1. Introduction

Sex hormones are implicated in immune response, with estrogens acting as enhancers [10]. Estrogens not only have anti-inflammatory but also pro-inflammatory roles depending upon different influencing factors [66]. Metabolism of estrogen within the body is a complex and important subject. Estrone and estradiol are biochemically interconvertable and yield the same family of estrogen metabolites. The metabolism of estrogen takes place primarily in the liver through Phase I (hydroxylation) and Phase II (methylation, glucuronidation, and sulfation) pathways with ultimate excretion in the urine and feces [26].

Cytochrome P-450 enzymes mediate the hydroxylation of estradiol and estrone, which is the major Phase I metabolic pathway for endogenous estrogens. Several extrahepatic target tissues or cultured cells from target tissue express estrogen-hydroxylating enzymes activities [28]. Each cytochrome P-450 favors the hydroxylation of specific carbons, altogether, these enzymes can hydroxylate virtually all carbons in the molecule, with the exception of the inaccessible carbons. Different functional groups produced by the action of P-450 at the specific sites of steroid nucleus markedly effects the biological properties of different estrogen metabolites. For example, different hydroxylation reaction give estrogenic or carcinogenic metabolites. Functionally, the important reactions catalyzed by cytochrome P-450 are at carbon number 2, 4 and 16.

In this chapter, we venture into the area of the ambiguous relationship between catecholestrogens (CEs) and rheumatoid arthritis (RA). This analysis is focused in a part on the possible role of CEs in the pathogenesis of RA and on exploring the mechanism behind the generation of autoantibodies taken consideration the role of CEs. It also explains a unique hypothesis, which showed that CEs formed in various tissues undergo oxidative metabolism (enzymatic or non-enzymatic) to produce reactive oxygen species (ROS), which could
modified DNA and alter it immunogenicity. This would lead to the induction and elevated levels of RA autoantibodies. In addition, chapter further explain the role of CEs in antigen-driven induction of RA autoantibodies that may unfold various aspects of RA.

2. Catecholestrogens (CEs)

Catecholestrogens (CEs) are biological active metabolites of estrogen which are synthesized by estrogen 2- and 4- hydroxylase in the liver, brain and other organ [29]. They are produced by the actions of genes encoded by CYP1A1, CYP1A2 and CYP1B1, which is an estrogen 4-hydroxylase (Figure 1).

![Figure 1. Pathway of estrogen metabolism including estradiol, estrone and various estrogen metabolites (including CEs).](image)

CEs also yield potent genotoxic molecules implicated in carcinogenesis. 4-Hydroxyestrone/estradiol was found to be carcinogenic in the male Syrian golden hamster kidney tumor model, whereas 2-hydroxylated metabolites were without activity [46]. 4-Hydroxyestrogen can be oxidized to quinone intermediates that react with purine base of DNA, resulting in depurinating adduct that generates highly mutagenic apurinic sites. Quinones derived from 2-hydroxyestrogens produced stable DNA adducts and are presumed to be less genotoxic [67]. As described elsewhere, estrogen may be oxidized by hepatic cytochrome P-450 enzymes to hydroxyl CEs and further oxidized to the semiquinone and quinone form [1]. The quinone formed from oxidation of 4-hydroxyestrogen has half-life of 12 minutes as compared with the short life of 2-hydroxyestrogen (t<sub>1/2</sub> = 47 s) [30]. Estrone and estradiol are oxidized to lesser amount to 2-hydroxycatechols by CYP3A4 in liver and by CYP1A in extrahepatic tissues or to 4-hydroxycatechols by CYP1B1 in extra-
hepatic sites, with the 2-hydroxycatechol being formed to a larger extent. Catechol-o-methyl transferases (COMTs) inactivated 2- and 4-hydroxycatechols to their respective methyl derivatives [69]. COMT, a phase II enzyme, catalyses the transfer of methyl group from S-adenosyl methionine to hydroxyl groups of a number of catechol substrates, including the catecholestrogens. CEs under normal condition are o-methylated by COMT to form 2- and 4-o-methylethers, which are excreted in the urine [56]. Small amount of CEs may also be converted by peroxidase-catalysed reactions to semiquinones or quinones that are capable of forming DNA adducts or generating ROS via redox cycling that could oxidize DNA bases [69]. Estrogen metabolites have indirect and direct genotoxicity. Hydroxylation is an important elimination step for estrogen to generate CEs. Microsomal estradiol hydroxylation in human breast cancer showed significantly higher 4-hydroxy:2-hydroxyestradiol ratio in tumor tissue than in adjacent normal breast tissue [48], but the breast cancer tissue samples contained fourfold higher levels of 4-hydroxyestradiol (4-OHE2) than in normal tissue from benign breast biopsies [57].

It has been observed that intra-tissue concentration of estrogens, hydroxyestrogens (16 α-hydroxyestrone, 2-hydroxyestrone, 2-hydroxyestradiol, 4-hydroxyestrone, and 4-hydroxyestradiol) and methoxyestrogens (2-methoxyestrone, 2-methoxyestradiol, 4-methoxyestrone, and 4-methoxyestradiol) in normal and malignant breast showed the highest concentration of 4-hydroxyestradiol in malignant tissue [4]. The concentration, as evaluated by combined high performance liquid chromatography (HPLC) and gas chromatography-mass spectrometry (GC-MS) was found to be more than twice as high as that of any other compound. CEs have been measured in rat brain and various endocrine tissues by sensitive radioenzymatic assay. The concentration of these CEs in the hypothalamus and pituitary are at least ten times higher than parent estrogen. CEs have potent endocrine effect i.e. they have an important role in neuroendocrine regulation [54].

