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TOWARD A 3D IN VITRO MODEL BASED ON DECELLULARIZED THYMUS TO MAINTAIN ADULT THYMIC EPHITELIAL CELLS FUNCTIONALITY

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1. INTRODUCTION

1.1 THYMUS

In mammals and man, the thymus is the primary lymphoid organ for T lymphocyte maturation, whereas the bone marrow serves as the primary lymphoid organ for the B lymphocyte compartment, although the gut-associated lymphoid tissue is its phylogenetic ancestor.

The thymus is the first of the lymphoid organs to be formed in all vertebrates, and grows considerably immediately after birth in response to postnatal antigen stimulation and the demand for large numbers of mature T cells. Genetic factors also influence the age of onset, rate and magnitude of thymus dependent immunological function. The thymus reaches maximal size by sexual maturity and then gradually involutes [1].

In humans, the thymus is located in the mediastinum, just anterior to the ascending aorta and inferior to the left brachiocephalic vein. If examined when its growth is most active, it consists of two lateral lobes placed in close contact along the middle line, situated partly in the thorax, partly in the neck, and extending from the fourth costal cartilage upward, as high as the lower border of the thyroid gland. It is covered by the sternum, and by the origins of the Sternohyoidei and Sternothyreoidei muscles. Below, it rests upon the pericardium, being separated from the aortic arch and great vessels by a layer of the middle cervical fascia. In the neck it lies on the front and sides of the trachea, behind the Sternohyoidei and Sternothyreoidei muscles.

The thymus is of a pinkish-gray color, soft, and lobulated on its surfaces. This superficial arrangement corresponds to a threedimensional (3D) organization of its parenchyma, specifically to 3D

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prismatic structures, hystologically constituted by two subcompartments, the cortex and the medulla. Each compartment contains distinct populations of thymic ephitelial cells (TEC), as weel as mesenchymal cells, endothelial cells and dendritic cells.



Fig. 1 A: Structure of the mammals thymus. The thymus is a lympho-ephitelial organ surrounded by a connective capsule. Each of its constituting units , the thymic lobule can be divided into a central medulla (m) region, which contains medullary thymic ephitelial cells (m TECs), and an outer cprtex (c), which contains cortical thymic ephitelial cells (c TECs). B: Histology of the adult mouse thymus. Hematoxylin and eosin-stained sagittal section highlighting the thymus subcompartments.

The thymus provides a unique microenvironment for the efficient production of a diverse T cell repertoire[2]. However, after puberty it begins to regress. This *involution*, which seems to occur in almost all vertebrates, rodents included, suggesting that this is an evolutionary ancient and conserved process. Age-associated thymic involution includes a decrease in tissue mass and cellularity, together with a loss of tissue organization, leading to a reduction in the cell output. This decline in the T cell output is believed to have a major impact on the properties of the peripheral T cell pool such that with increasing age, these cells exhibit alterations in phenotype and function, loss of diversity, and replicative senescence[3].

1.1.1 Anatomy of the thymus

In mammals and man, the thymus has a bilobed structure, it is classidfied as a lympho-ephitelial organ containing many developing lymphocytes, and is surrounded by a capsule of connective tissue. The two lobes generally differ in size; they are occasionally united, so as to form a single mass; sometimes they are separated by an intermediate lobe. Each lateral lobe is composed of numerous lobules, featured as 3D prismatic structures of prismatic geometry, held together by delicate areolar, stromal tissue; the entire gland being enclosed in an investing capsule of a similar but denser structure. The lobules vary in size and consists of a cortical and a medullary portion. The cortical portion is mainly composed of lymphoid cells, supported by a network of finely branched cells, the cortical thymic ephitelial cells (cTECs), which is continuous with a similar network in the medullary portion, formed by medullary TEC (mTECs). A network of reticular TECs forms an adventitia the blood vessels. In the medullary portion, this reticulum is around coarser than in the cortex, the lymphoid cells are relatively fewer in number, and there are found peculiar nest-like bodies, the concentric corpuscles of Hassall, originating from involution of local TECs appoases in concentric layers. These concentric corpuscles are characterized by a central mass, consisting of one or more granular elelments, and of a capsule which is formed of epithelioid cells.

Each lobule is surrounded by a vascular plexus, from which vessels pass into the interior, and radiate from the periphery toward the center, forming a second zone just within the margin of the medullary portion. In the center of the medullary portion there are very few vessels, and they are of minute size.

Arteries supplying the thymus are derived from the internal mammary, and from the superior and inferior thyroids. The veins end in the left innominate vein, and in the thyroid veins. The nerves are exceedingly minute; they are derived from the vagus and sympathetic chain. Branches from the descendens hypoglossus and phrenic nerves reach the investing capsule, but do not penetrate into the substance of the gland. The thymic arteries follow the course of the interlobular connective tissue septae, and enter the thymic parenchyma at the corticomedullary junction. The corticomedullary arterioles ramify into capillaries that extend into the cortex and medulla. In the cortex they form a complex of capillary arcades which together with perivascular lymphocytes, macrophages and peripheral reticular epithelial cells form the blood-thymus barrier[4]. Capillaries in the cortex are rarely fenestrated [5]. This restricts access of circulating antigenic molecules to developing cortical lymphocytes. By contrast, medullary capillaries are fenestrated and freely permeable to circulating antigens. Blood drains into the postcapillary venules, and finally returns to the corticomedullary junction in medullary veins [4].

Prothymocytes are thought to enter the thymic stroma through the large venules at the corticomedullary junction, and re-enter the circulation through the vascular lining of post-capilliary venules. These perivascular areas contain accumulations of phenotypically mature T cells that express the marker Mel 14, a receptor involved in the migration of lymphocytes through the lymphoid vasculature [6]. Efferent lymphatics drain into an adjacent pair of lymphnodes. The thymus has no afferent lymphatics. The

rodent thymus has an abundant noradrenergic innervation but some cholinergic innervation has also been demonstrated [7]. Nerves follow the vasculature, within the capsule and septae and the highest concentration of nerve fibres is in the corticomedullary junction.



Fig. 2 Schematic diagram outlining the pathway of T cell development. Aging can impact on variety of pathways during the development of T cells.

1.1.2 Development of the thymus

The epithelium of the thymus is derived from the embryonic endodermal layer, with possible contributions from the ectoderm, through a series of differentiating steps, each of which must be completed in order to provide the optimum environment for thymic development and function. The roof of the endodermal yolk sac is folded into the expanding embryo as the foregut (Fig.3). The embryonic pharynx is quite important for the development of many organs, including the thyroid and parathyroid glands as well as the thymus.



Fig.3 Representation of the foregut formation in the embryo development.

Alongside the thymocites, at different stages of maturation, the thymic environment is formed by ephitelial cells, which form a meshwork to provide mechanical support and stimuli for the proliferation and development of thymocites and by macrophages, dendritic cells, fibroblast and matrix molecules[8].

The developing pharyngeal region in contact with the ectodermal layer produces inclusions of ectodermal cells into the thymus, a

mechanism possibly explaining the heterogeneity of ephitelial cells in yhe mature thymus. The mass of endodermal cells from the 3rd and 4th faringeal pouch give rise to the primordial thymus (and forthcoming parathyroid glands), and attracts lymphoid precursor cells that are brought to this region through the bloodstream. The lymphoid precursor cells leave the blood vessels and migrate through the mesenchyme into the primitive organ bud. This migration starts between 11 and 12 gestation days (GD) [9]. In the rat, the process is less well known and according to the few existing reports, it starts between GD 13 and 15[10]. The earliest phase of organogenesis, culminates in the formation of two primordial buds, one of each antimeres, surrounded by a condensing mesenchymal capsule. By GD 13.5 for mice and 16.5 for rats the parathyroid and thymus are separated into physically distinct organs and soon afterwards they reach their approximate adult positions within the embryo (Fig.4).



Fig. 4 Photomicrograph showing sagittal sections of the rat embryo. A at 10 GD the thymus is not presetn; B at 14 GD it is visible at rudiment of the organ and C at 19 GD the two lobes can be distinguished. Hematoxylin-eosin staining.

In the embryonic thymic cortex, rat lymphocytes, in cortex, show a different antigenic pattern at the 16th day and at the 31th day after the birth. In adult rats, differences in CD4:CD8 cell ratio involve the major histocompatibility complex (MHC) haplotype[11], and peripheral CD4⁺ T cells predominate over CD8⁺ T cells but the origin of these differences 14

has not yet been elucidated.

Unlike adult thymic precursors, fetal thymic progenitors give rise to reduced numbers of CD4⁺ peripheral T cells in chimeric animals (CD4:CD8 cell ratio <1). This relative under- representation of CD4⁺ peripheral T cells was seen 4–16 weeks after either intrathymic injection of suspensions of foetal CD4⁻CD8⁻ thymocytes into adult thymuses, or transplantation of foetal thymic lobes under the kidney capsule of adult host rats. Then, foetal thymic precursors seem to be responsible for a relative underproduction of CD4⁺ peripheral T cells, regardless of the developmental age of the thymic stromal environment. However, the developmental age of the thymic stromal environment is the key regulatory element involved in the generation of distinct proportions of CD4⁺ and CD8⁺ thymocytes.

1.1.3 Thymic epithelial and dendritic cells in the adult thymus

The bulk of the supporting framework in the thymus is composed of a network of TECs [12]. In the subcapsular region of the thymus, TECs form a layer, 1 or 2 cells deep. In the outer cortex, TECs are thin and sheet like, but elsewhere they assume a stellate appearance.

TECs manifest expression of MHC class I antigens, neuropeptides, neurophysin, oxitocin/vasopressin and neuroendocrine markers. They are responsible for interleukins IL-6 and IL-7 secretion, and for secretion of thymic hormones. Superficial TECs demonstrate common phenotype traits with medullary traits [13]. TECs in the mid and deep cortex also have slender cytoplasmic processes providing a large surface area for cell–cell interactions with developing thymocytes[14].

Morphologically, TECs are characterized by electronlucent vacuoles with a small amount of moderate electrondense material, linked to secretory activity. The cortical TECs are distinguished by expression of both MHC class I and II molecules and protein gene product 9.5 (PGP 9.5). This latter is also a marker of a specific population of cortical TECs, known as thymic nurse cells (TNCs) [13].

In the outer cortex, some TECs appear to completely surround and enclose a large number (about 10–250) of thymocytes[15]. Moreover, TECs have been described in the subcapsular and outer cortical regions of the human [16] and rat thymus[17].

In contrast to cortical TECs, those contain a small number of thymocytes [18]. Scanning electron microscopic images of the mouse

thymus have suggested that thymocytes migrate in or out these lymphostromal complexes[17]. TECs are thought to provide a specialized microenvironment for T-cell differentiation[19]. Although the epithelium they constitute is thought to be a sessile component of the thymus[20], the subcapsular epithelium in particular may be a dynamic structure, since transitional forms have been observed between sheet-like TECs, basketlike TECs, and TECs depending on the type of cortical lympho- stromal interaction.

Most medullary TECs are closely packed displaying voluminous cytoplasm and short processes. They are commonly subdivided into two subsets, representing different stages of their differentiation[21,13].Cells of the first subset are defined "undifferentiated" and are around 1% of the total medullary population. They remain unrelatively unknown and are identified on the basis of ultrastructural traits in the outer medulla, as solitary cells or arranged in small groups of cells, which comprise forms of traits intermediate with those of the second subsubset of medullary TECs. The second type of TECs are polygonal, with a rich cytoplasm, polarized with respect to the position of organelles and having ultrastructural traits of endocrine cells. They express a wide range of peptide hormones and several co-stimulatory molecules including CD40.

In the adult mouse, dendritic cells are present mainly in the medulla; they are voluminous with electronlucent cytoplasm penetrating between surrounding lymphoid cells (Fig.5).



Fig.5 Electron micrograph showing thymic morphology in the rat. Dendritic cells are characterized by electron-lucent cytoplasm.

1.1.4 Composition of the thymus microenvironment

The thymus microenvironment forms a complex network of interaction that comprises non lymphoid cells, extracellular matrix elements (ECM), matrix metalloproteinases, cytokines, chemokines, and other soluble proteins. During their journey inside the thymus, thymocites interact with stromal cells and extracellular matrix proteins to receive appropriate signals for survival, proliferation and differentiation.

On day 13 of fetal lifea number of ECM components have been observed in the thymus. Collagen type I expression was restricted to a few cells in the capsule area, whereas laminin was present in the capsule and in some scattered stromal cell clusters. In contrast, a filamentous network observed throughout the organ shows positivity for Collagen type IV and fibronectin[25].

On day 15 of fetal life it is not possible to distinguish cortical and medullary regions by ECM expression patterns, and the entire organ displayed a thin network of laminin, fibronectin and collagen type IV. On day 17 a similar profile is apparent, although all these proteins were additionally detected at basement membranes and intralobular perivascular spaces. At this stage collagen type I expression is similar to that found in young adult animals. The complete adult pattern of thymic ECM distribution was achieved on day 19 of fetal life[25].

The distribution of ECM in the normal thymus does not follow a homogeneus pattern, and glycoproteins such as fibronectin, laminin and collagen type IV are not restrictly located at typical basement membrane sites. In the medullary region of the thymic lobules, they form a rather thick medullary network, whereas very thin ECM fibres are found within the cortex.



Fig.6 Intrathymic distribution of fibronectin in human thymus, as revealed by confocal mycroscopy. The fibronectincontaining network is denser in the medulla (right part on the panel) if the thymic lobule, as compared to the cortex (left part of the panel). Additionally, Fibronectin is seen around vessels.

The distribution of collagen type I is limited to the capsule, intraseptal and perivascular spaces[22].

The fibronectin is located throughtout the all thymic parenchima and outlines the cortico-medullary junction of the thymic lobules[23]. Different isoform of the laminin could be found into the thymus. Laminin 2 is restricted to cortical ephitelial cells whereas laminin 5 is located surrounding small blood vessels[24]. Laminin 1 and 2 can modulate thymocyte migration within TNCs complexs. In the human thymus a large number of fibers in the interior of the connective tissue septae are made of collagen type I whereas only some fibers are labelled with collagen type IV. Instead the basement membrane bordering these septae contained large amount of collagen type IV. The interior of interlobular septae cointain laminin and fibronectin; in contrast the connective fibres situated at the corticomedullary junction are made up of collagen type IV, type I

and fibronectin.



