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inflammation of the synovial membrane, characterized by hyperplasia, neo-angiogenesis, and immune cell infiltration that drives local inflammation and, if untreated, can lead to joint destruction and disability. In

parallel to the well-known clinical heterogeneity, the underlying synovitis can also be significantly heterogeneous. In particular, in about 40% of patients with RA, synovitis is characterized by a dense lymphocytic infiltrate that can acquire the features of fully functional tertiary lymphoid organs (TLO). These structures amplify autoimmunity and inflammation locally associated with worse prognosis and potential implications for treatment response. Here, we will review the current knowledge on TLO in RA, with a focus on their pathogenetic and clinical relevance.

Tertiary Lymphoid Organs in Rheumatoid Arthritis

Felice Rivellese, Elena Pontarini, and Costantino Pitzalis

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Abstract Rheumatoid Arthritis (RA) is a chronic systemic autoimmune disease. 24 RA mainly affects the joints, with inflammation of the synovial membrane, char-25 acterized by hyperplasia, neo-angiogenesis, and immune cell infiltration that drives 26 local inflammation and, if untreated, can lead to joint destruction and disability. In 27 parallel to the well-known clinical heterogeneity, the underlying synovitis can also 28 be significantly heterogeneous. In particular, in about 40% of patients with RA, 29 synovitis is characterized by a dense lymphocytic infiltrate that can acquire the 30 features of fully functional tertiary lymphoid organs (TLO). These structures 31 amplify autoimmunity and inflammation locally associated with worse prognosis 32 and potential implications for treatment response. Here, we will review the current 33 knowledge on TLO in RA, with a focus on their pathogenetic and clinical 34 relevance. 36

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1 Introduction

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Rheumatoid Arthritis (RA) is the most common autoimmune disease, affecting up 38 to 1% of the population worldwide (Smolen et al. 2016). Although RA is well 39 recognized as a systemic disease, its main feature is the chronic inflammation of the 40 synovial membrane, which is characterized by infiltration of immune cells, cellular 41 hyperplasia, and neo-angiogenesis (McInnes and Schett 2011, 2017). Ongoing 42 synovitis and its corresponding clinical features of joint pain and swelling are the 43 main causes of functional disability in patients with RA. Despite the availability of 44 effective medications, in a large proportion of patients, the treatments fail to control 45 the inflammatory response. When un-optimally controlled, synovial inflammation 46 can progress and ultimately lead to joint destruction and permanent disability. Such 47 inconsistent response to treatment has been attributed at least in part to the clinical 48 and physiopathological heterogeneity of RA. In fact, similar to other autoimmune 49 diseases, under the umbrella of RA, we are grouping a diverse spectrum of patients 50 with different clinical features, which are mirrored by significant differences in 51 terms of pathogenesis and, therefore, variable response to targeted treatments. For 52 example, it is well recognized that the positivity for anti-citrullinated protein 53 antibodies (ACPA) identifies a group of patients-around 70%- with a clinical 54 phenotype of highly aggressive and destructive disease (Willemze et al. 2012). In 55 line with its marked clinical heterogeneity, a variable degree of immune cell 56 infiltration has been described in the synovia of RA patients and has been recently 57 linked to distinct clinical features, including disease severity, progression, and 58 treatment response. 59

⁶⁰ 2 The Synovial Membrane as Site of Inflammation in RA

The main physiopathological feature of RA is the inflammation of the synovial 61 membrane (SM). In physiological condition, the SM is a composed by an intimal 62 layer formed of synoviocytes, also known as fibroblast-like synoviocytes (FLS), 63 which are specialized fibroblast-like cells with the main function of producing the 64 synovial fluid that lubricates and nourish the avascular articular surfaces. Below the 65 thin layer of FLS, there is a sub-intimal layer composed by connective tissues, 66 scattered infiltrating macrophage-like cells, and blood vessels. During RA, the 67 synovial membrane undergoes the following changes: (i) infiltration of immune 68 cells, including cells of innate (e.g., macrophages, natural killer [NK] cells, innate 69 lymphoid cells, dendritic cells, mast cells) and adaptive immunity (e.g., B and T 70 lymphocytes, plasma cells); (ii) proliferation of FLS, leading to the thickening of 71 the intimal layer, and (iii) growth of new blood vessels (neo-angiogenesis) which 72

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further sustains the infiltration of immune cells, thus facilitating the perpetuation of 73 the inflammatory response. Despite the enormous advancements in our under-74 standing of the pathogenesis of RA, leading us to recognize a number of genetic 75 and environmental factors contributing to its pathogenesis, the initial trigger of 76 synovial inflammation is currently unknown. Also, we do not know whether the 77 first hit happens directly in the joints or somewhere else, such as the lungs or other 78 organs. However, once the inflammatory response is triggered and gets perpetuated. 79 synovitis represents the main feature of RA, thus the study of synovial inflamma-80 tion is of utmost importance to improve our understanding of RA (Pitzalis et al. 81 2013). 82

