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1	Thymic B cell-mediated attack of thymic stroma precedes Type 1
2	Diabetes Development.
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7	

8 Abstract

9 Type 1 diabetes results from a co-ordinated autoimmune attack of insulin producing beta 10 cells in the pancreas by the innate and adaptive immune systems, beta cell death being 11 predominantly T cell-mediated. In addition to T cells, peripheral B cells are important in type 12 1 diabetes progression. The thymus of mice and man also contain B cells, and lately they 13 have been linked to central tolerance of T cells. The role of thymic B cells in type 1 diabetes 14 is undefined. Here we show there are abnormalities in the thymic B cell compartment prior to 15 beta cell destruction and type 1 diabetes manifestation.

Using non-obese diabetic (NOD) mice, we document that preceding type 1 diabetes 16 17 development, there is significant accumulation of thymic B cells-partly through in situ 18 development- and the putative formation of ectopic germinal centres. In addition, in NOD mice we quantify thymic plasma cells and observe *in situ* binding of immunoglobulins to 19 20 undefined antigens on a significant proportion of medullary thymic epithelial cells. In 21 contrast, no ectopic germinal centres, or pronounced intrathymic autoantibodies are 22 detectable in animals not genetically predisposed to developing type 1 diabetes. Binding of autoantibodies to thymic stroma correlates with apoptosis of medullary thymic epithelial 23 24 cells, including insulin-expressing cells. In contrast, apoptosis of medullary thymic epithelial cells was decreased by 50% in B cell deficient NOD mice suggesting intrathymic 25 autoantibodies may selectively target certain medullary thymic epithelial cells for destruction. 26 Futhermore, we observe that these thymic B cell-associated events correlated with an 27 increased prevalence of premature thymic emigration of T cells. 28

Together our data suggests that the thymus may be a principal autoimmune target in type 1diabetes and contributes to disease progression.

31

32 Introduction

33 The thymus is a primary lymphoid organ involved in shaping the T cell repertoire. Sequential compartmentalisation of developing T cells into the cortical region of the thymus, and 34 35 subsequently the medulla enable the effective positive and negative selection events, respectively, that are integral in generating an immature T cell repertoire enriched to respond 36 to pathogens but not self-tissue- termed central tolerance [1]. Central to this role for the 37 thymus, are the medullary thymic epithelial cells (mTECS) [2; 3]; capable of autoimmune 38 regulator (AIRE) driven expression of peripheral tissue specific antigens (TSAs) [4] and 39 presentation in the context of MHC class I or class II molecules, they trigger events that lead 40 41 to apoptosis of developing T cells bearing high affinity receptors for self-peptides. 42 Although extensive studies have documented the importance of mTECs for negative selection of autoreactive T cells [5], other antigen presenting cell (APC) populations within the thymus 43 have also been shown to participate in T cell negative selection, particularly dendritic cells 44 45 [6]. A newer member of this family of APCs involved in negative selection are the thymic B 46 cells [7; 8], although it is still not clear how significant their role is in negative selection with respect to that of mTECs and thymic DCs [9]. Thymic B cells are present both in man and 47 48 mice; constituting a minor population of the thymic cellular pool, they are detectable in foetal

through to adult mammalians [10; 11]. Thymic B cells have a similar phenotype to peripheral

50 B2 cells [12; 13], and their thymic frequency is stable from birth onwards. Interestingly

51 expansion of thymic B cells in Myasthenia and Systemic Lupus Erythematosus (SLE)

52 patients [14; 15], or animal models of SLE have been linked to disease progression,

53 suggesting thymic B cells may have a potential role in breakdown of central tolerance [16].

54 Type 1 Diabetes (T1D) is an autoimmune condition where insulin secreting β cells in the

55 islets of Langerhans are destroyed through co-ordinated attack by both the innate and

56 adaptive immune systems; the final assault being perpetuated by CD8⁺ cytotoxic T cells [17; 18: 19]. Defects in central tolerance is linked to emergence of a β cell-specific T cell 57 repertoire [20], yet definitive understanding of the mechanisms underlying defective central 58 59 tolerance are unclear. Much of our understanding of the immunological events leading to β cell pathology has been derived from the non-obese diabetic (NOD) mouse, a murine model 60 that spontaneously develops T1D with many similarities to those seen in man [21]. Studies in 61 62 NOD mice show T1D is a progressive condition, with priming of the T cell repertoire to β cell antigens in early life followed by infiltration of islets with immune cells (termed 63 64 insulitis), a period of regulation of the autoreactive response, but ultimately an aggressive and sustained attack on the β cells. It is not clear what immunological event triggers this final 65 stage of the disease. 66

67 B cells, too, are known to be important in the T1D process both in man and in NOD mice; 68 abnormally high numbers of islet-infiltrating B cells is linked to rapid progression to T1D in young children [22], and increasing diversity of serum antibodies for β cell antigens increase 69 70 substantially the risk factor of developing T1D genetically-predisposed children [23]. In NOD mice, genetic or immunological ablation of B cells protects against T1D development 71 [24; 25], and in both diabetic NOD mice and diabetic patients, depletion of B cells can 72 73 resolve the condition albeit transiently [26; 27]. To date, the role for B cells in T1D progression has been linked to their peripheral APC function- their ability to present β cell 74 antigens to β -reactive CD4⁺ T cells [28] enhances CD4⁺ T helper cell activation of CD8⁺ T 75 cells, and in islets B cells provide survival signals for activated CD8⁺ T cells enabling a 76 sustained cytotoxic attack on β cells [29]. 77

Here we provide that the thymus of diabetes-prone NOD mice displays evidence of
autoreactivity prior to type 1 diabetes development. We show that the post-insulitic/prediabetic phase is characterised by abnormally high thymic B cell development, B cell

accumulation in follicles at the cortical-medullary junction and the emergence of ectopic
germinal centres. Intrathymic autoantibodies bind to undefined antigens on selective mTECs.
Subsequently increased mTECs apoptosis, including insulin-expressing mTECs occurs.
These events correlate with increased levels of peripheral T cells that have a RAG-GFP
phenotype akin to thymocytes that have yet to undergo negative selection, suggesting in NOD
mice thymic B cells may contribute to decreased efficacy of negative selection of
autoreactive T cells.

Our data provides new insights into thymic abnormalities that precede β cell destruction and
highlight the importance of focusing research on these unique thymic B cells as mediators of
this chronic condition.