Fig.7 Frozen section of adult mouse thymus showing the distribution pattern of a collagen type I, b type III and C type IV and D fibronectin. A part of cortex (C) of a thymic lobule , as well as septum containing a blodd vessel(V) coul be seen. Bar $50\mu m$

The extracellular localization of the laminin is mainly medullary and also medullary TECs contain and produce this ECM protein.



Fig.8 Frozen section of adult mouse thymus showing the distribution pattern of fibronectin within the thymic lobules. (B) Higher magnification of the fibronectin-positive network occurring in the medullary region. Bars: a- 50µm b-20µm

In the mouse fibronectin and laminin are strongly detected at the basement membranes, not only in the connective tissue septa but also in the interlobular perivascular spaces. In addition, filaments containing fibronectin, laminin and few collagen type IV have been found within the thymic medulla.



Fig.9 Frozen section of adult mouse thymus showing the distribution pattern of fibronectin within the thymic lobules. (B) Higher magnification of the fibronectin-positive network occurring in the medullary region. Bars: a- 50μm b-20μm

In normal young adult animals, the distribution patterns of both interstitial and basement membrane proteins were similar to those previously described for the infant human thymus[22].



Fig. 10 Extracellular matrix labelling in the fetal thymus. a) fibronectin and b) laminin labeling patterns in a 13- day fetal thymus. c) the fibronectin network observed in a 15.day fetal thymus. Bar-20µm

All ECM components are produced by thymic microenvironmental cells, which also drive thymocyte differentiation. Signals triggered by ECM are conveyed into thymocytes or microenvironmental cells through specific membrane receptors, and most of them belong to the type, such

as the VLA-3, VLA-4, VLA-5 and VLA-6. In vitro studies revealed that adhesion of thymocytes to thymic integrin microenvironmental cells is mediated by extracellular matrix. Such an adhesion is preferentially done by immature thymocytes[26].

Cultured TECs produce laminin, fibronectin and type IV collagen, fibroblasts and major histocompatibility complex (MHC) class II⁺ phagocytic cells of the thymic reticulum also produce these ECM components[27,28,29].

Fibronectin, recognized by VLA-5 (very late antigen-5) through its RGD motif, is located throughout the thymic parenchyma, whereas the isoform derived from alternative splicing of fibronectin mRNA (being recognized by VLA-4 through the LEDV motif) seems to be restricted to the medulla[30].

In addition, there are some isoforms of laminin. Laminin-5, formed from the $\alpha 3\beta 3\gamma 2$ heterotrimer, was detected also in the human thymus, being produced by medullary TECs and able to trigger outside-in signals to thymocytes[31].In the mouse system, anti-laminin-5 monoclonal antibodies significantly blocked thymocyte expansion, as well as double negative(DN) \rightarrow double positive (DP) differentiation[32].

In the human thymus, laminin-1 seems to be distributed widely within lobules and septae, whereas laminin-2 is restricted to cortical TECs.

Laminin-1 and laminin-2 can modulate thymocyte migration within TNC complexes and in addition to the DN \rightarrow DP progression, the migration of DP thymocytes is also influenced by laminin(s).

Cells of the thymic microenvironment express fibronectin receptor

(VLA-4 and VLA-5), as well as laminin receptors (VLA-3 and VLA-6) [33]. These receptors are also expressed by differentiating thymocytes. ECM proteins form molecular bridges between thymocytes and microenvironmental cells, a biological event that potentially provides signals to both cell types.

Thymocyte adhesion to cultured TECs is enhanced in the presence of ECM components, such as fibronectin and/or laminin, whereas antibodies against these adhesive molecules promote an opposite effect.

Thymocyte release from TNCs is enhanced by fibronectin and laminin and diminished by corresponding anti-ECM or anti-ECM receptor antibodies.

ECM-mediated epithelial-thymocyte interactions play a role in the traffic of thymocytes within TNCs, affecting both the entrance and exit of lymphocytes in this particular microenvironmental niche.

The expression of thymic ECM molecules in aging normal mice (50 week-old) is more intense than that observed in young animals.

In addition to a higher expression of basement membrane components in the medullary region, there is a formation of a cortical network of fibronectin, laminin and collagen type IV.

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1.1.5 Physiological function of the Thymus

The thymus supports the differentiation and selection of T cells. The thymic development of T cells consists of several processes that require the dynamic relocation of developing lymphocytes into, within and out of the multiple environments of the thymus trought distinct cell-cell and cell-matrix interactions. These processes include:1) the entry of lymphoid progenitor cells into the thymus; 2) the generation of CD4+CD8+ double-positive (DP) thymocytes at the outer cortex of the thymus;3) the positive and negative selection of DP thymocytes in the cortex; fourth, the interaction of positively selected thymocyte development and ensure central tolerance; 4) the export of mature T cells from the thymus [34-39] (FIG.11).

Chemokines might act in concert with extracellular matrix (ECM), resulting in the migration of a given cell subset, either within the thymus or at the entrance into and/or exit from the organ [40].



Fig. 11 Traffic of thymocytes for T-cell development and selection. \mathbf{a} | In the postnatal thymus, circulating Tlymphoid progenitor cells migrate into the thymic parenchyma through the vasculatures that are enriched around the corticomedullary junction. \mathbf{b} | The outward migration of CD4⁻CD8⁻ double-negative (DN) thymocytes to the capsule is regulated by chemokine signals through CXC- chemokine receptor 4 (CXCR4) and CC-chemokine receptor 7 (CCR7). \mathbf{c} | Further outward migration of the DN thymocytes to the subcapsular region is mediated by CCR9 signals. \mathbf{d} | CD4⁺CD8⁺ double-positive (DP) thymocytes generated in the outer cortex are motile, interacting with stromal cells that are localized in the cortex for positive and negative selection. \mathbf{e} | Positively selected DP thymocytes that gain the capability to survive and differentiate into CD4 or CD8 single- positive (SP) thymocytes show an increase in the surface expression of CCR7, through which the cells are attracted to the medulla, which expresses CCR7 ligands. \mathbf{f} | In the medulla, further selection of SP thymocytes express sphingosine-1phosphate receptor 1 (SP1)through which the cells are attracted back to the circulation that contains a high concentration of sphingosine-1-phosphate. cTEC, cortical thymic epithelial cell; DC, dendritic cell; mTEC, medullary thymic epithelial cell; PSGL1, platelet-selectin glycoprotein ligand 1.

The seeding of the thymus with lymphoid progenitor cells occurs as early as embryonic day 11.5 (E11.5) in mice at the eight week of gestation in humans [41-42], and is mediated by at least two different pathways: the vasculature-independent pathway, which probably occurs during the early stage of embryonic development before vascularization of the thymus, and the vasculature-dependent pathway, which probably occurs in the late stage of embryogenesis and postnatally after vascularization.

It is thought that vasculature-independent colonization of the fetal thymus is regulated by the chemotactic attraction of lymphoid progenitor cells to the thymic primordium (FIG. 12a). The partial but significant roles of two chemokines, CC-chemokine ligand 21 (CCL21) and CCL25, in this early stage of fetal thymus colonization have been reported. CCL21 and CCL25 are expressed by the fetal thymus primordium, along with several other chemokines[43-44]. Ccl25 and Cxcl12 regulate the hematopoietic precursor homing process [45]and have also chemotactic functions.

CCL21 and CCL25 are partially involved in fetal thymus colonization. CCL25 is expressed by TECs and dendritic cells and attracts immature thymocytes. The CC-chemokine receptor 9 (CCR9) receptor for CCL25 is expressed at various stages of murine thymocytes differentiation, particularly in DP and CD8+ single positive cells[40].

Interestingly, the entry of lymphoid progenitor cells into the thymus is not a continuous event but an intermittent and gated event that occurs in waves during embryogenesis and in adulthood[46-48]. During embryogenesis, distinct waves of thymus-colonized cells give rise to distinct progenies of $\gamma\delta$ T cells with different usages of T-cell receptor (TCR) -V γ and V δ chains, indicating that T-lymphoid progenitor cells that colonize the fetal thymus differ in developmental potential from Tlymphoid progenitor cells that enter the postnatal thymus[48-52].

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Following entry into the thymus, lymphoid progenitor cells begin their development into T cells through the developmental pathway that is commonly identified by the expression profiles of CD25 and CD44, until the developmental checkpoint at the CD4–CD8– CD25+CD44– (double-negative 3, DN3) stage[53-54].

Only the cells that succeed in in-frame rearrangement of the gene encoding the TCR β -chain are selected for further differentiation beyond this DN3 stage. The initial thymocyte development until the DN3 stage is promoted by Notch-mediated signals delivered by binding of Delta ligands [55-56] and is supported by signals delivered by interleukin-7 (IL-7) [57-58]: the signals derived from cortical thymic epithelial cells (cTECs) (FIG. 12c). Along this developmental pathway, immature DN thymocytes promote the differentiation of thymic stromal cells and trigger the formation of the cortical-epithelial environment in the thymus[59-62].

The differentiation of thymocytes from the DN1 stage to the DN3 stage regulates the differentiation of TEC precursor cells into cTECs that form the cortical environment in the thymus (FIG. 12b).

Concomitantly, DN thymocytes relocate outwards from the corticomedullary junction to the subcapsular region of the thymic cortex [63] (FIG. 11b). Several chemokine receptors, including CXCR4, CCR7 and CCR9, have been suggested to be involved in this movement of immature thymocytes[64-66].

The chemokine-dependent outward localization of the thymusseeded progenitor cells from the cortico-medullary junction to the outer cortex seems to be essential for optimal initiation of thymocyte development, although the accumulation of DN3 thymocytes in the subcapsular region might not be required for further thymocyte development.

In the thymic cortex, on their way to the subcapsular region, DN thymocytes begin to rearrange their *Tcrb* locus and the cells that succeed in generating the in-frame *Tcrb* rearrangement begin assembling TCR^β and pre-TCR α (pT α) chains to form the cell-surface pre-TCR complex[67-68]. The successful expression of the pre-TCR complex on the cell surface[68-69], along with the Delta-Notch interaction [70], initiates the signals for further development to DP thymocytes that express TCR $\alpha\beta$ antigen receptors. This is the first checkpoint of T-cell development at the DN3 stage which censors the cells that have succeeded in in-frame TCRB rearrangement and allows further development of thymocytes beyond the DN3 stage. It has been shown that the subcapsular region is rich in transforming growth factor- β (TGF β), which retards the cell-cycle progression of pre-DP thymocytes and negatively regulates the generation of DP thymocytes[71], indicating that the migration of pre-DP thymocytes to the subcapsular region might have a role in regulating the rate of production of DP thymocytes (FIG. 11c).

DP thymocytes that are newly generated in the cortex express low levels of the TCR $\alpha\beta$ antigen–receptor complex. The cortical DP-thymocyte population contains the unselected repertoire of T cells, making this population the main target of the most rigorous positive and negative selection[72-73].

Most thymocytes in the cortex of the adult thymus are highly motile and they pause to interact through their TCR with peptide–MHC complexes that are expressed by stromal cells, such as cTECs, and dendritic cells in the cortex[74]. Following TCR recognition of peptide– MHC ligands at low-avidity interactions, DP thymocytes are induced to receive signals for survival and further differentia- tion into single-positive (SP) thymocytes.

High-avidity interactions elicit signals that lead to the deletion of thymocytes. Apoptosis is the mechanism of negative selection. The process of negative selection contributes to the deletion of self-reactive T cells, thereby avoiding autoimmunity. In addition to the negatively selected thymocytes, most cortical DP thymocytes fail to receive TCR signals and are also destined to die at this stage. Therefore, only 3–5% of developing thymocytes survive this checkpoint of T-cell development at the cortical DP-thymocyte stage(FIG. 11d; FIG. 12d) [75-76].

Positively selected DP thymocytes then begin relocating from the cortex to the medulla[77]. The expression of the chemokine receptor CCR7 by developing thymocytes is associated with the phenotypic stage of cortex-to-medulla migration during the development of immature DP thymocytes to mature SP thymocytes[78-79]. Following TCR ligation, cortical DP thymocytes show an increase in cell-surface expression of CCR7. By contrast, CCR7 ligands (CCL19 and CCL21) in the postnatal thymus are predominantly produced by mTECs[80]. Therefore, cortical DP thymocytes that have received TCR-mediated signals are attracted to the medulla through CCR7-mediated chemotaxis (FIGS 11e).

However, the migration of developing thymocytes to the medulla cannot be solely mediated by the passive inward flow, present in the postnatal thymus from the capsular and subcapsular areas to the medullary area[81-82].

The positive selection of thymocytes is a prerequisite for both the formation of the medulla and the migration of positively selected thymocytes to the medulla (FIG. 12e).

The signals of the nuclear factor- κ B (NF- κ B)-mediated signal pathway, occuring through lymphotoxin- β receptor, tumour-necrosis-factor-receptor-associated factor 6 (TRAF6), NF- κ B-inducing kinase (NIK) and the NF- κ B subunit RelB are crucial for the development of the thymic medulla[83-86].

Positively selected DP thymocytes are induced to differentiate into SP (that is, CD4+CD8- or CD4-CD8+) thymocytes and relocate to the medulla. The SP thymocytes are assumed to spend approximately 12 days in the medulla before being exported from the thymus [87]. During this period, the SP thymocytes go through a maturation process that is commonly identified by the expression profiles of CD62 ligand (CD62L; also known as lymphocyte (L)-selectin) and CD69, and with the acquisition of functional capability of these cells as mature but naive T cells[88-90]. The newly generated SP thymocytes are CD62LlowCD69hi semi-mature cells that are func- tionally incompetent and susceptible to various apoptotic signals, including dexamethasone, and undergo further maturation to become mature SP thymocytes that are functional, dexamethasone-resistant and CD62LhiCD69low. The process of SPthymocyte maturation presumably occurs in the medulla and is accompanied by further deletion of self-reactive thymocytes that have escaped negative selection in the cortex (FIGS 11f). Such additional deletion in the medulla seems to be particularly important in establishing central tolerance to tissue- specific antigens, as mTECs express tissuespecific anti- gens promiscuously[91]. The expression of tissue-specific antigens by mTECs is at least partially dependent on the transcriptional factor autoimmune regulator (AIRE) [92-93].