In addition, brain is capable of 2-hydroxylation and consequently forms CEs from estrone and estradiol. Increased availability of estrogen and estradiol for binding and hypothalamic sites would facilitate formation of CEs. 2-Hydroxylated and 4-hydroxylated estrogen exerts biologic responses on the cyclic secretion of lutenising hormone (LH) perhaps even on follicle stimulating hormone (FSH) and prolactin. Catecholestradiol competes with estradiol for estrogen binding sites in the anterior pituitary gland and hypothalamus and dopamine binding sites on anterior pituitary membranes [41].

Mitogenicity associated with estrogen receptor (ER)-mediated cellular events was believed to be the mechanism by which estrogens contributed to carcinogenesis [59]. The role of CEs as genotoxic chemical procarcinogenesis, independent of ER mediation, has been well recognized [70]. Although oxidative metabolism of estrogen to CEs is generally thought to terminate the estrogenic signal, but CEs retain some binding affinity to the ER. Exposure of MCF-7 cells with 2- and 4-hydroxyestradiol increased the rate of cell proliferation and the expression of estrogen-inducible gene such as progesterone receptor (PR) gene and pS2. Compare to estradiol, 2- and 4-hydroxyestradiol increased proliferation rate, level of PR protein as well as pS2 mRNA expression to certain fold [62].
As described previously, estrogens are metabolized via two pathways, i.e., formation of CEs and to a lesser extent, 16α-hydroxylation. The catechols are formed as 2- and 4-hydroxylated estrogen. These two CEs can be inactivated by the enzyme COMT through α-methylation [69]. Other possible mechanisms of inactivation include conjugation of CEs by glucuronidation and sulphation. High concentration of 4-hydroxylated metabolites caused insufficient production of methyl, glucuronide or sulphate conjugate which in turn results in CE toxicity in cell and consequently competitive catalytic oxidation to semiquinone (CE-SQ) and quinone (CE-Q). CE-SQ and CE-Q may conjugate with glutathione (GSH), catalyse by S-transferase. CE-Q may also react with DNA to form stable and de-purinating adduct if this inactivating process is incomplete [47].

3. CEs and DNA

Endogenous estrogens can become carcinogenic via formation of catecholestrogen quinones, which react with DNA to form specific estrogen-DNA adducts. The mutations resulting from these adducts can cause cell transformation and initiation of breast cancer [58]. The 4-hydroxyestrogen generates free radicals from reductive-oxidative cycling with the corresponding CE-SQ/CE-Q forms, thus causing DNA damage extensively. In comparison to 4-hydroxyestrogen, 2-hydroxyestrogen is not carcinogenic and has potent inhibitory effect on the growth of tumor cells and on angiogenesis [17]. The DNA adducts generated by 2-hydroxyestrogen and 4-hydroxyestrogen can form stable modifications that remain in the DNA unless they are removed by repair. On the other hand, the modification bases can be released from DNA by destabilization of the glycosydic bond and result in the formation of depurinated or depyrimidinated sites [7].

Earlier studies from our lab showed that CE modified DNA brings about single and double strand breaks, hyperchromicity, damage at the restriction sites and modification of different bases (Table 1) [40, 36, 37, 38]. It has been observed that CE caused more damage to the DNA in presence of copper in comparison to nitric oxide (NO) as the extent of base modification was greatest for thymine followed by guanine, adenine and cytosine [40]. We proposed that CE and NO formed in different tissue may react with each other to produce CE-Q.

The quinone/semiquinone redox system produces superoxide ions (O$_2^-$) which can react with NO to form peroxynitrite, which could cause DNA damage [38]. Quenching studies of DNA modified with CEs in presence of C(II) showed that catalase and bathocuproine strongly (~ 80%) inhibited the modification by CEs plus Cu(II) indicating the involvement of H$_2$O$_2$ and Cu(I) during the modification [36, 37]. These results demonstrate that CEs lead to the production of potent ROS, capable of causing DNA damage, thus playing important role not only in carcinogenesis [36] but also in systemic lupus erythematosus (SLE) [40, 37, 39] and RA [38].
<table>
<thead>
<tr>
<th>Parameter</th>
<th>Native DNA</th>
<th>CE-modified DNA</th>
</tr>
</thead>
<tbody>
<tr>
<td>Hyperchromicity</td>
<td>—</td>
<td>32.2</td>
</tr>
<tr>
<td>Absorbance ratio ($A_{260}/A_{280}$)</td>
<td>1.86</td>
<td>1.52</td>
</tr>
<tr>
<td>Melting temperature ($T_m$) °C</td>
<td>88</td>
<td>80.5</td>
</tr>
<tr>
<td>S1 nuclease action</td>
<td>No digestion</td>
<td>Digestion</td>
</tr>
</tbody>
</table>

**Restriction enzymes**

<table>
<thead>
<tr>
<th>Restriction enzyme</th>
<th>Digestion</th>
<th>No digestion</th>
</tr>
</thead>
<tbody>
<tr>
<td>PvuI (CGAT↓CG)</td>
<td>Digestion</td>
<td>No digestion</td>
</tr>
<tr>
<td>PvuII (CAG↓CTG)</td>
<td>Digestion</td>
<td>No digestion</td>
</tr>
<tr>
<td>SspI (AAT↓ATT)</td>
<td>Digestion</td>
<td>Digestion</td>
</tr>
</tbody>
</table>