⁸³ 2.1 Histological Patterns of Synovial Inflammation

The infiltration of immune cells is one of the main features of RA synovitis. In line 84 with the clinical heterogeneity of the disease, a variable degree of immune cell 85 infiltration in synovia has been described. Despite the complexity and partial 86 overlap of immune cell infiltration, the parallel study of large numbers of synovial 87 samples from patients with early untreated RA (Humby et al. 2019) has allowed to 88 describe three distinct groups based on the patterns of immune cell infiltration in 89 synovia: (1) lympho-myeloid, dominated by lymphoid lineage infiltration (T cells, 90 B cells, plasma cells) in addition to myeloid cells; (2) diffuse-myeloid, with myeloid 91 lineage predominance but poor in B cells/plasma cells and (3) pauci-immune, 92 characterized by scanty immune cells and prevalent stromal cells. Within the 93 lympho-myeloid group, the infiltrating B cells, T cells, and plasma cells often 94 organize into aggregates that resemble the lymphoid follicles of secondary lym-95 phoid organs, acquiring features such as segregation of T cells and B cells, the 96 presence of high endothelial venules (HEVs), and follicular dendritic cells (FDCs) 97 networks. Although TLO can also be detected at extra-articular sites, including the 98 lungs (Barone et al. 2015) and bone marrow (Bugatti et al. 2005) of RA patients, 99 they mainly form within the sublining of the synovial tissue, where they have been 100 described in about 40% of patients with early untreated RA (Pitzalis et al. 2013). 101 A representative example of TLO is offered in Fig. 1a-c, including a schematic 102 representation of their organization in Fig. 1d, with additional details in Fig. 2. In 103 the next paragraphs, we will describe the ontogeny of tertiary lymphoid organs in 104 RA, their functions, and their correlation with clinical features and disease prog-105 nosis, including response to treatment. 106



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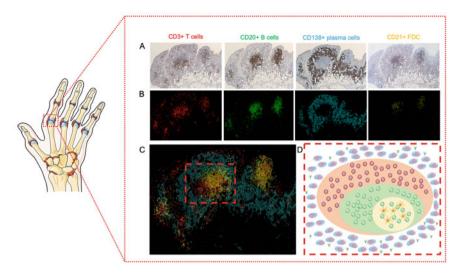


Fig. 1 Tertiary lymphoid organs in synovia. **a** Immunohistochemical staining of synovial membrane, **b** color deconvolution of the images in (**a**), **c** overlap of the above images, and **d** schematic representation of the organization of TLO in synovia, with FDC in yellow, B cells in green, T cells in red, and plasma cells in blue

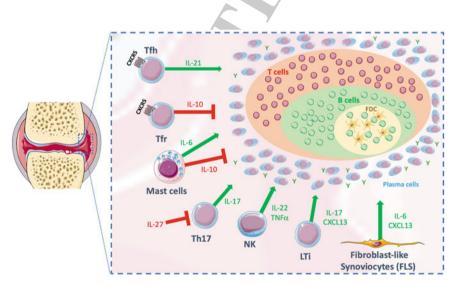


Fig. 2 Schematic representation of synovial TLOs and immune cells contributing to their development. Tfh = T follicular helper cells; Tfr = T follicular regulatory cells, Th17 = T helper cells 17, NK = Natural killer cells; LTi = Lymphoid tissue inducer



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3 Synovial Tertiary Lymphoid Organs in RA

3.1 The Development and Regulation of Synovial Tertiary Lymphoid Organs in RA

110 3.1.1 Chemokine and Lymphotoxin Beta

One of the initial steps in the formation of TLO is the infiltration of lymphoid cells 111 into the synovia, which is driven by the inflammatory milieu produced by FLS and 112 innate immunity cells. As the inflammatory process becomes chronic, however, a 113 number of specific mediators are required for the formation of TLO, such as 114 lymphotoxin- β (LT β), CXCL13, CCL19, and CCL21 (Corsiero et al. 2012). The 115 development of TLO largely mirrors the ontogeny of secondary lymphoid organs, 116 thus most of our knowledge on TLO development is derived from the study of 117 secondary lymphoid organs, where animal models have identified a number of 118 stimuli which are essential for the development of secondary lymphoid organs 119 (Randall et al. 2008; Drayton et al. 2006). Among these stimuli, the primum movens 120 has been recognized to be the production of lymphotoxin- β from so-called lym-121 phoid tissue inducer (LTi) cells (Bar-Ephraim and Mebius 2016), which in turns 122 leads to the production of lymphoid chemokines (CXCL13, CCL19, and CCL21) 123 from lymphoid tissue organizers and mesenchymal cells. Although the presence of 124 many of these lymphogenic stimuli has been confirmed in TLO in rheumatoid 125 synovium (Bugatti et al. 2014; Manzo et al. 2007), the initial trigger of TLO 126 formation in RA has not been identified. Several immune cells have been shown to 127 be a source of lymphoid chemokines, and some of these key cells are represented in 128 Fig. 2b. Among the various mediators, CXCL13 produced by follicular dendritic 129 cells (Takemura et al. 2001) and other immune cells plays a pivotal role in deter-130 mining the spatial organization of TLO, inducing the segregation of B cells within 131 the germinal center, which is an essential drive for affinity maturation and antigenic 132 selection (De Silva and Klein 2015). In line with its pivotal role, serum levels of 133 CXCL13 have been associated with the presence of synovial TLO in patients with 134 RA (Bugatti et al. 2014; Dennis et al. 2014). 135

3.1.2 T Follicular Helper Cells

In recent years, a specialized class of T helper cells, named T follicular helper cells
 (Tfh), has been recognized for their central role in sustaining B cell activation and
 differentiation in the germinal center (GC) reactions in secondary lymphoid organs.
 Tfh cells are specialized T helper cells that upon priming by antigen presenting cells
 (APCs) acquire the expression of CXCR5, the receptor for CXCL13, enabling them
 to migrate into the B cell area of GC.
 The ectopic expression of CXCL13 has been described in RA synovium (Manzo