FIG. 12 Crosstalk between thymocytes and thymic stromal cells. Thymic stromal cells, including the common progenitor thymic epithelial cells (pTECs), attract the entry of T-lymphoid progenitor cells to the thymus (signal **a**). The developing double-negative (DN) thymocytes are required for pTECs to generate cortical thymic epithelial cells (cTECs) (signal **b**) that form the cortical environment that is needed to promote the generation of double-positive (DP) thymocytes (signal **c**) and the positive selection of DP thymocytes (signal **d**). The generation of single-positive (SP) thymocytes by the positive selection of DP thymocytes is required for the development of mature medullary thymic epithelial cells (mTECs) (signal **e**) that form the medullary environment to support the maturation, further selection and export of mature SP thymocytes (signal **f**) to supply the peripheral T-cell pool. Red arrows indicate crosstalk signals.

This process of medullary tolerance is shown to occur, at least in part, through the deletion of self-reactive T cells[94-96].

The medulla is thought to be the place for the production of regulatory T cells[97-99]. It has been shown that most forkhead box P3 (FOXP3)- expressing regulatory T cells within the thymus are found in the medulla[98]. Therefore, the period of maturation in the medulla seems to be essential for SP thymocytes to establish central tolerance by ensuring the deletion of T cells that are reactive to tissue-specific antigens and by producing regulatory T cells (FIGS 12f).

Hassall's corpuscles express TSLP, which activates thymic dendritic cells that in turn induce the generation of regulatory T cells[100]. Independent studies have indicated that CCL17 is expressed by dendritic cells that are mostly localized in the medulla[100-101], and CCL22 is expressed by the outer walls of Hassall's corpuscles[102-103], whereas CCR4 (a receptor for CCL22 and CCL17) is expressed by semi- mature SP thymocytes[79,102,103]. These chemokine signals might direct medullary semi-mature SP thymocytes to Hassall's corpuscles and have a role in establishing central tolerance by enhancing the generation of regulatory T cells and/or optimizing negative selection.

In the postnatal thymus, lymphoid progenitor cells that have just entered the thymic parenchyma are found mainly in the area close to the cortico–medullary junction, where the vasculature is well developed, indicating that the lymphoid progenitor cells enter the postnatal thymus by transmigrating from the blood to the thymic parenchyma, mainly through the area around the cortico–medullary junction (FIG. 11a) [63].

Thymus seeding of the adult thymus is regulated by the adhesive interaction between platelet (P)-selectin glycoprotein ligand 1 (PSGL1), which is expressed by circulating lymphoid progenitor cells, and P-selectin, which is expressed by the thymic endothelium[104]. Thymocyte emigration is an active process controlled by signals mediated by G-protein-coupled receptors [105] phingosine-1-phosphate receptor 1 (of the S1P receptors, is expressed by mature SP thymocytes and that S1P is required for the egress of T cells from the adult thymus[106]. During maturation from DP thymocytes, SP thymocytes acquire the expression of S1P [106-107].

The chemoattraction of mature thymocytes to circulating S1P regulates the exit of mature T cells from the postnatal thymus(FIG. 11g). CCL19-mediated CCR7 signals contribute to thymocyte emigration in newborn mice but not in adult mice[108], indicating that there might be a developmental switch between CCR7-mediated emigration and S1P-mediated emigration during ontogeny. The repulsive movement of SP thymocytes repelled from a high con- centration of CXCL12 has a role in the emigration of T cells from artificial thymus organoids[109].

The export of mature thymocytes from the thymic parenchyma to the circulation is thought to occur through the perivascular space, which is channelled to post-capillary venules, arterioles and lymphatics[110-111].
1.2 The Natural Scaffold: the extracellular matrix

End-stage organ failure is a key challenge for the medical community because of the ageing population and the severe shortage of suitable donor organs available. It is a serious, growing, and costly issue. Every year in the USA alone, 120 000 people die from chronic lung disease[112], 112 000 die from kidney failure[113], 27 000 die from end-stage liver disease[114], and 425 000 die from coronary heart disease[115]. Equally, injuries to or congenital absence of complex tissues such as the trachea, oesophagus, or skeletal muscle have few therapeutic options.

At present, definitive treatment for end-stage organ failure is allogeneic transplantation. It is a well-established procedure but is a victim of its own success in that there is a growing disparity in numbers between the donor organ pool available for transplantation and the patients eligible for such a procedure, with high mortality rate in those who are on the waiting list[116].

Patients fortunate enough to receive a donor organ endure life-long immunosuppressive therapy with its associated morbidity and also are at risk of acute or chronic organ rejection[117].

Recent advances in tissue engineering, an interdisciplinary field that applies the principles of engineering and life sciences toward the development of functional replacement tissue for clinical use, and regenerative medicine have established a foundation on which the functional replacement of whole organs and complex tissues such as skeletal muscle, trachea, and oesophagus seems possible. Many developments employ the seeding and cultivation of cells onto acellular scaffolds[116]. These scaffolds can act as an inductive template for functional tissue and organ reconstruction after recellularisation with autologous stem cells or differentiated cells. This approach involves the use of naturally occurring extracellular matrix, obtained by the decellularisation of allogeneic or xenogeneic whole organs or tissues (Figure 13). The ready availability of an off-the-shelf xenogeneic scaffold that could subsequently be recellularised with autologous cells is a potential solution to the donor shortage that exists for allogeneic whole-organ transplantation. In addition, this approach would obviate the need for immunosuppression.



Fig: 13 Organ and tissue Bioengineering

An advantage of utilizing the ECM in its native state as a scaffold for tissue repair is the presence of all of the attendant growth factors (and their inhibitors) in the relative amounts that exist in nature and, perhaps most importantly, in their native three-dimensional ultrastructure. It is reasonable, therefore, to consider ECM scaffolds as temporary controlled-release vehicles for naturally occurring growth factors[118].

Another key advantage of this approach is the ready availability of an intact vascular network in the decellularised organs with appropriately sized inflow and outflow conduits for anastomosis to the recipient circulation and thus perfusion with nutrients and appropriate cues for cell behaviour.

The feasibility of a therapeutic strategy based on cellular repopulation of an intact extracellular matrix has evolved as a result of an understanding of improved cell-matrix interactions[119-120], development of methods for isolation of tissue-specific and organ-specific native extracellular matrix with little change in native structure and composition[121], rapid advancements of stem cell and progenitor cell biology (including the potential use of inducible pluripotent stem cells) [122-123], and the integration of the principles of developmental biology into regenerative medicine[120]. Although notable scientific and ethical challenges remain as this approach advances to clinical use, successful proof of principle for organs such as liver[124-125], heart[126], and lung [127-130]and complex tissues such as the trachea[131], oesophagus[132], and skeletal muscle[133] has been shown. Functional restoration of such tissues and organs with matrices might become a

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viable and practical therapeutic approach to meet future demand after organ failure[124-128,131].

Biologic scaffold materials composed of ECM are commonly used to facilitate the constructive remodelling of a variety of tissue both in preclinical animal studies and in human clinical applications[134-137].

The ECM possesses the ideal characteristics of the tissueengineered scaffold or biomaterial, being biocompatible, a supportive medium for blood vessels, nerves, and lymphatics and for the diffusion of nutrients from the blood, as well as being able to undergo constructive remodeling and degradation within the body's own systems[138-140].

All tissues and organs are made up of cells and associated extracellular matrix (ECM), a secreted product of the resident cells consisting of a unique, tissue-specific three-dimensional environment of structural and functional molecules. It is custom designed and manufactured by the resident cells of each tissue and organ and is in a state of dynamic equilibrium with its surrounding microenviromnment[141].

The extracellular matrix is a multicomponent structure, a dynamic protein network that provides cells with positional and environmental information, serving as a tissue-specific structural framework that controls cell function. The composition of ECM is represented by a complex mixture of functional and structural molecules that affect a variety of cell activities[142]. These molecules are arranged in a unique, tissue specific, three-dimensional ultrastructure and are ideally suited to the tissue or organ from which the ECM is harvested. The structure and the composition of the ECM are constantly changing in response to the

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current metabolic activity of the resident cell population, the mechanical demands of the tissue, and the prevailing microenvironmental niche conditions. This concept of "dynamic reciprocity" between the ECM and the resident cell population is a major advantage for the use of ECM scaffold materials and emphasize the importance of maintaining as much of the native composition and ultrastructure as possible during the preparation of these three-dimensional scaffolds. Matrix organization and its connection to intracellular pathways are important regulators of cell function, [143]defects in which can have significant pathological consequences [144].

Individual components of the ECM are common throughout most tissues such as collagen, laminin, fibronectin, and hyaluronic acid.

Collagen is the most abundant protein within the ECM. More than 20 distinct types of collagen have been identified and the most prevalent form found in mammalian tissues is type I collagen. Collagen has been well characterized and is ubiquitous across both the animal and plant kingdoms [145]. Collagen has maintained a highly conserved amino acid sequence through the course of evolution and for this reason allogeneic and xenogeneic sources of type I collagen have been long recognized as effective biologic scaffolds for tissue repair with low antigenic potential. Collagen types other than type I exist in the native ECM, altought in much lower quantities. These alternative collagen types each provide distinct mechanical and physical properties to the ECM and contribute to the utility of the intact ECM (as opposed to isolated components of the ECM) as a scaffold for tissue repair. By way of examples: type IV collagen is present within the basement membrane of all vascular structures and is an

important ligand for endothelial cells, type VII collagen is the principal component of the anchoring fibrils of keratinocytes to the underlying basement membrane of the epidermis, and type VI collagen functions as a connector of functional proteins and glycosaminoglycans to larger structural proteins such as type I collagen, helping to provide a gel-like consistency to the ECM. Type III collagen exists within selected submucosal ECMs, such as the submucosal ECM of the urinary bladder, where nonrigid structure is de- manded for appropriate function.Most collagen types cause differentiation of progenitor cell populations. It is common, therefore, to find little if any collagen in the ECM surrounding ESC populations.

Fibronectin represents an important noncollagenous component of the ECM. Fibronectin exists both in soluble and tissue isoforms and possesses many desirable properties of a tissue repair scaffold, including ligands for adhesion of many cell types[142,146]. In fact, this protein is the first example of a protein recognized to have dual functions, i.e., both struc- tural and functional properties. Fibronectin exists within the ECM of both submucosal structures and basement membrane structures [147,142]. Fibronectin, on the other hand, has been shown to play a central role in the development of vascular structures and can be found at an early stage in developing embryos. Fibronectin has also been found to be critical for the maintenance of hematopoietic stem cell activity ex vivo, which is another indication of the important role for this molecule in the support of the stem cell phenotype.

Laminin is a complex adhesion protein found within the ECM, especially within the basement membrane form of ECM[142]. This trimeric

crosslinked polypeptide exists in numerous forms dependent on the particular mixture of peptide chains[148,149]. The prominent role of laminin in the formation and maintenance of vascular structures is particularly noteworthy when considering the ECM as a scaffold for tissue repair[150,151] . Laminin can be found within basement membrane structures and is perhaps the best studied of the ECM protein found in ESC-derived embryoid bodies[152]. Laminin plays a prominent role in the formation and the maintenance of vascular structures and is among the first and most critical of the ECM components in the process of cell and tissue differentiation.

Glycosaminoglycans (GAGs) are important components of ECM and play important roles in the binding and preservation of growth factors and cytokines, water retention, and the gel properties of the ECM. The heparin binding properties of numerous cell surface receptors and of many growth factors (e.g., FGF family, VEGF) make the heparin-rich GAGs extremely desirable components of scaffolds for tissue repair[118].

The composition of the ECM includes a variety of bioactive molecules admixed with the binding molecules such as decorin and biglycan. Although cytokines and growth factors are present within ECM in extremely small quantities, these molecules act as potent modulators of cell behavior.

The ability of the ECM scaffold to facilitate and integrate cellular and external environmental cues is the consequence of its bioinductive properties, allowing the important constructive remodeling of tissue after the in vivo implantation of ECM scaffolds[153-155]. This remodeling cannot be solely attributed to characteristics such as viscoelastic

behavior, biomechanical properties, or host cell attachment through collagen, fibronectin, and laminin ligands within the ECM. Angiogenesis, abundant host cell infiltration, mitogenesis, and deposition and organization of new host ECM are common events during the remodeling of ECM scaffolds. Component growth factors such as vascular endothelial cell growth factor (VEGF), basic fibroblast growth factor (bFGF), and transforming growth factor beta (TGF-b) are released during scaffold degradation and exert their biologic effects as they are dissociated from their binding proteins and activated [156-161]. These growth factors promote angiogenesis, mitogenesis and cellular differentiation during the remodeling process. The rapid degradation of the native ECM scaffold material is mediated by enzymatic and cellular processes and may be considered as a mechanism for controlled release of the ECM constituent molecules. The degradation process itself, which is mediated by enzymatic and cellular processes, may be considered as a mechanism for controlled release of the ECM constituent molecules. Indeed, the process of scaffold degradation and growth factor release continue until the scaffold is completely degraded. Perhaps more importantly, degradation products of the parent molecules that constitute the ECM appear to mediate a subsequent series of remodeling events. Cryptic peptides released by the degradation process initiate and sustain the recruitment of circulating, bone-marrow-derived cells that actively parti- cipate in longterm tissue remodeling[162-163].

