## Thymus and T cells

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#### 1. Introduction

Evidence for intimate interconnections between the three major systems of cell communication, the nervous, endocrine and immune systems, has opened important novel research perspectives. Neuroimmune-endocrine interactions are now established as crucial factors for the control of body development and homeostasis. In distant species and invertebrates, the foundations of both the neuroendocrine system and innate immunity were coexisting until now without any apparent problem. Some 300 millions years ago, in a relatively short period after agnathan fishes (e.g., hagfish and lamprey), adaptive immunity emerged in the first gnathostomes, cartilaginous fishes (e.g., shark and ray). Somatic recombination machinery characterizes adaptive immunity and is responsible for the random generation of the huge *diversity* of immune receptors able to recognize infectious antigens. The emergence of this novel form of immune defenses exerted a so potent pressure that structures and mechanisms developed along the paths of lymphocyte traffic to impose immunological self-tolerance, that is, the inability of the immune system to attack the host organism. Together with the generation of diversity and memory, self-tolerance constitutes a fundamental property of the immune system. The progressive rise in the level of immune diversity and complexity also explains why self-tolerance failures (i.e., organ-specific autoimmune diseases) were increasingly detected during evolution, the maximum being currently observed in the human species. The first thymus appeared in cartilaginous fishes (chondrichthyes), concomitantly with the emergence of rudimental forms of adaptive immunity. Though some forms of tolerance induction already takes place in primary hemopoietic sites (fetal liver and bone marrow), antigen-dependent B-cell tolerance is primarily due to an absence of T-cell help. Among all lymphoid structures, the thymus is the only organ specialized in the establishment of central self-tolerance. The thymus crucially stands at the crossroad between the immune and neuroendocrine systems. In this organ responsible for thymopoiesis—T-cell generation—(Kong et al., 1998), the neuroendocrine system regulates the process of T-cell differentiation from the very early stages. In addition, T lymphocytes undergo inside the thymus a complex educative process that establishes central T-cell self-tolerance of neuroendocrine principles (Geenen et al., 1992; Martens et al., 1996). Within the thymus, a confrontation permanently occurs between previously established neuroendocrine principles and a recent system equipped with recombination machinery promoting stochastic generation of response diversity. Contrary to a previous assumption, the thymus functions throughout life (Poulin et al., 1999; Geenen et al., 2003) and plays a fundamental role in the recovery of a competent T-cell repertoire after intensive chemotherapy or during highly active antiretroviral therapy (Mackall et al., 1995; Douek et al., 1998).

## 2. Development and cytoarchitecture of the thymus

In the human species, the thymus is located in the anterior mediastinum, overlying the heart and the basal blood vessels. Its general shape resembles the leaves of the thyme plant. It is

divided into two lobes surrounded by a connective capsule. By sending incomplete septa, the capsule delineates thymic lobules further divided into distinct zones: subcapsular cortex, cortex, cortico-medullary junction, and medulla. The generation of a peripheral CD4<sup>+</sup> and CD8<sup>+</sup> T-cell pool occurs in the thymic organ that provides the appropriate cellular microenvironments. Thymocyte development is not a pure autonomous genetic program, but it closely depends on interactions of pre-T cells with the complex cellular network of the thymus.

Thymic epithelial cells (TECs) constitute the major component of the thymus stroma and mainly derive from the endoderm of the third branchial pouch. In this endoderm, recent studies have identified epithelial stem cells able to form a thymus with normal phenotype and function in vivo (Bennett et al., 2002; Gill et al., 2002). Further development of this primitive epithelial rudiment depends upon an important mesenchymal contribution from the cephalic neural crest (Bockman, 1997). Some human diseases (and animal models) include a defective thymus development leading to primary immune deficiencies. DiGeorge's syndrome associates congenital absence (or hypoplasia) of thymus, parathyroids, and defects in the heart and truncal vessels. This syndrome results from a migration failure of the cephalic neural crest. Mice in whom the homeobox gene *Hoxa3* has been disrupted present thymic aplasia, parathyroid hypoplasia, and frequent defects in heart and great vessels. Wild animals with immune deficiencies most closely related to DiGeorge's syndrome are nude mice with hairlessness and lack of thymic development resulting from defect in epithelial cells. The nude phenotype is caused by mutations in *nude* gene on murine chromosome 11 encoding the transcription factor winged-helix nude (whn or Foxn1). In the absence of whn, the thymus anlage still develops but is filled with primitive epithelial cells that do not specialize and segregate into subregions. Other transcription factors (Pax-1, Pax-9) also contribute to the ontogeny of thymic epithelium.