**Circular dichroism (mdeg) Wavelength (nm)**

<table>
<thead>
<tr>
<th>Wavelength (nm)</th>
<th>+274</th>
<th>−246</th>
<th>−210</th>
</tr>
</thead>
<tbody>
<tr>
<td>mdeg</td>
<td>0.77</td>
<td>−1.3</td>
<td>−0.85</td>
</tr>
<tr>
<td>mdeg</td>
<td>0.52</td>
<td>−7.60</td>
<td>−0.58</td>
</tr>
</tbody>
</table>

**Scavengers**

<table>
<thead>
<tr>
<th>Scavenger</th>
<th>inhibition of modification</th>
</tr>
</thead>
<tbody>
<tr>
<td>Catalase</td>
<td>82</td>
</tr>
<tr>
<td>Sodium azide</td>
<td>21</td>
</tr>
<tr>
<td>Bathocuproine</td>
<td>80</td>
</tr>
<tr>
<td>Ascorbic acid</td>
<td>35</td>
</tr>
</tbody>
</table>

**Base modification**

<table>
<thead>
<tr>
<th>Base</th>
<th>—</th>
<th>16.9%</th>
</tr>
</thead>
<tbody>
<tr>
<td>Adenine</td>
<td></td>
<td>45.5%</td>
</tr>
<tr>
<td>Guanine</td>
<td></td>
<td>12.6%</td>
</tr>
<tr>
<td>Cytosine</td>
<td></td>
<td>47.1%</td>
</tr>
<tr>
<td>Thymine</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

Table 1. Characterization of Native and CE-modified DNA under identical experimental conditions. (Adapted from [36]).

Over the last 2 decades, only two studies have been reported that explain the presence of CE adducts in human breast tissue. First study involves the analysis of CE adducts in the DNA from malignant breast tumor. DNA sample showed the presence of deoxyguanosine adducts of 4-hydroxyestradiol and 4-hydroxyoestrone in almost every samples [21]. Other study showed the presence of CE quinone-derived DNA adduct in two breast samples that were collected from one women with and one woman without breast cancer [51]. The catechol quinone-derived adducts identified were 4-hydroxyestradiol-1-N3-adenosine, 4-hydroxyestrone-1-N3-adenosine and 4-hydroxyestradiol-1-N7-guanine. Animal experimentation involves the injection of 4-hydroxyestradiol (or estradiol-3,4-quinone) into the mammary glands of female ACI rats which resulted in the formation of the depurinating adducts, 4-
hydroxyestradiol-1-N3-adenosine and 4-hydroxyestradiol-1-N7-guanine. [47]. In addition to these adducts, 4-hydroxyestradiol-GSH conjugates was also detected. Another study demonstrates the presence of 4-hydroxy catecholestrogen conjugates with GSH or its hydrolytic product (cysteine and N-acetylcysteine) both in tumors and hyperplastic mammary tissues in ERKO/Wnt-1 mice [18]. Estradiol-3,4-quinone reacted rapidly to form 4-hydroxyestradiol-1-N3-adenosine adducts that are considered to be as depurinating adducts.

It has been shown that CEs can induce DNA lesions not only in plasmid DNA [40, 36, 37, 38] but also in calf thymus DNA [49]. In short, quinone intermediates produced by oxidation of the CE 4-hydroxyestradiol or 4-hydroxyestrone may react with purine bases of DNA to form depurinating adducts that generate highly mutagenic apurinic sites. In contrast, quinone of 2-hydroxylated estrogen produce less harmful and stable DNA adducts [7]. The CE metabolites may also produced potentially mutagenic oxygen radicals by metabolic redox cycling that can also caused DNA damage. The abasic sites resulting from the spontaneous depurination-depyrimidination of the modified bases and the aldehydic base and sugar lesions resulting from the oxidative damage to deoxyribose moieties in the DNA molecules [49], disturb the structural integrity of DNA and destabilize the double helix. This would alter it property and render DNA immunogenic.

4. CEs and Immune Response

Despite a large number of reports that may explain the role of estrogen (or estradiol) in modulation of immune response [10, 66], the role of CEs in the immune response is lacking. It has been observed that the conversion from 2-hydroxyestrogens to 4-hydroxyestrogen might be an additional pro-inflammatory signal because these endogenous metabolites can be oxidized to 3,4-quinone, which lead to depurination and mutation [49]. The 3,4-semiquinone and quinone produce ROS, which stimulate inflammation, whereas methylation and sulfation are detoxification pathways, which involve the release of detoxifying agents in the urine [56]. Other pro-inflammatory role of 4- hydroxyestrogen can be explain by the fact that 4-hydroxyestradiol undergoes 2-electron oxidation to quinone via semiquinone and during this process, ROS can be generated to cause DNA damage and cell death [8]. A similar effect was observed in our studies that explained extensive DNA damage and base modification [40, 36, 37, 38, 39]. In contrast to the pro-oxidant activities of 4-hydroxyestradiol, 2-hydroxyestradiol is a antioxidant that show no carcinogenic activities and has potent inhibitory effect on the growth of tumor cells and on angiogenesis [17]. Thus, 4-hydroxyestradiol in comparison to 2-hydroxyestradiol induce carcinogenesis and have pro-inflammatory effects [20]. But now, we are unaware of the fact that whether pro-inflammatory mediators induce a shift from 2-hydroxyestrogen to 4- and 16-hydroxylated estrogens. From all these studies there is a clear evidence that 2-methoxyestradiol is a pro-apoptotic and cytostatic endogenous compound, which can inhibit not only angiogenesis [17] but also inflammation in animal models. In contrast, 4-hydroxylated estrogens might exert pro-inflammatory roles by inducing ROS and DNA damage. Just like conversion from 2-hydroxyestrogens to 4-hydroxy-
yestrogens, a shift to 16 α-hydroxylated forms of estrogen can be an important pro-inflammatory and pro-proliferative signal [66].