91 Methods

92 *Mice*

C57BL/6 (B6), FVB.RAGp2-GFP reporter mice [30] and NOD.µMT^{-/-} mice [25] have been 93 described elsewhere. FVB.RAGp2-GFP reporter mice were backcrossed 20 generations to 94 either non-obese diabetic (NOD) mice (NOD.RAGp2-GFP) or NOD.µMT^{-/-} mice (NOD.µMT^{-/-} 95 ¹.RAGp2-GFP mice). All mice used in this study were maintained under specific-pathogen free 96 97 conditions with a 12 hour light-dark cycle and fed normal chow. All animal experimental procedures were carried out in accordance with the Animals and Scientific Procedures Act 98 1986 were approved by the University of York Animal Welfare and Ethics Review Board and 99 conducted under UK Home Office License approval conforming to ARRIVE guidelines 100 101 (https://www.nc3rs.org.uk/arrive-guidelines). Diabetes development was determined by assessing urine glucose levels using Diastix (Bayer, Inc). All animals used in this study were 102 103 not diabetic. In addition, only female mice were used.

104

105 Antibodies and Flow Cytometry

All antibodies, unless otherwise stated, were purchased from e-Bioscience. Single-cell 106 suspensions were incubated with antibodies against CD16/32 unconjugated (93), CD3 FITC 107 (145-2C11), CD3 BV421 (17A2; Biolegend), CD4 eFluor450 (RM4-5), CD4 PE (RM4-5), 108 CD4 BV650 (GK1.5; Biolegend), CD8a FITC (53-6.7), CD8B PE-Cy7 (H35-17.2), CD19 109 eFluor450 (6D5), CD19 PE (6D5), CD19 BV421 (1D3; Biolegend), CD21/CD35 PE (4E3), 110 CD23 PE-Cy7 (B3B4), IgM APC (II/41), IgD eFluor450 (11-26c), IgE BV650 (R35-72; BD-111 Biosciences), IgG PerCP-eFluor 710 (Polyclonal), IgA PE (11-44-2), biotinylated-insulin 112 (ibtsystems), CD45 PerCPCy5 (30-F11), CD45 BV510 (30-F11; Biolegend); PD1 (29F.1A12; 113

114 Biolegend), ICOS APC (C398.4A), CD138 BV650 (281-2; Biolegend), CD11b FITC (M1/70) , CD11c PE (N418), B220 eFluor450 (RA3-6B2), BCL-6 PerCP-eFluor 710 (BCL-DWN), 115 CXCR5 PE (SPRCL5), IL-21 PE (mhalx21) and Ki67 PE (B5; BD Biosciences). Intracellular 116 labelling of Ki67 was performed using eBioscience kit following manufacturer's guidelines 117 (catalogue number 00-5523-00). Cells were acquired using a BD LSR Fortessa X-20 (BD 118 Biosciences) and data analysed using FlowJo software® (Tree Star). Doublets were excluded 119 using forward light-scatter gating (FSC-A versus FSC-W) followed by gating on cells based 120 on FSC-SSC. Dead cells were excluded by gating on LIVE/DEAD® Fixable Dead Cell 121 122 Staining (ThermoFisher) negative cells. The gating strategies are described in the paper in the main figures and supplementary figures, explicit in the axis or described in detail in figures 123 legends. The gates were defined using fluorescence minus one and isotype controls: Rat IgG2a 124 125 eF450 (eBR2a), Rat IgG2a FITC (eBR2a), Rat IgG2a PE (eBR2a), Rat IgG2a PE-Cy7 (eBR2a), Rat IgG2a APC (eBR2a), Rat IgG2a BV421 (RTK2758, Biolegend), Rat IgG2a 126 BV650 (RTK2758, Biolegend), Rat IgG2a PerCP-eFluor 710 (eBR2a), Rat IgG2b PE (10H5), 127 Armenian Hamster IgG APC (eBio299Arm), Rat IgG1 Biotin (eBRG1) and Rat IgG1 BV650 128 (RTK2071; Biolegend). 129

130

131 Detection of thymic B cells bearing insulin-specific receptors.

132 The detection of B cells with receptors that bind insulin has been previously described [31]. Briefly single cell suspensions isolated from the thymus were incubated overnight at 4°C in 133 PBS supplemented with 1% foetal bovine serum, 1% anti-CD16/32 antibodies (eBiosciences) 134 and biotinylated insulin (0.1 μ g/10⁶ cells, ibtsystems). Bound insulin was detected with 135 fluorochrome-labelled streptavidin Alexa6470(Invitrogen) for 30 minutes at 4°C. The cells 136 were subsequently incubated with anti-CD19 PE (6D5; eBiosciences), B220 eFluor450 (RA3-137 6B2; eBiosciences), - CD4 BV650 (GK1.5; Biolegend), CD8β PE-Cv7 (H35-17.2; 138 eBiosciences), -CD45 PerCPCy5.5 (30-F11; eBiosciences) antibodies and LIVE/DEAD 139

Fixable Dead Cell Stains (Thermo Fisher Scientific) for 30 min at 4°C, following which the cells were analysed by flow cytometry. B cell gates were defined following exclusion of dead cells and T cells (dump channel). All samples were stained with insulin-biotin followed by streptavidin or with streptavidin only, frequencies of B cells insulin⁺ were calculated subtracting the background calculated in sample-matched streptavidin only control.

145

146 Soluble tissue extracts and Enzyme Linked Immunosorbent Assay (ELISA)

Cell-free supernatants from thymic and splenic tissue were prepared as described [32]. Briefly, 147 single cell suspension were centrifuged at 300g for 10 mins, 4 °C then 15 mins, 4 °C at 3000g. 148 Cell-free supernatants were collected and stored at -20°C until analysis.IL-2 and IL-21 149 cytokines were detected using mouse IL-2 ELISA Ready-SET-Go and mouse IL-21 ELISA 150 151 Ready-SET-Go ELISA kits following manufacturer guidelines (eBioscience). Isotype 152 classification of immunoglobulins in thymic cell-free supernatants or serum was achieved using a rapid ELISA Mouse mAb isotyping kit (Thermofisher) following manufacturer's 153 154 guidelines.

155

156 *Cultures*

Bone marrow derived dendritic cells (BM-DCs) were prepared from the appropriate mice by 157 standard methodology. Immature DCs were pulsed for 16 hours with whole insulin (Sigma; 5 158 µg/ml), LPS (Sigma; 10ng/ml), or pro-insulin peptide pB15-23 peptide (Thermo Fisher; p4878-159 1; 5 μ g/ml). Thymocytes were prepared from mice described in the results and 1x10⁶ 160 thymocytes were co-cultured in complete RPMI media (10% FCS, 50µmol/L β-161 mercaptoethanol, L-glutamine, 50 units/ml penicillin and streptomycin (Life-Sciences)) with 162 $3x10^4$ BM-DCs only, or $3x10^4$ BM-DCs pulsed with insulin or $3x10^4$ BM-DCs pulsed with 163 B15-23 peptide, or anti-CD3 (5µg/ml) (eBioscience) and anti-CD28 (2.5µg/ml) antibodies 164 (eBioscience). The co-cultures were incubated at 37°C, 5% CO₂ for 72 hrs, following which 165 cell proliferation was assessed by flow cytometry. The stimulation index was calculated 166

dividing the frequency of T cells in active proliferation (Ki67⁺) in cells following antigen
stimulation by the frequency of T cells in active proliferation in paired non-stimulated culture
(background).