Thymic nurse cells (TNCs) are large epithelial cells located in the subcapsular and outer cortex. They contain the subcellular equipment and the enzymatic machinery required for antigen processing and presentation. Though they express various neuroendocrine markers and synthesize neuropeptides, their precise embryonic origin remains unknown.

Thymic dendritic cells (DCs), mainly located at the cortico-medullary junction, and macrophages, without any precise topography, derive from bone marrow and constitute the other components of the thymic stroma.

The thymus also receives sympathetic autonomic innervation (Bulloch, 1985). A number of neurotransmitters have been identified by immunocytochemistry within thymic sensory nerve fibers.

The thymus is classically described as a lymphoepithelial structure. However, it is important to understand that the lymphoid compartment is a mobile population that originate as T-cell progenitors in primitive hemopoietic sites, migrate into the thymus, and differentiate through close contacts with the surrounding TECs/TNCs, DCs and macrophages.

## 3. The dual role of the thymus in T-cell differentiation

The pioneering studies conducted by Miller and Osoba (1967) have demonstrated the active role played by the thymus in the generation of peripheral T lymphocytes, and its previously assumed function as an endocrine gland was less and less considered. From the primary sites

of hemopoiesis (fetal yolk sac, liver, and then bone marrow), possibly under the control of some chemokines, T-cell precursors enter into the thymus through capillaries at the corticomedullary junction, then migrate to the subcapsular cortex (Fig. 1). An intense proliferation of T-cell blasts takes place in the outer cortex. The transmembrane molecule Notch-1 plays an important role in the induction of T-lineage restriction and commitment. Then, intimate contacts with TECs/TNCs promote the activity of the recombination-activating genes (RAG 1 and RAG 2). RAG-catalyzed generation of T cell diversity occurs through the random recombination of gene segments encoding the variable parts of the T-cell antigen receptor (TCR) (Fugmann et al., 2000). TCR recombination and expression by thymocytes (pre-T cells) constitutes a pivotal event because, among the number of about 10<sup>8</sup> possible combinations, many of them are able to recognize self-antigens presented by the major histocompatibility complex (MHC) proteins expressed by TECs/TNCs, DCs and macrophages. The so-called thymic negative selection represents the massive deletion of Tcell clones expressing self-reactive TCR and results either from programmed cell death or developmental arrest. Thymus censorship is responsible for the establishment of central Tcell self-tolerance (Nossal, 1994). This fundamental process is very powerful since, from 100 T-cell precursors, only 1-2 T cells will leave the thymus in a state of self-tolerance and competence against foreign antigens.

The concomitant occurrence within the same organ of opposite events such as T- cell negative selection and T-cell life/diversity generation represents a prominent problem in modern biology. A number of models have been proposed to explain the paradox of thymus physiology. A first model attributed different functional roles to the cells of thymic stroma: TECs/TNCs would be mainly responsible for inducing T-cell proliferation and development, whereas thymic DCs and macrophages, as dedicated antigen-presenting cells (APCs), would mediate T-cell negative selection. Although DCs and macrophages are very efficient in inducing T-cell deletion, this model is no more valid since it is now demonstrated that TECs/TNCs are also able to present self-antigens and provoke T-cell negative selection. Medullary TECs in particular are highly competent in achieving self-tolerance. The most popular model is called the avidity model (Ashton-Rickardt et al., 1994): T cells bearing a TCR with a high affinity for self-antigen at high density will undergo clonal deletion, while those with a low-affinity TCR and/or in presence of low-density self-antigen survive and further develop. However, the affinity of a given TCR for its cognate antigen is rather low (with a  $K_D$  around  $10^{-7}$  M), and thus one may question the physiological meaning of lower affinities. The model of the dual role played by the repertoire of thymic neuroendocrinerelated genes recapitulates at the molecular level the paradox of thymus physiology. According to this model, thymic neuroendocrine-related precursors are the source of two distinct types of interactions with immature T cells. On the one hand these precursors deliver ligands that are not secreted but targeted to the outer surface of TECs/TNCs from where they can bind to neuroendocrine receptors expressed by immature T cells. A high affinity (K<sub>D</sub> ranging from 10<sup>-11</sup> to 10<sup>-9</sup> M) and a low specificity characterize this cryptocrine type of signaling. On the other hand the same precursors undergo antigen processing and are the source of *neuroendocrine self-peptides* presented by the thymic MHC machinery. A negative signal would result from the low-affinity but highly specific binding of the complex MHC/neuroendocrine self-antigen to its cognate TCR. According to the theory of clonal deletion, the central T-cell self-tolerance of neuroendocrine principles results from the intrathymic presentation of neuroendocrine self-antigens (Fig. 2).