The phenotype of the resident cells, including their active genetic profile, proteome, and functionality, is influenced by conditions of the microenvironmental niche including factors such as oxygen concentration, pH, mechanical forces, and biochemical milieu. In turn, resident cells secrete the appropriate molecules in which to survive, function effectively, and communicate with neighbouring cells. Through this reciprocal interaction, the ECM has an essential role in prenatal development and postnatal maintenance of healthy function[119,120,164,165]. In addition to the matrix's role in development and adult homoeostasis, it can be used as an inductive scaffold to promote a constructive tissue remodelling response after injury. Mechanisms of this response include release of cryptic peptides that are mitogenic and chemotactic for endogenous stem and progenitor cells[166-167], modulation of the innate immune response[169], and provision of tissue-specific molecular cues that support cell phenotype and function[129,169-171].

1.2.1 Principles and methods of decellularisation

The preparation of a matrix from a large, intact mammalian organ requires several sophisticated processing steps. The selection of decellularization reagents is critically important because these agents can damage the structure and composition of the resultant matrix[172].

Several factors can affect the ability to decellularize tissues/organs which include: (a) the cellular density of a particular tissue or organ, (b) the specific density of the tissue/organ, (c) lipid content, (d) thickness of the tissue/organ and (e) properties of the selected decellularization agent(s) [121]. The most effective agents for decellularization of each tissue and organ will depend upon many factors, including the tissue's cellularity (e.g. liver vs. tendon), density (e.g. dermis vs. adipose tissue), lipid content (e.g. brain vs. urinary bladder), and thickness (e.g. dermis vs.

pericardium). It should be understood that every cell removal agent and method will alter ECM composition and cause some degree of ultrastructure disruption. Minimization of these undesirable effects rather than complete avoidance is the objective of decellularization.

Detergents, enzymes and chemicals used for whole organ decellularization have been used in decellularization for various organs. Since decellularization agents can also negatively impact the native ECM composition and structure during the decellularization process, selection of these agents, along with the development of new protocols, have been considered to be critical steps of the process. To achieve this goal, several different groups of decellularization agents have been explored and such agents include: (a) non-ionic (Triton-X), ionic (sodium dodecyl sulfate: SDS) and zwitterionic detergents, (b) enzymatic agents, (c) physical agents and (d) direct application of force[121].

- Chemical agents

Acids and bases cause or catalyze hydrolytic degradation of biomolecules. Peracetic acid is a common disinfection agent that doubles as a decellularization agent by removing residual nucleic acids with minimal effect on the ECM composition and structure[173-174] . Acetic acid damages and removes collagens with a corresponding reduction in ECM strength, but it does not affect sulfated glycosaminoglycans (sGAG) [175]. Bases (e.g. calcium hydroxide, sodium sulphide, and sodium hydroxide) are harsh enough and commonly used to remove hair from dermis samples during the early stages of decellularization[176-177] . However, bases can completely eliminate growth factors from the matrix and decrease ECM mechanical properties more significantly than

chemical and enzymatic agents[177]. The primary mechanism by which bases such as sodium hydroxide reduce mechanical properties is the cleavage of collagen fibrils and disruption of collagen crosslinks [178].

Hypertonic saline dissociates DNA from proteins [180]. Hypotonic solutions can readily cause cell lysis by simple osmotic effects with

minimal changes in matrix molecules and architecture . For maximum osmotic effect, it is common for the tissues to be immersed alternately in hyper- and hypotonic solutions through several cycles. Hypertonic and hypotonic solutions also help rinse cell residue from within tissue following lysis.

non-ionic, zwitterionic detergents lonic, and solubilize cell membranes and dissociate DNA from proteins, and they are therefore effective in removing cellular material from tissue [179-181]. However, these agents also disrupt and dissociate proteins in the ECM as evidenced by their use in protein extraction procedures in tissue proteomics [182-183]. The removal of ECM proteins and DNA by detergents increases with exposure time[184-185] and varies with organ subunit, tissue type, and donor age[186-188]. Combining multiple detergents increases ECM protein loss [187] but also allows for more complete detergent removal from ECM after decellularization [188-189]. Triton X-100 can effectively remove cell residues from thick tissues such as valve conduits where enzymatic and osmotic methods are insufficient, with concomitant ECM protein loss accompanied by decreased adverse immune response in vivo[191]. Sodium dodecyl sulfate (SDS) appears more effective than Triton X-100 for removing nuclei from dense tissues while preserving tissue mechanics [186,192]. The addition of a detergent such as SDS to a decellularization protocol can make the difference between complete and incomplete cell nuclei removal [193] but has the associated drawback of ultrastructure disruption[194-195] and growth factor elimination [177]. The zwitterionic detergent 3-[(3cholamidopropyl)dimethylammo- nio]-1-propanesulfonate (CHAPS) is most effective for cell removal from thinner tissues such as lung and may be ineffective for decellularization of thicker tissues, even when used in combination with SDS[197] or alone for relatively acellular tissues[185]. A blinded categorical comparison of detergents for peripheral nerve decellu- larization showed better preservation of ultrastructure by nonionic and zwitterionic detergents but better cell removal by ionic detergents[198]. However, the low concentration of cell bodies in peripheral nerve and the size of the tissue may limit the translation of these results to other tissues.

Care must be taken to flush residual chemicals from ECM after decellularization, particularly detergents such as SDS that penetrate into thick or dense tissues. Cytotoxicity is possible even at reduced agent concentrations and will inhibit or completely negate the beneficial properties of a cell-free ECM scaffold[199-200]. Even thin tissues such as valve leaflets require multiple (more than six) agitated washes to completely remove detergents[197].

Alcohols such as glycerol aid in tissue decellularization by dehydrating and lysing cells instead isopropanol, ethanol, and methanol are more effective than lipase in removing lipids from tissue and are capable of rendering adipose tissue lipid-free in a relatively brief period[201-202]. Methanol in combination with chloroform has been used during delipidation of tissues[176].. Caution should be used in treating tissues with alcohols such as ethanol and methanol due to their use as tissue fixatives in histology, their ability to precipitate proteins[203], and the damage they cause to ECM ultrastructure[178-202-203].

Acetone can also be used to remove lipids during decellularization[192]. However, like alcohols, the use of acetone as a tissue fixative [203] and its damage of ECM ultrastructure[204] warrant conservative use, especially for biological scaffolds used in load-bearing clinical applications[178]. In comparison to detergent treatments, acetone crosslinks ECM to produce stiffer scaffolds with mechanical properties further removed from those of native tissue[192].

-Biologic agents

Enzymes reported in tissue decellularization protocols include nucleases, trypsin, collagenase, lipase, dispase, thermolysin, and agalactosidase. Enzymes can provide high specificity for removal of cell residues or undesirable ECM constituents. However, complete cell removal by enzymatic treatment alone is difficult and enzyme residues may impair recellularization or evoke an adverse immune response.

Nucleases (e.g. DNases and RNases) cleave nucleic acid sequences and can therefore aid in removal of nucleotides after cell lysis in tissues[169,184,193,205]. Endonucleases such as benzonase[169] may be more effective than exonucleases because they cleave nucleotides mid-sequence and thereby more effectively fragment DNA in preparation for its removal. Likewise, non-restriction endonucleases will more effectively fragment DNA compared to their sequence-dependent

counterparts.

Trypsin is a serine protease commonly used as an enzymatic decellularization agent. However, ECM proteins such as collagens have limited resistance to trypsin cleavage[206] and tissue exposure to trypsin should therefore be used with caution. In comparison to detergents, trypsin is more disruptive to elastin and collagen and slower to remove cells but shows better preservation of GAG content[195, 188, 207,208]. Trypsin disruption of ECM can be correlated to changes in mechanical properties[208]. Removal of cells and ECM constituents by trypsin is time-dependent, and complete decellularization by trypsin alone may require lengthy incubation even for thinner tissues such as valve leaflets[209]. Trypsin can be used effectively to disrupt tissue ultrastructure and improve penetration of subsequent decellularization agents; therefore, exposure to trypsin as the initial step in a tissue decellularization protocol may be desirable or even necessary, particularly for complete removal of cell nuclei from dense tissues[184].

Collagenase may be used during decellularization, but only when ultrastructure preservation and maximum collagen retention are not critical to the intended clinical application of the resultant ECM.

Lipase aids in delipidation but is typically insufficient to remove all lipids when used alone[201-202].

Dispase and trypsin can be used successively to improve cell removal from thicker tissues such as dermis if used in combination with detergents, and repeated treatments with dispase may further improve decellularization[210]. Using an enzyme such as dispase or thermolysin as the sole decellularization agent is only effective for removing cells on the surface of a tissue and is likely to require mechanical abrasion for complete cell removal[211]. With regard to the underlying basement membrane and ECM, thermolysin is less disruptive compared to dispase[211].

Decellularized xenogeneic tissues can be treated with α galactosidase to reduce the immunogenic cell surface antigen galactose- α - (1,3)-galactose (Gal epitope) [212], although the immunomodulatory effect of Gal epitope does not adversely affect in vivo remodeling of xenogeneic ECM[213].

-Non-enzymatic agents

Chelating agents such as ethylenediaminetetraacetic acid (EDTA) and ethylene glycol tetraacetic acid (EGTA) aid in cell dissociation from ECM proteins by sequestering metal ions[214-215]. It is likely that chelating agents contribute to subtle disruptions in protein-protein interactions by the same mechanism[216]. Chelating agents alone are insufficient for superficial cell removal even with agitation, and they are therefore typically used in combination with enzymes such as trypsin[193, 184,205] or detergents[177,193,196,205]. The combined efficacy of chelating agents and simple hyper- or hypotonic solutions for decellularization is unknown[193,205,217].

Serum associates with nucleic acid fragments to aid in their removal from tissue, but it does not remove some immunogenic constituents such as phospholipids[197]. Additionally, xenogeneic serum has the disadvantage of introducing immunogenic constituents that may associate with the ECM, thus potentiating a downstream adverse host response. The use of serum in preparing ECM for clinical applications is therefore limited.

Serine protease inhibitors such as phenylmethylsulfonyl fluoride (PMSF) [193,201,217], aprotonin, and leupeptin[218] prevent undesirable damage to ECM that might otherwise result from release of intracellular proteases during cell lysis.

-Physical and miscellaneous agents

Freeze-thaw processing effectively lyses cells within tissues and organs, but the resulting membranous and intracellular contents remain unless removed by subsequent processing. A single freeze- thaw cycle can reduce adverse immune responses such as leukocyte infiltration in vascular ECM scaffolds[217]. Multiple freeze-thaw cycles may be used during decellularization [201-219]and do not significantly increase the loss of ECM proteins from tissue. The effect of freeze- thaw processing on mechanical properties is minimal for load- bearing, mechanically robust tissues[188].

Cells on the surface of a tissue or organ (e.g., urinary bladder, small intestine, skin, amnion) can be effectively removed by mechanical abrasion in combination with enzymes[211], hypertonic saline, or chelating agents, all of which facilitate dissociation of cells from their subjacent basement membrane. However, underlying ultrastructure and basement membrane integrity are invariably damaged by any direct application of mechanical force[211].

Hydrostatic pressure requires relatively little time and can be more effective than detergents or enzymes for removing cells from blood vessel and corneal tissues, although the baric formation of ice crystals may disrupt ECM ultrastructure. [205] Increased temperature during pressure decellularization prevents ice crystal formation[205] but may disrupt ECM due to the associated increase in entropy, which can be mitigated by colloids such as dextran.

Agent/Technique	Mode of action	Effects on ECM
Chemical Agents	and a second second second	
Acids and bases	Solubilizms cytoplasmic	May damage collagen.
	components of cells.	GAG, and growth factors
	disrupts michaic acids, trend	
	In departure proteins	
Divostonic and	Cell hais by perpetic shock	Effectively losses cells, but does not effectively
Investigate adultant	discost DMB, contain interactions	remain cellular anidure
ubble mear and more	mucht over brother spiratrous	Termine central resolution
Non-ionic detergents	Disrupt DNA-protein interactions,	
	disrupt lipid-lipid and	
	linid contein interactions	
	and to a lesser degree	
	pentele-protein interactions	
- Triton X-100	because become association	Mixed sesuits with efficacy dependent on tissue
		more effective cell removal from this tissues.
		some disputting of all rationating and removal
		of CAC, best effective than \$26
lacic determents	Solubilize cell and nucleic membranes	on on the series concerned control boost
and a second second	tend to denature proteins	
- Sodiare dodersi	tions to decision provident	Effectively permoves nuclear remnants and
will de (NIN)		cutoolaumic nonteins from Aeros tissues
manufact (specify		bands to discuss of matricellum company CAP
		and mounth further and damages collision
Rodings descriptions		Allowed another with affective dependent on three
- Sodiara deorgenolate		when mous will employ dependent of time
		thickness, some disruption of ultrastructure
		and removal of CAG
Trime X 200		More effectively remains cells from this
- 101000 A-200		Hours but with meater discertion of
		classes and wate greater complete of
Partmenter in Astronomics	Publishis proposation of some locality	senanoacture comparen so ouser netergents
ownineer oregens	and louis determents	
- CHAPS	and once or rear from	Effectively removes cells with mild disputtion
		of ultrastructure in this tissues
- Sulfabetaine-10 and		Effectively removes cells with mild domestion.
10 /08,10 58,167		of ultrastructure in this tissues
Coharate		AN ARTICLE DURING AN ARTICLE
Hanhala	Out have be debudention	Effectively semicory role from dance down
- Alcohota	can ryan by detiyoration,	LINULAWEY PERSONEN CEUM STOLEN CHEMICAL
	sonation and remove against	tissues and mactivates pyrogens, out
		crussions and precipitates proteins,
	COLUMN AND AND	including collagen
- Acetone	cell tysis by deligitration,	Effectively removes cells from dense dense
	solutilizes and certoves lipids	tissues and inactivates pyrogens, but
		crosolinks and precipitates proteins,
		including collagen
 Tributyl phosphate (TBP) 	Forms stable complexes with metals,	Mixed results with efficacy dependent on tissue.
	disrupts protein-protein interactions	dense tissues lost collages but impact on
		mechanical properties was minimal

Biologic Agents		
- Nucleuses	Gatalyze the hydrolysis of ribonacleotide and deoxyribonacleotide chains	Difficult to remove from the tissue, could invoke an immune response
- Trypsin	Cleaves peptide bonds on the C-side of Arg and Lys	Prolonged exposure can disrupt ECM ultrastructure, removes ECM constituents such as collagen, laminin, Ebronectin, elastin, and GAG, slower removal of GAG compared to detregents
- Dispase	Cleaves specific peptides, mainly fibronectin and collagen IV	Prolonged exposure can disrupt DCM ultrastructure, removes DCM components such as fibronectin and collagen IV
Chelating Agents (EDITA, EGTA)	Chelating agents bind metallic ions. thereby disrupting cell adhesion to ECM	Typically used with enzymatic methods (e.g. trypsin) but can be used with other agents, ineffective when used alone
Physical and Miscelianeous Agent		
Temperature (freezing and thawing)	Intracellular lor crystals discupt cell membrane	for crystal formation can disrupt or fracture ECM
Direct application of force	Removal of tissue eliminates cells and force can burst remaining cells	Force can directly damage ECM
Pressure	Porssure can burst cells and aid in removal of cellular material	Pressure can disrupt BCM
Dectroporation	Pulsed electrical fields discupt cell membranes	Electrical field oscillation can disrupt DCM

Fig. 14 scheme of the different agents used in the decellularization protocols.

