

Immunoglobulin E response in health and disease beyond allergic disorders

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2	disorders
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24	Lists of abbreviations:

Ag	Antigen
AID	Activation Induced cytidine Deaminase
Aka	Also Known As
APE1	APurinic-apyrimidinic Endonuclease 1
BSA	Bovine serum albumin
CDx	Cluster of Differentiation n°x
CSR	Class Switch Recombination
DMBA	7,12-dimethylbenz[a]anthracene
FAP	Facilitated antigen presentation
FcERI	Receptor I for constant chain of immunoglobulin E
FceRII	Receptor II for constant chain of immunoglobulin E
GC	Germinal Center
GF	Germ Free
GLTs or I _x	Germ Line Transcript
IDO	indole-2,3-diamine oxygenase
IgE	Immunoglobulin of isotype E
IgG	Immunoglobulin G
IgND	Immunoglobulin Not Determined
IgY	Immunoglobulin Y
MALT	Mucosal Associated Lymphoid Tissues
MHC	Major Histocompatibility Complex
PBMC	Peripheral blood mononuclear cells
PID	Primary Immune Deficiency
SHM	Somatic HyperMutation

SLO	Secondary Lymphoid Organs
srIg	Self-reactive immunoglobulin
TPA	12-O-tetradecanoylphorbol-13-acetate
TCR	T cell receptor
Tfh	T follicular helper cell
Tfr	Follicular regulatory T cell
TNP	Trinitrophenyl
UNG	uracil-DNA glycosylase

26 **Running title:**

- 27 IgE involvement in health and non-allergic diseases
- 28

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47

48 Abstract:

49 Immunoglobulin E is the latest discovered of immunoglobulin family and has been long 50 associated with anaphylaxis and worm expulsion. Immunoglobulin E, along with mast-cells, basophils and eosinophils, is also a hallmark of type 2 immunity which is dysregulated in 51 52 numerous diseases such as asthma, rhinitis, atopic dermatitis, eosinophilic esophagitis in addition to anaphylaxis as aforementioned. However, recent advances have shed light on IgE 53 54 regulation and memory explaining the low level of free IgE, the scarcity of IgE plasma cells that are mainly short live and the absence of IgE memory B cells in homeostatic conditions. 55 56 Futhermore, IgE was implicated in inflammatory conditions beyond allergic disorders where 57 IgE-mediated facilitated antigen presentation can enhance cellular and humoral response against autoantigens in systemic lupus or chronic urticaria leading to more severe disease and 58 59 even against neoantigen facilitating tumor cell lysis. At last, IgE was unexpectedly associated 60 with allograft rejection or atheromatous cardiovascular diseases where precise mechanisms remain to be deciphered. The purpose of this review is to summarize these recent advances in 61 IgE regulation, biology and physiopathology beyond allergic diseases opening whole new fields 62 63 of IgE biology to explore.

64

65 Key words: IgE, Type 2 immunity, Facilitated-antigen presentation, FcεRI, CD23

66 INTRODUCTION

67

Immunoglobulins of isotype E (IgE) were initially reported as a "reagin" in the early 68 1920's by Coca and Cooke who worked on a classification of "the phenomena of 69 hypersensitiviness"¹. Little work occurred on "reagin" from the 1920's until the 1960's, until 70 71 newer techniques to better detect proteins were developed. In 1966, Kimishige and Teruko 72 Ishizaka described an anti-serum that was able to block type 1 hypersensitivity and called this 73 molecule γ E-globulin². This globulin did not bind complement nor induced a precipitin 74 reaction like other immunoglobulins. Simultaneously, Bennich and Johansson discovered a paraprotein in a patient affected by myeloma that did not belong to any of the known 75 immunoglobulins. He called it "Immunoglobulin Not Determined" (IgND) and found that it 76 had similar properties to reagin³. Both IgND and γ E-globulin initiated the Prausnitz-Kustner 77 78 test (i.e., passive cutaneous anaphylaxis) and the authors discovered that "reagin", IgND and γ E-globulin were finally the same immunoglobulin ⁴. In 1968, γ E-globulin and IgND were 79 80 officially named Immunoglobulin E by the World Health Organization International Reference 81 Center for Immunoglobulins ⁵.

Phylogenetically talking, IgG and IgE are thought to have emerged from a reptilian 82 common ancestor called Immunoglobulin Y (IgY)⁶. On one hand, IgE and IgG share 83 84 similarities. Both are monomeric immunoglobulins composed of two identical heavy and light 85 chains ending with same variable domain. They adopt a 3D structure depending on their FC receptors binding profiles and thus on their biological function. They exert their effector 86 functions through interactions with Fc receptors⁷. On the other hand, IgG and IgE evolved 87 differently on several points in mammals. Indeed, they are respectively the most and the least 88 89 abundant immunoglobulin classes in human serum. IgE heavy chain is longer than IgG with 4 constant domains and contains seven N-linked glycosylation sites (three on $C\varepsilon_1$, one on $C\varepsilon_2$ 90

and three on C ε 3) while a single N-linked glycosylation site is present in IgG^{7,8}. This high glycosylation rate may contribute to increase its solubility whereas deglycosylated IgE had a tendancy to aggregate⁸.

To date, IgE is mostly known as an anaphylactic immunoglobulin and studied in allergic diseases such allergic rhinitis & asthma and food allergy where specific IgE-allergen complex plays a major part in the physiopathology both in acute and chronic phases of allergic inflammation⁹. The main mechanism relies on allergen cross-linking with specific IgE bound on effector cells (mast-cells, eosinophils, basophils) via its high affinity receptor, Fc ϵ RI, inducing secretion of inflammatory cytokines and mediators such as histamine, heparin, tryptase, prostaglandins⁴.