The ectopic expression of CXCL13 has been described in RA synovium (Manzo
 et al. 2005, 2008) and has been shown to induce TLO formation and recruits B cells

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to non-lymphoid tissues in mice (Luther et al. 2000). In fully formed GC, Tfh cells support somatic hypermutation of auto-reactive B cells and plasmablast generation 146 directly in the diseased tissues mainly through the production of IL-21. The latter is 147 Tfh signature cytokine, known to be a potent cofactor for B cell survival, prolif-148 eration and plasma cell differentiation, in particular in the context of CD40 149 co-stimulation and in synergy with B cell activating factor (Karnell and Ettinger 150 2012: Liu et al. 2015).

Importantly, because of the role of IL-21 and Tfh cells in supporting GC 152 response, they have been implicated in the development of TLO in rheumatic 153 autoimmune diseases, including RA, as represented in Fig. 2. 154

- Data in animal models of arthritis identified a number of Tfh-associated markers 155 during the development of inflammatory arthritis. In particular, CXCR5 has been 156 shown to be an essential factor for the development of inflammatory arthritis: 157 CXCR5-deficient animals or lacking CXCR5 on T cells are resistant to RA, 158 showing impaired GC response (Moschovakis et al. 2017). Also, selective defi-150 ciency in T helper cells of SLAM-associated protein (SAP), required for the B/T 160 cell interaction, thus essential for Tfh differentiation, protects mice from RA, further 161 supporting the pathogenic role of ectopic GC formation (McCausland et al. 2014). 162 In parallel, IL-21 and its receptor are highly expressed in synovial tissue of 163 patients with RA (Jüngel et al. 2004; Kwok et al. 2012), and increased IL-21 164 expression is associated with synovial TLO (Jones et al. 2015). IL-21R 165 up-regulation is mainly described on macrophages and fibroblast with an acti-166 vated phenotype (Jüngel et al. 2004), and IL-21 has been involved in the devel-167 opment of articular damage by promoting both osteoclastogenesis (Kwok et al. 168 2012) and metalloproteinase release by fibroblast-like synoviocytes (Xing et al. 169 2016). Finally, Tfh cells are also enriched in the synovia of patients with RA, while 170 almost absent in osteoarthritis and normal synovium (Penatti et al. 2017; Chu et al. 171 2014). 172
- In addition to conventional CXCR5+ Tfh cells, a population of T helper cells 173 lacking CXCR5 expression and producing CXCL13 has been also described in the 174 synovia of RA patients (Manzo et al. 2008). A recent breakthrough publication has 175 shed new light on these cells, which have been re-named as PD1+ CXCR5-T 176 peripheral helper cells (Tph), since they have been found in the synovia but also in 177 the peripheral blood of patients with RA and their ability to induce the activation of 178 B cells has been confirmed in vitro (Rao et al. 2017). Similar to GC-Tfh, these cells 179 are an important source of CXCL13, support synovial B cell proliferation and 180 activation through IL-21 production and SLAMF5 receptor ligation, and co-localize 181 with B cells in synovial TLO (Rao et al. 2017). Although Tfh and Tph cells share 182 the main markers, the tissue localization, and the ability to support B cell activation, 183 it is unclear whether Tph in RA are Tfh cells with impaired CXCR5 expression, or a 184 more distantly related cell type. Despite the evidence of Tfh and Tph contribution to 185 the pathogenesis of RA, the functional link between these cells and TLO formation 186 remains to be elucidated, as well as their contribution to the local production of 187 autoantibodies within TLO. 188

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Additionally, although the enrichment of Tfh in RA synovium has been well 189 described, there are conflicting data regarding circulating Tfh cell frequency 190 [comprehensively reviewed in (Gensous et al. 2018)]. Some authors reported IL-21 191 directly correlating with the frequency of Tfh-like cells, with IL-21 level and 192 number of Tfh-like cells associated with higher titer of anti-CCP antibodies and 193 disease activity score in RA (Ma et al. 2012). The circulating counterpart shares 194 phenotypic and functional features with tissue Tfh cells, except for the expression 195 of prototypical Tfh transcription factor B cell lymphoma protein 6 (Bcl-6), but their 196 biology is still poorly defined. Data from SAP-deficient mice show how these cells 197 are committed to Tfh lineage and are generated prior the GC response (He et al. 198 2013; Tsai and Yu 2014). Moreover, it is still unclear if circulating Tfh can reflect 199 an ongoing humoral activity. 200

201 3.1.3 Other Pro-inflammatory Cytokines and Cells

It is now clear that a number of other pro-inflammatory cytokines, such as IL-17, IL-21, IL-22, IL-23, and TNF α , are also critical for lymphoid neogenesis in autoimmune diseases (Jones and Jones 2016).

The IL-23–IL-17 pathway has been involved in the initiation and perpetuation of TLO, and several cells of the innate and adaptive immunity are able to produce IL-17. In particular, a subset of adult innate lymphoid cells [type-3 innate lymphoid cells (ILC3 cells)] can produce IL-17 in the initial phases of TLO formation (Sawa et al. 2010). Accordingly, IL-17 positive cells are observed in the proximity of TLO in RA synovia (Chabaud et al. 1999), and the activation of the IL-23–IL-17 pathway correlates with the presence of synovial TLO (Cañete et al. 2011).