Unlike agitation and diffusion-only techniques used for the decellularization of simple and thin tissues, solid organs require dynamic decellularization techniques in order to efficiently remove cellular components from the entire organ system. In particular, the mechanical forces incorporated with dynamic decellularization techniques are important for efficient decellularization. Such mechanical parameters can be pressure, shear stress and flow rate of decellularization agents. Different levels of mechanical forces need to be applied to decellularization processes depending on the size and types of organs. Antegrade and retrograde perfusion techniques of the decellularization agent(s) through the native vascular system of the organ has been the most commonly used method to produce acellular whole organ scaffolds[169,172,177,184]. In this approach, perfused decellularizing agent(s) are evenly distributed throughout the organ and, in addition, the application of a certain amount of pressures during the perfusion process has shown to effectively achieve tissue permeation from large blood vessels down to the capillary level. Perfusion of decellularization solution also allows cellular debris to be excreted through the venous system. [172]

1.1.2 Characterization of the decellularized tissue

After the decellularization process, it is necessary to evaluate the resulting scaffolds to determine the extent of decellularization and whether or not the scaffold is damaged. Assays used for this purpose include qualitative and quantitative assays. Qualitative evaluation involves demonstrating that cellular material has in fact been removed from the matrix as well as demonstrating that the main ECM proteins have been retained. This is performed using several different histological and immunofluorescent methods. The removal of cellular components by the decellularization process can be confirmed by observing the lack of visible nuclear material in tissue sections stained with 4, 6-diamidino- 2phenylindole (DAPI) or hematoxylin & eosin (H&E). The retention of ECM components is visualized by several staining methods, such as Masson's trichrome and Herovici staining for collagens, Alcian blue/PAS staining for glycosoaminoglycans (GAGs) and Movat's staining for collagen and elastin fibers. In addition, electron microscopic studies using a scanning electron microscope (SEM) and/or transmission electron microscope (TEM) are often conducted to examine the decellularized scaffolds on a very fine scale to identify ultrastructure within the scaffolds[121].

However, these techniques are not sensitive enough to identify and quantitatively analyze the residual DNA content within the ECM[176]. The presence of DNA is directly correlated to adverse host reactions[188,191].

Hence, interest in the quantification of the residual nucleic material after decellularization has been justified[176]. Based upon findings from in vivo studies in terms of constructive remodeling and adverse host responses using two-dimensional decellularized tissues, the following minimal criteria for acceptable amounts of residual DNA after decellularization have been created[176]: (a) <50 ng double strand DNA per mg dry weight, and (b) <200 bp DNA fragment length[188].

Likewise, the biological activities of ECM components need to be considered in the process of whole organ bioengineering. Following the decellularization of solid organs, the resulting acellular organ scaffolds will be combined with a cellularization technique for further application in in vivo studies, and the biological activities of many matrix proteins are required for the recellularization step to be successful. First, cell adhesion to the acellular organ matrix is a pre-requisite step for the recellularization process. In many cases, natural cell-adhesion domain sequences in the matrix (e.g., RGD) [204] can bind with members of the integrin family, which are expressed on many cell types. This binding is crucial to allow cells to adhere to acellular organ matrices. These types of cell adhesion peptide motifs exist on ECM components such as collagen and fibronectin. Collagen is particularly important because it retains the RGD motif as well as bound growth factors. Fibronectin is a representative glycoprotein in ECM that plays roles in cell adhesion, growth, migration and differentiation [195,203].

The presence of collagen, fibronectin and laminin is crucial for cells seeded onto a decellularized organ to attach to the matrix and begin recellularizing the organ.

Once seeded cells are attached to the ECM components within acellular scaffolds, the behaviors of the cells will be influenced by growth factors and bioactive molecules that confer inherent bioactivity within the matrix. Generally, growth factors easily degrade after secretion from the cells that produce them. However, within the ECM, GAGs are critically important in the protection of free growth factors from proteolytic degradation[196]. Due to the existence of heparan sulfate residues on GAGs and the low isoelectric point (IEP \approx 5) of heparan sulfate, GAGs are able to bind growth factors that have an IEP of ≈10, such as fibroblast growth factor-2 (FGF-2), vascular endothelial growth factor (VEGF), transforming growth factor- β (TGF- β) and platelet-derived growth factor (PDGF). Such growth factors become anchored to GAGs and can thus be protected from enzymatic degradation. This process allows them to retain biological activity, and they can affect cells that bind the various ECM compounds during the recellularization of a decellularized organ scaffold. Another role of GAGs for the recellularization process is of particular importance in morphogenesis which promotes cellular adhesion permissive cell shape changes and motility as well as the maintenance of original phenotypes of repopulated cells[221]. Hyaluronic acid (HA), one subset of GAGs, is such a molecule. HA is considered to be important for several aspects of cellular behaviors such as the direction of cellular shape changes and the induction of cytoskeletal rearrangements within cells. However, during current decellularization processes, 50 to 90% amounts of GAGs in various organs [222,223] were reported to be removed when compared to that of original organs. The reduction of GAG amounts may cause the inactivation of endogenous growth factors and

also adversely influence the efficient recellularization process due to lack of primary components of GAGs. Further investigations may be needed into the modification of decellularization method for the maintenance of GAG architectures after decellularization.

Just as solid organs have an intricate vascular tree within them, acellular organ scaffolds should maintain continuous and intact vascular structures after the decellularization process. This vascular matrix can be used to reseed the matrix for efficient whole organ bioengineering. This vascular system can be used as a template for cell seeding with organspecific types of parenchymal cells or it can be directly seeded with endothelial cells. This is especially important because a continuous and functional re-endothelialized vascular system within a recellularized whole organ can reduce thrombosis and blood clots when the organ is integrated with the vasculature of the host during the implantation procedure.

1.2.3 State of the art of the different decellularization scaffolds

Intact three-dimensional extracellular matrix scaffolds from different allogeneic and xenogeneic tissues and organs have been manufactured effectively for regenerative medicine.



Fig.15 Some reported perfusion protocols for decellularization of heart, liver and lung. Abbreviations: DMEM, Dulbeccos's Modified Eagle's Medium; EDTA, ethylene diamine traacetic acid; FBS, fetal bovine serum; PBS, phosphate buffered saline; SDS, sodium dodecyl sulfate.

The heart

The first report in which whole-organ decellularization was successfully performed was on the heart[222]. The aorta of a rat heart was cannulated to allow retrograde coronary perfusion with heparinized phosphate-buffered saline with adenosine, 1% sodium dodecyl sulfate (SDS), and 1% Triton X-100 in sequence alternating with rinsing with

deionized water between steps. This produced a decellularized scaffold that was structurally similar to the heart, but had a translucent white appearance throughout (Fig. 16). Reimplantation of the acellular heart scaffold onto the aorta demonstrated an intact and patent vascular network .

Recellularization of the heart scaffold with cardiomyocytes under electro- physiological stimulation formed a construct capable of muscular contraction[126].



Fig. 16 The heart in stages of increasing decellularization over time with 1% sodium dodecyl sulfate as the decellularizing agent, with the end result of a translucent ap- pearing heart-like scaffold structure. The panel on the right shows histological evidence (H&E staining) of decellu- larization. Scale bar = 200 mm.

The lungs

A number of groups have investigated the use of perfusion decellularization in the lungs[128,129,194,223,224], which contain two accessible compartments separated by a short diffusion distance, that is, the vascular and the airway systems, by using varying protocols of vascular perfusion alone [194,224], or in combination with endotracheal instillation[128,223], or perfusion of the airway compartment[129]. The decellularizing agents used were also far more varied from protocol to protocol, from SDS at different concentrations [129,198,224] to 8mM of Triton CHAPS[128], to а combination X-100/deoxycholate/ DNAse/bleach[223]. All groups demonstrated preservation of the major components of the ECM and the micro- and macroarchitecture of the lung.

Recellularization was achieved with all the reports, although Petersen et al. mimicked physiological conditions within the bioreactor by ventilation of the airway compartment at negative pressure at 1 breath/min, and maintained pulmonary artery pressure at 20 mmHg or below. Orthotopic implantation of these tissue-engineered lung constructs demonstrated active gas exchange occurring via sampling of pulmonary arterial and venous blood.

However, results in this study and Ott et al. showed vascular leakage and pulmonary edema after a few hours of implantation, indicating damage to the microvasculature from the decellularization process. In a follow-up study to Ott et al., Song et al. demonstrated further optimization of graft preservation and oxygenative function postimplantation for as long as 7 days[224]. Recellularization was performed with mixtures of whole-lung cell isolates in four cases[128,194,223,224], but one group used a homogeneous mouse embryonic stem cell population[129], which showed that the matrix was capable of promoting site-appropriate differentiation without any other specific differentiation cues.



Fig. 17 Perfusion decellularization of whole rat lungs. (a) Photographs of a cadaveric rat lung, mounted on a decellularization apparatus allowing antegrade pulmonary arterial perfusion. pa, pulmonary artery; pv, pulmonary vein; tr, trachea; RUL, right upper lobe; RML, right middle lobe; LL, left lobe. Freshly isolated lung (left), after 60 min of SDS perfusion (middle), and after 120 min of SDS perfusion (right). The lung becomes more translucent as cellular material is washed out first from apical segments, then from the middle segments and finally from the basal segments.

The liver

Liver decellularization has been performed by antegrade perfusion via the portal vein with SDS and Triton X-100, either alone or in combination [124.] 225-227]. although another study used trypsin/EGTA[185], and Baptista et al. used Triton X-100 in combination with ammonium hydroxide [177, 228]. Decellularization was demonstrated with histology and evidence of DNA removal, and retention of key ECM constituents with preservation of microvasculature and ECM ultrastructure. Hepatocytes were seeded onto the acellular liver scaffold via portal vein perfusion and were shown to be functional in four studies [177,225,227-228]bv showing evidence of synthesis of lactate dehydrogenase, albumin, and urea after heterotopic implantation. In particular, two studies attempt to scale-up this approach to further approximate human- scale engineered organ constructs by use of ferret[177] and porcine[227] livers, and both studies seeded the bioscaffolds with human liver cells, which were shown to be compatible on these xenogeneic scaffolds.



Fig. 18 Decellularization of ischemic rat livers. (a–e) Representative images of ischemic rat livers during decellularization process at 0 h (a), 18 h (b), 48 h (c), 52 h (d) and 72 h (e).

The kidney

Several groups have decellularized whole kidneys by the perfusion method with preservation of the vascular network and complete cell removal[228-230]. Decellularizing agents vary from 1% Triton or SDS[228-229] to 3% Triton with DNase and additional 4% SDS[230]. The preservation and presence of ECM constituents such as laminin and collagen IV were also demonstrated, as well as collagen I, fibronectin, and heparin sulfate in a separate study, which did not employ perfusion[208]. The latter study juxtaposed fetal cells in fresh renal explants with the ECM scaffolds by layering fetal kidney tissue against the ECM, which supported developmental renal phenotypes in these cells after they migrated into the ECM. In the study by Ross et al., decellularized rat kidneys scaffolds were reseeded with murine embryonic stem cells, and the renal ECM was shown to support the renal differentiation of the embryonic stem cells.

The last but not the least Ott et al. decellularized rat, porcine and human kidneys by detergent perfusion, yielding acellular scaffolds with vascular, cortical and medullary architecture, a collecting system and ureters.

To regenerate functional tissue, they seeded rat kidney scaffolds with epithelial and endothelial cells and perfused these cell- seeded constructs in a whole-organ bioreactor. The resulting grafts produced rudimentary urine *in vitro* when perfused through their intrinsic vascular bed. When transplanted in an orthotopic position in rat, the grafts were perfused by the recipient's circulation and produced urine through the ureteral conduit *in vivo*. [231]

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Fig. 19 Perfusion decellularization of whole rat kidneys. (a) Time-lapse photographs of a cadaveric rat kidney undergoing antegrade renal arterial perfusion decellularization. Shown are a freshly isolated kidney (left) and the same kidney after 6 h (middle) and 12 h (right) of SDS perfusion. Ra, renal artery;Rv, renal vein; U, ureter.

Trachea

Macchiarini et al. repopulated a decellularized human windpipe in a bioreactor whit cultured auologus respiratory ephitelial cells and autologus chondrocytes of bone marrow-derived mesenchymal stromal cell origin, and used the resultant graft to replace a terminally diseased left main bronchus.

For the decellularization technique they used 25 cycle of distilled water for 72 h, then incubated in 4% sodium deoxycholate and 2000 kU deoxyribonuclease I in 1 mmol/L sodium chloride (Sigma Chemicals, Barcelona, Spain). [131]



Fig. 20 The graft obtained after the decellularization process and immediately before the implantation.