101 Beyond allergic inflammation and related diseases, specific IgE secretion is a soluble mediator 102 and also a hallmark of "type 2 immunity" along with eosinophil, mast-cell and basophils 103 infiltration in inflamed tissue and peripheral blood. So called "type 2 immunity" or "type 2 104 response" is triggered by a large panel of microbial (virus, bacteria, parasite) and non-microbial (venom, allergens, synthetic or natural adjuvants) stimuli ranging from nanometer to several 105 106 meters long suggesting a multiples variants of type 2 responses either protective or pathologic where the precise roles of IgE remains poorly understood¹⁰. However, several advances in 107 108 recent years have begun to shed light on IgE regulation and memory, on its role in homeostatic 109 and inflammatory conditions beyond allergic disorders such as helminth infection, neoplasia, 110 auto-immunity/auto-allergy, vascular disease and solid organ transplantation. The purpose of 111 this review is to summarize these advances in IgE physiopathology.

113 IMMUNOGLOBULIN E REGULATION

114

115 Concentrations of free serum IgE are \sim 50–200 ng per ml of blood in healthy humans 116 without any atopic background compared with around 1-10 mg per ml of blood for other immunoglobulin isotypes¹¹. In addition, the serum half-life of IgE is the shortest of all 117 immunoglobulin isotypes in human to ~ 2 days compared to 20 days for IgG¹¹. Of note, IgE 118 half-life is markedly prolonged around 9-12 weeks when bound to its high affinity receptors on 119 mast-cells and basophils $(Fc \in RI)^{12}$. At last, IgE production can occur at a loco-regional 120 level^{13,14}. These observations suggest that IgE production is tightly regulated at low levels under 121 122 homeostatic condition to avoid potential life-threatening condition such as anaphylaxis or 123 deleterious type-2 hyperresponsiveness.

124

125 Intrinsic regulation of IgE

126 IgE class switch recombination

127 Class switch recombination (CSR) occurs in naive B cells (CD19+, CD24+, CD38+, 128 IgM+ and IgD+) in secondary lymphoid organs (SLO) such as lymph nodes, spleen or mucosal 129 associated lymphoid tissues (MALT) under BCR activation, CD40 ligation and cognate CD4+ T cell help at the T-B zone before organizing into germinal centers¹⁵. CSR is an 130 131 intrachromosomal rearrangement of the immunoglobulin heavy-chain locus resulting in isotype 132 switching from IgM-IgD to either IgG₁₋₄ or IgA or IgE that differ in effector functions without altering the specificity for the antigen ¹⁶. CSR relies on activation of several key enzymes such 133 134 as Activation Induced cytidine Deaminase (AID), uracil-DNA glycosylase (UNG) and 135 APurinic-apyrimidinic Endonuclease 1 (APE1) that bind to specifically intronic regions called donor and acceptor switch regions (S_d and S_a respectively) ^{17–19}. DNA recombination between 136 S_d and S_a as well as the transcription of germ line transcript (GLTs aka I_x) which are crucial 137

regulators of CSR via specific transcription factor binding sites²⁰ under the dependence of
super-enhancer 3'RR²¹. Selection of IgE isotype is driven by cytokine micro-environment, IL4, IL-13 and IL-9²².

Ide IgE CSR has unique features. On a first hand, S ε structure is shorter (around 2kb) than other S region making more difficult DNA recombination all the more so chromatin is condensed according to epigenetic information^{23,24}. On a second hand, IgE CSR can occur directly from S μ to S ε or indirectly from S μ to S γ 1 and from S γ 1 to S ε ²⁵. Recently, a probabilistic model suggested that direct CSR occur more frequently with a low activation threshold generating low affinity IgE ²⁶ mostly from naive B cells in SLO ²⁷ whereas indirect CSR would be more frequently associated with high affinity specific IgE ^{26,28} (Figure 1).

148

149 *IgE BCR impacts on B cell fate*

IgE BCR itself induces B cell fate toward short life. First of all, IgE mRNA transcript lack the 150 canonical polyadenylation tail and thus a lower ratio of IgE BCR at the cell membrane²⁹. 151 Moreover, IgE B entry in germinal center (GC) is impaired due to lower mobility preventing 152 153 memory B cells or long-live plasma-cells differentiation all the more so recent findings have revealed that CSR occurs preferentially before GC formation¹⁵. At last, membrane IgE BCR 154 induces a suicide locus conferring to IgE+ B cells a higher propensity to apoptosis³⁰. Those 155 156 observations argue that IgE+ B cells are fragile, prone to apoptosis and to become short-live 157 plasma cells consistent with low but not null IgE secretion (Figure 1).

158

159 Extrinsic regulation of IgE

160 *IgE CSR is tightly regulated in SLO.*

As aforementioned, indirect class switch recombination results more frequently in high affinity
 IgE^{26,28}. Though for a same rate of somatic hypermutation, IgE affinity remains lower

compared to IgG1 affinity suggesting that T follicular helper cells (Tfh) regulate IgE CSR and 163 affinity maturation in SLO. A new population of Tfh, "Tfh-13", was described in several mice 164 165 models of food allergy and allergic asthma promoting high affinity IgE maturation compared to non-allergic or helminth-infected mice³¹. These particular Tfh expressed canonical master 166 167 genes for type 2 inflammation (GATA3) and follicular polarization (Bcl-6). They also 168 expressed high amount IL-4 and IL-13 and low amount of IL-21 leading to indirect CSR. Indeed, IL-21, IFN- γ and IL-10 are known to restrain IgE CSR all the more so they act in 169 170 synergy $^{32-34}$. At last, Tfh-13 are associated with IgG1 memory B cells that held IgE memory 31 . Clement et al described a novel follicular regulatory T cells (Tfr) producing neuritin that were 171 able to constrain IgE-CSR and Tfh-13 formation preventing IgE high affinity maturation^{35,36}. 172 Of note, human IgE repertoire was shown to have less SHM in healthy individuals^{37,38}. 173 174 Furthermore, Shade et al demonstrated that several glycosylation patterns on IgE correlated 175 with its function. More precisely, N384-linked oligomannose on C ε 3 was essential to enhance FceRI binding^{39,40} whereas a higher number of sialic acid residues on N168 (C ε 1) and N265 176 (C ϵ 2) was essential to promote mast-cell degranulation in food allergic patients⁴¹. Those 177 modification were probably acquired after affinity maturation in the germinal center⁴². IgE 178 binds a C-type lectin domain on its low affinity receptor (CD23 or Fc ε RII; Kd ~ 0.1–1 μ M)⁴³. 179 180 Interestingly, IgE binding on its monomeric form (soluble or membrane-anchored) acts as a 181 negative feedback loop on B cells, suppressing IgE synthesis when interacting with CD21 44(p21),45 182