The direct role of CEs in mediating immune response can be explain in our studies that demonstrate the production of antibodies to CE-modified plasmid DNA [40, 36, 37, 39, 38]. As reported earlier, CE-modified DNA (CE-DNA) was found to be highly immunogenic (> 1:12800) in animal model [36]. These antibodies were effectively used as probe for detecting oxidative lesions in human genomic DNA as well as for the estimation of 8-hydroxy-2’-deoxyguanosine (8-OHG) level in the urine of cancer patients [36] and RA patients [38]. Antibodies evaluated from competition ELISA showed strong specificity for the CE-DNA (immunogen) causing about 92 % inhibition in antibody binding at 20 μg/ml of the immunogen (competitor) concentration. Fifty percent inhibition was observed at 3.5 μg/ml of the immunogen. In addition, these antibodies showed a very high degree of binding toward 8-OHG that may explain the role of ROS in mediating various immune responses including inflammation. CE-modified form of different bases (adenine, guanine, cytosine, thymine), human and calf thymus DNA, ROS-DNA, poly nucleotides (dT, dA, dC, dG, dA-dT) were also recognized by these antibodies. Another evidence that explain the role of CEs in mediating immune responses comes from our previous study that showed that CE-DNA was not only recognized highly by cancer autoantibodies [36] but also by anti-DNA autoantibodies in SLE [40, 37] and RA [38]. Results from all these studies demonstrate that CEs somehow produce some antigenic stimulus that may contribute to the generation of antibodies which may directly or indirectly affect the immune system.

Dual pro-inflammatory and anti-inflammatory role of estrogens has been described earlier [66], which depend on the immune stimulus, subsequent antigen-specific immune response, the cell types involved during different disease phase, the specific microenvironment, timing of estrogen administration in relation to the disease course, the concentration of estrogen, the variability in expression of estrogen receptor depending on the microenvironment and the cell type and intracellular metabolism of estrogen leading to important biologically active metabolites (CEs) with quite different anti- and pro-inflammatory function. In addition, one of the striking property of estrogen is its ability to differentiate T and B cells, increase immunoglobulin production and increase immune complex-mediated disease such as SLE [9]. It has been found that women, who are diagnosed with breast cancer, have significantly elevated anti-HMdU autoantibodies [23]. This study is in agreement with our previous finding that explains the presence of circulating antibodies against CE-modified DNA in cancer patients [36]. The modified DNA presents unique epitopes which may be one of the factors for autoantibody induction in cancer. The presence of high levels of these antibodies proves to be the pro-oxidant conditions that have led to the oxidation of bases in genomic DNA and have stimulate an immune response. Since native DNA is a known weak immunogen, it appear that DNA damage by CE render it immunogenic leading to the induction of cancer autoantibodies. It has been suggested that the oxidative DNA damage and the immunologic responses it evokes (autoantibodies) start occurring not only in the carcinogenic process [36] but also in various autoimmune diseases [40, 36, 37, 39, 38]. Estrogen also modulate the function of PMNs (poly morphonuclear leukocytes, neutrophils, granulocytes),
which in turn cause the production of ROS on their stimulation. ROS generated by PMNs cause DNA modification as well as oxidation of protein and lipid peroxidation. It is interesting to note that 2-hydroxylated estrogens works as powerful inhibitors of PMNs activity, which shows one of the protective property of the 2-hydroxylated CEs. Therefore, estrogens (and estrogen metabolites like CEs) affect immunological responses and in turn, their activities are affected by the immunological products. In addition, estrogen in physiological concentration serves to enhance immune response by inducing the production of various types of cytokines and interleukins.

5. CEs and Cytokines

The sex hormones are likely to directly modulate the function of cells involved in the immune response [42]. The pro- and anti-inflammatory effects of estrogen and their metabolites (CEs) on secretion of pro-inflammatory cytokines have been a matter of debate for two or three decades. In fact, at physiological dose, estradiol induce IL-1α, a cytokine that can initiate a cascade of other cytokines, chemotactic and growth factors [12]. It is interesting to note that estradiol also inhibits IL-1α-induced IL-6 production. Therefore, estradiol not only results in increased human epithelial cell proliferation (a process important in tumor growth) but also inhibits the activity of natural killer cells, thus allowing tumor growth [52].

Estrogen and their metabolites (such as CEs) stabilized or increased cytokine secretion whereas testosterone inhibited this secretion [32]. This study explains that down stream estrogen, 2-hydroxyestradiol (2-OHE₂), 4-OHE₂ and 16-hydroxyestradiol (16-OHE₂) did not stimulate TNF secretion. However, the combination of 16-OHE₂ and 2-OHE₂ or 4-OHE₂ markedly stimulated TNF secretion that was observed in presence of cortisol. Therefore, at physiological concentration, estradiol and a combination of downstream estrogens (like 2-OHE₂, 4-OHE₂ and 16-OHE₂) increased (or stabilized) immune stimuli-induced TNF secretion. In addition, estradiol also caused the production of IL-6, IL-2, IFNγ, IL-4 and IL-10, caused by the pro-inflammatory influence of applied immune stimuli [32]. The study of Jaenele et al, 2006,] demonstrates that ratio of 16-hydroxylated estrogen in relation to 2-/4-hydroxylated estrogen is important for TNF secretion. It has been found that the ratio of 10:1 of 16-OHE₂ in relation to 2-/4- hydroxylated estrogen markedly stimulated TNF secretion in presence of cortisol (Figure 2).