SIS (small intestinal mucosa)

Totonelli et al in 2012, using a detergent-enzymatic treatment (DET), optimized in rats a protocol that creates a natural intestinal scaffold, as a base for developing functional intestinal tissue. After 1 cycle of DET, histological examination and SEM and TEM analyses showed removal of cellular elements with preservation of the native architecture and connective tissue components. This DET consists of deionized water at 4 °C for 24 h, 4% sodium deoxycholate (Sigma) at room temperature (RT) for 4 h, and 2000kU DNase-I (Sigma) in 1 M NaCI (Sigma) at RT for 3 h, as previously descrive also for trachea. Maintenance of biomechanical, adhesion and angiogenic properties were also demonstrated strengthen the idea that matrices obtained using DET may represent a valid support for intestinal regeneration[131].



Fig.21 Decellularization of rat small intestine with detergent-enzymatic treatment. Macroscopic pictures prior (A) and following (B) one cycle of decellularization.

Thyroid

Working in the laboratory of Regenerative Morphology and Bioartificial Structures at the S.Bi.BI.T Department of the University of Parma, under the supervision of Prof. Roberto Toni, I have been engaged in developing a protocol for decellularization of the rat thyroid. We settled a protocol based on freeze and thaw, 0.02% of trypsin, 3% of Triton X-100 in PBS, 4% of Sodium deoxycholate in PBS and final sterilization with Peracetic Acid. After tissue process the structure of the matrix remained intact , in terms of 3D-structure, and at the same time the matrix obtained was very well cleaned in DNA.

We also collected preliminary results on the recellularization of this scaffold, using a culture model with cells derived from the same animal source, the rat thyroid. As a result, we observed that after 7 days of culture the thyroid matrix provided attachement of the seeded cells, and maintenance of their differentiation state [233].



Fig.22 Decellularization of rat thyroid with detergent-enzymatic treatment. Macroscopic pictures prior in situ (A) and ex situ (B) following (C) decellularization protocol.

1.3 Culture of thymic epithelial cells

Culturing thymic epithelial cells in vitro, freshly obtained from the tissue, is a problem of substantial interest.

In vivo, Thymic ephitelial cells undergo a rapid expansion during the early stages of thymus development, but as lymphopoiesis progresses, the more mature ephitelial cells exit the cell cycle to leave only thymic ephitelial stem cell pool.

Monolayer cultures of thymic stromal cells have been used for in vitro assays of stromal cell function. Immortalization techniques have been used in conjuction with monolayer culture to generate long term stromal cell lines and it has been shown that monolayer cultures are capable of inducing the maturation of CD4⁻ CD8⁺ CD3⁻ cells from their CD4⁻ CD8⁻ CD3⁻ precursors.

Thymic ephitelial cell lines have also been used to induce a limited growth and differentiation of early thymic precursors cells in coculture.

A number of studies have shown that thymic stromal cells cultured as monolayer rapidly loose their ability to support T-cell development, suggesting a change in their normal phenotype.

They change their normal phenotype in a serum free medium, altought fetal calf serum or human serum has been used as a necessary constituent of the culture medium but this has several disadvantages such as the addiction of unknown factors (hormones, endotoxins, viruses and other macromolecules) [234], which may vary from batch to batch. This creates problems of reproducibility but also makes it difficult to define growth requirements for the cells, and to isolate and figure out the meaning of biological activities executed by molecules released by the cells into the culture medium [235].

It has been reported that for growing of murine TEC is essential the ephidermal growth factor, insulin and hydrocortisone [236] instead, for human TEC, it's important the supplement of selenium, transferrin and insulin to the culture media. [237-238]

There are also some different way to culture thymic ephitelial cells using not only a specific media but doing coculture with other cell type that could support TEC.

In 1986, Ehmann at al., cultured TEC, using a tissue culture plastic, were plated togheter with letally irradiated cells of the LA7 rat mammary tumor line that gave a support for growing. Ephitelial cells from the normal mouse thymus could be cultured and subcultured for at least 8 passages in this way. As the irradiated LA7 cells slowly decreased in number TEC proliferated concomitantly to form a confluent monolayer like show in figure 23. [239]



Fig.23 An image of a colony of distinct thymus cells with LA7 cells near the top of the image. Magnification X89 .

1.3.1 Three dimensional culture of thymic epithelial cells

Culturing thymic ephitelial cells in vitro preserves the ability of these cells to promote thymocytes maturation so long as they are cultured as 3-dimensional aggregates. The 3-dimensional structure of the thymic ephitelial cells is essential to maintain their function within the intact thymus.

The correct 3-dimensional structure therefore appears to be a prerequisite for full ephitelial cell function in vitro since the 3D thymic architecture contributes to critical cell-cell interactions.

3D matrices were created by culturing thymic fragments on the surface of disks of CellFoam (Cytomatrix) and this resulted in the colonization with thymic stromal cells reaching 80% of confluency [240].

This matrices were then used with the addition of progenitor cells CD34⁺ or AC133⁺ and a coculture for 14 days demonstrated that these matrices were capable to induce a differentiation in the progenitor cells and they were able to high proliferate in the presence of interleukin 2 and phytohemagglutinin [241].

There was also some way of doing a coculture in a 3D structure like a three dimensional organotypic coculture model. This was used for the culture of isolated mTECs to mimic a dermal equivalent in vitro. It consisted of dermal fibroblast that were embedded in a inert scaffold consistuted by fibrin gel like shown in Fig. mTEC seeded on top of this fibrin gel proliferated and underwent terminal differentiation. This three dimensional model preserves key features of mTECs: proliferation and terminal differentiation of CD80^{lo}, Aire⁻ mTECs into CD80^{lo}, Aire⁺ mTECs, responsiveness to RANKL and sustained expression of FoxN1, Aire and tissue-restricted genes into CD80^{hi} mTECs [242].



Fig.24 A schematic rapresentation of the 3D organotypic coculture.

Less it has been done for the reconstruction of the thymus, in vitro and then in vivo, since it has been demonstrated that after injection of IL-22 after chemioterapic treatment is enough in the regeneration of thymic epithelium.

IL-22 has been identified as an important player in the thymic regeneration. It appears to be mainly produced by thymic innate lymphoid cells in response to tissue damage, and this process is at least partly regulated by dendritic cells throught the production of IL-23. The receptor for IL-22 is expressed on TEC and seems to induce their survival and proliferation[243].

In 2002, Gill et al, identified a population of MT24+ cells, a subpopulation of TEC. These cells contained epithelial progenitor cells that were competent and sufficient to fully reconstitute the complex thymic ephitelial microenvironment that supported normal T cell development. This population has been cultured for 24 hours and then, after reaggregation, it has been implanted under the kidney capsule of congenic recipient mice. These reaggregates were able, after 3 weeks, to induce the maturation of pre T-lymphocytes in a comparable situation to
the normal thymic tissue[241].

In 2012, Piccinini et al, developed an in vitro 3D module to culture TEC. They were cultured in different 3D shape with a scaffold made up fibrin and they identified like a best shape a simple drop in which cells are embedded. In this configuration TEC were able to maintain their differentiated state like underlined by the expression of a series of marker that were expressed in a comparable quantity to the in vivo situation. They maintened their phenotype after 14 days in vitro and, after coculture with pre-T lymphocytes, they were able also to induce the maturation of them in a more differentiated phenotype[244].

2.AIM OF THE THESIS

2. AIM

The aim of my experimental work has been to develop an innovative technique to reproduce in vitro the 3D thymic microenvironment, to be eventually used for growth and differentiation of thymocytes, and possible transplantation replacement in conditions of depressed thymic immune regulation. The work has been developed in the laboratory of Tissue Engineering at the University Hospital in Basel, Switzerland, under the tutorship of Prof.Ivan Martin, Director of this research Center.

Since a number of studies have suggested that the 3D structure of the thymic microenvironment might play a key role in regulating the survival and functional competence of thymocytes, I've focused my effort on the isolation and purification of the extracellular matrix of the mouse thymus, choosen as a suitable research model. Specifically, based on the assumption that TEC can favour the differentiation of pre-T lymphocytes, I've developed a specific decellularization protocol to obtain the intact, and DNA-free extracellular matrix of the adult mouse thymus. In this manner, I have tried to prevent the cytotoxic effect of contaminating nucleic acids on seeded TECs, and to provide a functional 3D structure rich in intrinsic proteins for TEC growth and assembly.

This 3D matrix, then, has been used as a natural scaffold for culturing TEC in static conditions. In particular, I have analyzed a number of surface markers indicating TEC capability to maintain their own phenotype, and eventually promote the differentiation of pre-T lymphocytes, similar to what previously demonstrated with other 3D structure like the fibrin gel.

3.MATERIALS AND METHODS

3.1 TECs isolation

3.1.1 Thymi digestion

TECs are isolated from B6 mice, 4-6 weeks old.

Thymus isolation and cleaning is performed on the lab bench, while the digestion is performed in sterile conditions under the hood.

Mice are sacrificed by cutting the head to avoid CO2 and excessive bleeding in the thymic area.

The mouse is open cutting the peritoneum, cutting the membrane separating the chest from the peritoneum, cutting the ribs and exposing heart and thymus.

The thymus is isolated by pulling it, without cutting, and after it is placed in a Petri dish on ice containing PBS + 2%Fetal Bovine Serum.

After the explant and before the digestion, Thymi are cleaned, under the microscope, by fat and capsule to facilitate enzymes' action in a solution of

The thymi are cleaned under the microscope. First lobes are separated and then each lobe is cleaned from fat and connective tissue. Finally to favor the digestion small holes are done in the capsule.

The digestion is carried out in a water bath at 37°C in an enzymatic solution made of PBS, Liberase (Product Number:05401127001 Roche Diagnostic) and DNase (Product Number:11284932001 Roche Diagnostic).

For 6 thymi the solution is made of:

Liberase: 50mg/20 ml H2O sterile 500µl DNase: 10mg/1 ml H2O sterile 50µl PBS: 5,5 ml

The thymi are left at 37°C for ten minutes. Then the solution with pieces are pipetted up and down with a tip of $1000\mu l$, cut before to favor the digestion. This is done for two times.

The pellet is allowed to sit down, the supernatant with cells is collected and enzymes are inactivated added to them IMDM+ 2%Fetal Bovine Serum.

3ml of the digest solution is added to pellet and left at 37°C for ten minutes. Then the solution with pieces are pipetted up and down with a tip of $1000\mu l$, cut before to favor the digestion. This is done for two times.

After these steps all pieces are digested and enzymes are inactivated added to solution IMDM+ 2%Fetal Bovine Serum.

Two solutions, coming from first and second digestion respectively, are pooled down together at 1500 rpm 5'.

Cells are resuspended in 10ml of PBS + 2%Fetal Bovine Serum and filtered through a 70 μm strainer.

 $10\mu l$ of the solution is used to count cells at the Cell Coulter.

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Fig.25 An image of the cell coulter.

For the counting $10\mu l$ of solution with cells is added to 10ml of PBS and 3 drops of Zap-oglobin.

The result obtained on Cell Coulter is multiplied per 2 and per 10^3 per the ml of the solution.

The solution is adjusted to bring cells to a concentration of 100×10^{6} cells.

3.1.2 CD45 depletion

The cells' solution freshly obtained is used for CD45 depletion. CD45 beads (Product Number:130-052-301 Miltenyi Biotec) are added to cells at a concentration of $5\mu l/10^7$ cells.

The solution is left 15' at 4°C in the dark.

After that cells are washed 2 times with PBS + 2%Fetal Bovine Serum spinning down at 1500rpm 5'.

Cells are filtered through a 70 μm strainer and then distributed in FACS tubes (2ml each tube) after another filtration through a 45 μm

strainer attached to a 2,5 ml syringe.



Fig.26 A image prior the filtration of the cell suspension.

The enrichment can start with the AutoMACS Pro Separator System(Miltenyi Biotec) using the protocol **Deplet_s + QRinse**.



Fig.27 A image of the AutoMacs used for the CD45 depletion.

Cells of interest will be in the CD45⁻ fraction. These cells are spinned down at 1500rpm 5' and the pellet is resuspeded in 1ml of

Iscove's Modified Dulbecco's Medium(IMDM) (Gibco; Invitrogen) containing 10% FCS, β -mercaptoethanol and Kanmycin (Gibco; Invitrogen) and again counted as described above.

3.1.3 TECs purity

75 μ l of total cells obtained are used to check the purity.

The 75 μ l of solution are brought to 1ml with PBS + 2% FBS. To the tube are added directly the two antibodies:

EpCam Alexa 647(1:2000) (Product Number:118211 BioLegend) CD45 Alexa 700 (1:20) (Product Number:103127 BioLegend)

The solution is incubated 15' a 4°C, then centrifugated and resuspended in $300\mu l$ of PBS + 2% FBS. A filtration is performed through a 45 μm strainer attached to a 2,5 ml syringe and then $100\mu l$ of DAPI (Product Number:422801 BioLegend) are added and all is acquired at FACS Aria.



Fig.28 FACS Aria used for FACS analysis.

TECs purity usually goes from 20% to 30% of total cells obtained.

3.1.4 Culture medium and supplements

TEC were cultured in CnT-57 cell culture medium (CELLnTEC, Bern, Switzerland). CnT-57 is a progenitor cell target liquid culture medium with low calcium concentration (0.07mM) and low concentrated Bovine Pituitary extract (BPE) (6μ gr/ml). CnT-57 is a protein free medium containing a variety of amino acids, minerals, vitamins and other factors such as Fibroblast Grow Factor (FGF), Epidermal Growth Factor (EGF), and Insulin.

Additionally, the culture medium was supplemented with 10nM dihydrochloride monohydrate, a cell permeable anti-apoptotic selective Rho-associated Kinase (ROCK) Inhibitor (10nM; Calbiochem, Merk) to enhance viability. Kanamycin was supplemented to the medium to reduced potential hazard of contamination.

3.2 DNA Quantification

Samples were dried onto sterile paper and weighted. Samples were then frozen at -20°C.

The day before the analysis, samples were put in a solution of Proteinase K at 56°C, until complete digestion. Usually overnight is enough to digest the entire sample.