For the detection of IL-21, single cell suspensions from the appropriate tissues were prepared placed in RPMI media (as above) supplemented with 50ng/ml PMA and 1 μ g/ml ionomycin for a total of 5 hours at 37°C, 5% CO₂. Brefeldin A (SIGMA) was added to the cultures at a concentration of 0.4mg/ml 1 hr after the initiation of the culture.

174

175 Immunofluorescence Analysis.

Thymi frozen in OCT compound were sectioned (~8 µm) on a cryostat. Sections were fixed in 176 4% paraformaldehyde or ice-cold acetone then blocked in PBS supplemented with 0.5% BSA. 177 The sections were incubated with unconjugated primary antibodies rabbit-anti mouse IgG 178 (Abcam), rabbit anti-mouse insulin (Abcam) or rabbit anti-mouse cytokeratin V (Abcam) 179 overnight at 4 °C. Detection of bound antibody was achieved with goat anti-rabbit IgG-Alexa 180 647 or goat anti-rabbit Ig-Alexa488 (Invitrogen) or goat anti-rat IgG Alexa 488 (Invitrogen). 181 Anti-B220 directly conjugated with Alexa 647 was incubated for 45 minutes at room 182 temperature. For detection of apoptosis, following incubation with the secondary antibody an 183 in situ apoptosis kit was used (Click-iTTM Plus TUNEL Assay, Alexa FluorTM 647 dye; 184 Thermofisher) according with the manufacturer instructions. Sections were counterstained with 185 DAPI (Molecular Probes) and mounted in Prolong Gold anti-fade or Prolong Diamond 186 (Invitrogen). Confocal microscopy was undertaken using Zen software on a Zeiss LSM 710 187 fitted on an Axioimager using a 63x (1.4) planApochromat or 20x (0.6) Neofluor. Binding of 188 autoreactive immunoglobulin and TUNEL in microscopy images was quantified using 189 StrataQuest V64 software. Individual nuclei were counted and the data was presented as 190

scatterplots of mean fluorescence intensity of DAPI versus mean fluorescence intensity of Igor TUNEL positive cells.

193

194 RNA Isolation and real-time RT-PCR analysis

Thymic tissues were stored at -80°C in RLT. Samples were allowed to thaw and RNA were 195 carried out using the RNeasy mini kits (Qiagen, Manchester, UK), according to the 196 197 manufacturer's instructions. On-column DNase digestion was carried out to remove any contaminating genomic DNA using the RNAse-free DNase set (Qiagen, Manchester, UK) 198 according to the manufacturer's instructions. The cDNA syntheses were performed with the 199 Superscript II reverse transcriptase system (Invitrogen), according to manufacture's 200 instructions. The qRT-PCR of aicda mRNA expression (AID gene) in total thymus was 201 202 performed with the Taqman qPCR Kit (Applied Biosystems, Warrington, UK)). mRNA expression levels were normalized to HPRT1 housekeeping gene using $\Delta\Delta$ Ct calculations. 203 Mean relative mRNA expression levels between control and experimental groups were 204 calculated using the $2^{-\Delta\Delta Ct}$ calculations. 205

206

207 Statistical analysis

Statistical analyses were performed by parametric or non-parametric tests, selected based on the distribution of the raw data. The comparisons between experimental groups were performed using student unpaired t-test, Mann-Whitney and one-way ANOVA as appropriate. The statistical analyses for fold-changes were performed using Wilcoxon signed-rank test. All analyses were conducted using GraphPad InStat (version 5) software (GraphPad).

214 **Results**

215 T1D progression correlates with increased intrathymic B cell numbers in NOD mice

216 Thymic B cells normally constitute a small population of cells within the murine and human

thymus in normal individuals. Abnormality in thymic B cell numbers has been linked to

certain autoimmune conditions [14; 15]. To determine whether thymic B cell populations

219 differ between diabetes-prone or non-prone mice, we performed time-course flow cytometric

studies of age-matched, sex-match NOD and control C57BL/6 (B6) mice.

Diabetes incidence in our female NOD mouse colony is 95%, approximately 3% of mice 221 222 develop T1D at 12-14 weeks of age, 85% at 18-20 weeks of age with the remaining 7% of females progressing to T1D by 23 weeks of age. Animals not diabetic by 23 weeks of age 223 224 rarely develop T1D. The data is based on a cohort of 200 animals (Supplementary Figure 1a). 225 This diabetes incidence, combined with the insulitis score- that is the degree of immune cell infiltration of islets and degree of β cell destruction- as mice age (Supplementary Figure 1b) 226 highlight that 12-14 weeks of age in our colony represents late insulitic-preultimate diabetic 227 stage, a critical time when immunoregulation of the autoreactive response starts to 228 229 breakdown.Thus, in our initial studies we focused on two major time points; the pre-early insulitic phase (4-6 weeks) and the post-insulitic/pre-diabetic phase (12-14 weeks) to assess 230 the presence of CD19⁺ thymic B cells. Representative flow cytometry plots for the respective 231 232 mice are shown in Figure 1a. Although absolute numbers of CD19⁺ B cells remained relatively static in the thymi of control B6 mice at the time points investigated (Figure 1b), 233 with perhaps a slight increase at 12-14 weeks of age, the absolute numbers of CD19⁺ B cells 234 increased significantly in the later age group of NOD mice in comparison to numbers seen 235 either at 4-6 week old NOD mice or 12-14 week old B6 mice. Importantly, the number of 236 237 thymic B cells at 4-6 weeks of age was comparable between NOD and control B6 mice.

This increased number of thymic B cells in 12-14 week old NOD mice was not related to increased B cell development in the bone marrow, as frequencies of CD19⁺ B cells in this primary lymphoid tissue was comparable between the two strains of mice at both time points investigated (data not shown). These data show that inappropriate accumulation of thymic B cells precedes the overt β cell destruction phase of T1D.

243

244 Intrathymic signals trigger enhanced B cell development in NOD mice.

Although previous studies have documented the ability of the thymic environment to enable 245 246 B cell development in non-autoimmune-prone mice, other reports suggest thymic B cells accumulate via periphery B cell migration to the thymus [16; 33]. To determine whether the 247 NOD mouse thymus promotes B cell development we used recombination activation gene 248 249 green fluorescent protein (RAG2p-GFP) reporter mice on a non-T1D-prone FVB background (hereafter called FVB-RAG-GFP), or on the NOD background (hereafter called NOD-RAG-250 GFP). In RAG2p-GFP reporter mice, highest GFP expression occurs when RAG genes are 251 active [30]. Once recombination of the B cell receptors and T cell receptors is complete and 252 RAG activity is silenced, GFP expression decreases over a 54 hour period [30]. As such, 253 254 newly developed B cells can be identified from thymic resident/recirculatory B cells based on the expression of GFP. 255

Since our control RAG2p-GFP transgenic mice are on a FVB background, we compared
thymic B cell frequencies and numbers of this alternative control murine strain to control B6
mice or NOD mice. Although frequencies and absolute numbers of thymic B cells in the FVB
strain were higher than the B6 strain, the NOD strain demonstrated significantly greater
thymic B cell frequencies and numbers to the FVB strain (Supplementary Figure 2a-b).

