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Metabolic control of T-cell activation and death in SLE

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Abstract

Systemic lupus erythematosus (SLE) is characterized by abnormal T-cell activation and death, processes which are crucially dependent on the controlled production of reactive oxygen intermediates (ROI) and of ATP in mitochondria. The mitochondrial transmembrane potential $(\Delta\psi_m)$ has conclusively emerged as a critical checkpoint of ATP synthesis and cell death. Lupus T cells exhibit persistent elevation of $\Delta\psi_m$ or mitochondrial hyperpolarization (MHP) as well as depletion of ATP and glutathione which decrease activation-induced apoptosis and instead predispose T cells for necrosis, thus stimulating inflammation in SLE. NO-induced mitochondrial biogenesis in normal T cells accelerates the rapid phase and reduces the plateau of Ca²⁺ influx upon CD3/CD28 co-stimulation, thus mimicking the Ca²⁺ signaling profile of lupus T cells. Treatment of SLE patients with rapamycin improves disease activity, normalizes CD3/CD28-induced Ca²⁺ fluxing but fails to affect MHP, suggesting that altered Ca²⁺ fluxing is downstream or independent of mitochondrial dysfunction. Understanding the molecular basis and consequences of MHP is essential for controlling T-cell activation and death signaling in SLE.

- Lupus T cells exhibit mitochondrial dysfunction
- Mitochondrial hyperpolarization (MHP) and ATP depletion predispose lupus T cells to death by necrosis which is pro-inflammatory
- MHP is caused by depletion of glutathione and exposure to nitric oxide (NO)
- NO-induced mitochondrial biogenesis regenerates the Ca²⁺ signaling profile of lupus T cells
- Rapamycin treatment normalizes Ca²⁺ fluxing but not MHP, suggesting that the mammalian target of rapamycin, acts as a sensor and effector of MHP in SLE

Keywords

lupus; mitochondria; nitric oxide; glutathione; calcium

Introduction

Abnormal T cell activation and cell death underlie the pathology of SLE (1). Potentially autoreactive T and B lymphocytes are removed by apoptosis during development and after completion of an immune response. Paradoxically, lupus T cells exhibit both enhanced

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spontaneous apoptosis and defective activation-induced cell death (2). Increased spontaneous apoptosis has been linked to chronic lymphopenia (2) and compartmentalized release of nuclear autoantigens in patients with SLE (3). By contrast, defective activation-induced cell death (AICD) may be responsible for persistence of autoreactive cells (2).

Both cell proliferation and apoptosis are energy-dependent processes. Energy in the form of ATP is provided through glycolysis and oxidative phosphorylation. The mitochondrion, the site of oxidative phosphorylation, has long been identified as a source of energy and cell survival (2). The synthesis of ATP is driven by an electrochemical gradient across the inner mitochondrial membrane maintained by an electron transport chain and the membrane potential $(\Delta \psi_m, \text{ negative inside and positive outside})$. A small fraction of electrons react directly with oxygen and form reactive oxygen intermediates (ROI). The $\Delta \psi_m$ is regulated by the supply of reducing equivalents (NADH/NAD + NADPH/NADP + GSH) and the production of ROI and nitric oxide (NO) (4). Regeneration of GSH by glutathione reductase from its oxidized form, GSSG, and synthesis of NO depend on NADPH produced by the pentose phosphate pathway (PPP) (2). While disruption of the mitochondrial membrane potential $\Delta \psi_m$ has been proposed as the point of no return in apoptosis, elevation of $\Delta \psi_m$ or mitochondrial hyperpolarization (MHP) occurs prior to activation of caspases, phosphatidylserine (PS) externalization and disruption of $\Delta \psi_m$ in Fas- (5), H₂O₂- (6), and NO-induced apoptosis (7). MHP is also triggered by T-cell receptor (TCR) stimulation that is associated with transient inhibition of F_0F_1 -ATPase, ATP depletion, and sensitization to necrosis, suggesting that $\Delta \psi_m$ elevation is a critical checkpoint of T cell fate decisions (8). Importantly, lupus T cells exhibit persistent MHP and ATP depletion which causes predisposition to death by necrosis that is highly pro-inflammatory (2). The mammalian target of rapamycin (mTOR) is located in the outer mitochondrial membrane and serves as a sensor of the $\Delta \psi_m$ in T cells (9). Focused on targeting MHP for treatment of lupus patients resistant or intolerant to conventional immunosuppressants, rapamycin improved disease activity and normalized baseline and T cell activation-induced Ca^{2+} fluxing without affecting MHP(10). These findings suggested that mTOR represents a gate keeper between MHP and altered Ca²⁺ signaling in SLE. The present review will focus on establishing the hierarchy of the metabolic pathways that underlie and mediate the consequences MHP and identify checkpoints that can be targeted for therapeutic intervention in SLE (Fig. 1).