In addition to deletion of self-reactive T-cell clones, the establishment of central thymic self-tolerance also includes the generation of regulatory T cells ( $T_{REG}$ ) that are enriched in the

CD4<sup>+</sup>CD25<sup>+</sup> T cell subset (Shevach, 2000; Sakagushi, 2001). It remains to be determined how far the thymic self-antigen repertoire contributes to the generation of T<sub>REG</sub> lymphocytes.

## 4. The nature of neuroendocrine self (Table I)

From the investigation of intrathymic expression of neuroendocrine-related self-peptide precursor genes, a series of properties were drawn to define the nature of neuroendocrine self. (1) Neuroendocrine self-antigens usually correspond to peptide sequences that have been highly conserved throughout the evolution of one given family. (2) A hierarchy characterizes their expression pattern. In the neurohypophysial family, oxytocin (OT) is the dominant peptide synthesized by TECs/TNCs from different species (Geenen et al., 1986). The binding of OT to OT receptor (OTR) expressed by pre-T cells induces a very rapid phosphorylation of focal adhesion related kinases. This event should play a major role in the promotion of immunological "synapses" between immature T lymphocytes and thymic APCs (TECs/TNCs, macrophages and DCs). With regard to tachykinins, neurokinin A (NKA)—but not substance P (SP)—is the peptide generated from the processing by TECs of the preprotachykinin A (PPT-A) gene product (Ericsson et al., 1990). All the genes of the insulin family are expressed according to a precise hierarchy and topography: IGF2 (TECs/TNCs) > IGF1 (thymic macrophages) >> INS (thymic DCs and/or medullary TECs) (Geenen and Lefèbvre, 1998; Derbinski et al., 2001). This hierarchical pattern is meaningful since the tolerogenic response primarily concerns the dominant epitopes of a protein family (Sercarz et al., 1993). In fetal thymic organ cultures, the blockade of thymic IGF-mediated signaling, at the level of IGF ligands (in particular IGF-2) or IGF receptors, severely interferes with early stages of T-cell differentiation and proliferation. On the contrary, the treatment of fetal thymic organ cultures with antibodies against (pro)insulin does not exert any significant effect on T-cell differentiation (Kecha et al., 2000). (3) Neuroendocrine precursors are not processed in a classic fashion, but they undergo antigenic processing for presentation by, or in association with, thymic MHC proteins. (4) Some differences exist in the processing between thymic APCs and dedicated peripheral APCs. At least for some neuroendocrine self-antigens (OT and neurotensin), those differences imply that their presentation by thymic APCs is not restricted by MHC alleles like presentation of foreign antigens by peripheral APCs (Vanneste et al., 1997; Geenen et al., 1999).

