

Supplementary Material

LAMTOR2 (p14) controls B cell differentiation by orchestrating endosomal BCR trafficking

Marcin Łyszkiewicz^{1,2,*}, Daniel Kotlarz^{2,*}, Natalia Ziętara^{1,2,*}, Gudrun Brandes³, Jana Diestelhorst², Silke Glage⁴, Elias Hobeika⁵, Michael Reth⁶, Lukas A. Huber⁷, Andreas Krueger^{1,8,#}, Christoph Klein^{2,#}

Correspondence: andreas.krueger@kgu.de (A.K.); Christoph.Klein@med.uni-muenchen.de (C.K.)

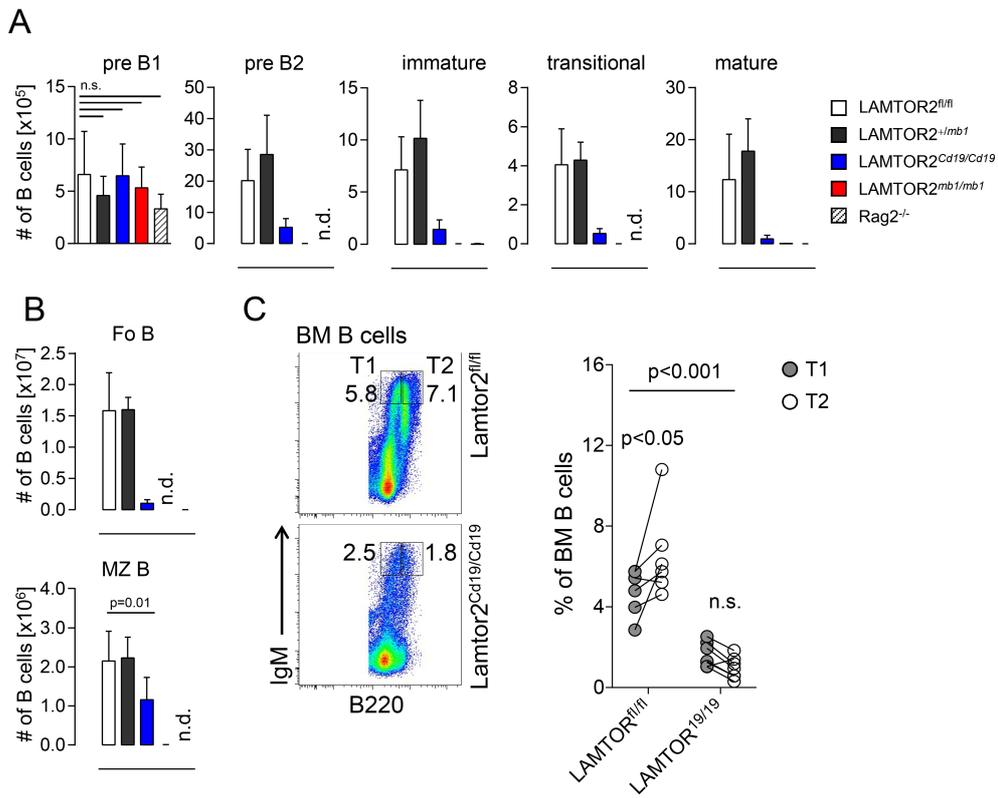


Figure S1 (related to Figure 2). Absolute numbers of B cells in spleen and BM of mice sufficient or deficient for LAMTOR2. (A) Total number of B cell progenitors in BM and (B) main B cell populations in spleen. Pooled data of one or two independent experiments, $n = 5-11$, Rag2^{-/-} $n = 3$; n.d. – not detectable. (C) Representative pseudo-colour plots (left) and quantification (right) of transitional 1 (T1) and T2 B cells in BM. Statistical significance was assessed with 2-way ANOVA (effect for genotype is shown) followed by Bonferroni post-test. Pooled data of two independent experiments is shown. Each pair of dots represents data from one mouse.

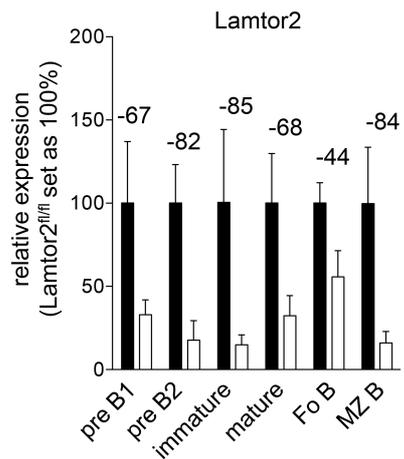


Figure S2. qRT-PCR analysis of CD19-cre mediated deletion of LAMTOR2. Purified populations of BM and splenic B cells of LAMTOR2^{fl/fl} (black bars) and LAMTOR2^{Cd19/Cd19} mice (open bars) were tested for expression of LAMTOR2 mRNA. Average expression in wild-type cells was set as 100%. Numbers above the bars indicate difference in expression between sufficient and deficient cells. Data of one experiment with n=3 for both genotypes is shown.