Metabolic control of T cell activation and Ca²⁺ Fluxing

ROI modulate T cell activation, cytokine production, and proliferation at multiple levels (2). The antigen-binding $\alpha\beta$ or $\gamma\delta$ TCR is associated with a multimeric receptor module comprised of the CD3 $\gamma\delta\epsilon$ and ζ chains. The cytoplasmic domain of CD3 ζ chain contains an immunoglobulin receptor family tyrosine-based activation motif (ITAM) which is crucial for coupling of intracellular tyrosine kinases (11). Expression of CD3ζ is suppressed by ROI (12). Binding of p56^{lck} to CD4 or CD8 attracts this kinase to the TCR-CD3 complex, leading to phosphorylation of ITAM. Phosphorylation of both tyrosines of each ITAM is required for SH-2-mediated binding by zeta-associated protein-70 (ZAP-70) or the related SYK. ZAP-70 is activated through phosphorylation by p56^{lck}. Activated ZAP-70 and SYK target two key adaptor proteins LAT and SLP-76 (11). Phosphorylated LAT binds directly to phospholipase C-y1 that controls hydrolysis of phosphatydilinositol-4,5-biphosphate (PIP2) to inositol-1,4,5triphosphate (IP₃) and diacylglycerol (DAG). Phosphorylation of inositol lipid second messengers is mediated by phosphatidylinositol 3'hydroxyl kinase (PI3K). The stimulatory effect of the TCR alone on PI3K activity is small. Concurrent triggering of the CD28 costimulatory molecule by its ligands CD80 or CD86 is required for optimal PI3K activation. IP3 binds to its receptors in the endoplasmic reticulum (ER), opening Ca²⁺ channels that release Ca^{2+} to the cytosol. Decreased ER Ca^{2+} concentration activates the Ca^{2+} release-activated Ca^{2+} channel (CRAC) in the cell membrane. The resultant Ca^{2+} influx activates the

phosphatase calcineurin which dephosphorylates the transcription factor, NFAT. Dephosphorylated NFAT can translocate to the nucleus where it promotes transcription of IL-2 in concert with AP-1, NF κ B, and Oct-1. While activities of AP-1 and NF κ B are increased by oxidative stress (13), both thiol insufficiency and H₂O₂ treatment suppresses calcineurinmediated activation of NFAT (14). Thus, expression of cytokines, i.e. IL-2 (with AP-1 and NFAT motif-containg promoter) and IL-4 (with AP-1-less NFAT enhancer), can be selectively regulated by oxidative stress depending on the relative expression level of transcription factors involved (Fig. 1).

The calcium signal is dysregulated in SLE T-cells in a number of ways. SLE T-cells have elevated intracellular and mitochondrial calcium at baseline (15), although the amount of Ca^{2+} present in the endoplasmic reticulum is normal (16). And while the T-cell activation-induced calcium flux is elevated initially, the plateau phase is reduced relative to activated control T-cells (15). Dysregulation of NO may play a role, since normal T cells pre-treated with NO donors recapitulate the Ca^{2+} fluxing abnormalities observed in SLE T cells (15). Considering that the Ca^{2+} is important for activation of PKC, calcineurin, NFAT, and production of IL-2, the altered Ca^{2+} fluxing may account for the inappropriate activation of T cells in SLE, or conversely, may contribute to their inability to produce adequate amounts of IL-2 (1).

Metabolic control of cell death

Programmed cell death (PCD) or apoptosis is a physiological mechanism for elimination of autoreactive lymphocytes during development. Signaling through the Fas or structurally related TNF family of cell surface death receptors represent dominant pathways in elimination of unwanted cells under physiological and disease conditions (2). Fas stimulation leads to sequential activation of caspases resulting in the cleavage of cellular substrates and the characteristic morphologic and biochemical changes of apoptosis. Apically, trimerization of cell surface Fas receptors through formation of disulfide bonds is required for activation of the death-inducing signaling complex and cleavage of caspase 8 (2). Importantly, GSH is required for activity of caspases and its depletion can prevent Fas-dependent apoptosis (17). ATP is also required for apoptosis and its deficiency predisposes to necrosis (18). Autophagy mediates the bulk degradation of cytoplasmic contents, proteins and organelles including mitochondria, in lysosomes; this process is induced by rapamycin through inactivating mTOR (19). Interestingly, rapamycin treatment in vivo did not affect MHP or mitochondrial mass of lupus T cells (10).