Another important aspect is the potential plasticity between other T helper subsets and the Tfh. In fact, several other subsets, including Th17 cells, Th1 and Th2, have been described to acquire Tfh-like phenotype (Ueno et al. 2015). For example, Tfh2 and Tfh17, but not Tfh1, are able to secrete IL-21 and induce naïve B cells to secrete class-switched immunoglobulin (Ig) (Morita et al. 2011).

Within RA synovium, proliferation of fibroblast-like synoviocytes is sustained 217 by IL-22, a cytokine required for the development and maintenance of TLO. IL-22 218 role in ectopic lymphoneogenesis comes from data in experimental models of 219 inducible TLO in salivary glands, mimicking TLO formation in Sjogren's syn-220 drome salivary glands. In this animal model, IL-22 is able to directly induce 221 CXCL13 production in a subset of GP38+ stromal cells through phosphorylation of 222 signal transduced and activator of transcription 3 (STAT3) (Barone et al. 2015). 223 Once lymphocytes are recruited, IL-22, together with LTa1B2, supports also pro-224 liferation of a population of podoplanin (pdpn)-positive stromal cells, 225 over-expressing IL21R, into a network of immunofibroblasts that are able to sup-226 port the earliest phases of TLS establishment (Navar et al. 2019) in the same model. 227 In RA synovium, IL-22 expression and IL-22 receptor on fibroblast-like synovio-228 cytes have been reported (Ikeuchi et al. 2005), suggesting its contribution to the 229 maintenance of TLO. In particular, IL-22 expression is increased in cells expressing 230

the long isoform of complement receptor type 2 (Cr2, also known as CD21) (Cañete 231 et al. 2011), usually present in networks of stromal-derived follicular dendritic cells 232 (FDCs), that contribute to the presentation of immune complexes necessary to 233 generate activated B cells, in TLO. In synovial tissue, IL-22 is also produced by NK 234 cells (Zhu et al. 2015). NK cells are innate immune lymphocytes with cytolytic and 235 immune-regulatory activities representing a significant proportion (8-25%) of 236 immune infiltration in synovial fluid of RA patients, identified in the joints in the 237 early stage of RA development (Tak et al. 1994). Initially, NK cells were described 238 in RA pathogenesis for their production of cytotoxic serin protease granzyme-A and 239 B and pro-inflammatory cytokines, such as IL-1 and TNF α as dominant mediators 240 of proliferative synovitis in RA (Klimiuk et al. 1997), supporting ocleoclastogen-241 esis and thus involved in the development of articular damage (Kotake et al. 2001). 242 In fact, increased production of IFNy and TNFa characterizes synovial fluid NK 243 cells of erosive RA patients with joint damage in comparison with non-erosive RA 244 (Yamin et al. 2019). Recent evidence suggests that NK cells may support TLO 245 maintenance within RA synovium as a subset of NK cells expressing a natural 246 cytotoxicity receptor NKp44 which is able to produce IL-22 (Zhu et al. 2015). 247 NKp44+ NK cells are enriched in both peripheral blood and synovium of RA 248 patients secreting IL-22 and TNFa, which in vitro studies showed to support 249 RA FLS proliferation (Ren et al. 2011), through the activation of STAT3 pathway 250 (Zhu et al. 2015). IL-22 induced proliferation of synovial fibroblast, an effect that 251 was inhibited by neutralizing antibodies targeting IL-22 and TNFa (Ren et al. 252 2011). Thus, NK cells may participate in TLO organization supporting the prolif-253 eration of synovial fibroblasts responsible for the local secretion of chemoattractant 254 molecules and, as consequence, lymphocytes recruitment. 255

In addition to cells of the adaptive immunity, many other innate immunity cells
 and the stromal compartment have been shown to contribute to the development of
 synovial TLO (Barone et al. 2016).

Fibroblast-like synoviocytes (FLS), for example, have been shown to produce the T cell/dendritic cell chemoattractant CCL21 (Manzo et al. 2007) and express CXCL12 and IL-7, involved in immune cell retention and lymphoid-like microanatomical organization (Timmer et al. 2007; Bradfield et al. 2003).

Recently, we have also shown a strong association between synovial mast cells 263 (MCs) and the presence of TLO in a large cohort of patients with early RA 264 (Rivellese et al. 2018). MCs were also found to induce B cell activation and 265 differentiation in vitro, including the production of ACPA autoantibodies. Finally, 266 in animal models of inducible TLO (IL27R knockout), we confirmed the associa-267 tion of MCs with TLO. Overall, this points out to the relevance of MCs as potential 268 contributors to the formation of TLO, although additional studies are needed to 269 confirm their functional relevance (Rivellese et al. 2017, 2019b). 270



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3.1.4 Negative Regulators of TLO Including Tfr

In addition to the mediators and pathways acting as positive regulators of TLO, several cells and cytokines have been characterized as negative regulators of TLO development.