In addition, in absence of cortisol, the combination of 16-hydroxylated and 2-hydroxylated estrogens even strongly inhibited TNF secretion. During chronic inflammatory conditions, administration of therapeutic estrogens increased 16-hydroxylated estrogens. In addition, during chronic inflammation in the situation of rheumatic diseases (example RA) the balance is switch to estrogen and particularly to 16-hydroxylated estrogen. In conclusion, one should assume that therapeutic administration of estrogen would enhance 16-OHE₂ over 2-/4-hydroxylated estrogens, which would support secretion of pro-inflammatory cytokines.
It has been observed that estradiol increased IgG and IgM production by peripheral blood mononuclear cells (PBMC) in patients of SLE. This led to elevated levels of polyclonal IgG (including IgG anti-dsDNA) by enhancing B-cell activity by way of IL-10 [33]. It should be important to check these results in the presence of 16 α-hydroxyestrone (16-OHE1) and the naturally occurring 2-hydroxylated CE. Previous study shown that decrease activity in SLE patients correlated negatively with urinary concentration of 2-hydroxylated estrogens [68]. Interestingly, estradiol increase IL-10 production by monocytes and exogenous IL-10 further enhances estradiol-induced increase in antibody production by B cells. Therefore, in SLE patients, estradiol increases IL-10 production by patient’s monocytes and exogenous IL-10 acts additively with estradiol to increase antibody production of B cells. It is hypothesized that estradiol-estradiol receptor complexes may enhance the transcription of IL-10 genes by binding to estrogen-responsive elements or by binding to some transcription factor that directly bind those elements [24].

High doses of estrogen (1500-2000 pg/ml) administration to non-overiectomized mice ameliorated disease systems and significantly decreased the percentage of TNF α or IFNγ producing CNS antigen-specific T Cells in both the CNS and spleen [31]. However, similar studies in mice lacking IFNγ, IL-4 and IL-10 genes provide evidence that these cytokines may not be necessary for estradiol-mediated improvement of disease severity, discrediting the notion that an estrogen-induced shift towards Th 2 cytokine involved [31]. Estrogen also stimulates secretion of IL-4, -5, -6 and -10 by T_{H2} lymphocytes. These cytokines are potent stimulators of B-cell proliferation, maturation into plasma cells and synthesis of antibody. Interleukins 4, 5, 6 and 10 are expressed in greater amount in an estrogen-dominant hormonal milieu [53]. SLE patients have increased level of IL-6 and IL-10, product of T_{H2} lymphocytes and macrophages, which can be directly correlated with clinical disease severity [53]. Patients with lupus also have an inherent defect in interferon gamma production. Under normal condition, increased estrogen leads to an increase interferon gamma concentration but in SLE patients, the estrogen-associated increase in interferon gamma secretion is absent.

Figure 2. Effect of estrogen and their catechol metabolites (CEs) on the secretion of pro-inflammatory cytokines in RA.
In addition, interesting changes of serum estrogens that correlated with cytokine variation have been found during pregnancy in SLE patients [19]. The local effect of sex hormones in autoimmune rheumatoid disease (RA) seem to consist mainly of altered cell proliferation (i.e. estrogen enhance) and cytokine production.

6. Reminding the concept of rheumatoid arthritis (RA)

Rheumatoid arthritis (RA) is an autoimmune disease whose aetiology and progression are multifactorial, including a range of immune, neuroendocrine and psychosocial variable [14]. How these variable interact with one another and how they ultimately influence the disease process in RA is only partially know. A growing body of evidence showed that the stress system and its interaction with the immune system, play a vital role in RA [14]. RA is more prevalent among women than men and 80 % of the total cases occurring between the age of 35 and 50 [34]. The primary presenting symptoms are pain, stiffness and swelling of the joints resulting in impaired physical function. Synovial inflammation underlines the cardinal manifestations of this disease, which include pain, swelling and tenderness followed by cartilage destruction, bone erosion and subsequent joint deformities. In RA, joint involvement is typically symmetric, a characteristic usually not found in other forms of arthritis [50]. Early theories on the pathogenesis of RA focused on autoantibodies and immune complexes, T cell-mediated antigen-specific responses, T cell-independent cytokine network and aggressive tumor-like behavior of synovium have also been implicated [22]. Recently the contribution of autoantibodies in RA has come in the front line and specific therapeutic interventions can be designed to suppress synovial inflammation and joint destruction in RA [22].