261 We performed time-course, flow cytometry studies of the two strains of mice at the ages shown in Figure 1c, and quantified the number of GFP^{hi} B cells. Representative flow 262 cytometry plots showing the gating strategy for CD19⁺GFP^{hi} B cells is shown in 263 Supplementary Figure 1c. Recently developed CD19⁺GFP^{hi} B cells were readily detectable in 264 both strains of mice at all time points analysed (Figure 1c). In control FVB-GFP mice, there 265 was no significant changes in B cell development as mice aged. In the NOD strain, although 266 there was no significant change in B cell development when the two age groups were 267 compared, it was clear that thymic B cell development is enhanced as mice enter the late 268 269 insulitic-prediabetic phase of the T1D pathway.

In light of evidence that the late insulitic-prediabetic phase is characterised by increased B 270 cell development, we asked if homeostatic proliferation of thymic B cells is also affected as 271 mice enter the late insulitic-prediabetic phase. We performed comparative flow cytometric 272 studies between NOD and control B6 mice, assessing for Ki67 expression as a marker for 273 homeostatic proliferation. Interestingly for both strains of mice, the highest level of 274 275 homeostatic proliferation of thymic B cells is an early event, with CD19⁺Ki67⁺ B cell 276 frequencies higher in younger mice when compared to older mice (Figure 1d). Further, this decrease in homeostatic proliferation in the 12-14 week old group was more pronounced in 277 278 NOD mice, although the decrease was not significant.

279

280 The NOD thymic environment has ectopic germinal centre formation potentiality.

To further investigate the phenotype of thymic B cells in NOD mice we assessed their surface markers. B cells undergo a series of transitions from the immature stage developing follicular or marginal zone properties. Thus, we qualified the phenotype of thymic B cells assessing for follicular (IgM^{lo}IgD⁺CD21/35⁺,CD23⁺) versus marginal zone (IgM⁺IgD^{lo}CD23⁻CD21/35⁺).

We focused our studies on 11-14 week old mice due to the evidence that at this age B cell development is enhanced as are thymic B cell numbers in NOD mice when compared to control mice. Representative flow cytometric plots for our gating strategies are shown in Supplementary Figure 2d.

As shown in Figure 2, the frequency of follicular B cells within the thymic B cell pool was significantly higher in NOD mice compared to control B6 mice (Figure 2a). This enhancement in follicular B cells in the NOD mouse thymus was recapitulated when absolute number of follicular B cells was calculated (Figure 2b). In contrast, although the frequency of B cells with a marginal zone phenotype were significantly decreased in the NOD mouse thymus compared to control B6 mouse thymus, the absolute numbers of these cells was similar between the two strains of mice.

296 The increased numbers of FO B cells in NOD mice with respect to B6 control mice led us to investigate whether the thymic B cell form follicle-like structures. Immunohistochemical 297 298 studies revealed B cell follicle-like structures form only in NOD mice (Figure 2c). Initially B 299 cells are detectable at the cortical-medullary junction at 9 weeks of age in NOD mice (not shown) with pronounced accumulation of B cells into follicle-like structures in this location 300 301 by 11 weeks of age. The presence and location of B cell follicle-like structures was identical irrespective of whether we used anti-B220 or anti-CD19 antibodies to identify B cells (not 302 shown) confirming the accumulating B220⁺ cells are not plasmacytoid DCs. We quantified 303 the number of B cell follicle-like structures in the thymus of 9-11 week old NOD mice; of 15 304 individual sections assessed, 90% contained 1 follicle, 5% two follicles and 5% no follicles. 305 The presence of follicle-like structures in the thymus of late insulitic-pre-diabetic NOD mice, 306

but not control B6 mice, lead us to ask if the thymic environment could support germinal
centre formation. Of interest was the relationship between IL-2 and IL-21, the latter being a

309	key mediator of germinal centre formation; the cytokine promotes B cell somatic
310	hypermutation and class switching, and the development and maintenance of T follicular
311	helper cells (TfH cells) [34]. In NOD mice IL-21 has been associated with T1D progression
312	[35; 36] and CD4 ⁺ CD45R ⁻ T cells isolated from T1D patients secrete greater quantities of IL-
313	21 than quantified from normal individuals [37]. We prepared cell-free supernatants [32]
314	from the thymi of NOD and control B6 mice at the ages shown in Figure 3a and performed
315	ELISA assays. As a comparison, we analysed cell-supernatants from the spleens of the same
316	mice. The results were tabulated as ratio of IL-21:IL-2. No differences were seen in IL-21:IL-
317	2 ratios in splenic preparations from the two strains of mice. However, the NOD mouse
318	thymus had a significant bias in IL-21 concentrations in comparison to B6 mice.
319	In light of this IL-21 bias, we quantified the frequency and absolute numbers of CD4SP cells
320	that expressed a TfH cell phenotype in the thymus of the two strains of mice. As shown in
321	Figure 3b-c, NOD mice exhibited a significant increase in frequencies and absolute numbers
322	of CD4SPPD1 ^{hi} ICOS ⁺ T cells, and these cells also expressed transcription factor Bcl-6
323	(Figure 3d) and CxCR5 (Supplementary Figure 3a). In addition, approximately 5% of NOD
324	putative thymic TfH cells secreted IL-21, a frequency that was comparable to that seen for
325	splenic TfH cells from the same mice (Supplementary Figure 3b). Furthermore, this increase
326	in thymic TfH cells in NOD mice in comparison to B6 control mice correlated with an
327	increased number of CD4 ⁻ CD8 ⁻ B220 ^{low/-} CD138 ⁺ plasma cells in the NOD mouse thymus,
328	although this increased number was not significant (Figure 3e, Supplementary Figure 3c).
329	Together, these data suggested that ectopic germinal centres could be present in the NOD
330	mouse thymus, but absent in control B6 mouse thymus. To support this hypothesis we looked
331	for a bone-fide germinal centre marker; the enzyme activation-induced cytidine deaminase
332	(AID). RNA was prepared from thymi isolated from NOD mice or control B6 mice and
333	quantitative real-time RT-PCR performed. As an additional control we included thymic

mRNA isolated from age-matched, sex-matched NOD- μ MT^{-/-} mice. The relative expression of transcripts for AID in NOD mice was normalised to control B6 mice. As shown in Figure 31, the NOD mouse thymi has enhanced AID expression in comparison to control B6 mice. Thus, ectopic germinal centre formation is likely a feature of the NOD thymus and precedes the preultimate β cell destruction phase of T1D.