Taken together, those observations suggest that, in SLO, IgE production and memory is
constitutively constrained at low level with a relative low affinity to their cognate antigen. This
is mostly due to Tfr balancing with Tfh-13 and CD21/CD23-mediated negative feeback loop.
These concepts are summed up in Figure 2.

188 IgE function is actively constrained out of SLO and in peripheral tissues.

In 2018 Shan et al described an novel mechanism constraining IgE-driven basophils 189 190 degranulation in epithelia and inflamed tissues via IgD and galectin 9. Briefly, in an ovalbumin 191 (OVA)-sensitized mice model, OVA-specific IgD could bound basophils via galectin 9 and 192 CD44 preventing their degranulation and enhanced type 2 inflammation. Galectin 9 could also bind IgE subsequently inhibiting $Fc \in RI$ mast-cells degranulation⁴⁶. Finally, no intrinsic defect 193 194 in activating nor regulatory signals were found in human basophils either from allergic patients or healthy volunteers⁴⁷. These data strongly suggest that anaphylactic IgE are tuned down even 195 196 when type 2 immunity is triggered.

197 FccRI (aka high affinity receptor - Kd ~ 1nm) is found as a tetramer, $\alpha\beta\gamma2$, on mast cells and basophils, and as a trimer, $\alpha \gamma 2$, on other cells such as monocytes, dendritic cells, eosinophils, 198 and platelets. FceRI interacts with IgE through Ce348-50. A mathematical modelisation of 199 200 immunoglobulin metabolism found that IgE undergoes substantial catabolism at extravascular sites related to IgE/FceRI interaction in the peripheral tissues⁵¹. This prediction was confirmed 201 202 in vitro (human dendritic cells/monocytes) and in vivo (transgenic mice humanized with human 203 trimeric form of FceRI). Briefly, IgE is quickly endocytosed in a FceRI-dependent manner promoting serum IgE clearance and participate in IgE homeostatis⁵². 204

At last, IgG_{1-4} auto-antibodies against IgE were described in atopic/allergic diseases and autoimmune disorders and healthy volunteers ^{53–55}. Those anti-IgE auto-IgG are able to recognize free and FccR-bound IgE in contrast to Omalizumab that can bind only free IgE^{54,56}. Recently, Chan et al showed that anti-IgE auto-IgG could either activate or inhibit in vitro basophil degranulation test to grass pollen recombinant protein⁵⁷ suggesting an inhibitory role preventing type 2 inflammation though mechanism is still unclear.

Altogether, those data highlight that in peripheral tissues, IgE function are constitutively
 tracked down by active mechanisms (IgD mediated inhibition of basophils degranulation +/-

inhibitory auto-IgG against IgE and constant IgE catabolism) in addition to previously detailed
mechanisms in SLO (Figure 3).

215

216 *IgE production is a basal trait of mucosal and imbalanced immunity.*

217 More systemic and integrative hypothesis of IgE homeostasis arose from germ-free 218 (GF) mice studies and monogenic primary immunodeficiencies in humans (PID). Such 219 deficiencies result in hyper-IgE and atopic/allergic-like clinical manifestations along with 220 susceptibility to infections and/or auto-immune manifestations. Those PID could affect either 221 regulatory T cell compartment (FoxP3 mutation loss of function), TCR/cytokine signaling (ZAP-70 hypomorphic mutation loss of function as an example), TCR repertoire (restrained 222 oligo-clonal repertoire), glycosylation protein pathway (phosphoglucomutase 3 deficiency), 223 224 transduction pathways (STAT3 or DOCK8 deficiency) or even barrier function (Netherton syndrome caused by SPINK5 mutation loss of function)⁵⁸. Despite precise mechanisms leading 225 to hyper-IgE are still poorly understood, a common mechanism would be that the 226 imbalanced/impaired type 1 or type 17 immunity unmask type 2 immunity⁵⁹. 227

228 Indeed, a high amount of IgE in GF mice occurred early in life (before 5 weeks of age) in secondary lymphoid structure at mucosal sites. It was dependent on IL-4 produced by CD4+ 229 230 T cells which precise lineage is not known. IgE production could be abolished by neonatal 231 microbiota colonization or (before 5 week of age) by fecal microbiota transplantation from conventional-housed mice whereas in adults the process was stabilized⁶⁰. In addition, food 232 233 antigens in non-sensitized GF mice induced sustained specific IgE production at mucosal site 234 but also in SLO. The IgE synthesis was dependent on CD40L+ ICOS+ Tfh in secondary lymphoid structures⁶¹. In both studies, GF mice developed aberrant Ag-specific IgE bound to 235 236 mast-cells that could induce anaphylactic symptoms after second Ag exposure.

Taken together, these data suggest that IgE production (Tfh dependent) is set by default either at mucosal sites in absence of physiological microbiota colonization and/or by an imbalanced type 1 and/or type 17 immunity during early life. This makes an echo to birthdelivery mode that impact on neonatal gut microbiota colonization and diversity leading to miseducation of immune system. Indeed, neonates with low gut microbiota diversity have a higher risk to develop atopic/allergic diseases consistent with IgE dysregulation ^{62,63} (Figure 4).