RFs have been the hallmark autoantibody found in RA [71]. Autoantibodies to the major cartilaginous collagen have been found in the sera of some RA patients. In addition, cross-reactive natural autoantibodies (IgM) have been described in the sera of RA patients [25]. These antibodies are predominantly directed against histone moieties or against double-stranded DNA. Anti-ds DNA has also been reported following treatment of RA with IFNγ. Beside that, autoantibodies have been found in the sera of RA patients that showed preferentially high binding of CE-modified DNA [38]. In addition, the antibodies have been shown to represent a alternative immunochemical probe to detect oxidative lesions in DNA (genomic) as well as for the estimation of 8-OHdG level in different body fluid of RA patients [38], which may be used as marker in the diagnosis of the disease. Many autoantibody system that could participate in inflammatory joint disease are now recognized in RA, including rheumatoid factor, type II collagen, immunoglobulin heavy gene binding protein, heat shock proteins and hn RNP-33 [64]. Anti-collagen antibody and rheumatoid factor are produced by rheumatoid synovial B cells. The ability of ubiquitous antigen (glucose-6-phosphoisomerase) to induce synovial inflammation is probably related to their adherence to the cartilage surfaces. The presentation of immobilized antigen-antibody complexes on cartilage provides an exceptionally good substrate for complement fixation, similar to rheumatoid factor embedded in rheumatoid cartilage.
Macrophage-derived cytokines (like tumor necrosis factor alpha, TNF α) know to play an important role in the induction and perpetuation of the chronic inflammatory processes in rheumatoid joints as well as in the systemic manifestation of this disease [27]. Over production of this cytokines in the joints of RA patients triggers increase in synoviocyte proliferation and a cascade of secondary mediators involved in the recruitment of inflammatory cells and in the process of joint destruction [3]. It has been found that increase inflammatory markers are responsible for disease progression and joint destruction in early RA. Infect the rate of cartilage and joint destruction is correlated with plasma elevation in inflammatory acute phase reactants (C-reactive protein and vascular endothelial growth factor) and in the synovial concentration of matrix metalloproteinase, matrix digesting enzymes directly responsible for joint damage [2].

In recent years the treatment of RA has undergone somewhat of a revolution, with a strong consensus emerging in favor of early, aggressive therapy [63]. There is growing evidence that early treatment of the disease has a beneficial impact on treatment outcome. The goal to be achieved in managing RA are to explain the underline mechanism behind the generation of antibodies to cartilaginous collagen as well as other oxidative stress condition generated in the synovial fluid, which may be used as marker in the early diagnosis of the disease.

7. Role of CEs in the aetiopathogenesis of RA

Estrogen and their CE metabolites know to play an important role in autoimmune rheumatic diseases [38, 68, 15]. A role for estrogen in the pathogenesis of RA has been review for the last few years [66, 70, 38, 12, 32, 68, 15] but the exact patho-aetiology remains elusive. The evidences concerning the possibility of CEs in the development of RA are very limited and preliminary. It has been observed that the ratio of 16-OHE1/4-OHE2 in synovial fluid (SF) of RA was found to be significantly higher than control [6]. In addition, SF level of 4-OHE2 was significantly increased in RA patients compared to controls [5]. Interestingly, urinary concentration of 2-hydroxylated estrogen were 10 times lower in patients with RA than in healthy individuals but the ratio of 16-OHE1/2-hydroxylated was 20 times in RA patients as compared to control [68]. These finding suggests that the magnitude of conversion to the mitogenic 16-OHE1 is extremely upregulated in RA which most likely contributes to the maintenance of the disease. Furthermore, peripheral estrogen hydroxylation was found to be increased in both men and women with SLE and estrogenic metabolites were reported to increase B cell differentiation and activate T cells [44].

Both RA and SLE are associated with an altered sex hormone balance characterized by lower amount of immunosuppressive androgens and higher amount of immune enhancing estrogens [15]. Increased estrogen to androgen ratios have been observed in RA patient’s SF because of increased aromatase expression by inflammatory macrophages infiltrating synovial tissue [5]. Increased estrogen concentration in RA SF from patients of both sex likely results from the pro-inflammatory cytokines TNFα, IL-1β and IL-6, which accelerate the metabolic conversion of estrogen from androgen by inducing the synovial tissue aroma-
tase (Figure 3). As a result, increased estrogen level might exert activating effects on synovial cell proliferation including macrophages and fibroblast [35]. Interestingly, renal excretion studies showed that the enhanced estrogen metabolism observed in both RA and SLE patients resulted in elevation of pro-inflammatory metabolites derived from estrone or estradiol such as 16-OHE, which exert mitogenic effects and may induce proliferation of synovial fibroblast [68, 13]. In addition, RA SF showed reduced amounts of anti-inflammatory estrogens metabolites such as 2-hydroxyestrogen that inhibits the growth promoting effects of estrogen. In contrast, 16-OHE metabolites considered as enhancer of cell proliferation. In RA synovial tissue, biological effects of estrogen metabolites as a consequence of altered peripheral sex hormone synthesis mainly results in stimulation of cell proliferation and cytokine production [11].

Figure 3. Immune modulation by estrogen and their metabolites in the synovial tissue of RA patients.

It has been observed that RA synovial cells mainly produce the cell pro-proliferation 16-OHE, which in addition to 16-OHE, is the downstream estrogen metabolites that interferes with monocyte proliferation. In addition, urinary concentration and total urinary loss of 2-hydroxyestrogen was found to be 10 times higher in healthy subjects compared to RA [11]. Peripheral metabolic conversion of androgens to potent metabolites of estrogen that promote immune cell proliferation and activation may have role in the female predisposition to autoimmunity.