339

340 Thymic immunoglubulins binding selective mTECs correlates with mTEC apoptosis.

The presence of AID and enhanced plasma cell frequencies in the NOD thymus with respect 341 342 to control B6 mice, made us query the immunoglobulin isotype of the thymic B cells and secreted antibodies. Since we previously had investigated the IgM⁺ B cell thymic subtype 343 (Figure 2), this time we focused on class-switched IgM⁻ cells. The number of IgM⁻IgD⁻IgA⁺ 344 345 and IgM⁻IgD⁻IgG⁺ B cells were similar in the thymus of both NOD and control B6 mice as determined by flow cytometry (Figure 4a). In contrast, the number of IgM⁻IgD⁻IgE⁺ B cells 346 were significantly increased in the NOD mouse thymus with respect to control mice. 347 Interesting, a unique population of IgM⁻IgD⁺ B cells (similar to those reported in T1D 348 patients [31]) was detectable in the thymic tissue. These IgM⁻IgD⁻ B cells dually expressed 349 IgG, IgE or IgA with the number of dual expressing IgD⁺IgA⁺ and IgD⁺IgG⁺ B cells being 350 significantly higher in the NOD mouse thymus than the B6 control mouse thymus, the most 351 significant being the IgD⁺IgG⁺ isotype (Figure 4b, Supplementary Figure 4a). In contrast, no 352 differences in IgD⁺IgE⁺ B cell numbers were seen between the two strains of mice. We next 353 354 assessed the isotype of soluble thymic immunoglobulin by ELISA, in comparison to serum immunoglobulin. Only the IgG1 and IgA immunoglobulin isotypes were enhanced in the 355 356 thymi of NOD mice in comparison to control B6 mice (Figures 4c,e). In contrast, IgG2a, IgG2b and IgM antibody levels were similar in both strains of mice, with IgG3 antibody 357

358 levels being slightly lower in NOD mice than in the thymus of B6 control mice. Interestingly, in NOD mice thymus, B cells predominately used the kappa light chain, there being a 359 significant decrease in the presence of lambda light chains when compared to B6 control 360 361 mice (Supplementary Figure 4c). In addition, this isotype pattern documented in the NOD mouse thymus seemed unique for this tissue, as similar ELISA-based isotyping of 362 immunoglobulins in the serum of the two strains of mice revealed little difference in levels of 363 364 each isotype assessed (Figure 4d, f). However, similar to the thymus, in the serum there was a significant decrease in lambda light chain usage in NOD mice in comparison to B6 controls 365 366 (Supplementary Figure 4b). Quantification of the thymic Ig isotypes supported the data that IgG1 and IgA are significantly greater in the thymus of NOD mice with respect to control B6 367 mice (Supplementary Figure 5a). 368

We decided to explore further these thymic B cells to determine whether they harboured 369 370 receptors specific for islet autoantigens, focusing on their specificity for insulin [31]. Representative gating strategy for identifying insulin-reactive B cells is shown in 371 372 Supplementary Figure 5b. Although the frequency of cells bearing receptors specific for 373 insulin is significantly less in the thymus of NOD mice with respect to control B6 mice within the B cell fraction, absolute numbers of insulin reactive B cells was similar in NOD 374 mice and B6 control mice (Supplementary Figure 5c). Thus, insulin-reactive B cell numbers 375 do not correlate with T1D susceptibility at this time point. Due to this finding, we decided to 376 ask whether thymic B cells produce antibodies that target, as yet, undefined antigens on 377 thymic stroma. Thymic tissue sections from 11 week old NOD mice were incubated with 378 anti-mouse antibodies that would detect any mouse immunoglobulin bound to thymic stroma 379 *in situ* and bound antibodies were detected by confocal microscopy. To qualify whether any 380 381 immunoglobulins that bound to thymic stroma interacted with mTECs, we included mTECbinding anti-cytokeratin V antibodies in the assay. As shown in Figure 5a, there was 382

383 detectable binding of murine immunoglobulins to thymic stroma, suggesting these cells had murine immunoglobulins bound to them *in situ*. Interestingly, intrathymic immunoglobulins 384 were bound almost exclusively to cytokeratin V⁺ mTECs and it appeared that only a 385 386 proportion of mTECs were being targeted by the immunoglobulins. In contrast to NOD mice, there was substantially less intrathymic immunoglobulin in control B6 mice interacted with 387 thymic stroma, particularly cytokeratin V⁺ mTECs (Figure 5b). Further, there was no 388 evidence of intrathymic immunoglobulins bound to thymic stroma, including cytokeratin V⁺ 389 mTECs in B cell deficient NOD-µMT^{-/-} mice confirming the specificity of the anti-mouse 390 391 antibodies for mouse immunoglobulins (Supplementary Figure 6). We quantified the frequency of cells with murine immunoglobulin bound in the thymus of 11-14 week old NOD 392 and B6 mice. We selected images that had comparative frequencies of cytokeratin V⁺ mTECs 393 and counted 2-3x10⁴ DAPI⁺ cells/mm². As shown in Figure 5c, approximately 7% of cells 394 had bound murine immunoglobulin (Ig) in the NOD mouse thymus. In contrast, the 395 frequency of cells bound by murine immunoglobulins in B6 mice was so low as to be 396 397 undetectable.

398 Finally we queried the significance of *in situ* binding of thymic stroma by immunoglobulins, particularly the potential that a selective number of mTECs underwent apoptosis. In this 399 regard, we incubated thymic tissue sections from 11 week old, female NOD mice with 400 antibodies specific to cytokeratin V^+ and assessed for apoptosis by confocal microscopy 401 following Tunel staining (Figure 6a). As controls, we similarly analysed thymic tissue 402 sections from control B6 mice and NOD-µMT^{-/-} mice. The inclusion of NOD-µMT^{-/-} mice 403 404 was important to determine whether the diabetes-associated MHC class II molecules unique to NOD mice was sufficient to trigger mTEC apoptosis via non-B cell-mediated mechanisms. 405 406 Thymic tissue sections from NOD mice had clear evidence of apoptosis, and such apoptotic cells were almost exclusively cytokeratin V⁺ mTECs. Apoptosis of cytokeratin V⁺ mTECs 407

was also evident in NOD- μ MT^{-/-} mice, although the proportion of apoptotic cells seemed 408 lower than that for B cell sufficient NOD mice. In contrast to the NOD strains, we could not 409 see any apoptotic cells in the B6 control mouse thymic tissue section. We quantified the 410 411 frequency of apoptotic cells in the thymic sections of the respective strains of mice (Figure 6). We counted a total of 4×10^4 DAPI⁺ cells/mm² per section, ascertaining similar frequencies 412 of cytokeratin V⁺ mTECs for each tissue sections examined. As shown in Figure 6b, ~6% of 413 DAPI⁺ cells were apoptotic in NOD mice. This frequency of apoptosis was two-fold higher 414 than seen for NOD- μ MT^{-/-} mice (~3%). In contrast to the NOD strains, <1% of cells were 415 apoptotic in control B6 mice. We were curious to determine if apoptotic cytokeratin V+ 416 mTECs in NOD mice expressed insulin. Thymic tissue sections from 11 week old female 417 NOD mice were incubated with anti-cytokeratin V⁺ and anti-insulin antibodies and apoptosis 418 419 determined by Tunel staining as before. As a control, we similarly analysed thymic tissue sections from age-matched, female B6 mice. Interestingly, within the apoptotic cytokeratin 420 V⁺ mTEC pool in NOD mice resided cytokeratin V⁺ mTECs that expressed insulin, although 421 it is important to note that some insulin⁺ cytokeratin V⁺ mTECs were not apoptotic 422 suggesting there is not a complete loss of insulin⁺ cytokeratin V⁺ mTECs but a reduction in 423 424 their numbers. Similarly, some apoptotic mTECs did not express insulin (Supplementary Figure 7). 425