MHP and ATP depletion predispose lupus T cells to necrosis

The mitochondrion is the site of ATP synthesis via oxidative phosphorylation. The synthesis of ATP is driven by an electrochemical gradient across the inner mitochondrial membrane maintained by an electron transport chain and the $\Delta \psi_m$. Activity of caspases require ATP to the extent that depletion of ATP by inhibition of F₀F₁-ATPase with oligomycin (18) or exhaustion of intracellular ATP stores by prior apoptosis signals, Fas stimulation or H₂O₂ pretreatment, leads to necrosis. Thus, intracellular ATP concentration is a key switch in the cell's decision to die via apoptosis or necrosis (18).

Apoptosis is a physiological process that results in nuclear condensation and break-up of the cell into membrane-enclosed apoptotic bodies suitable for phagocytosis by macrophages thus preventing inflammation. By contrast, necrosis is a pathological process that results in cellular swelling, followed by lysis and release of proteases, oxidizing molecules, and other proinflammatory and chemotactic factors resulting in inflammation and tissue damage (20). Swollen lymph nodes of patients with SLE harbour increased numbers of necrotic T lymphocytes and dendritic cells (DC) (21). Necrotic, but not apoptotic, cell death generates

inflammatory signals necessary for the activation and maturation of DCs, the most potent antigen-presenting cells (2;22). High mobility group B1 (HMGB1) protein, an abundant DNAbinding protein, remains immobilized on chromatin of apoptotic bodies, however, it is released from necrotic cells (2). Necrotic but not apoptotic cells also release heat shock proteins (HSPs), HSPgp96, hsp90. hsp70, and calreticulin. Mature DCs express high levels of the DC-restricted markers, CD83 and lysosome-associated membrane glycoprotein (DC-LAMP) and the costimulatory molecules CD40 and CD86 (2), which may contribute to altered intercellular signaling in SLE (Fig. 1). Their activation is driven by circulating interferon- α (IFN- α) that may come from one of the DC subsets, i.e., plasmacytoid dendritic cells (PDC) that infiltrate lupus skin lesions. SLE patients harbour activated PDC which may be responsible for increased production of IFN- α in SLE (23).

Targeting of metabolic checkpoints of T cell activation for therapeutic intervention in SLE

Depletion of intracellular glutathione

Reduced glutathione (GSH) levels are profoundly depleted in lymphocytes SLE patients (8). Low GSH in T cells over-expressing transaldolase predispose to MHP (5). GSH depletion is robust trigger of MHP via S-nitrosylation of complex I upon exposure to NO (24). Thus, the effect of NO on MHP is tightly related to GSH levels. Diminished production of GSH in face of MHP and increased ROI production is suggestive of a metabolic defect in de novo GSH synthesis or maintenance of its reduced state due to deficiency of NADPH (25). Recent studies showed diminished GSH/GSSG ratios in the kidneys of 8-month-old versus 4 month-old (NZB \times NZW) F1 mice; treatment with N-acetylcysteine (NAC), a precursor of GSH and stimulator of its de novo biosynthesis, prevented the decline of GSH/GSSG ratios, reduced autoantibody production and development of glomerulopnephritis (GN) and prolonged the survival of (NZB \times NZW) F1 mice (26). Oral NAC has been used to treat oxidative stress in patients with idiopathic pulmonary fibrosis (IPF) (27). In a one-year study of IPF patients treated with prednisone and azathioprine, addition of NAC ($3 \times 600 \text{ mg/d}$) improved vital capacity and reduced myelotoxicity in comparison to placebo. Therefore, prospective clinical studies appear justified to assess whether NAC treatment can reverse GSH depletion, correct T-cell signaling defects and provide clinical benefit to patients with lupus.

Inhibiting NO production

NO is a particularly interesting molecule in this context because it provides a link between seemingly dissociated features of T cell activation and mitochondrial function. NO is produced by nitric oxide synthases (NOS) that require Ca^{2+} to function and use NADPH and arginine as substrates. Thee isoforms exist: endothelial NOS, neuronal NOS, and inducible NOS, of which T cells express the former two (28). NO induces MHP and mitochondrial biogenesis, increases Ca^{2+} in the cytosol and mitochondria of normal T cells, and recapitulates the enhanced CD3/CD28-induced Ca^{2+} fluxing of lupus T cells (15). As recently reported, eNOS is recruited to the site of T-cell receptor engagement, locally increasing NO at the immunological synapse in a Ca^{2+} and PI3K-dependent manner, resulting in reduced IL-2 production(29) which is characteristic of SLE (1).

NO contributes to the development of GN in the MRL/lpr lupus mouse model (30). Inactivation of iNOS does not block the development of lupus (31), suggesting a role for eNOS and nNOS isoforms expressed in T cells. However, given the widespread expression of these isoforms in vascular smooth muscle and brain, it will be necessary to develop T-cell-specific approaches for inhibiting NOS to avoid potentially deleterious side effects.