244 IMMUNOGLOBULIN E BIOLOGY

245

246 Does natural IgE production exist?

Several mechanisms are at play to tune down IgE production. Though, IgE titer is not 247 248 null under homeostatic condition suggesting that a natural secretion of IgE does exist despite 249 the absence of long lived IgE plasma cells. One hypothesis would stand that IgE memory is kept by IgG1 memory B cells^{64(p1)}. Thus, constant IgE secretion would suggest a low-to-250 251 intermediate affinity to avoid uncontrolled type 2 inflammation and a constant/regular 252 stimulation by (auto-)antigens that remain to be determined. Another hypothesis relies on the secretion of natural IgE independent of major histocompatibility complex (MHC) cognate T-253 254 cell help in secondary lymphoid structures in a GF and T-cell-deficient mice models. Those IgE had a low rate of SHM suggesting a low-to-intermediate affinity for antigens. At last, no 255 evidence of deleterious type 2 activation was observed ⁶⁵. 256

Whatever their origin, the role of natural IgE remains poorly understood. An hypothesis comes 257 258 from old clinical observations where tissular damages (burns or in immediate post-operative 259 period) resulted in a rapid polyclonal increase of circulating IgE without deleterious type 2 inflammation ^{66,67}. A recent study by Crawford et al demonstrated that a carcinogenic 260 261 environmental xenobiotic called 7,12-dimethylbenz[a]anthracene (DMBA) triggered a strong 262 polyclonal IgE response with self-reactivity when applied on skin. This IgE response, which was dependent on $Fc \in RI$ binding, prevented epithelial damage⁶⁸. Although, their exact 263 specificities remain to be determined, those IgE could be implied in maintaining immune 264 tolerance to self-antigens from damage tissues¹⁰. This is supported by two recent studies in mice 265 266 humanized for FcERI alpha subunit which demonstrated anti-inflammatory properties of IgE-FceRI activated dendritic cells. Platzer et al showed that antigen/specific IgE complex 267 dampened systemic inflammation in an allergic asthma mice model in a $Fc \in RI$ -dependent 268

manner⁶⁹. Baravalle et al showed that monovalent antigen covalently bound to IgE first induced 269 CD4+ T cell proliferation, quickly followed by a systemic clonal deletion of the antigen specific 270 CD4+ T cells⁷⁰. In addition, IgE-Fc*e*RI activated DC promoted a IL-10 dependent retro-control 271 loop leading to effector T-cell deletion and regulatory T cell activation via indole-2,3-diamine 272 oxygenase (IDO) activation ^{71–73}. A FceRI soluble form was detected in human serum ⁷⁴ and 273 exhibited an inhibitory effect on mast cell degranulation⁷⁵. Altogether, those data argue for an 274 immuno-modulatory role of natural IgE on innate immunity and indirectly on adaptative 275 276 immunity. Of note soluble form of $Fc \in Rs$ could be an add-on level of regulation. Yet, further 277 investigations are needed to understand the complex inter-play between IgE, FcERs and immune tolerance (Figure 5). 278

279

280 IgE facilitated antigen presentation – a pivotal role in unmasking IgE 281 response/inflammation

IgE-mediated response/inflammation is somehow necessary¹⁰. The IgE facilitated antigen presentation (IgE-FAP) aka IgE antigen focusing plays a pivotal role in IgE inflammation. This process has been mainly studied in immediate allergic diseases where it set up and amplified the allergic inflammation against allergens⁷⁶.

In 1993, Heyman et al showed that the rate of IgG anti-bovine serum albumin (BSA) 286 and BSA-specific IgG+ B cells was higher when mice were immunized with trinitrophenyl 287 (TNP) – BSA and TNP-specific IgE^{77,78}. Another description of IgE-FAP came from the "toxin 288 hypothesis" that stood a protective role of IgE against venom⁷⁹. Later, it was demonstrated that 289 290 bee-venom immunized mice were more resistant to IgE-mediated anaphylaxis when rechallenged with a lethal dose of bee or viper venom⁸⁰. In the same line, a recent study 291 demonstrated that a pre-existing IgE sensitization against Staphylococcus aureus (SA) toxin 292 enhanced SA IgG immunization and infection clearance in a mice model of pulmonary 293

infection⁸¹. These data suggest that one role of IgE-FAP would be to enhance humoral andcellular response against non-self-antigen.

There are two main mechanisms known so far for IgE-FAP implying either FcERI or 296 FceRII. Multivalent antigen/IgE complexes were shown to activate dendritic cells ⁸² mostly 297 toward a type 2 inflammation but not exclusively⁸³. IgE-FAP activated dendritic cells were also 298 showed to boost specific CD4+ effector T cells via MHC-II presentation with a 100 to 1000 299 300 fold higher efficiency than canonical MHC-II presentation (low dose of antigen)⁸². Moreover, IgE-FAP activated basophils were more efficient at MHC-II cross presentation than dendritic 301 cells^{84,85}. At last, FceRI-driven IgE-FAP demonstrated in vitro a broad spectrum of action on 302 human mast cells ranging from self-survival, pro-inflammatory cytokines secretion to complete 303 degranulation according to the nature of IgE and the antigen⁸⁶ (Figure 6a). 304

305 CD23 has three major known ligands: CD21, MHC-II and IgE with different interaction sites respectively⁸⁷ especially with B cells and monocytes/macrophages⁸⁸. Noteworthy, CD23 306 307 has two isoforms: CD23a is constitutively expressed and up-regulated after IL-4 stimulation whereas CD23b is expressed only after IL-4 stimulation⁸⁹. Only CD23a under its trimeric form 308 is implied in antigen processing whereas CD23b is more related to phagocytosis⁹⁰. Conversely 309 to FcERI-driven IgE-FAP, several studies demonstrated that CD23a IgE-FAP enhanced 310 311 humoral response (IgG and IgM) as well as the specific cellular response (CD4+ T cells) against 312 a multivalent antigen (TNP-OVA) in mice models either adjuvanted with TNP-specific IgE or transgenic mice over-expressing CD23^{91,92}. The magnitude of the response was higher with 313 antigen/IgE complex and low dose of antigen⁹². Those data were confirmed in human by in 314 vitro models where B cells from allergic patients expressed a high density of CD23 arguing for 315 a trimeric form^{93,94}. Trimeric soluble CD23 is also detectable in human serum which function 316 seems to be identical to the membrane form ⁹⁵ (Figure 6b). 317