426 Taken together, these data suggest that B cell-mediated autoimmune targeting of cytokeratin 427 V^+ mTECs results in the loss of a distinct population of cytokeratin V^+ mTECs, some of 428 which express insulin, and this key feature occurs before sustained autoimmune attack in the 429 pancreas.

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Thymic B cells enhance premature egress of T cells from the thymus.

The evidence that thymic stroma had bound autoantibodies and the presence of these 433 autoantibodies correlated with increased apoptosis of thymic stroma, including some insulin⁺ 434 mTECs, we investigated the impact this may have on thymocytes capable of responding to 435 islet antigen, particularly insulin. We isolated the thymocytes from NOD mice thymi and 436 cultured the cells in the presence of bone-marrow derived dendritic cells and either whole 437 insulin or proinsulin peptide15:23 [38]. The proliferative response to the CD4SP and CD8SP 438 thymocytes to the respective stimulants was assessed by flow cytometric analysis of Ki67 439 (Figure 7, Supplementary Figure 8a). As controls we included B6 mice stimulated with whole 440 insulin, and B cell-deficient NOD-µMT^{-/-} mice stimulated with whole insulin or proinsulin 441 peptide 15:23. For CD4SP cells only those isolated from B cell sufficient NOD mice 442 responded to both whole insulin, although the response was not significant in comparison to 443 444 control mice. (Figure 7a). The responses to whole insulin for thymocytes from B6 and NODµMT^{-/-} mice being close to baseline. In contrast, CD4SP thymocytes from NOD mice 445 exhibited a significantly increased response to proinsulin P15:23 with respect to NOD-µMT^{-/-} 446 mice. The responses of CD8SP thymocytes was slightly different; whereas thymocytes 447 isolated from NOD mice responded to the whole insulin molecule, the responses for 448 individual mice was quite diverse- some responded well, others' response close to baseline 449 levels for B6 control mice (Figure 7b). Similarly, CD8SP thymocytes from NOD-µMT^{-/-} mice 450 had some diversity in responsiveness to whole insulin, although it was noted that even the 451 best responders still responded weaker than that seen for NOD mice. In contrast, CD8SP 452 thymocytes from NOD mice responded far better to proinsulin P15:23, than those from NOD-453 -µMT^{-/-} mice, although the response was not significantly enhanced. In these same mice the 454 responses to the proinsulin peptide were less diverse and above baseline levels. 455

456 We initially wondered whether this increased response for insulin and proinsulin peptide by NOD thymocytes was representative of increased survival of autoreactive T cells, and thus a 457 breakdown in negative selection. In particularly, we queried whether thymocytes that had 458 459 very recently rearranged their TcR escaped from the thymus before completing negative selection. If this held true, we expected an increase in RAG-GFPhi T cells in the blood; RAG-460 GFP levels normally fall during negative selection due to the time to complete the process 461 and as such, peripheral T cells are usually RAG-GFP^{int} [30]. To test this hypothesis we 462 performed flow cytometry analysis of total GFP levels of T cells in the peripheral blood of 463 464 NOD-RAG-GFP mice in comparison to control FVB-RAG-GFP mice and B cell deficient NOD-µMT^{-/-}-RAG-GFP mice. Representative flow cytometry plots showing the gating 465 strategy is shown in Supplementary Figure 8b. As shown in Figure 7c, in the NOD murine 466 467 strains the frequency of total GFP⁺ T cells in peripheral blood was greater than seen for the 468 control FVB strain, for NOD mice this increase being significant. Furthermore, this increased frequency of GFP⁺ T cells in the peripheral blood of the NOD strains was almost entirely due 469 to GFP^{hi} cells, as GFP^{int} cells were only slightly increased in frequency in comparison to 470 control FVB-RAG-GFP mice, again NOD mice showing a significant increase. Importantly, 471 although not significant, it was clear that the frequency of RAG-GFP^{hi} cells in B cell 472 sufficient NOD-RAG-GFP mice was higher than in B cell deficient NOD-µMT^{-/-}-RAG-GFP 473 highlighting the importance of B cells in the early release of T cells from the thymus prior to 474 475 negative selection.

476 **Discussion**

Ablation of efficient purging of autoreactive T cells in the thymus and the role of B cells in
T1D seem two distinct entities in understanding how immunological tolerance is broken in
this chronic autoimmune condition. Here, we establish that inappropriate accumulation of B
cells in the NOD mouse thymus is a unique feature of the disease process, and these thymic B
cells may play a role in the egress of pre-negatively selected T cells.

T1D progression in both man and NOD mice occurs over time. The initial stages of T1D, 482 where priming of the immune response to islet antigen occurs but not overt β cell destruction, 483 is characterised by autoantibodies to β antigens [39]. It is accepted that following priming of 484 the autoreactive T cell repertoire to β cell antigens, the activity of the autoreactive T cells is 485 486 kept in check by regulatory mechanisms. Ultimately, such regulation fails, and leading to β cell destruction. Little is known as to why regulation of autoreactive T cells fails over time, 487 although paucity of, or dysfunction of, T regulatory cells is speculated to contribute to the 488 489 phenomenon [40; 41; 42]. Our data adds a new dimension to our understanding of the 490 immunological changes that occur at the late insulitic- pre-diabetic phase that may tip the autoreactive T cell response in favour of β cell destruction; targeted thymic B cell 491 autoimmune attack of thymic stroma expressing β cell antigens. 492

B cells are present in the thymus of mammals from fetal age to adulthood, their numbers
remaining relatively static in ontogeny and equating to those of thymic dendritic cells [5; 11;
13]. Previous studies in NOD mouse strains documented B cell accumulation in the thymus
of aged mice [43; 44]. Here we extended on these early studies showing that in NOD mice,
thymic B cell numbers are not static, their numbers significantly increase at the late insuliticpre-diabetic phase suggesting the restricted B cell niche normally present expands. This
change in B cell numbers occurs at the same time as increased numbers of RAG⁺ B cells are

500 detected in the thymus, but decreased homeostatic proliferation. Together these findings suggest that permissiveness of B cell development that can normally occur within the thymus 501 [33; 45; 46; 47] is enhanced in NOD mice as they age, and the increase in B cell numbers 502 503 potentially reflects this increased rate of development rather than *in situ* proliferation. Although we cannot exclusively discount that peripheral B cells migrating to the thymus 504 contribute to the thymic B cell population, we, like others, have found peripheral B cells have 505 506 little propensity to traffic to the thymus (data not shown, [47]). Future studies in how the NOD mouse thymic environment potentially nutures B cell development and retention will be 507 508 informative.