This lysis solution contains:

Proteinase K: 1mg/ml in 50mM Tris pH 7.6 (SIGMA-ALDRICH Product Number:P2308)

Pepstatin A: (SIGMA-ALDRICH Product Number: P5318)

EDTA 100mM+lodoacetamide 100mM (Siegfried)

TRIS: (SIGMA-ALDRICH Product Number:A5456-3)

DMSO: (FLUKA Product Number:41640)

The day after the digestion:

Thaw the CyQUANT® Cell Proliferation Assay Kit (InVitrogen Product Number:C7026)

a. Preparation of the working buffer

Dilute Component B (20x): 20 ml = 1 ml Component B + 19 ml

H2O

Add Component A (400x) to the same tube 50 μl / 20ml

The component A is light sensitive so you have to wrap the tube in an aluminum sheet

b. Preparation of DNA working solution

Bovine DNA: Stock solution 10 μg/ml

DNA working solution : 1 μg/ml Dilution 1/10

c. Preparation of the standards

For each standard we need: 3 x 200 μ l so we prepare 640 μ l

d. Preparation of the standard curve: (use the black plates)

Each standard will be done in triplicate: Pipette 200 $\mu\text{l}/\text{well}$ for each standard

e. Preparation of the samples

Dilution 1/200: 5 µl sample directly to the well

+ 195 μl working Buffer directly to the well

When all the samples are pipetted, cover the plate with an aluminium paper

f. Reading of the values

Use the Perking Elmer HTS 700 apparatus, and a protocol with a wavelength of 480/520nm.

Results were normalized for the weight, and compared using the bar graph.

3.3 SEM Analysis

Samples were fixed and dehydrated following this sequence:

- Rinse in PBS for 10 min
- Formaldehyde 4% overnight 4 ° C
- 2X rinse in PBS for 10 min
- 3X 20 min 30% Ethanol
- 3X 20 min 50% Ethanol
- 3X 20 min 70% Ethanol
- 3X 20 min 90% Ethanol

Tissues were transferred to 100% ethanol, subjected to Critical Point Drying, and covered with gold particles.

3.4 Istochemistry

3.4.1Hematoxylin and Eosin Staining

Samples were fixed by immersion with Paraformaldehyde 4% for 12 hours at 4°C, and then put in Sucrose(Sigma Aldrich Product Number:

S7903) 30% in Milli Q water.

After that, samples were put in the embedding machine for the alcohol dehydratation and the final step in liquid paraffin at 60°C.

When samples were in liquid paraffin, were embedded and then cut with the microtome in slides of 6µm using polylisinate glasses.

Then slides were dried at 37°C overnight and put in the staining machine.

The staining includes these phases:

- Ultraclear 90s (Biosystem Product Number:41-5213-00)
- Ultraclear 90s
- Ultraclear 90s
- Ethanol 100% 90s
- Ethanol 100% 90s
- Ethanol 96% 90s
- Ethanol 70% 90s
- Ethanol 50% 90s
- H2O distilled 90s
- HE mayer 90s (Biosystem Product Number:38-7010-00)

- HE mayer 90s
- HE mayer 90s
- H2O tap 90s
- HCI -Alcohol 0.2% 90s
- H2O tap + NH4 90s
- Eosin 0.2% 90s (Biosystem Product Number:38-7410-00)
- Eosin 0.2% 90 s
- Eosin 0.2% 90s
- Ethanol 96% 90s
- Ethanol 96% 90s
- Ethanol 100% 90s
- Ultraclear 90s
- Ultraclear 90s
- Ultraclear 90s

Slides were mounted with Mounting medium (Pertex Product Number:41-4012-00), and acquired at different magnification with an Olympus DP80 light microscope.

3.4.2 Immunohistochemistry

The immunoistochemisty it has been used to evaluate the presence of specific proteins in tissues analized:

Antibody anti-Collagen type IV (AbCam Product Number:ab6586) Antibody anti-Fibronectin (AbCam Product Number:ab2413)

- Ultraclear 10m (Biosystem Product Number:41-5213-00)
- Ultraclear 10m
- Ethanol 100% 5m
- Ethanol 95% 1m
- Ethanol 70% 1m
- Ethanol 35% 1m
- Rinse with buffer TBS 1X 1m
- Rinse with buffer TBS 1X 1m
- HEAT MEDIATED ANTIGEN RETRIEVAL
 - Retrieval solution (Dako Product Number:S1700) 10m at 95°C
 - Let it cool down for 10-15m
 - Rinse with buffer TBS 1X 5m
 - Rinse with buffer TBS 1X 5m
- BLOCKING
 - Goat serum (Invitrogen Product Number:16210-064) 10m
 - Rinse with buffer TBS 1X 5m
- Primary antibody diluted in TBS 1%BSA overnight at 4°C
- Rinse with buffer TBS 1X 5m
- Rinse with buffer TBS 1X 5m
- Secondary antibody diluted in TBS 1%BSA 1:200 45m (polyclonal Goat anti Rabbit Immunoglobulins/HRP E0432)
- Rinse with buffer TBS 1X 5m
- Rinse with buffer TBS 1X 5m
- DAB 5 m (1 tablet in 15ml of TBS1X plus 12 μl of 30% H2O2) (Sigma Aldrich Product Number:D5905)

- Rinse with buffer TBS 1X 5m
- Rinse with buffer TBS 1X 5m
- Rinse with tap water 1m
- Hematoxylin 20s
- Rinse with tap water 1m
- Ethanol 96% 90s
- Ethanol 96% 90s
- Ethanol 100% 90s
- Ultraclear 90s
- Ultraclear 90s
- Ultraclear 90s

Slides were mounted with Mounting medium (Pertex Product Number:41-4012-00), and acquired at different magnification with an Olympus DP80 light microscope.

Antibody anti-Laminin (AbCam Product Number:ab128053)

- Ultraclear 10m (Biosystem Product Number:41-5213-00)
- Ultraclear 10m
- Ethanol 100% 5m
- Ethanol 95% 1m
- Ethanol 70% 1m
- Ethanol 35% 1m
- Rinse with buffer TBS 1X 1m
- Rinse with buffer TBS 1X 1m
- ENZYMATIC DIGESTION

- Trypsin 0,1% in TBS (Gibco Product Number:15090-046)
- BLOCKING
 - Goat serum (Invitrogen Product Number:16210-064) 10m
 - Rinse with buffer TBS 1X 5m
- Primary antibody diluted in TBS 1%BSA overnight at 4°C
- Rinse with buffer TBS 1X 5m
- Rinse with buffer TBS 1X 5m
- Secondary antibody diluted in TBS 1%BSA 1:200 45m (polyclonal Goat anti Rabbit Immunoglobulins/biotinylated E0432)
- Rinse with buffer TBS 1X 5m
- Rinse with buffer TBS 1X 5m
- Vectastain ABC/AP kit (Linaris Product Number:AK-5000) 45 m in a humid chamber
- Rinse with buffer TBS 1X 5m
- Rinse with buffer TBS 1X 5m
- Vector Red AP substrate + Levamisole (Linaris Product Number: SK-5100)
- Rinse with buffer TBS 1X 1m
- Rinse with tap water 1m
- Hematoxylin 20s
- Rinse with tap water 1m
- Ethanol 96% 90s
- Ethanol 96% 90s
- Ethanol 100% 90s
- Ultraclear 90s

- Ultraclear 90s
- Ultraclear 90s

Slides were mounted with Mounting medium (Pertex Product Number:41-4012-00), and acquired at different magnification with an Olympus DP80 light microscope.

3.5 Reagent for the decellularization process

- Triton X-100 (Sigma Aldrich Product Number: T8532)
- Sodium Deoxycholat (Sigma Aldrich Product Number:30970)
- DNAse I from Bovine Pancreas (Sigma Aldrich Product Number:D4263)

3.6 MTT assay

- The yellow tetrazolium salt (MTT) (Sigma Aldrich Product Number: M5655) is reduced in metabolically active cells to form insoluble purple formazan crystal, which are solubilized by the addition of a detergent.
 - 1. Weight 12.5 mg
 - 2. Dilute it in 25 mL culture medium w/o phenol red.
 - 3. Filter sterilize = *MTT* stock solution (0.5mg/ml), storage in the freezer

4. Dilute 1/10 of MTT stock solution with culture medium w/o phenol red = *MTT working solution (0.05mg/ml=50ug/ml)*

- 5. Remove culture medium from your wells
- 6. Rinse with PBS
- 7. Add *MTT working solution* (to cover the sample)
- 8. Keep in incubator for about 2 hours
- 9. Remove MTT working solution
- 10. Rinse with PBS
- 11. Look at the distribution of the cells in the scaffold

3.7 Scaffold preparation

- Dry the scaffold on paper to let it dry for the disruption
- The scaffold is put into cryovial with metal beads inside given with the machine itself (Mikro-Dismembrator).
- The vials were freezed in dry ice to let the scaffold hard and more easy to disrupt
- After 10 minutes the vials were transferred into the Mikro-Dismembrator S
- The scaffold were disrupted at 3000 rpm for 1 minute.
- the product of dismembration is a powder that is subsequently dissolved in 200 µl per thymus of MilliQ water
- 50 µl of the solution are distributed for each well of 96 well
- the 96 well is directly freezed at -20°C
- · after 2 hours the weel is transferred into the lhyophilizator
- it is set at a pressure of 0.0030mbar and a temperature of -50°C
- · the sample are left overnight for drying

4. RESULTS AND DISCUSSION

4.1 STUDY OF A PROTOCOL FOR THYMUS DECELLULARIZATION

To obtain an efficient protocol to decellularized the murine thymus, I started to apply a protocol, previously developed for the rat thyroid. It consists of different steps, detailed in the table below.

Freze and thaw		
Trypsin 0.05%	30 minutes at 37°C	
Triton X-100 3%	48 hours	
Deoxycholic acid 4%	24 hours	
PBS	30 minutes at 4°C	
Peracetic acid 0.1 %	3 hours at RT	
PBS	48 hours	

Table 1. protocol of thyroid decellularization applied to mouse thymus

The obtained matrices were analyzed to determine the extent of decellularization, if the cellular material had been thouroughly removed and wheter or not the scaffold tissue was damaged. For this purpose Haematoxylin and Eosin (H&E) staining and SEM analysis were performed.



Fig.29 H&E staining of decellularized thymus at 10X and 20X of magnification



Fig.30 SEM analysis of the matrix obtained at two different magnification

These analysis showed that the protocol initially is too aggressive for the mouse thymus and it damaged the structure and the compositions of its ECM when compared to the native one.



Fig.31 H&E staining of native thymus at 4X and 20X of magnification



Fig.32 SEM analysis of the native thymus at two different magnification. At higher magnification is possible to see some limphocytes on the surface of the organ itself.

Starting from this point, I've analyzed the role played by different agents used in the protocol and their activity on the mouse thymus.

Thymic matrices were decellularized using the same protocol but eliminating one by one the different solutions to evaluate the action of the different agents on the ECM.

The different agents evaluated has been: Triton X-100 3%, Deoxyxholic Acid 4%, Trypsin 0,05% and 0,01%. The matrix structure was evaluated by H&E staining, SEM and DNA quantification to figure out the influence of these agents on the ECM of the thymus and its 3D structure.

In the following table is described the first modified protocol tested on the mouse thymus to evaluate the action of the Deoxycholic acid on the ECM.

Freze and thaw			
Trypsin 0.05%	30 minutes at 37°C		
Deoxycholic acid 4%	24 hours		
PBS	30 minutes at 4°C		
Peracetic acid 0.1 %	3 hours at RT		
PBS	48 hours		

 Table 2. Protocol of thymus decellularization without the use of Triton X-100

The matrix obtained was evaluated performing Haematoxylin and Eosin (H&E) staining and SEM analysis.



Fig.33 H&E staining of decellularized thymus without the use of the Triton X-100 at 10X and 20X of magnification



Fig.34 SEM analysis of the matrix obtained without the use of the Triton X-100 at two different magnification

In the following is described table the second modified protocol tested on the mouse thymus to evaluate the action of the Triton X-100 on the ECM.

Freze and thaw	
Trypsin 0.05%	30 minutes at 37°C
Triton X-100 3%	48 hours
PBS	30 minutes at 4°C
Peracetic acid 0.1 %	3 hours at RT
PBS	48 hours

Table 3. Protocol of thymus decellularization without the use of Deoxycholic Acid



Fig.35 H&E staining of decellularized thymus without the use of the Deoxycholic acid at 10X and 20X of magnification



Fig.36 SEM analysis of the matrix obtained without the use of the Deoxycholic acid at two different magnification

In the following table is described the third modified protocol tested on the mouse thymus to evaluate the action of the Deoxycholic acid on the ECM.

Freze and thaw		
Triton X-100 3%	48 hours	
Deoxycholic acid 4%	24 hours	
PBS	30 minutes at 4°C	
Peracetic acid 0.1 %	3 hours at RT	
PBS	48 hours	

Table 4. Protocol of thymus decellularization without the use of Trypsin



Fig.37 H&E staining of decellularized thymus without the use of the Trypsin at 20X of magnification



Fig.38 SEM analysis of the matrix obtained without the use of the Trypsin at two different magnification

Freze and thaw			
Trypsin 0.01%	30 minutes at 37°C		
Triton X-100 3%	48 hours		
Deoxycholic acid 4%	24 hours		
PBS	30 minutes at 4°C		
Peracetic acid 0.1 %	3 hours at RT		
PBS	48 hours		

 Table 5. Protocol of thymus decellularization with Trypsin diluited at 0,01%



Fig.39 H&E staining of decellularized thymus with the use of the Trypsin 0,01% at 10X and 20X of magnification



Fig.40 SEM analysis of the matrix obtained with the use of the Trypsin 0,01% at two different magnification



FIG.41 DNA quantification of decellularized thymi in absolute value.