Altogether, those data suggest another role of IgE-FAP which could be 1) the detection of low amount of antigen and 2) the onset of an adaptative type 2 immune response mainly but not exclusively. This suggests that IgE takes part to immuno-surveillance to non-self. As well, IgE-FAP seems to trigger an inflammatory response with multivalent antigens. This makes an echo to a recent study where the more the IgE interacting epitopes were numerous and close to each-other the more the IgE complexes were able to cluster IgE receptors (Fc ϵ RI) and trigger mast cell activation⁹⁶.

325 POTENTIAL ROLES OF IMMUNOGLOBULIN E IN HEALTH AND NON-

326 ALLERGIC DISEASES

327 IgE and venom toxicity – a host defence mechanism?

328 The hypothesis that IgE could play a protective role against toxins after arthropods or reptile bite was proposed by James Stebbings in 1974⁹⁷ and Margie Profet in the early 1990's⁷⁹. Since, 329 330 important works from Stephen Galli's group have get credit to this hypothesis suggesting that IgE and its interaction with mast cells can enhance host resistance to venoms. More precisely, 331 332 honeybee venom resistance to a lethal dose can be transferred to a naive mice by systemic serum therapy from a honeybee immunized mice⁸⁰. This protective effect was lost either after 333 334 depletion or inactivation of IgE in the serum or when Fc ϵ RI α or γ chain were knocked out in the naïve recipient mice^{80,98}. Those result suggested that IgE-mediated resistance to venom is 335 336 linked to a FccRI-IgE FAP on effector cells. Indeed, mice lacking mast cell and/or basophils were unable to acquire venom resistance after immunization⁹⁹. Furthermore, local FccRI-IgE 337 FAP mast cell degranulation enhanced venom detoxification^{99,100} thanks to proteases contained 338 in their granule such as carboxypeptidase 3 and chymase^{101,102}, which degrade venom peptides 339 340 (sarafotoxin and helodermin respectively) with structure similarity to endogenous mammalian peptide (endothelin 1 and vasoactive intestinal peptide respectively)¹⁰³. Interestingly in those 341 mice model of envenomation, mast cell, basophil or even innate lymphoid cells type 2 were not 342 required to induce IgG1 and IgE immunization against venom antigen^{98,99}. Last but not least, 343 344 previous IgE immunization against an irrelevant allergen added to venom resulted in enhanced resistance to the venom despite prior immunization⁹⁹. Altogether, those data demonstrate that 345 detoxification to a complex mixture of antigen such as venom involve specific IgE sensitization 346 347 to a small fraction of antigen that induce the release of proteases due to FccRI-IgE local degranulation of mast cells. Furthermore, those results suggest that venom itself induce specific 348 349 IgE immunization independent of effector cells (Figure 7).

351 IgE and helminth – who is protecting who?

352 High IgE titer is one hallmark of active helminthic infection with eosinophilia reflecting type-353 2 inflammation. In human, several studies associated resistance to (re)-infection to specific IgE against Schistosoma mansoni or haematobium^{104,105} whereas the underlying mechanisms 354 355 associated with IgE and parasite resistance remain unclear. So far, mice models of helminthic 356 infection used to better decipher the role of IgE during helminthic infection in has led to 357 conflicting results. An explanation, at least in part, could be the use of inadequate model to study IgE response such as AID-deficient mice, B cell deficient mice or FcyR deficient 358 359 (reviewed in¹⁰⁶). Table 1 summarizes different studies that used IgE-deficient or FccRIdeficient mice to better decipher their role during primary or secondary helminthic infections. 360 361 Several interesting patterns draw according to worm burden mostly and to a lesser extent on 362 granuloma in target organs. It is worth noticing that the time after infection inversely correlates 363 with worm burden: the longer it is, the smaller the difference in worm burden is observed. In 364 most studies, IgE or FceRI deficiency is associated with an increased worm burden in the early 365 phase (before 2 weeks) of the infection either during the first exposure (Schistosoma mansoni, Strongyloides venezuelensis or Brugia malayi) or the second exposure (Nippostrongylus 366 brasiliensis, Heligmosomoides polygyrus)¹⁰⁷⁻¹¹². Interestingly, IgE or FceRI deficiency was 367 associated with a reduced specific IgG1 response during the first exposure^{110,112}. The 368 consequence of IgE or FccRI deficiency in repeated exposures to helminths which can be 369 370 modelized by the presence of granuloma in target tissue remains unclear. Though, IgE 371 deficiency seems to be associated with smaller granuloma (i.e smaller inflammatory infiltrate)^{112,113} and FccRI deficiency is associated with increased damages and inflammatory 372 infiltrate in the target tissues ^{108,114}.Furthermore, basophils activation by IgE/worm antigens 373 complexes were essentials for granuloma formation and limit tissue damages^{108,109}. Altogether, 374

those data suggest that IgE response might have different impact in helminthic infection. Upon 375 acute exposure, IgE would play a role in worm clearance 1)thanks to enhancement of specific 376 377 IgG response, 2)FccRI-driven IgE degranulation of effector cells such as mast cells, basophils, platelets and eosinophils ^{106,115}. Upon repeated exposure (i.e, worm clearance failure), specific 378 379 IgE and their interaction with FccRI on basophils would take part in granuloma formation to constrain the worm, avoid target tissue damages¹¹⁶ and tissue healing making an echo to the 380 concept of natural IgE implied in tolerance⁶⁵. Nevertheless, this latter point remains speculative 381 382 since to date no clear role for IgE in wound healing has been documented (Figure 8).