509 The phenotype of thymic B cells in NOD mice resembles that of thymic B cells in nonautoimmune strains of mice; they predominantly express B2 follicular cell markers, and have 510 a predominantly activated phenotype with high MHC and costimulatory molecule expression 511 512 (not shown, [45; 48]). The location of thymic B cells in NOD mice is also reminiscent of reports in other murine strains- positioned predominantly at the cortico-medullary junction-513 514 but in contrast to non-autoimmune prone mice, large B cell follicles form and this is age-515 dependent. Furthermore, the hallmarks of germinal centres are readily detectable in the NOD thymus; IL-21 and T follicular helper (TfH) cells. Abnormalities in levels of IL-21 and TfH 516 517 cells in peripheral tissues, and blood, have been strongly associated with T1D [37; 49]. Here, we show similar abnormalities exist in the thymus occurring specifically at the late insulitic-518 pre-diabetic phase of the T1D condition. In addition the thymus of NOD mice has enhanced 519 levels of AID mRNA transcripts, suggesting increased in situ somatic hypermutation and 520 class switching of the B cell repertoire activity. Plasma cells are also increased in the thymus 521 of NOD mice with respect to control animals which taken all this information together 522 implies ectopic germinal centres are a feature of the NOD thymus, not just their pancreas 523 [50]. Our evidence that the NOD mouse thymus is populated with significantly increased 524

numbers of B cells with IgG, IgA and IgE receptors with respect to non-autoimmune prone
mice, as well as enhanced levels of soluble IgG1 and IgA antibodies supports our rationale of
ectopic germinal centre formation in this primary lymphoid tissue.

The significance of these unique changes in the NOD mouse thymus as mice progress along 528 the T1D pathway, we believe, is that they have the potential to impact on the capacity of 529 negative selection of autoreactive T cells to occur effectively. The importance of mTEC 530 expression of TSAs for efficient deletion of developing T cells bearing autoreactive T cell 531 receptors is well established [51]. Our evidence that a selective population of mTECs have 532 autoantibodies bound in situ, and in the presence of thymic B cells a proportion of mTECs 533 534 undergo apoptosis, a number of which express insulin, is likely to have implications on negative selection of islet-reactive T cells. The antigenic specificity of the intrathymic 535 autoantibodies target is unknown, and we do not believe that they must recognise insulin to 536 537 impact of T1D progression. It is possible that the intrathymic autoantibodies recognise and promote apoptosis of particular mTECs that express certain TSAs that are associated with 538 539 other autoimmune conditions NOD mice develop [52; 53; 54]. It follows that reduction in 540 insulin expressing mTECs may happen inadvertently. Alternatively, or in addition, it is possible that thymic cognate B-T cell interactions promote survival of developing 541 542 autoreactive T cells as opposed to their deletion [55].

Our data is supportive of the rationale that pre-negatively selected T cells are potentially released from the thymus prematurely. Two photon microscopy has documented that developing T cells reside in medulla for 3-5 days to complete negative selection [56]. In RAG2p-GFP reporter mice, this duration in the medulla equates to decreased GFP intensity due to the 54 hour half-life of the molecule [30]. Our evidence that in NOD mice CD3⁺GFP^{hi} cells are significantly enhanced in peripheral blood with respective to non-autoimmune prone mice suggests an aborted time, or failed entry into, the medulla of GFP^{hi} T cells, and as a

550 consequence failed negative selection. It follows that the increased export of non-negatively 551 selected T cells could overpower waning regulatory mechanisms in the islets leading to the 552 final sustained attack of the β cells.

The fledgling field of thymic B cell research is starting to unravel the importance of this
unique population of cells in the immune system. Our data highlight a new relationship
between thymic B cells and type 1 diabetes development. Future studies that define the *in situ*developmental pathway and receptor specificity of thymic B cells will be important for
identifying key therapeutic strategies for type 1 diabetes and other autoimmune conditions in
which thymic B cells make a contribution.

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564 Authors contributions:

AP, JS, MK, KH and EAG performed the experiments and analysed data; AP and EAG wrotethe paper.

567 Conflict of Interest Statement

568 All authors declare there was no conflict of interest.

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740 Figure Legends

Figure 1. Intrathymic B cell accumulation precedes ß cell destruction. Single cell 741 suspensions were prepared from the respective thymi and all data analyzed on a single cell, live 742 gate. (a) Representative dot plots of thymic CD19⁺ cells: (I) Isotype control for CD19 antibody; 743 (II) 12 week old female B6 mouse; and (III) 12 week old female NOD mouse. (b) Number of 744 B cells in the thymus of 4-6 week old female B6 mice (n=11), 12-14 week old female B6 mice 745 (n=25), 4-6 week old female NOD mice (n= 13) and 12-14 week old female NOD mice (n= 746 25). (c) Number of RAG-GFP^{hi} B cells in the thymus of 4-6 week old female FVB-RAG-GFP 747 mice (n=3), 12-14 week old female FVB-RAG-GFP mice (n=8), 4-6 week old female NOD-748 RAG-GFP mice (n=4) and 12-14 week old female NOD (n=8). (d) Frequency of Ki67⁺ B cells 749 in the thymus of 4-6 week old female B6 mice (n=10), 12-14 week old female B6 mice (n=10), 750 4-6 week old female NOD (n=9) and 12-14 week old female NOD mice (n=8). Data presented 751 752 as scatter plot, each dot equating to a mouse, the bar representing the mean value. Statistical significance determined using the non-parametric Mann-Whitney U-test, significant P values 753 754 are shown, ns= not significant.

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Figure 2. B cells form follicle-like structures at the cortical-medullary junction in NOD 756 thymi. Flow cytometric analysis of the (a) frequency of B cells displaying a Follicular (FO) or 757 758 Marginal-zone (MZ) phenotype in the thymus of B6 (n=10) or NOD mice (n=13) and (b) absolute number of B cells displaying FO or MZ phenotypes in the thymus of B6 (n=5) or 759 760 NOD mice (n=6). Comparisons made between aged-matched, female11-14 week old mice in a single cell, live gate. Data acquired from at least two independent experiments and is presented 761 as scatter plot; P values were calculated using the Mann-Whitney U test analysis, ns= not 762 763 significant. (c) Representative confocal immunofluorescence microscopy images of thymi sections examined for B220 (yellow), cytokeratin V (green) expression and the DNAintercalating dye DAPI identified nuclei (blue) from 11 week old female NOD or B6 mice. A total of 14 sections from eight NOD mice and a total of six section from three B6 mice were analyzed, and there was consistency in the data obtained from the appropriate strains of mice. Confocal fluorescent images were obtained with a Plan-Apochromat 20x objective.