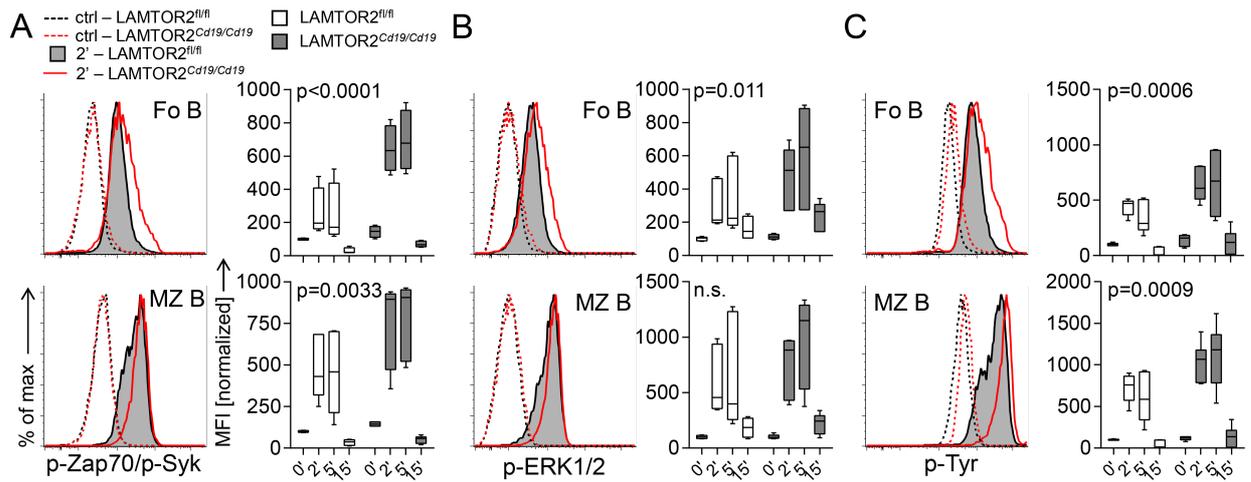


Figure S3 (related to Figure 4). Hyper phosphorylation of BCR-associated kinases in LAMTOR2-deficient B cells. Flow cytometric analysis of (A) pZap70 (pY319)/pSyk (pY352), (B) pERK1/2 (pT202/pY352), and (C) total pY in splenic B cells isolated from LAMTOR2^{fl/fl} or LAMTOR2^{Cd19/Cd19} mice. Starved splenocytes were left untreated (ctrl) or stimulated with anti-IgM F(ab')₂ and H₂O₂ for 2, 5 and 15 minutes. Representative histograms of three independent experiments are shown, graphs show summarized data of two independent experiments, n = 6 for each genotype, whiskers indicate min. to max. range of data, horizontal bars show mean value. Median fluorescence values (MFI) were normalized to ctrl (set as 100%). Statistical analysis was performed using two-way ANOVA (p-values for effect of genotype).

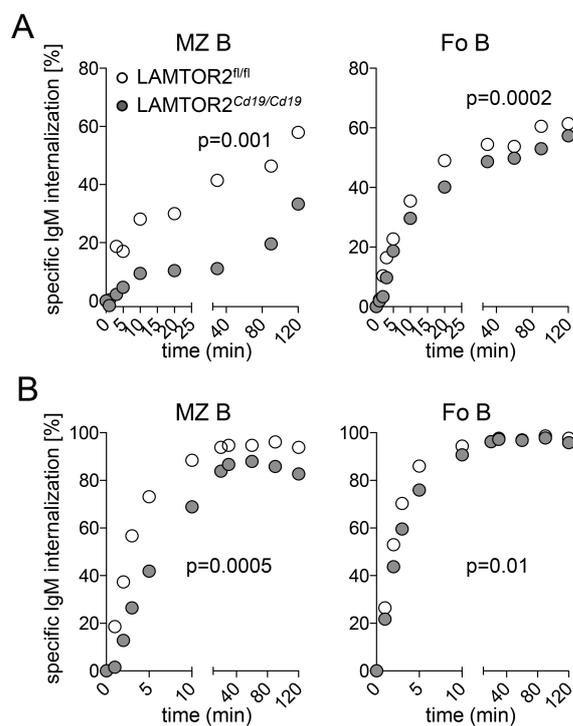


Figure S4 (related to Figure 6). Altered internalization of BCR in LAMTOR2-deficient B cells. Full-time data of the assays presented in Figure 6. (A) Passive BCR internalization in MZ B and Fo B cells or (B) ligand-induced BCR internalization in MZ B or Fo B cells. Purified splenic B cells were labelled in an ice-cold environment with either monovalent (A) or bivalent (B) biotin-labeled anti-IgM Fab or F(ab')₂ fragments, respectively, and then BCR internalization was assessed over time at 37°C. Plots for electronically gated cells are representative for two independent experiments. Charts summarize data of one representative experiment out of two.