For example, IL-27, an heterodimeric cytokines part of the IL-12 family 275 (Yoshida and Hunter 2015), has been recently identified as a negative regulator of 276 TLO. In fact, animals with knockout of the IL27Ra develop a severe form of 277 antigen-induced arthritis, including the development of synovial TLO (Jones et al. 278 2015). Importantly, synovial TLO are not normally produced in animal models of 279 arthritis; thus, the identification of these structures in IL-27R α knockout animals 280 points to the relevance of IL-27 as a regulator of TLO development. Accordingly, in 281 patients with RA, IL-27 was found to be inversely correlated with TLO and with 282 TLO-related gene signatures. Finally, both in clinical and experimental arthritis, 283 synovial TLO coincided with an increased local expression of cytokines and 284 transcription factors of the Th17 and T follicular helper (Tfh) cell lineages, where 285 IL-27 is able to inhibit the differentiation of Th17 cells, in line with previous 286 evidence (Stumhofer et al. 2006). 287

As local counterpart of the circulating T regulatory cells, T follicular regulatory cells (Tfr) have been recently described within GCs, including GCs in TLO. Tfr cells are able to prevent the differentiation of auto-reactive B cells (Wu et al. 2016; Botta et al. 2017), by regulating Tfh cells, but also by directly inhibiting B cell activation (Wing et al. 2014).

Although the relevance of Tfr cells in the regulation of GCs in animal models is well established (Linterman et al. 2011), the involvement of Tfr cells in human autoimmune disease, including RA, is still unclear.

Several studies have reported decreased levels of Tfr in patients with active RA 296 and, accordingly, negative correlations with autoantibodies and disease activity 297 (Romão et al. 2018; Niu et al. 2018). On the other hand, increased levels of Tfr 298 were found in patients who were in remission (Liu et al. 2018). Using animal 299 models of autoimmunity with spontaneous development of GCs, IL-21 was shown 300 to induce an unbalance between Tfh and Tfr, increasing the formation of GCs, 301 while administration of Tfr was able to restore Tfh:Tfr ration and suppress GC 302 responses (Ding et al. 2014). 303

Another group found that the resolution of collagen-induced arthritis following administration of intravenous immunoglobulins was accompanied by an increase of Tfr cells (Lee et al. 2014). Taken together, this suggests that the reduction of circulating Tfr cells is associated to the development of RA and that restoration of Tfr cells could potentially improve autoimmune responses.

In line with this, monitoring the ratio between Tfr and Tfr could be useful in patients with RA, as confirmed by a several observations (Niu et al. 2018; Wang et al. 2019).

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As for the function of Tfr in RA, these cells have been shown to have sup-312 pressive effects in vitro, which were enhanced in patients in remission (Liu et al. 2018). However, it has also been speculated that Tfr in autoimmune diseases might 314 be functionally deficient (Fonseca et al. 2017). 315

3.2 The Function of Tertiary Lymphoid Organs in RA 316

As TLO mirrors secondary lymphoid organs in their ontogeny and maturation, it is 317 expected that they also recapitulate the main functions of secondary lymphoid 318 organs, which is supporting germinal centers (GC) reaction toward maturation of B 319 cells and antibody production. 320

Within a considerable proportion of TLO forming in rheumatoid synovium, 321 ectopic GC reactions take place similar to secondary lymphoid organs (Bombardieri 322 et al. 2017). Many of RA-associated autoantibodies are high affinity IgG (e.g., 323 ACPA) (van Delft and Huizinga 2020), and B cells forming TLO are auto-reactive 324 and somatically mutated (Humby et al. 2009), indicating the involvement of a GC 325 response in RA progression. Indeed, TLO in RA synovium can display functional 326 features of germinal centers, like the expression of the enzyme activation-induced 327 cytidine deaminase (AID) involved in in situ B cell affinity maturation and clonal 328 selection (Humby et al. 2009). 329

Accordingly, the analysis of B cells isolated from the synovia of patients with 330 RA has confirmed the generation of synovial plasma cells from locally activated B 331 cells (Scheel et al. 2011), and the local production of class-switched autoantibodies 332 in rheumatoid synovium has been demonstrated (Humby et al. 2009). Also, we 333 have recently demonstrated that the presence of synovial TLO in early untreated 334 RA is associated with autoantibody positivity (Humby et al. 2019). Interestingly, 335 this is in contrast with previous data that failed to show an association between TLO 336 and autoantibody positivity (Thurlings et al. 2008). Recently, comparing two large 337 cohorts of patients with early and established RA, we were able to confirm the 338 strong association between TLO and autoantibody positivity in early RA that could 339 not be observed in established RA, thus explaining the previous findings, possibly 340 because of treatment effect or other biases from long-standing diseases (Rivellese 341 et al. 2019a). 342

Importantly, the initiation of a germinal center reaction requires antigen pre-343 sentation to B cells. In RA, the aberrant immune response against citrullinated 344 proteins culminating in the production of anti-citrullinated protein antibodies 345 (ACPA) is well recognized as a key pathogenetic feature (Derksen et al. 2017). 346

Accordingly, citrullinated proteins have been described in the synovia of RA 347 patients (Baeten et al. 2001) together with PAD enzymes, which are responsible for 348 citrullination (De Rycke et al. 2005). The specificity of synovial citrullinated pro-349 tein has been challenged (Vossenaar et al. 2004), but this does not come as a 350 surprise since citrullination and other post-translational modifications of proteins 351 are recognized as physiological processes (Trouw et al. 2017). On the contrary, the 352

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aberrant immune response to modified proteins represents the hallmark of RA, and
accordingly, the local production of ACPA in synovia has been confirmed (Humby
et al. 2009; Amara et al. 2013; Masson-Bessière et al. 2000). Finally, several groups
have been able to isolate ACPA-producing B cell clones from the synovia and
synovial fluid of patients with RA (Germar et al. 2019; Corsiero et al. 2016, 2018).