During ontogeny of Balb/c mice, *OT* transcripts are detected on fetal day 13 (E13) both in thymus and brain, whereas vasopressin (*VP*) gene transcription starts on E14 in the brain and is clearly detected in the thymus only on E15. The earlier *OT* expression in the thymus supports the role of thymic OT in the induction of central T-cell tolerance of the neurohypophysial peptides before their appearance in the hypothalamic magnocellular neurons (Fig. 3). The expression of the neurohypophysial receptor genes (*OTR*, *V1R*, *V2R* and *V3R*) was investigated on murine thymic CD4<sup>-</sup> CD8<sup>-</sup> lymphoma cell lines, as well as on murine thymic T-cell subsets. *OTR* transcripts are detected in CD4<sup>-</sup>CD8<sup>-</sup>, CD4<sup>+</sup> CD8<sup>+</sup> and CD8<sup>+</sup> cells, while a faint *V3R* expression is restricted to CD4<sup>+</sup> CD8<sup>+</sup> and CD8<sup>+</sup> cells. *V1R* and *V2R* expression could not be detected on any T cell subset. In fetal thymic organ cultures, a specific OTR antagonist increases late T-cell apoptosis confirming the involvement of OT/OTR-signaling in T-cell proliferation/survival (Hansenne et al., in press).

## 5. The development of autoimmunity as a defect in thymic self-tolerance establishment

In 1992, a defect in thymic T-cell education to recognize and to tolerate self-antigens was hypothesized to play a pivotal role in the development of hypothalamus-specific

autoimmunity (Geenen et al., 1992; Robert et al., 1992). As already hypothesized by Burnet (1973), the pathogenesis of autoimmune diseases could result from the appearance of "forbidden" self-reactive clones in the peripheral lymphocyte repertoire. Since the thymus is the primary site for induction of self-tolerance, thorough investigation of a defective thymic censorship should provide the scientific community with important keys to understand the mechanisms underlying the development of autoimmune responses. A number of abnormalities of thymic morphology and cytoarchitecture have been described for several autoimmune disorders. The expression of the transcription factor autoimmune regulator (AIRE) gene is maximal in murine medullary TECs, but it is absent in the thymus epithelium of diabetic NOD mice (Heino et al., 2000). Type 1 diabetes (juvenile or insulin-dependent diabetes) is a chronic devastating disease resulting from an autoimmune response specifically oriented against pancreatic islet B cells, the only cells secreting insulin according to the endocrine model. In accordance with the above hypothesis, INS transcripts were measured at lower levels in the thymus of human fetuses with short class I VNTR (variable number of tandem repeats) alleles, a genetic trait of type 1 diabetes susceptibility (Vafiadis et al., 1997; Pugliese et al., 1997). The expression of insulin-related genes was also analyzed in thymus, liver and brain of a common animal model of type 1 diabetes, the bio-breeding (BB rat). A defect of Igf2 expression was evidenced in the thymus of more than 80% of diabetes-prone BB rats, in close accordance with the incidence of diabetes in this rat strain (Kecha-Kamoun et al., 2001). In these rats, this tissue-specific defect in *Igf2* transcription could contribute both to lymphopenia (including a lack of antigen-specific T<sub>REG</sub> cells that control autoimmune diabetes), as well as to the absence of central self-tolerance of insulin family. As a consequence of the defective thymic censorship, self-reactive T cells bearing TCR oriented against dominant epitopes of insulin-related peptides continuously migrate from the thymus and enrich the peripheral pool with self-reactive T cells exhibiting a potential cytotoxic power against islet ß cells. Under certain environmental influences, a molecular bridge could be installed between the target autoantigenic epitopes leading to activation of the self-reactive Tcell pool and subsequent \(\beta\)-cell destruction (Fig. 4). Diabetes-prone NOD mice also display a defect in central tolerance (negative selection) of T lymphocytes (Kishimoto and Sprent, 2001). The transcription of genes encoding tissue-specific antigens (including Ot, Igf2, Ins2, and Npy) is markedly down-regulated in the thymus epithelium of Aire<sup>-/-</sup> mice (Anderson et al., 2002). Moreover, Aire deficiency results in an almost complete thymic failure to delete CD4<sup>+</sup> T cells with high affinity for a pancreas-specific antigen (Liston et al., 2003). Thus, more and more arguments provide strong evidence that a defect in thymic self-tolerance plays a pivotal role in the pathogenesis and development of organ-specific autoimmunity. The origin of systemic (non-organ specific) autoimmunity is still a matter of debate and could be elucidated through further investigations of the non-specific, innate immune system.