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770 Figure 3. The NOD thymus has the hallmarks of ectopic GC development. (a) Evaluation of IL-2/IL-21 ratio in cell-free supernatants from spleen and thymic tissue from 11-15 week 771 old B6 (n=5) or NOD (n=4) mice. The data shown is representative of two individual 772 experiments showing similar results. (b) Frequency of CD4⁺ T follicular helper cells (TfH) in 773 the thymus of B6 (n=12) and NOD mice (n=17). (c) Number of CD4⁺ T follicular helper cells 774 775 in the thymus of B6 (n=10) and NOD mice (n=11). (d) Representative histogram of BCL-6 expression in CD8⁻ CD4⁺ PD1^{hi} ICOS⁺ cells in thymus of NOD (n=5) and B6 (n=5) mice. (e) 776 Number of plasma cells in the thymus of B6 (n=15) and NOD mice (n=15). For (b-e) 777 comparisons made between female, age-matched 10-14 week old B6 and NOD mice. The 778 779 analysis was performed on a single cell, live gate, and the data is presented as a scatterplot, each dot equating to a mouse, the bar represents mean value. P values were calculated using 780 the Mann-Whitney U test analysis and are shown in the figure, ns= not significant. The data is 781 pooled from at least two independent experiments giving similar results. (f) Quantitative PCR 782 (qPCR) analysis of AID mRNA (aicda expression) levels in the whole thymus. Data was 783 784 normalized to HPRT mRNA as described in methods, and fold change in NOD mice AID mRNA when compared to normalised AID mRNA levels for B6 mice. All mice were 11-14 785 786 weeks of age, a total of 5 female B6 mice were compared to 5 female NOD mice. One thymic sample from a female B cell-deficient NODµMT^{-/-} mouse was used as a negative control. The 787 data is pooled from two independent experiments and is presented mean \pm standard error mean 788

(SEM). P values were calculated using the Mann-Whitney U-test and are shown in the figure,
ns= not significant.

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Figure 4. The NOD thymus harbours a unique pattern of immunoglobulin isotypes (a) 792 Number of IgM⁻ IgD⁻ IgA⁺, IgM⁻ IgD⁻ IgE⁺, IgM⁻ IgD⁻ IgG⁺ B cells in the thymus of 11-14 week 793 794 old female B6 (n=10) or female NOD mice (n=10). (b) Number of IgM⁻ IgD⁺ IgA⁺, IgM⁻ IgD⁺IgE⁺, IgM⁻ IgD⁺ IgG⁺ B cells in the thymus of 11-14 week old female B6 (n=10) and 795 female NOD mice (n=10). (c-f) Optical density (OD) values of the respective immunoglobulins 796 in cell free tissue supernatants (c,e) or serum (d,f). A total of 6 female B6 and 6 female NOD 797 mice were assessed in two independent experiments. Data is presented as scatter plot, each dot 798 799 equating to one mouse and bar representing the mean. P values were calculated using the Mann-Whitney U-test analysis and are shown in the figure, ns= not significant. 800

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802 Figure 5. IgGs bind to thymic stromal components in NOD mice. (a and b) Representative confocal immunofluorescence microscopy images of thymi sections of NOD (aI-III) and B6 803 mice (bI-II) examined for cytokeratin V (red), murine IgG (green), and the DNA-intercalating 804 dye DAPI (white). A total of six 11 week old NOD mice and five 11 week old B6 mice, two 805 sections per mouse were examined. (a)I,II and II are derived from different NOD mice. The 806 confocal fluorescent image in AI was obtained with a Plan-Apochromat 20x objective to give 807 a broader view of the extent of immunoglobulin bound to thymic stroma, arrows indicating 808 809 some of the cells co-positive for cytokeratin V and mouse IgG. The confocal fluorescent images in AII and AIII were obtained with a Plan-Apochromat 63x objective. For (b) the confocal 810 811 fluorescent image was obtained using a Plan-Apochromat 20x objective. (c) Quantification of 812 murine Ig-bound to stromal cells of age-matched 11 week old, female NOD or B6 mice.

Confocal immunofluorescence microscopy images were subjected to StrataQuest V64 analysis, a total of $2-3x10^4$ DAPI⁺ cells/mm² were counted and the mean fluorescence intensity of DAPI⁺ cells versus mean fluorescence intensity of anti-Ig is presented as a scattergram. The data shown, is representative of two independent mice examined giving similar results.

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Figure 6. Increase in thymic B cells was associated to increased apoptosis of stromal cells. 818 (a) Representative confocal immunofluorescence microscopy images of thymi sections from 819 9-14 week old female NOD, and NOD-µMT^{-/-} and B6 mice examined for cytokeratin V 820 (yellow), apoptosis (red), and the DNA-intercalating dye DAPI (white) expression. The data is 821 representative of similar data acquired from 6 female NOD, 6 female NOD-µMT^{-/-} and 4 822 female B6 mice, three sections per mouse were examined. In all cases, the confocal fluorescent 823 images were obtained with a Plan-Apochromat 63x objective. Bar represents 20µm. (b) 824 Quantification of Tunel⁺ stromal cells of age-matched 11 week old, female NOD, NOD-µMT⁻ 825 ¹⁻ or B6 mice. Confocal immunofluorescence microscopy images were subjected to StrataQuest 826 V64 analysis, a total of $4x10^4$ DAPI⁺ cells/mm² were counted and the mean fluorescence 827 intensity of DAPI⁺ cells versus mean fluorescence intensity of Tunel is presented as a 828 scattergram. 829

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Figure 7. B cells promote premature thymic-release of T cells prior to negative selection.

(a-b) Thymocytes from 11-12 week old B6, NOD and NOD NOD- μ MTKO mice were stimulated with insulin or B15:23 peptide (NOD and NOD- μ MTKO mice, only) for 72 hours and Ki67 expression in CD4SP (a) or CD8SP (b) cells as a measure of proliferation was determined by flow cytometry. The frequency of Ki67⁺ cells for stimulated samples was normalized against the frequency of Ki67⁺ cells in unstimulated samples. (c) Frequency of

- 837 RTEs (RAG-GFP^{hi}) in peripheral blood of 11-12 week old FVB-GFP (n=8), NOD-GFP (n=8)
- 838 or NOD- μ MT^{-/-}-GFP (n=5) mice. Data is pooled from two independent experiments and cells
- 839 were analysed on a live, single, $CD3^+$ T cell gate. The data is presented as scatter plot, the bar
- 840 representing the mean value;P values were calculated using the two-way Anova followed by
- 841 Tukey multi comparison test and are shown the figure, ns= not significant.