383

IgE in tumor immuno-surveillance – a kind friend?

385 Selective IgE deficiency is defined as total circulating IgE < 2.5 KU/L after excluding common variable immune deficiency¹¹⁷. Recently, three studies demonstrated either retrospectively or 386 387 prospectively a significant association between selective IgE deficiency and the onset of 388 malignant diseases in adult and children. Interestingly, the demographic data of neoplasm 389 diseases were identical to the target population: solid organ cancer in adulthood and hematologic cancer in childhood^{118–120}. These data suggest that IgE-FAP could play a role in 390 tumor immune-surveillance. Indeed, a recent study by Crawford et al demonstrated that DMBA 391 induced skin epithelial carcinogenesis triggered $\gamma\delta T$ cell activation and local self-reactive IgE 392 393 production via IL-4 in mice. Those IgE took part in preventing epithelial carcinogenesis in a $Fc \in RI$ dependent manner⁶⁸. This assumption is supported by several observations in human 394 where non anaphylactic specific IgE immunization against tumors could trigger specific 395 cytotoxic responses in patients with colorectal, pancreatic, prostatic and breast cancer^{121–124} in 396 a FcERI-dependent manner¹²⁵. On the opposite, two recent observations demonstrated a 397 398 detrimental role of IgE in a mice model of skin cancer. More precisely, Taniguchi et al demonstrated that skin tumors were infiltrated with IL-33 induced FccRI+ macrophages that 399

promoted tumor growth suggesting a role of IgE on those tumor-infiltrating macrophages¹²⁶. 400 401 Also, Hayes et al demonstrated that tumor growth was promoted by an increase of polyclonal 402 "natural" IgE after 12-O-tetradecanoylphorbol-13-acetate (TPA) repeated applications. Tumor growth was also dependent on FceRI IgE-FAP on basophils that infiltrated the tumor. 403 Interestingly, selective IgE deficiency in those mice model abrogated tumor growth¹²⁷. 404 405 Altogether those data suggest that IgE would prevent tumor growth in the early phase of 406 carcinogenesis whereas IgE would promote tumor growth at an advanced stage (Figure 9). The 407 distinct IgE repertoires induced by DMBA and TPA (self-reactive versus polyclonal natural 408 respectively) could explain, at least in part, the janus-faced role of IgE in cancer immunopathology despite type 2 immunity induction in both case^{68,127}. In conclusion, further 409 410 researches are needed 1)to consider selective IgE deficiency as a predictive biomarker of cancer onset¹²⁸ and 2) to better characterize the role of IgE at molecular (IgE repertoire, antigen 411 412 identification, receptors) and cellular (CD8+ T cells, NK cells, $\gamma\delta T$ cells) levels in the tumor micro-environmment (enhancing or preventing tumor growth) in order to achieve new add-on 413 therapeutic strategies^{129,130}. 414

415

416 IgE self-reactivity: from autoallergy to autoimmunity – friend or foe?

The concept of autoallergy implies the presence of self-reactive IgE (srIgE) and was introduced 417 by the end of 1950's^{131,132} without any understanding of its role. To date, this concept has 418 419 evolved into IgE autoimmunity with a growing spectrum of autoantigens identified in systemic 420 erythematous lupus (double strand DNA, SSA/B, RNP, nucleosome, acidic ribosomal P2 421 protein among others), chronic urticaria (thyroperoxidase, double strand DNA, IL-24), bullous pemphigoid (BP180 or BP230) and atopic dermatitis (> 140 autoantigens identified) mostly¹³³. 422 srIgE were associated with disease severity in systemic erythematous lupus^{134,135}, bullous 423 pemphigoid¹³⁶ and chronic urticaria¹³⁷. They are probably important in atopic dermatitis 424

pathogenesis since a high percentage of patients have $srIgE^{138,139}$, though the correlation 425 between srIgE and disease severity remains debated requiring further investigations¹³³. The 426 427 cellular and molecular effects of srIgE were mainly deciphered in lupus where they enhance 428 disease activity through two main pathways. On one hand, FceRI-driven srIgE-FAP on plasmacytoid dendritic cells enhance srIgG/srIgE through B cells expansion and plasma cell 429 differentiation in SLO in a TLR9-dependent manner¹³⁴. On the other hand, MHC-II cross-430 431 presentation of autoantigen (double strand DNA) via FceRI-driven srIgE-FAP by activated basophils enhanced autoantibodies and B cell differentiation through type-2 inflammation in 432 SLO in a BAFF-dependent manner¹⁴⁰ (reviewed in¹⁴¹). In bullous pemphigoid, cellular and 433 molecular effect of autoreactive IgE are not that well deciphered¹³³. In chronic urticaria, the 434 435 importance of auto-IgE has been strongly suggested by the great efficiency of omalizumab on disease activity¹⁴² and also by passive anaphylaxis transfer with sera from patients with chronic 436 urticaria¹⁴³. Though, the precise mechanisms remains poorly understood¹³³. In conclusion, the 437 438 data strongly argue for a deleterious implication of srIgE through $Fc \in RI$ -driven FAP. It also 439 suggests that srIgE are of high affinity (indirect CSR) able to cross-link multivalent 440 autoantigens and activate effector cells. Though, how autoreactive IgE and type 2 immunity are 441 triggered leading to complex and different clinical features remains poorly understood.

442

443 Unexpected associations between IgE and diseases

Implication of IgE was suggested in cardiovascular diseases thanks to mice models of aorta
aneurysm where FcɛRI alpha subunit knock-out partially protected from smooth muscle
senescence¹⁴⁴, neutrophils and macrophage infiltration and IL-6 mediated inflammation^{144,145}.
Yet, those results were not translated to human. Another observation suggested that IgE
sensitization against alpha-gal oligosaccharide without any anaphylactic features was
associated with a higher risk of coronary atheromatous disease at a younger age¹⁴⁶ which

precise mechanisms remain to be determined. Altogether, those observations open a whole newfield of IgE biology in cardiovascular homeostasis and disease.