Rapamycin

The mammalian target of rapamycin (mTOR) is associated with the outer mitochondrial membrane and senses mitochondrial dysfunction and changes in the $\Delta \psi_m$ of T cells (9). Rapamycin normalized T cell mitogen-stimulated splenocyte proliferation and IL-2 production, prevented the typical rise in anti-double-stranded DNA antibody and urinary albumin levels and GN, and prolonged survival of lupus-prone MRL/lpr mice (32). With a focus on mitochondrial dysfunction, we began to utilize rapamycin for treatment of SLE patients resistant or intolerant to conventional medications. We observed normalization of baseline Ca²⁺ levels in the cytosol and mitochondria and of CD3/CD28-induced Ca²⁺ fluxing as well as persistence of MHP (10), indicating that increased Ca^{2+} fluxing is downstream or independent of MHP in the pathogenesis of T-cell dysfunction in SLE. The effectiveness of rapamycin in murine and human SLE suggest that mTOR is a potential sensor and down-stream effector of MHP and mediator of T cell dysfunction and autoimmunity in SLE. Rapamycin can selective expand CD4+/CD25+/Foxp3+ regulatory T cells (33) which appear to be deficient in patients with SLE (34). Therefore, understanding the mechanism of persistent MHP that leads to mTOR activation and enhanced Ca^{2+} fluxing may be fundamental to the pathogenesis of T cell dysfunction and autoimmunity in SLE.

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Fig 1.

A) Schematic outline of the metabolic pathways controlling (PPP,GSH, NO) and sensing (mTOR) mitochondrial hyperpolarization (MHP) of lupus T cells. In normal T cells, MHP and mitochondrial biogenesis is mediated via production of NO by eNOS or nNOS (28) and up-regulation of transcription factors PGC-1 α , Tfam, and ALAS (39). NO production by eNOS may be compartmentalized to the T cell synapse (29). NO causes transient MHP via reversible inhibition of complex IV/cytochrome c oxidase (7) and persistent MHP via S-nitrosylation of complex I of the ETC in a state of GSH depletion (24). The PPP regulates the $\Delta \psi_m$ by producing 1) NADPH that serves as a reducing equivalent for GSH regeneration from its oxidized form GSSG and for production of NO by NOS and 2) ribose 5-phosphate for biosynthesis of

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nucleotides, ADP, ATP, NAD, NAADP, (c)ADP-ribose, and cGMP, the latter is a second messenger of NO. NAADP and (c) ADP-ribose induce Ca^{2+} release from the endoplasmic reticulum (ER) via ryanodine receptors (RyR). mTOR senses $\Delta \psi_m$ and regulates IP3R-mediated Ca^{2+} release (40). The intracellular rapamycin receptor FKBP12 directly binds the RyR. The Bcl-2 family proteins control permeability of the outer mitochondrial membrane and release of apoptosis-inducing factors. Necrosis-prone T cells release oxidized DNA and HMGB1 which stimulate B cells, macrophages, and dendritic cells (DC). In turn, B cells produce IL-10 and macrophages and DC produce NO which stimulate MHP of T cells. B) Proposed hierarchy of metabolic pathways upstream and downstream of MHP in lupus T cells. Broken line demarcates checkpoints affected by rapamycin.

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Table 1

Signaling alterations underlying abnormal cell death of T cells in patients with SLE

Signal	Effect	Reference
Spontaneous apoptosis [↑]	Compartmentalized autoantigen release, disease activity \uparrow	(3;8)
AICD ↓	Persistence of autoreactive cells	(35;36)
FasL↑	Spontaneous apoptosis [↑]	(37)
IL-10 [↑]	Selective induction of apoptosis in SLE	(36;37)
$\Delta\psi_m \uparrow$	ROI $^{\uparrow}$, ATP $^{\downarrow}$	(8)
GSH ↓	ROI $^{\uparrow}$, Spontaneous apoptosis $^{\uparrow}$	(8;38)
ATP ↓	AICD \downarrow , Predisposes for necrosis \uparrow	(8;18)
NO $^{\uparrow}$	Disease activity $^{\uparrow}, \Delta \psi_m^{\uparrow}$, mitochondrial biogenesis $^{\uparrow}$, enhanced baseline and activation-induced Ca ²⁺ flux	(15;28)
ROI [↑]	Spontaneous apoptosis \uparrow , IL-10 production \uparrow	(8;36)
IL-10 blockade	Spontaneous apoptosis \downarrow , ROI \downarrow	(36;37)
IL-12	Spontaneous apoptosis \downarrow , ROI \downarrow	(36)

[↑]increase;

↓ decrease;