358 **3.3** The Clinical Relevance of Tertiary Lymphoid Organs 359 in RA

360 3.3.1 TLO and Disease Severity

Early studies on the analysis of synovial membrane relied on the use arthroscopy to 361 obtain synovial samples. These analyses pointed out a marked heterogeneity in 362 terms of synovial inflammation, particularly in the degree of immune cell infiltra-363 tion, with the description of aggregates of lymphoid cells in a proportion of patients. 364 However, when looking for an association with clinical features, these studies 365 vielded contradictory results: some found an association of lymphoid aggregates 366 with disease severity and autoantibody positivity (Humby et al. 2019; Bugatti et al. 367 2014; Orr et al. 2017) and others did not (Thurlings et al. 2008; Cantaert et al. 2008; 368 Van De Sande et al. 2011) (Table 1). These inconsistencies could be explained by a 369 number of biases: (i) the exclusive analysis of large joints, in which there can be 370 commonly overlapping osteoarthritis and are not the most representative of the 371 inflammatory process in RA (Linn-Rasker et al. 2007) (ii) the inclusion of patients 372 with long-standing disease, with the obvious bias of treatment and disease duration, 373 and (iii) the lack of a gold standard for the histological assessment of immune cell 374 infiltration (Humby et al. 2016). 375

The development of minimally invasive techniques such as ultrasound-guided 376 synovial biopsies has overcome most of these limitations, as it made possible to 377 obtain synovial tissues from small joints of a large cohort of patients with early RA 378 and, very importantly, prior to treatment star. Thus, it is not surprising that the 379 recently published analyses on this cohort highlighted a strong association with 380 disease severity and autoantibody positivity (Humby et al. 2019). Interestingly, a 381 direct comparison of early and established RA, using a validated semi-quantitative 382 score for the assessment of B cells, showed that while in early RA the presence of B 383 cell-rich synovitis was associated with disease severity, this was not the case in 384 established RA, possibly explaining the discrepancies from previous studies ana-385 lyzing patient with different disease duration Rivellese et al. (2019a). 386

When analyzing exclusively patients with early untreated RA, our group has recently shown that patients with a synovial lympho-myeloid pathotype, characterized by the presence of B and T cell aggregates, have significantly higher disease severity, autoantibody positivity, and baseline erosive load (Humby et al. 2019). Furthermore, molecular analyses showed that myeloid- and lymphoid-associated Table 1 Association of TLO with disease severity and clinical phenotype in RA

Author and	References	Population	Joints	Treatment	Time	Analyses	Results
year			biopsied	(if any)	points		
			and				
			proceedie				
Van	Van	57 RA	Knee	N.a.	Biopsy at	IHC	ACPA + patients showed higher mean
Oosterhout	Oosterhout		arthroscopy		time 0		number of infiltrating lymphocytes and
2008	et al. (2008)						higher rate of local joint destruction
Van de Sande	Van De Sande	93 (24 RA)	Knee	sDMARDs	Biopsy at	IHC	Lymphoid neogenesis present in 36% of all
	et al. (2011)		arthroscopy		0 (93) and		patients, associated with the degree of
					6 months		synovial inflammation, but not specific of
					(17)		RA. No relationship between the presence
							of lymphocyte aggregates at baseline and
							definitive diagnosis or clinical outcome
							after follow-up
De Hair 2013	De Hair et al.	55 seropositive	Knee	N.a.	Biopsy at	IHC	CD3 T cell numbers in the biopsy tissue
	(2013)	individuals	arthroscopy		time 0		showed a borderline association with
		without clinical					subsequent development of clinically
		evidence of					manifest arthritis. CD8 T cells were
		arthritis					associated with ACPA positivity
Gómez-Puerta	Gómez-Puerta	83 RA	Knee	N.a.	at	IHC	No significant differences in clinical
2013	et al. (2013)		arthroscopy		time 0		variables, acute phase reactants, synovial
							cell infiltrate or lymphoid neogenesis
						1	(LN) between ACPA positive and negative
							patients
Orr 2017	Orr et al.	123 RA	Knee	sDMARDs	Biopsy at	IHC	ACPA + RA patients were characterized by
	(2017)		arthroscopy	and	time 0		significantly higher levels of CD19+ B cells
				bDMARDs			and CD3+ and CD8+ T cells. Levels of
							lymphoid aggregates of CD19+ B cells and
							serum CXCL13 levels were significantly

