa Guerrero,^{*1,2} Tomasz Sewastianik,^{*3,4} Meng Jiang,^{3,5} Peter S. Dennis,³ Jianjun Zhao,³ Geraldine S. Pinkus,⁶ Petr Jarolim,⁶ Wong Kwok-Kin,^{2,7,8} Amanda Kofides,¹ Maria Demos,¹ Lian Xu,¹ Guang Yang,^{1,2} Zachary R. Hunter,^{1,2} Steven P. Treon,^{1,2} Ruben D. Carrasco.^{3,6}

¹Bing Center for Waldenström's Macroglobulinemia, Dana Farber Cancer Institute, Boston, MA, USA; ²Department of Medical Oncology, Dana-Farber Cancer Institute, Harvard Medical School, Boston, MA, USA; ³Department of Oncologic Pathology, Dana-Farber Cancer Institute, Harvard Medical School, Boston, MA, USA; ⁴Department of Experimental Hematology, Institute of Hematology and Transfusion Medicine, Warsaw, Poland; ⁵Department of Surgical Oncology, The Fourth Affiliated Hospital of Harbin Medical University, Harbin, China; ⁶Department of Pathology, Brigham & Women's Hospital, Boston, MA, USA; ^{7,2} Department of Medicine, Brigham and Women's Hospital and Harvard Medical School, Boston, MA, USA

³ Perlmutter Cancer Center, New York University, Langone Medical Center, New York, New York, USA Department of Medicine, Brigham and Women's Hospital and Harvard Medical School, Boston, MA, USA; ⁸Perlmutter Cancer Center, New York University, Langone Medical Center, New York, New York, USA.

*These authors contributed equally to the work

Waldenström's Macroglobulinemia (WM) is an incurable lymphoplasmacytic lymphoma characterized by bone marrow infiltration of IgM-secreting lymphoplasmacytic cells. Activating mutations in MYD88, present in 93-97% of WM patients, trigger the NF-κB pro-survival signaling. To this day, there is no bona fide mouse model for WM. We therefore generated conditional transgenic mice overexpressing the human MYD88 mutated and wild-type (WT) protein to investigate the oncogenic role of MYD88^{L265P} in WM. We generated compound transgenic mice by mating MYD88 transgenic mice with those expressing Cre recombinase under the control of Activation-Induced Cytidine Deaminase (*AID*) gene promoter to induce MYD88 expression in germinal center (GC) and post-GC B cells.

Transgenic mice were euthanized at different ages and lymphoid organs were rigorously examined. All *AID*^{Cre/-};*MYD88*^{WT/-} and *AID*^{Cre/-} mice were phenotypically and histologically unremarkable. Compared to controls, *AID*^{Cre/-};*MYD88*^{L265P/-} mice had significantly increased serum IgM levels. Histological analysis of 19 *AID*^{Cre/-};*MYD88*^{L265P/-} mice sacrificed frequently demonstrated a nodular lymphoid infiltrate of small atypical cells variously associated with CD138+ plasma cells in both lymphoid and non-lymphoid tissues, with enhanced expression of MYD88 and NF-κB activation (p65). Serum protein electrophoresis showed diffuse polyclonal

IWWM-10 Session 5: Thursday, October 11, 2018, Carrasco

increase in γ -globulins. Rouleaux formation was documented in the peripheral blood. Notably, 3 additional old AID^{Cre/-};MYD88^{L265P/-} mice developed a more aggressive monoclonal disease, resembling human large B cell lymphomas and showing a prominent lymphoplasmacytic infiltrate and enhanced MYD88 expression. Southern blot analysis revealed clonal bands of the IgH gene only in 2/3 mice harboring aggressive lymphomas. All three mice showed increased serum CXCL13 levels. Starting from 8 weeks of age, AID^{Cre/-};MYD88^{L265P/-} mice showed high frequency of submandibular dermatitis and/or alopecia, varying from mild to severe phenotype. Those sacrificed because of severe dermatitis had enlarged cervical lymph nodes and splenomegaly. Increased serum IL-6 levels (>49 pg/ml) were measured in 8 mice, 6 of which had intermediate or severe dermatitis. By IHC, the skin lesions revealed an infiltrate of macrophages/histiocytes and T cells.

These results indicate that MYD88L265P gives rise to a low-grade mature polyclonal B-cell lymphoproliferative disorder that shares many features of human WM that with time can transform into an aggressive lymphoma.