Estrogen represent a risk factor for the development of autoimmunity and therefore, their therapeutic use must be avoided in patients with active immune-mediated diseases specially RA [16]. In addition to estrogen, various CE metabolites know to increase the risk for the development of RA disease. The increased estrogen concentration has been observed in RA SF, where the hydroxylated forms make the major concentration in particular 16-OHE and 4-OHE whereas, the 2-OHE was found to be similar to the controls [6, 5]. As describe early, 16-OHE is a mitogenic and proliferative endogenous hormone which is responsible for the
proliferation of synovial fibroblast [13]. 16-OHE\textsubscript{1} is formed from upstream estrone and estradiol, which show biological responses by covalent linkage to the receptor [61]. The other conversion product of estrone and estradiol are the 2-hydroxylated estrogens such as 2-OHE\textsubscript{2} and 2-OHE\textsubscript{3}, which would classified as CE. In comparison to 16 hydroxylated estrogens, the 2-hydroxylated forms of estrogen inhibit growth-promoting effects of estradiol [61]. In one of the previous study, it has been observed that urinary levels of 2-hydroxylated estrogens were found 10 times lower in RA patients in comparison to normal subjects, whereas the ratio 16-OHE/2-hydroxyestrogens was found to be 20 times higher in RA patients than in healthy controls [68]. Decreased loss of 2-hydroxylated estrogen in relation to the mitogenic 16-OHE\textsubscript{1} might provide important stimulus to support proliferative state of the synovial cells in RA. Therefore, concentrated-related conversion to pro- and anti-inflammatory down stream metabolites of estrogen might support dual role of estrogen (pro- or anti-inflammatory) [13].

Estrogen at physiological concentration serves to enhance immune responses that may act as important driving force for human humoral immunity [65]. Estradiol increase IL-10 production by monocytes and exogenous IL-10 further enhance estradiol-induced increase in antibody production by B cells. In addition, estradiol enhances IgG and IgM production by PBMC, which leads to elevated levels of polyclonal IgG, including IgG anti-DNA in PBMC of SLE patients by enhancing B cell activity via interleukin 10 [33]. These studies are in agreement with our previous studies that describe the role of CEs in the etiopathogenesis of SLE, cancer and RA [40, 36, 37, 38, 39]. To have a better insight into the possible role of CEs in the aetiopathogenesis of RA, we have demonstrated that CE-modified DNA was highly recognized not only by SLE IgG [40, 37, 39] and cancer IgG [36] but also by RA IgG [38], pointing out the possible participation of modified DNA in the pathogenesis of various autoimmune diseases including RA, as it has been reported that CE-modified DNA bases cause DNA strand breakage and adduct formation in vivo and in vitro [7]. CE-modified DNA showed preferentially high binding with RA sera when compare with nDNA (p < 0.001). These studies clearly indicate that the modified DNA is an effective inhibitor showing substantial difference in the recognition of CE-modified DNA over the native form [38]. The results also demonstrate the pro-oxidant condition which is generated due to the oxidative metabolism of estrogen [40, 36]. Non-enzymatic oxidation of CE in the presence of NO [40] or Cu(II) [37] cause DNA modification and enhanced binding of these modified antigens to RA autoantibodies. To further confirm the role of CEs in RA, we evaluated the binding of autoantibodies by quantitative precipitin titration. The apparent association constant clearly indicates better recognition of CE-modified DNA over native by RA autoantibodies. The enhanced recognition of CE-modified DNA by RA IgG indicates possible participation of modified DNA in RA pathogenesis. The spontaneous production of autoantibodies in RA might arise as a consequence of antigenic change in DNA. Therefore, it could be possible that CE-modified bases of DNA might be one of the contributing factors towards the production of autoantibodies. However, the pro-inflammatory role of estrogen cannot be ruled out [65]. It might be possible that estrogen caused T and B cell differentiation and increase immunoglobulin production. These autoantibodies might be strongly bound with CE-modified DNA than native polymer. Further more, DNA isolated from sera and SF of different
RA patients caused appreciable inhibition in the activity of anti-CE-modified DNA antibodies clearly showing the presence of oxidatively altered epitopes on the isolated DNA molecules, which could be directly correlated to estrogen/radical mediated oxidative stress [38]. These studies show that hydroxylated estrogens (specially CEs) might have a role in the aetio-pathogenesis of RA and other autoimmune diseases.

Various CE metabolites (4-OHE$_2$, 4-OHE$_3$, 2-OHE$_2$, 2-OHE$_1$) known to secret various cytokines from human peripheral blood leukocytes in presence of cortisol [32]. It has been demonstrated that the ratio of 16-hydroxylated estrogen in relation to 2-/4-hydroxylated estrogen is important for TNF secretion. The study showed that the ratio of 10:1 of 16-OHE$_2$ in relation to 2-/4-hydroxylated estrogens stimulate TNF secretion in the presence of cortisol otherwise this effect was not observed in absence of this hormone. Furthermore, absence of cortisol strongly inhibited TNF secretion if the combination of 16-OHE$_1$ and 2-hydroxylated estrogens were used in this experiment [32]. In addition, it has been demonstrated that estrogens are able to enhance secretion of matrix metalloproteinase and IL-1β-induced IL-6 secretion from human fibroblast-like synoviocytes in RA [35]. However, with respect to serum levels of estrogen in RA patients, they are not changed which is in strict contrast to androgen levels [13]. High estrogen concentration has been found in particular in SF of RA patients. The presence in the RA SF of an altered sex hormone balance resulting in lower immunosuppressive androgen and higher immunoenhancing estrogens might determine a favourable condition for the development of the immune-mediated RA synovitis. The appropriated explanation for high concentration of estrogen (and their CE metabolites) can be originated from the study that showed that the inflammatory cytokines (TNF-α, IL-1, IL-6) are particularly increased in RA synovitis and stimulate aromatase activity in peripheral tissues [55]. Therefore, enzyme complex is responsible for the peripheral conversion of androgen to estrogen (estrone and estradiol). In addition, a significant correlation was found between the aromatase activity and IL-6 production and aromatase has also been found significantly in synoviocytes [45]. Therefore, the increased aromatase activity induced by locally produced inflammatory cytokines (TNF-α, IL-1, IL-6) might explain the altered balance resulting in lower androgens and higher estrogen in RA SF as well as their effects on synovial cells [15, 6]. In conclusion, an increase in 16-hydroxyestrogen relative to the sum of all 2- and 4-hydroxylated estrogens (CE) must be viewed as a pro-inflammatory signal, which is particularly evident in RA patients.