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Author and year References Population Joints biopsied Ti year biopsied (i) humby 2019 Humby et al. 144 early US-guided si Humby 2019 Humby et al. (<1 year) US-guided si KA RA RA biopsy si Lliso-Ribera Lliso-Ribera 2009 with uS-guided si Lliso-Ribera Lliso-Ribera 2009 with biopsy si		(
Humby et al. (2019) (2019) (<1 year) (<1 year)	uthor and	References	Population	Joints	Treatment	Time	Analyses	Results
Humby et al. (2019) (2019) treatment naïve RA Lliso-Ribera et al. (2019) with with synovial biopsy synovial biopsy synovial biopsy arthritis biopsy	ear	~		biopsied	(if any)	points		
Humby et al. (2019) (<1 year) (<1 ye				procedure				
Humby et al. 144 early US-guided (2019) (<1 year) biopsy reatment naïve RA RA RA Lliso-Ribera 200 early patients US-guided with biopsy arthritis biopsy								higher in ACPA + patients. EULAR
Humby et al. 144 early US-guided (2019) (<1 year) biopsy reatment naïve RA RA Lliso-Ribera 200 early patients US-guided with synovial inflammatory biopsy arthritis								response was significantly associated with
Humby et al. 144 early US-guided (2019) (<1 year) synovial treatment naïve biopsy RA RA Lliso-Ribera 200 early patients US-guided with synovial inflammatory biopsy arthritis								the level of CD3+ T cell infiltrates, while
Humby et al. 144 early US-guided (2019) (<1 year) synovial treatment naïve biopsy RA RA Lliso-Ribera 200 early patients US-guided with synovial inflammatory biopsy arthritis								CD68+ macrophage and CD8+ T cell
Humby et al.144 carly (2019)US-guided synovial treatment naïve RAUS-guided synovial biopsy BALliso-Ribera200 carly patients with arthritisUS-guided synovial biopsy								levels were predictive of the response to
Humby et al.144 earlyUS-guided(2019)(<1 year)								tumor necrosis factor inhibitors
(2019) (<1 year) synovial treatment naïve biopsy RA RA biopsy BA biopsy and treatment naïve biopsy biopsy arthritis synovial biopsy arthritis	fumby 2019	Humby et al.	144 early	US-guided	SDMARDs	Biopsy at	IHC and	Patients with a lympho-myeloid pathotype
Lliso-Ribera 200 early patients biopsy RA BA biopsy Lliso-Ribera 200 early patients US-guided with synovial inflammatory biopsy arthritis		(2019)	(<1 year)	synovial		time 0	nanostring	have significantly higher disease severity,
RA Lliso-Ribera 200 early patients US-guided et al. (2019) with synovial inflammatory biopsy arthritis				biopsy	,			autoantibody positivity, and baseline
Lliso-Ribera 200 early patients US-guided et al. (2019) with synovial inflammatory biopsy arthritis								erosive load. Myeloid- and
Lliso-Ribera 200 early patients US-guided et al. (2019) with synovial inflammatory biopsy arthritis								lymphoid-associated gene expression
Lliso-Ribera 200 early patients US-guided et al. (2019) with synovial inflammatory biopsy arthritis								strongly correlated with disease activity and
Lliso-Ribera 200 early patients US-guided et al. (2019) with synovial inflammatory biopsy arthritis								acute phase reactants
et al. (2019) with inflammatory arthritis	liso-Ribera	Lliso-Ribera		US-guided	sDMARDs Biopsy at		IHC	Patients fulfilling the 1987 RA criteria had
	019	et al. (2019)	with	synovial		time 0		significantly higher levels of disease
arthritis				biopsy				activity, histological synovitis, degree of
			arthritis					immune cell infiltration, and differential
								upregulation of genes involved in B and T
							7	cell activation/function compared with RA
								2010 criteria or UA, which shared similar
								clinical and pathobiological feature

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genes strongly correlate with disease activity and acute phase reactants. Another
 more recent publication in early RA has further highlighted the value of synovial
 tissue analyses in refining the diagnosis of RA vs undifferentiated arthritis.

Moreover, deep phenotyping of synovial tissue by molecular analyses has 395 identified specific gene signatures associated with clinical phenotype. In particular, 396 for example, peripheral blood interferon response genes were associated with the 397 lymhpo-myeloid pathotype, while synovial plasma cell signature was associated 398 with progression of structural damage (Lewis et al. 2019). Additional analyses from 399 the Accelerating Medicine Partners group, by integrating single cell RNA 400 sequencing and mass cytometry, have recently identified unique cell population 401 expanded in RA synovia that allow to distinguish the degree of synovial inflam-402 mation (Rao et al. 2017; Zhang et al. 2019). Specific cell populations included HY1 403 (CD90) +HLA-DRAhi sublining fibroblasts, IL1B+ pro-inflammatory monocytes, 404 ITGAX + TBX21 + autoimmune-associated B cells, and PDCD1+ peripheral 405 helper T (TPH) cells and follicular helper T (TFH) cells. The latter, in particular, are 406 essential for the formation of TLO and have been already discussed in the previous 407 paragraph. However, to date, little is known about the association of these cell types 408 with disease features, such as disease severity, progression, and response to treat-409 ment. In the near future, it will be of utmost importance to confirm the relevance of 410 these immune populations, by studying their association with clinical phenotype in 411 larger cohorts of patients with RA. 412

413 **3.3.2 TLO as Direct Therapeutic Targets**

Because of their well-established relevance in driving the pathogenesis of RA and
their association with worse disease outcomes, several strategies aiming at targeting
TLO in RA have also been tested.