Recently, functional anti-MHC IgE (ie. inducing mast cell degranulation) was detected in mice 452 models of skin and heart allograft serum. The author could also detect anti-HLA IgE in sera 453 from kidney transplanted patients with high titers of donor specific antibodies $(n=5)^{147}$. Another 454 recent study associated antibody-mediated rejection with anti-HLA IgE in the serum and IgE 455 deposit colocalizing with mast cell/basophil in the kidney transplant¹⁴⁸. Altogether, those data 456 457 suggest that the HLA/IgE complex could interact with mast cell/basophils in the transplant and 458 trigger deleterious type 2 immunity. It also questions about 1)the interaction of IgE with other cognate effector cells such as eosinophils¹⁴⁹; 2)the triggers of allo-immune response in solid 459 organ transplantation¹⁰. 460

461

463 CONCLUSION

464 Since its characterization in 1966, IgE has been extensively studied in allergic diseases such as 465 food or drug anaphylaxis, allergic rhinitis and allergic asthma where it has a pivotal role. 466 Furthermore, it would be finalistic to assume a simple pathological role for IgE especially since 467 IgE is only found in mammals⁷. Some evidence argues for a constitutive secretion of IgE in the body fluids with homeostatic or even immune-modulatory properties. However, their precise 468 469 roles and interactions with other immunoglobulins and immune cells remain to be determined. 470 Some evidence also argues for a pro-inflammatory role of IgE when type 2 inflammation is 471 triggered. In that case, IgE could enhance immune response against small amounts of antigen (IgE-FAP) with a wide spectrum of downstream responses modulated either by IgE affinity 472 473 and/or receptor binding (FceRI or FceRII) and/or the target cell (basophils, mast cells, eosinophils or even platelets). However, further studies are needed to decipher how these 474 475 responses are finely tuned. Additionally, breakthrough works have unraveled some 476 physiological roles of IgE such as resistance and detoxification to venom, worm clearance 477 during acute infection, prevention of tissue damage in case of helminthic chronic infection. 478 More surprising, there is a janus-faced role in cancer immunopathology where IgE seems to 479 take part in tumor clearance/immuno-surveillance at an early stage. On the opposite, IgE seems 480 to enhance tumor out-growth at an advance stage suggesting a diversion of IgE response which 481 mechanisms remains unclear to date. Last but not least, IgE was unexpectedly implicated 482 vascular disease or even rejection in solid organ transplantation where immunopathological 483 mechanisms remain to be determined. Altogether, those interesting developments argue for a 484 role of IgE response as an immunological adaptor to complex environmental changes which better characterization will lead to the development of innovating therapeutic strategies. 485

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926 <u>Figure 1:</u> Intrinsic regulation of IgE. Class switch recombination (CSR) to IgE is regulated by 927 the size of the ε acceptor region (S ε) which is smaller than all the others making more difficult 928 to access for transcription factors. Furthermore, CSR can occur either directly from μ to ε or 929 indirectly from μ to γ 1 to ε . In both case, the expression of IgE BCR leads to an increase rate 930 of apoptosis and a lower mobility to germinal centers. This figure was created with 931 BioRender.com.



932

933 Figure 2: Extrinsic regulation of IgE in secondary lymphoid organs. When a naïve B cell 934 commits into IgE CSR (direct or indirect), there are two main mechanisms limiting the 935 appearance of IgE long-live plasma cells. The first one is the presence of IL-10 or neuritin 936 secreted by regulatory follicular helper cells (Tfr). The second mechanism is the negative 937 feedback loop by IgE itself via monomeric CD23 (low affinity receptor) binding on centrocyte 938 or on plasma cell. This figure was created with BioRender.com.



940 <u>Figure 3</u>: Extrinsic regulation of IgE production in peripheral tissue. Under homeostatic 941 condition, IgE binds his high affinity receptor (FccRI) on mast cells and monocyte without 942 antigen leading to a rapid internalization of IgE and its receptor to lysosomes. Under 943 inflammatory condition, basophils/mast cell degranulation due to IgE cross-linking with 944 antigen is constrained 1)by the antigen/IgD complex binding to its receptor (CD44 and galectine 945 9) 2)by the auto-reactive IgG against IgE under its soluble and bound form. This figure was 946 created with BioRender.com.



947

948 Figure 4: Loco-regional IgE secretion at mucosal sites. Loco-regional IgE secretion is a basal
949 trait at mucosal sites in case of impaired type 1 or 17 immunity which can occur in case of 1)
950 abiotic or aberrant microbiota leading to barrier deficiency (i.e., leaky epithelium); 2) TCR
951 signaling or restrained repertoire; 3) tolerance impairment (mainly regulatory T cells); 4)
952 Deficient or aberrant N-glycosylation. This figure was created with BioRender.com.



954 Figure 5: Natural IgE secretion. There are two main hypotheses to explain natural IgE secretion 955 at steady state. 1)The first one consists of constant stimulation of naïve or IgG1 memory B cells 956 with (auto?)antigens without cognate T cells help leading to short live plasma cells. 2)The 957 second one consists as well in constant stimulation of IgG1 memory B cells by (auto?) antigens 958 with Tfh help in presence of IL-4 leading to shot live plasma cells. Those natural IgE could 959 exert an anti-inflammatory role by promoting active mechanisms of peripheral tolerance 960 (tolerogenic dendritic cells, regulatory T cells) but also preventing effector T cell activation. 961 This figure was created with BioRender.com.