## 6. Pharmacological applications and conclusions

The study of neuroendocrine gene expression and precursor processing in the thymus led to the identification of *neuroendocrine self-peptides*. With regard to insulin-related gene expression in the thymus, IGF-2—a prominent fetal growth factor—was identified as the dominant self-peptide precursor of the insulin family expressed in the thymus from different species, including man. This observation is in close accordance with the theory of self-recognition which, according to F.M. Burnet, is not an inherited property but is gradually acquired in the course of fetal life. Although the tolerogenic properties of neuroendocrine self-peptides remain to be further documented, they are suspected from what is known about the immunological tolerance of classic hormones. The development of specific antibodies by active immunization (i.e., experimental breakdown of self-tolerance) revealed that OT is

more tolerated than VP, and that IGF-2 is also more tolerated than IGF-1, and much more than insulin. Some cases of diabetes insipidus result from an autoimmune process against VP-producing hypothalamic neurons (infundibulohypothalamitis) (Scherbaum et al., 1985; Imura et al., 1993). Insulin is a primary autoantigen tackled by the autoimmune response observed in type 1 diabetes and this might result from its very low expression in the thymus. At the opposite, autoimmunity has never been observed against OT and IGF-2. The strong tolerance of these peptides, resulting from high expression of their genes in the thymus, may be considered as the consequence of evolutive pressure to protect fundamental processes such as species reproduction and individual ontogeny (Geenen et al., 1999). The putative pathogenic role of forbidden self-reactive T cells against IGF-2 deserves to be further analyzed through immunization of *Igf2*-/- mice.

Thus although VP and insulin behaves as immunogenic auto-antigens of their respective families, OT and IGF-2 may be viewed as the tolerogenic self-peptide precursors of neurohypophysial and insulin families, respectively. Quite provocatively, perhaps is it now appropriate to establish a distinction between an **auto**-antigen and a **self**-antigen (Fig. 5). According to this distinction, auto-antigens can be considered as "altered" self-antigens. Though they are highly homologous, i.e., evolutionarily related, they are not identical and this biochemical difference theoretically drives opposite immune responses (i.e., immunogenic vs. tolerogenic responses). This proved to be correct in a recent preclinical study. A dominant epitope of insulin (Ins B<sub>9-23</sub>) and the homologous sequence of IGF-2 (IGF-2 B<sub>11-25</sub>) compete for binding to DQ8, a MHC class II allele conferring major susceptibility to type 1 diabetes. In young DQ8<sup>+</sup> type 1 diabetic patients, Ins B<sub>9-23</sub> presentation by DQ8 elicits a dominant IFNγ secretion by isolated peripheral blood mononuclear cells, whereas presentation of IGF-2 B<sub>11-25</sub> promotes a dominant regulatory IL-10 secretion. The powerful tolerogenic properties of thymic self-antigens should be soon exploited to cure and/or prevent severe autoimmune diseases (such as type 1 diabetes) which constitute the heavy tribute, mainly paid by mankind, for the diversity, complexity, and efficiency of its system of immune defenses.

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## 8. Legends to figures

Fig. 1: Pathways of intrathymic T-cell differentiation.

From fetal liver then bone marrow, T-cell progenitors enter the thymus by capillaries at the cortico-medullary junction and migrate to the subcaspsular outer cortex where they divide intensively (T-cell blasts). Interactions between randomly rearranged TCR chains on thymocytes and self-antigen/MHC complexes presented by thymic stromal cells drive the programme of T-cell differentiation in the thymus. From 100 thymocyte precursors that enter the thymus, only 1-2 self-tolerant CD4+ helper and CD8+ cytotoxic emigrate from the thymus to establish the peripheral T-cell repertoire.