A number of studies have attempted to target mediators that are relevant in the 417 formation or maintenance of TLO. The modulation of the IL-21/IL-21R pathway as 418 a treatment strategy was first tested in experimental models of RA. IL-21R defi-419 ciency in the K/BxN mouse model of inflammatory arthritis (Kim et al. 2009) and 420 antigen-induced arthritis (Roeleveld et al. 2017) is sufficient to block RA initiation, 421 while the blockade of the IL-21/IL-21R pathway ameliorates disease in 422 collagen-induced arthritis models treated with murine IL-21R Fc fusion protein 423 (Young et al. 2007). However, there are still no data in patients with RA on the 424 blockade of IL-21/IL-21R. 425

Some other molecules have been already tested in patients, but results have not 426 been particularly striking, as in the case of inhibiting $LT\beta$, which did not show 427 clinical efficacy (Bienkowska et al. 2014). Similarly, drugs inhibiting IL-17 and 428 IL-12/IL-23 showed little or no differences compared with placebo in RA 429 (Kerschbaumer et al. 2019). This is in contrast with data on seronegative arthritis, 430 where inhibition of IL-17 and its axis proved to be extremely effective, although it 431 has been suggested that the analysis of targeted expression of these molecules could 432 potentially help in predicting treatment response (Boutet et al. 2018). 433

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Importantly, none of the above studies targeting mediators involved in TLO formation or maintenance in RA has stratified patients on the basis of TLO presence, which could have helped in selecting patients with higher chances of response.

3.3.3 TLO as Predictors of Treatment Response

As highlighted in the previous paragraphs, the presence of TLO is able to identify a 439 subset of RA patients with a specific disease phenotype, specifically higher disease 440 activity and higher prevalence of autoantibodies. Therefore, it is plausible to 441 hypothesize that the presence of TLO could help to predict treatment response. 442 A number of studies have explored the analyses of synovial tissues to predict 443 treatment response. However, relatively few included the systematic analysis of 444 TLO. Futhermore, because of the relatively small number of patients, the incon-445 sistency in the definition of TLO, and the use of different time points for repeated 446 biopsy, most of the results are fragmented and difficult to interpret. 447

Canete et al., for example, demonstrated significantly lower response in patients who were TLO positive despite a significantly higher use of anti-TNFa agents. (Cañete et al. 2009) By linear regression, TLO positive were found to predict lack of response to anti-TNFa. In this study, however, patients started sequential treatment with escalation to anti-TNFa in non-responders, and therefore, there could have been a selection of TLO + patients as the most severe, thus non-responders.

On the contrary, Klaasen et al., by analyzing synovial samples obtained before
 and after standardized treatment with infliximab in a cohort of 97 patients, found
 that the presence of TLO at baseline was a highly significant predictor of the
 clinical response to anti-TNF treatment (Klaasen et al. 2009).

More recently, Dennis et al. provided the molecular confirmation of the histological pathotypes previously described by histology. In addition, by analyzing the data from a previous cohort undergoing treatment with infliximab, they were able to identify TLO signature as predictor of response to TNFi (Dennis et al. 2014). The limitation of this manuscript consisted in the analysis of synovial samples obtained from arthroplasty, thus without standardization of treatment.

The observations published from our early RA cohort allowed to overcome such 464 limitations and have shown a reduction of lymphoid-associated genes in EULAR 465 good responders to csDMARDs (Humby et al. 2019). Similarly, molecular analyses 466 by RNAseq identified a number of cell modules, including B cells, in association 467 with B response to csDMARDs (Lewis et al. 2019). Importantly, these data come 468 from the analysis of synovial tissue obtained by US-guided synovial biopsies in 469 untreated patients with early Rheumatoid Arthritis, thus eliminating the bias of 470 long-standing disease, treatment or the exclusive inclusion of large joints in studies 471 based on arthroscopy. 472

In recent years, continuing on the same line, two international consortia have
 driven the delivery of the first two large-scale biopsy-driven RCTs in Rheumatoid
 Arthritis. As part of a study funded by the UK National Institute of Health

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Research, a randomized, open labeled study in anti-TNFa inadequate responders to 476 investigate the mechanisms for Response-resistance to rituximab versus tocili-477 zumab in RA (R4-RA), a total of 165 patients failing treatment with TNFi have 478 been recruited. Promising preliminary results were presented at the ACR 2019, 479 while the trial is currently being analyzed and final results will be soon published. 480 Similarly, as part of the MRC and versus arthritis-funded consortium maximizing 481 therapeutic utility in RA (MATURA), the stratifying therapies for rheumatoid 482 arthritis by pathobiology (STRAP) RCT has enrolled a total of 226 patients who 483 failed csDMARDs and is due to being completed in the last quarter of 2020. 484

These studies have been appropriately powered and thus will hopefully give clear answers on the utility of synovial biopsy analysis in predicting treatment response in RA. Specifically, the studies aimed at understanding if patients lacking synovial B cells have a lower response to B cell targeted treatment (Rituximab) as opposed to other treatments. At the same time, the studies will provide invaluable information to answer additional research questions, including the association of TLO with disease severity, progression, and treatment response.

492 4 Conclusions

Here, we offered a comprehensive review on the relevance of synovial TLO in RA. 493 The data presented indicate that the ontogeny of TLO resembles the development of 494 secondary lymphoid organs, since many of the mediators known to be involved in 495 lymphoneogenesis have been identified in the synotia of RA patients. Importantly, 496 these structures are fully functional, as they induce the local maturation of B cells 497 toward the production of autoantibodies. Their presence has been described in 498 about 40% of patients with RA from early disease stages and has been strongly 499 associated with disease severity and progression. Despite the availability of several 500 drug treatments that can directly or indirectly target TLO and their components, a 501 stratified medicine approach is needed to fully appreciate the potential effect of such 502 treatments. 503

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