Elevated serum concentration of 16-OHE$_1$ have been described in patients with SLE [43], indicating that abnormal pattern of estradiol metabolism may lead to increased estrogenic activity. Interestingly, similar phenomenon was observed in the synovial fluid of RA patients, where 16-OHE$_1$/4-OHE$_2$ was found to be significantly higher compared with control fluid [6, 5]. 4-Hydroxylated estrogens were found to be mitogenic and thus have pro-inflammatory effect [66, 68]. Importantly, all 2- and 4-hydroxylated estrogen demonstrated TNF-inhibitory effect, which was not observed with 16 α-hydroxylated estrogen. Thus the observed preference of 16 α-hydroxylated over 2-/4-hydroxylated estrogens must be considered an important pro-inflammatory signal, which occur particularly in RA patients. This study also describes methylation of hydroxylated estrogen metabolites (CEs) in addition to hydroxyla-
Catecholestrogens in Rheumatoid Arthritis (RA): Hidden Role

7. Conclusion

Remarkable progress have been made in understanding the role of estrogen in the etiology of RA but the role of CEs in RA is lacking. Catechol metabolites known to play an important role in RA but the exact patho-aetiology remain elusive. The evidence concerning the possibility of CEs in the development of RA is very limited and preliminary. It has been observed that oxidative reactions, often catalyzed by isoforms of the cytochrome P-450, can result in the formation of CEs from parent estrogen and subsequently, semiquinones and quinones derived from CEs, are capable of forming either stable or depurinating DNA adducts. Oxidation of CEs also leads to high amount of ROS that can generate extensive DNA damage. This would probably alter its immunogenicity leading to the induction and elevated levels of RA autoantibodies (Figure 4). Therefore, it is possible that the CE-modified bases of DNA might be one of the contributing factors towards production of SLE autoantibodies. In addition, estrogen not only induces DNA damage but also modulate immune response and immune mediated disease. Estrogen was found to increase IgG and IgM from PBMC, which let to elevated level of polyclonal IgG including IgG anti-ds DNA by enhancing B cell activity via IL-10. These autoantibodies could be strongly bound to DNA and serve as an immunological marker for the diagnosis of diseases. Estrogen is thought to play both pro- and anti-inflammatory role in chronic inflammatory diseases that were found to be related to low and high concentration. Increased estrogen to androgen ratio have been observed in RA patients SF, perhaps due to increase aromatase expression by inflammatory macrophages infiltrating synovial tissues. The discovery of high concentration in SF from RA patients of both sexes can also be explain by the fact that inflammatory cytokines (i.e. TNFα, IL-1, IL-6) are increased in RA synovitis and can markedly stimulate aromatase activity in peripheral tissue. But renal excretion studies showed that enhanced estrogen metabolism observed in RA resulted in elevation of pro-inflammatory metabolites derived from estrone and estradiol such as 16-OHE_1 (or 4-OHE_2 4-OHE_1) which exert mitogenic effect on different synovial fibroblast. In contrast, RA SF have reduced amount of anti-inflammatory estrogen metabolites such as 2-hydroxyestrogen that inhibit the growth-promoting effect of estradiol. An unwanted shift from 2- to 4-hydroxyestrogens might be an additional pro-inflammatory signal because these 4-hydroxylated metabolites can be converted to 3,4-quinone, which lead to depurination and mutation in DNA. In conclusion, an increase in 16-OHE_1 relative to the sum of all 2- and 4-hydroxylated estrogens must be considered as pro-inflammatory signal which is particularly evident in RA [60].
In light of these data, the review literature leads to the conclusion that CE (and other metabolites) might have a role in the pathogenesis of RA. Based on the preliminary report, it appears that the production of antibodies in RA involves the generation of IgG against CE-modified DNA through oxidative induced DNA damage. Results imply that oxidative damage to DNA alter it immunogenicity leading to the induction and elevated levels of RA antibodies. RA is a chronic, systemic, inflammatory condition that is characterized by increased production of inflammatory cytokines as well as alteration in estrogen metabolites (CEs). Only by understanding the complex interaction between various estrogen metabolites and mechanism of autoantibodies production, we will able to generate therapeutic interventions that can be designed to suppress synovial inflammation in RA.

Acknowledgements

Authors like to thank Prof. Farhatullah for his valuable contribution to the editing of this manuscript.

Author details

Wahid Ali Khan* and Mohd. Wajid Ali Khan

*Address all correspondence to: wahidalikhan@rediffmail.com
1 Department of Clinical Biochemistry, College of Medicine, King Khalid University, Kingdom of Saudi Arabia

2 Institute of Infection and Immunity, School of Medicine, Cardiff University, United Kingdom

References