963 Figure 6a: The pivotal role IgE facilitated antigen presentation (IgE-FAP) via FcERI. FcERI 964 IgE-FAP has a broad spectrum of effects on basophils (baso) or mast cells ranging from survival 965 in case of crosslinking with a monovalent antigen to degranulation, cytokine secretion or even 966 amplification of type 2 response in case of multivalent antigen cross-linking (indirect 967 presentation by class II major histocompatibility complex, (MHC-II)). FccRI IgE-FAP with a 968 multivalent antigen can also occur on dendritic cells (DC) where it enhances type 2 response 969 thanks to indirect MHC-II presentation of the antigen. This figure was created with 970 BioRender.com.



972 Figure 6b: The pivotal role IgE facilitated antigen presentation (IgE-FAP) via FccRII aka
973 CD23. There are two main effects of trimeric CD23 IgE-FAP with multivalent antigen that can
974 be mediated by B cells and probably by monocytes (Mono). The first one is to boost humoral
975 response with specific IgM and IgG immunization. The second one is to enhance clonal
976 expansion of antigen specific T cells. This figure was created with BioRender.com.





Figure 7: The role of IgE in venom detoxification. In case of venom inoculation, a specific IgG1
and IgE response will arise independently from basophils/mast cells against a small protein
fraction of the venom. These venom specific IgE will lead to basophils/mast-cells degranulation
releasing enzymes such as tryptase, chymase as example in the micro-environment that will
cleave venom proteins with deleterious biological activity. The IgG1 memory against venom
will warrant the specific IgE recall and venom detoxification in case of latter exposures. This
figure was created with BioRender.com.



985

Figure 8: The role of IgE in anti-helminthic immunity. In case of helminthic infection (oral 986 987 route mainly), the role of worm specific IgE is different in case of acute or chronic infection. 988 In case of acute infection, worm specific IgE will lead to basophils/mast cells or eosinophils or platelets degranulation of cytotoxic molecules to kill and expulse worms. In parallel, a specific 989 990 humoral immunity (IgM and IgG) will be boosted by worm specific IgE. In case of chronic 991 infection (i.e., failure of worm clearance), worm specific IgE will take part in the creation of 992 eosinophilic granuloma to constrain the worms and also prevent tissue damage. This figure 993 was created with BioRender.com.



994

995 Figure 9: The janus-faced role of IgE in cancer immunopathology. In the early phase of cancer 996 growth, self-reactive IgE will produced thanks to the help of IL-4 and resident $\gamma\delta$ T cells. Those 997 self-reactive IgE will activate the basophils thanks to FccRI IgE-FAP that will prevent tumor 998 growth and enhance anti-tumoral immunity. On the opposite at an advanced stage of cancer, 999 the pro-inflammatory micro-environment will lead to the local secretion of polyclonal "natural" 1000 IgE that will enhance tumor out-growth thanks to FccRI IgE-FAP on basophils and 1001 macrophages that infiltrate the tumor. This figure was created with BioRender.com.

<u>Table 1:</u> Impact of IgE or FccRI deficiency in mice models of helminthic infections based on
 time after infection, single or repeated exposure, worm burden, egg production and the presence
 of granuloma (i.e. chronic exposure). ND = not determined.

	Name of the parasite	Mice model	Prima ry infecti on	Second ary infectio n	Evaluat ion (time after infectio n	Worm burden	Eggs product ion	Granulo ma (if applicab le)	Referen ces
	Trichinella spiralis	BALB/ C IgE deficie nt mice	Yes	No	1 week to 4 weeks	Increas ed compar ed to WT mice	ND	Increase d compare d to WT mice	107
		SJA/9 mice	Yes	Yes	4 weeks	No differe nce for both infectio n	ND	No differenc e for both infection	150
	Angiostrong ylus costariecensi s	SJA/9 mice	Yes	Yes	6 weeks	No differe nce for both infectio n	ND	No differenc e for both infection	151
	Nippostrong ylus brasiliensis	SJA/9 mice	Yes	Yes	1 week & 2 weeks	No differe nce for both infectio n	ND	No differenc e for both infection	150
		BALB/ C FcεRIα - deficie nt mice	No	Yes	2-5 days	No differe nce for the first infectio n; Decrea sed in the skin & increas ed in the lung	ND	Decrease d number of granulo ma in the skin and larger lung injuries compare d to WT mice.	108

					for the second infectio n compar ed to WT mice.			
	BALB/ C IgE deficie nt mice	No	Yes	2 weeks	ND for the first infectio n; Increas ed number in the small intestin e compar ed to WT mice for the second infectio n	ND	ND	109
	BALB/ C IgE deficie nt mice	No	Yes	2 to 3 weeks	No differe nce	ND	No differenc e	116
Heligmosom oides polygyrus	BALB/ C IgE deficie nt mice	No	Yes	2 weeks	ND for the first infectio n; Increas ed number in the small intestin e compar ed to WT mice for the second	ND	ND	109

					infectio n			
Strongyloide s venezuelensi s	FcɛRI- /- mice	Yes	No	1,5 weeks	Increas ed compar ed to WT mice when FcγR are blocke d	ND	ND	110
Brugia malayi	BALB/ C IgE deficie nt mice	Yes	Yes	2 to 6 weeks	Increas ed compar ed to WT mice in primar y infectio n; No differe nce on second ary infectio n	ND	ND	111
Schistosoma mansoni	129/ter Sv IgE- deficie nt mice	Yes	Yes	8 weeks	Increas ed compar ed to WT mice in primar y infectio n with lower specific IgG1 respons e; No differe nce on	ND	Smaller granulo ma in the liver than WT mice	112

					second ary infectio n			
	BALB/ C FcεRIα - deficie nt mice	Yes	No	8 weeks	No differe nce	No differen ce	Larger granulo ma with higher liver fibrosis than WT mice.	114
	SJA/9 mice	Yes	No	8 weeks	No differe nce for both infectio n	No differen ce	ND	152
Schistosoma japonicum	SJA/9 mice	Yes	No	8 weeks	No differe nce	No differen ce	Smaller granulo ma in the liver than WT mice	113

1009 DISCLOSURE STATMENT:

1010 The authors of this manuscript have no conflicts of interest to disclose as described by Allergy.

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