Although the different solutions analyzed proved to be adequate to efficiently decellularize the thymic matrix, nevertheless they resulted too aggressive for maintainance of its 3D structure, and basic ECM components. In addition, using H&E a violet staining meaning was sometimes detected, indicating presence of cellular debrids and nucleic acids, able to bind to the hematoxylin. In particular, trypsin either at 0,05% or 0,01% was able to destroy both the cells and ECM, as a consequence of the low protein amount present in the mouse thymus. In addition, Triton X-100 and Deoxycolic acid resulted inadequate to efficiently clean the ECM from cell fragments and DNA contamination, although delicate enough to reasonable preserve the native structure of the thymus matrix.

I, therefore, proceeded to modify the original protocol by adding MilliQ water and DNAse, based on protocols previously established with other organ like trachea [131], with the aim of preserving the ECM structure and, at the same time to improve its cleaning from contaminating cellular residues.

MilliQ water was chosen for its ability to induce an osmotic stress to cells, and to eliminate them as floating fragments, as a result of the shaker movements forcing cellular membranes to pass from the organ into the solution.

DNase was chosen for its ability to degradate DNA. It was used at the end of the decellularization process, to digest the DNA remained into the matrix without damaging the ECM.

The new protocols tested are listed in the table below.

	Protocol	Protocol	Protocol	Protocol
	A	В	C	D
MilliQ water 72 hours at 4°C		\checkmark	\checkmark	$\sqrt{x 2}$ cycles
Triton X-100 3% 12 hours at RT	\checkmark	\checkmark		
Deoxycholic acid 4% at RT		$\sqrt{24 \text{ hours}}$	$\sqrt{4 \text{ hours}}$	$\sqrt{4}$ hours x 2 cycles
DNAse 3hours at RT				$\sqrt{x 2}$ cycles

 Table 5. Scheme of different protocols used for thymus decellularization

An Haematoxylin and Eosin (H&E) staining and SEM analysis were performed to analyse resulting scaffold tissues obtained. They are showed in pictures below.



Fig.42 H&E staining of decellularized thymus with the protocol A $\,$ at 10X and 20X of magnification



Fig.42 SEM analysis of the matrix obtained with the protocol A at two different magnification



Fig.43 H&E staining of decellularized thymus with the use of the protocol D at 10X and 20X of magnification



00133539 30 µm 2² M В Uni Basel

Fig.44 SEM analysis of surface and the inner part of the matrix obtained with the use of the protocol B at two different magnification.



Fig.45 H&E staining of decellularized thymus with the use of the protocol C at 10X and 20X of magnification





Fig.46 SEM analysis of the surface and the inner part of the matrix obtained with the protocol C at two different magnification



Fig.47 H&E staining of decellularized thymus with the the protocol D at 10X and 20X of magnification





Fig.48 SEM analysis of the surface and the inner part of the matrix obtained with the protocol D at two different magnification.

A DNA quantification was performed to evaluate the remaining cellular material in resulting scaffolds obtained.



FIG.49 DNA quantification of matrices obtained in absolute value

An immunoistochemistry for the mainly ECM components of the mouse thymus, collagen IV, fibronectin and laminin, was performed to evaluate changes in ECM proteins after the different protocols used. The native mouse thymus was used like positive control.


FIG.50 Immunohistochemical staining for collagen IV in native and decellularized thymi. A) native thymus; B) scaffold tissue obtained with the protocol A; C) scaffold tissue obtained with the protocol C; E) scaffold tissue obtained with the protocol D.



FIG.51 Immunohistochemical staining for fibronectin in native and decellularized thymi. A) native thymus; B) scaffold tissue obtained with the protocol A; C) scaffold tissue obtained with the protocol B; D) scaffold tissue obtained with the protocol C; E) scaffold tissue obtained with the protocol D.



FIG.52 Immunohistochemical staining laminin in native and decellularized thymi. A) native thymus; B) scaffold tissue obtained with the protocol A; C) scaffold tissue obtained with the protocol B; D) scaffold tissue obtained with the protocol C; E) scaffold tissue obtained with the protocol D.

Using these new protocols, a better cleaning and maintainance of the 3D structure of the thymic ECM mouse was achieved.

In particular, using the protocol A, based on Triton X-100, milliQ water and DNAse, I observed only a modest decrease in the quantity of cellular contamination, whereas the 3D structure of the matrix was maintained quite well. This was also confirmed by the immunohistochemical analysis, that showed presence of laminin,

collagen IV and fibronectin in distribution consistent with that of the native control tissue.

In contrast, using protocol B based on addition of Deoxycholic Acid, the quantity of DNA decreased in a significant manner, while no changes occurred to the 3D matrix structure. Specifically, using both H&E staining at light microscopic level, and SEM I noticed that the 3D structure of the thymic matrix was reasonably preserved, although some holes formed, possibly due to the removal of resident cells. In addition, ECM proteins were observed by immunocytochemistry in a pattern resembling that of protocol A, but clustered as a result of the absence in resident cells. Collectively, these results suggested that protocol B could provide a good scaffold system for homing and proliferation of seeded TEC.

Using protocol C, in which the Deoxycholic Acid was applied only for 4 hours, no changes in the 3D matrix structure were observed. However, an improvement in the size of the pores occurred, suggesting that the Deoxycholic Acid was able to efficiently remove cells even if used for a short time. In addition, the amount of DNA decreased to a reasonable extent, suggesting a primary decellularizing role for the Deoxycholic Acid. Finally, no relevant changes were found in the amount of ECM proteins.

At variance with previous protocols, using protocol D I found differences in DNA quantity, matrix size, and ECM components. Specifically, the amount of DNA was increased compared to protocol C, a reduction in matrix size occurred, likely depending on the osmotic action of MilliQ water, and the long time of shaking as well, whereas a reduction in fibronectin staining was equally apparent.

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In summary, the best protocols proved to be B and C, able to guarantee a very little amount of contaminating DNA, while maintaining the native 3D matrix structure, and protein compositions of the ECM.

4.2 ROLE OF DECELLULARIZED THYMIC MATRICES IN REGULATING THE IN VITRO BEHAVIOUR OF TEC

The decellularized, 3D thymic matrices obtained using the previously described protocol were used as natural, 3D scaffold to study the behaviour of seeded TECs.

Decellularized matrices could influence TEC by releasing different factors into the culture medium, by directing them to their 3D structure or by influencing them through the ECM composition.

Specifically, I started to evaluate the role played by decellularized matrices prepared with the protocol C, which seemed to have the best amount of ECM proteins and low quantity of DNA. To this purpose these matrices were transformed in a sort of "sponge", as detailed in Material and Methods, and their structure analysed to define the acquired, final geometry. SEM analyses were performed on scaffold obtained and the shape of its surface and their inner parts are showed in figures below.





Fig.53 SEM analysis of the surface of the matrix obtained with the protocol C at two different magnification.





Fig.54 SEM analysis of the inner part of the matrix obtained with the protocol C $\,$ at two different magnification.

Sem analyses revealed that these "3D scaffolds" had pores big enough, and with a regular distribution to ensure homing and possibly proliferation of eventually seeded TEC.

I, therefore, decided to verify whether the 3D scaffolds made with decellularized matrix might release soluble factors in the culture media able to influence the growth and the differentiation of seeded TECs.

To this aim, scaffolds were used to condition the culture media for 48 hours at 37°C. Media, then were diluited at 5% and filtered throught a filter of 0,2µm to select the soluble factors released getting rid of anymatrix debris.

At the same time, TECs were seeded on standard 48-well plates, at a density of 50.000 cells/well and conditioned medium was added 24 hours later the seeding, to allow attachment of seeded elements to the bottom of the plastic plate. Subsequently the medium change was done every 48 hours.

Cells were followed day by day taking pictures of them at every medium change having like positive control cells cultured with fresh media.

Pictures, taken every medium change for the positive control, are showed below.



Fig.55 Pictures taken with the light microscope in the control conditon during all the culture at every medium change.

Pictures, taken every medium change for cells cultured with medium conditioned at 5%, are showed in the figure below.



Fig.56 Pictures taken with the light microscope during all the culture, at every medium change, for the cells cultured with medium conditioned with scaffold obtained after disruption, at 5% diluition.



Fig.57 Pictures taken with the light microscope during all the culture, at every medium change, for the cells cultured with medium conditioned by decellularized matrix at 5% diluition.

Following initial seeding of culture cells in all conditions, I found that in the first days they were few, due to the death of differentiated cells, but from day 5 cells started to proliferate reaching confluence and cell clusters at day 9.

Their vitality and markers expression was confirmed by FACS analysis performed at day 12, underlying an expression of MHC II by TEC. FACS data are showed below for the positive control and for different culture conditions.



Fig.58 Cell surface marker characterization of the 2D positive control of TEC as determined by flow cytometric analysis. TEC were analyzed for: A) size; B) DAPI staining for the vitality; C) elimination of doublets; D) EpCam staining; E) MHC II staining.



Fig.59 Cell surface marker characterization of TEC, cultured with medium conditioned with scaffold obtained after disruption of decellualrized tissue, as determined by flow cytometric analysis. TEC were analyzed for: A) size; B) DAPI staining for the vitality; C) elimination of doublets; D) EpCam staining; E) MHC II staining.



Fig.60 Cell surface marker characterization of TEC, cultured with medium conditioned with decellualrized tissue, as determined by flow cytometric analysis. TEC were analyzed for: A) size; B) DAPI staining for the vitality; C) elimination of doublets; D) EpCam staining; E) MHC II staining.

In comparing cell shape, cell number and TECs marker expression, I found that only MHC II antigens resulted slighly higher in the cells treated with the conditioned medium compared to controls, suggesting a favourable effect of matrix products on TECs differentiation state.

In a following step, these culture conditions were applied to cells cultured in fibrin gel. This with the aim to determine whether the changes in the above mentioned parameters could be detected also in cells grown in a 3D environment, known to induce MHC II expression as in vivo conditions.

FACS data are showed below for two different positive controls in two different configuration (2D and 3D) and for different culture conditions.



Fig.61 Cell surface marker characterization of 2D positive control of TEC as determined by flow cytometric analysis. TEC were analyzed for: A) size; B) DAPI staining for the vitality; C) elimination of doublets; D) EpCam staining; E) MHC II staining.



Fig. 62 Cell surface marker characterization of TEC, cultured in 3D fibrin gel, as determined by flow cytometric analysis. TEC were analyzed for: A) size; B) DAPI staining for the vitality; C) elimination of doublets; D) EpCam staining; E) MHC II staining.



Fig.63 Cell surface marker characterization of TEC, cultured in 3D fibrin gel with medium conditioned with scaffold obtained after disruption of decellualrized tissue, as determined by flow cytometric analysis. TEC were analyzed for: A) size; B) DAPI staining for the vitality; C) elimination of doublets; D) EpCam staining; E) MHC II staining.



Fig.64 Cell surface marker characterization of TEC, cultured in 3D fibrin gel with medium conditioned with decellualrized tissue, as determined by flow cytometric analysis. TEC were analyzed for: A) size; B) DAPI staining for the vitality; C) elimination of doublets; D) EpCam staining; E) MHC II staining.

In preliminary studies still in progress, after 12 days in 3D fibrin culture and conditioned medium, seeded TECs depicted increased expression of MHC II antigens, suggesting a direct stimulatory role of the scaffold products on differentiation parameters for TECs.

Similarly, preliminary studies have also been conducted to assess the role of TEC seeding directly on the decellularized matrix, but limited trapping of the cell suspension by the matrix requires further studies. However, cell viability assays, based on MTT, proved that, at day 3 of culture cells were able to attach and proliferate on the decellularized matrix (fig 65), and that they survived and proliferate up to 10 days in static culture (figure 66).



Fig.65 Pictures of the MTT assay after 3 days of culture. The first one taken through the steromicroscope and the other one with a normal photocamera. Cells alive, that metabolize the MTT, are stained in violet.



Fig.65 Pictures of the MTT assay after 10 days of culture. The first one taken through the steromicroscope and the other one with a normal photocamera. Cells alive, that metabolize the MTT, are stained in violet.

These results are very encouraging, suggesting that TECs seeded on native, decellularized matrixes are metabolically active and, likely favourably respond to the 3D geometry of the substrate by preserving their proliferation potential.

5. CONCLUSIONS AND FUTURE PERSPECTIVES

During my PhD tenure at the Tissue Engineering Laboratory of the University of Basel I have investigated the possibility to obtain an efficient protocol for the decellularization of the mouse thymus, and to use it as a scaffold for culturing Thymic Ephitelial Cells. Since TEC in normal 2D condition on plastic loose their differentiation state, I investigated a suitable 3D alternative, in particular a 3D scaffold able to support their growth, and maintain their differentiation state.

Due to its anatomy, the thymus is a very difficult organ to decellularize. Indeed, it is served only by small vessels, that cannot be easily used to perform a perfusion-decellularization process. For this reason, I developed an original decellularization process by immersion. Two different protocols satisfied the main characteristics of a decellularized matrix, according to qualitative and quantitative assays. In particular, the quantity of DNA was less than 10% in absolute value, no positive staining for cells was found after the decellularization, and the 3D structure and composition of the ECM were maintained. In addition, I was able to prove that the decellularized matrixes were not cytotoxic for the cells, and were able to increase expression of MHC II antigens with respect to control cells grown in standard conditions.

Further analysis are required to study whether and to which gene level these effects take place, which soluble factors are released by the decellularized matrix in the culture media, and their possible effects on the differentiation state of TECs.

Improvements will be done in seeding efficiency of the cells directly on the matrix, possibly also exploiting appropriate bioreactor conditions, and in the choice of appropriate markers showing the level of differentiation reached by the seeded TECs.

After a complete characterization of the culture system, these innovative natural scaffolds could be used to improve the standard culture conditions of TEC, to study in vitro the action of different factors on their differentiation genes, and to test the ability of TECs to induce in vitro maturation of seeded T lymphocytes.

In a near future, I also envisage the use of a completely recellularized, thymic matrix as a tool to test in vivo lymphocyte maturation after its implant under the kidney capsule of the athymic mice, and possibly as an innovative therapeutic tool to rescue conditions of immunodepression in specific animal models of immunodeficiency.

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