- Fig. 2: The dual role in the thymus of neuroendocrine-related self-peptide genes. Neuroendocrine gene-encoded precursors are the source of two types of interactions with pre-T cells. (1) They deliver *ligands*, which are not secreted but targeted to the outer surface of thymic plasma cell membranes. Those ligands bind to T-cell neuroendocrine receptors leading to activation of transductory pathways such as increase of inositol triphosphate (IP<sub>3</sub>) and phosphorylation of focal adhesion kinases (FAK). (2) The same precursors are processed into *self-antigens* presented by—or in association with—thymic MHC molecules. This presentation will lead to clonal deletion and/or developmental arrest T cells bearing a TCR oriented against neuroendocrine self-antigens and MHC complexes.
- Fig. 3: Ontogeny of *OT*, *VP* and *OTR* expression in thymus and brain of Balb/c mouse. *OT* transcription is already detectable on E13 in both murine thymus and brain, whereas *VP* transcripts are detected in the brain later (E14), and are detected in the thymus only on E15. Though not quantitative, the profile of *OTR* expression seems to decrease in the thymus and to increase in the brain during fetal development.
- Fig. 4: Role of the thymus in the development of the autoimmune response against islet insulin-secreting  $\beta$  cells.

Together with abnormalities of thymic cytoarchitecture, the thymus-specific defect of IGF2 expression contributes both to lymphopenia (with a decreased number of  $T_{REG}$  cells) and to the absence of central T-cell self-tolerance of the insulin family. The influence of the environmental factors could be exerted at the level of thymic T-cell differentiation, as well as

in the periphery through the establishment of a bridge between the  $\beta$ -cell auto-antigen and forbidden self-reactive T cells. Coxsackievirus B4 (CVB4) infection is a major environmental factor implicated in the pathogenesis of type 1 diabetes. Three mechanisms are currently proposed to explain the CVB4 diabetogenic influence: (1) molecular mimicry between sequences of glutamic acid decarboxylase (GAD) and a structural CVB4 protein; (2) direct toxic action of CVB4 toward islet  $\beta$  cells and bystander activation of autoreactive T cells; and (3) CVB4 infection of the thymic epithelium and interference with the establishment of central T-cell self-tolerance (Brilot et al., 2002).

Fig. 5: Opposite immune responses driven by insulin as an auto-antigen vs. IGF-2 as a self-antigen precursor.

As evidenced by numerous authors, insulin (Ins) is a major  $\beta$ -cell specific auto-antigen tackled by the diabetogenic autoimmune response. Ins is immunogenic and the sequence  $B_{9-23}$  is a dominant epitope recognized by pathogenic CD4<sup>+</sup> T cells. IGF-2 is the dominant member of the insulin family expressed in TECs of different species and this may explain why the immune tolerance of IGF-2 is so hard to break. Both Ins  $B_{9-23}$  and the homologous sequence IGF-2  $B_{11-25}$  bind with the same affinity to DQ8, one HLA class II allele conferring major susceptibility to type 1 diabetes. DQ8 presentation of IGF-2  $B_{11-25}$  to peripheral blood mononuclear cells of type 1 diabetic patients elicits a dominant IL-10 secretion as revealed by Elispot analyses. The regulatory/tolerogenic properties of IGF-2 (or IGF-2  $B_{11-25}$ ) could be exploited in a vaccination strategy to reprogram immune self-tolerance of the insulin family.

**Table I.** The organization of the thymic repertoire of neuroendocrine-related self-peptide precursors/genes

Function	Endocrine factors	Paracrine/ autocrine factors	Thymic self-peptide precursors
Reproduction	Oxytocin (OT)	ОТ	ОТ
Water homeostasis	Vasopressin (VP)	VP	OT >> VP
Sodium homeostasis	Natriuretic peptides (ANP, BNP, CNP)	ANP, BNP, CNP	ANP
Pain	-	Neurokinins (SP, NKA, NKB)	NKA
Glucose metabolism	Insulin (Ins)	IGF-1 IGF-2	IGF-2 > IGF-1 >> Ins

The fundamental principle is based upon the close homology between peripheral hormones and related thymic self-peptide genes/precursors. Tolerogenic neuroendocrine self-peptide precursors are predominantly conserved throughout evolution of their family and are the dominant members of one family that are expressed in the thymus.

## 9. Further reading

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#### 11. Related websites

- www.nature.com/ni/special\_focus/autoimmunity/index.html
- www.endo-society.org
- www.diabetes.org
- www.easd.org
- www.isnim.org
- www.isni2004.org
- www.pnirs.org
- www.mindbody.org