

Keto Lean Call #1

# Homework

**Article:** Insulin, glucose and beta-hydroxybutyrate responses to a medium-chain triglyceride-based sports supplement: A pilot study

**Apps:**

Zero

Insight Meditation Timer

**Wearable(s):**

CGM

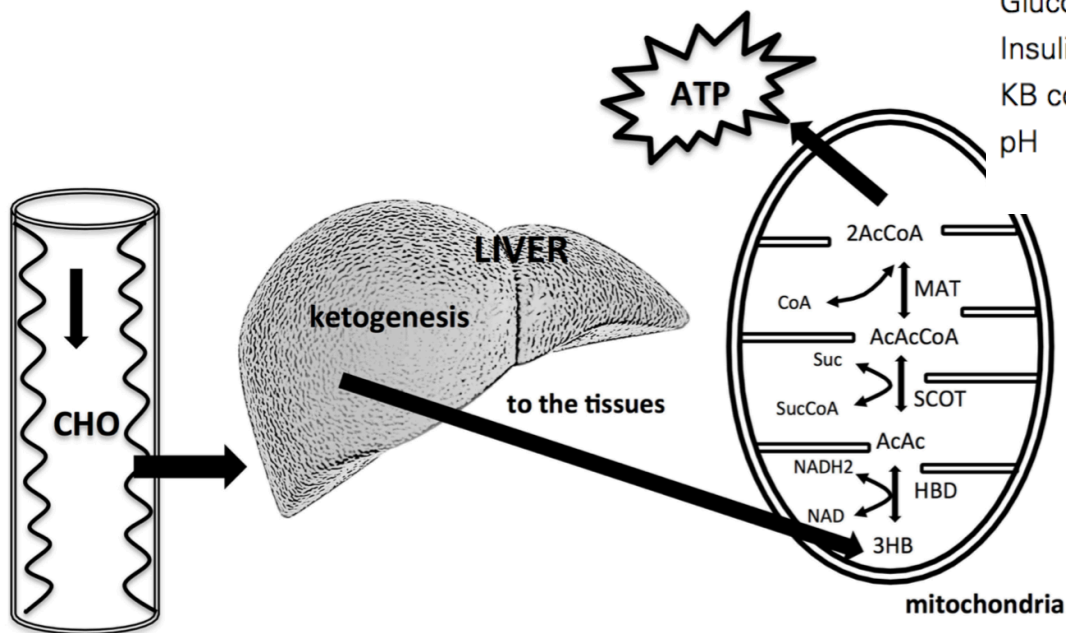
Glucometer

# Review

**Basics:** liver produces ketone bodies: (Beta-hydroxybutyrate, acetoacetate and acetone) from fatty acids

**Table 1 | Blood levels during a normal diet, ketogenic diet, and diabetic ketoacidosis (Paoli et al., 2012).**

Blood levels	Normal diet	Ketogenic diet	Diabetic ketoacidosis
Glucose (mg/dL)	80–120	65–80	>300
Insulin ( $\mu$ U/L)	6–23	6.6–9.4	$\cong 0$
KB conc (mmol/L)	0.1	7–8	>25
pH	7.4	7.4	<7.3

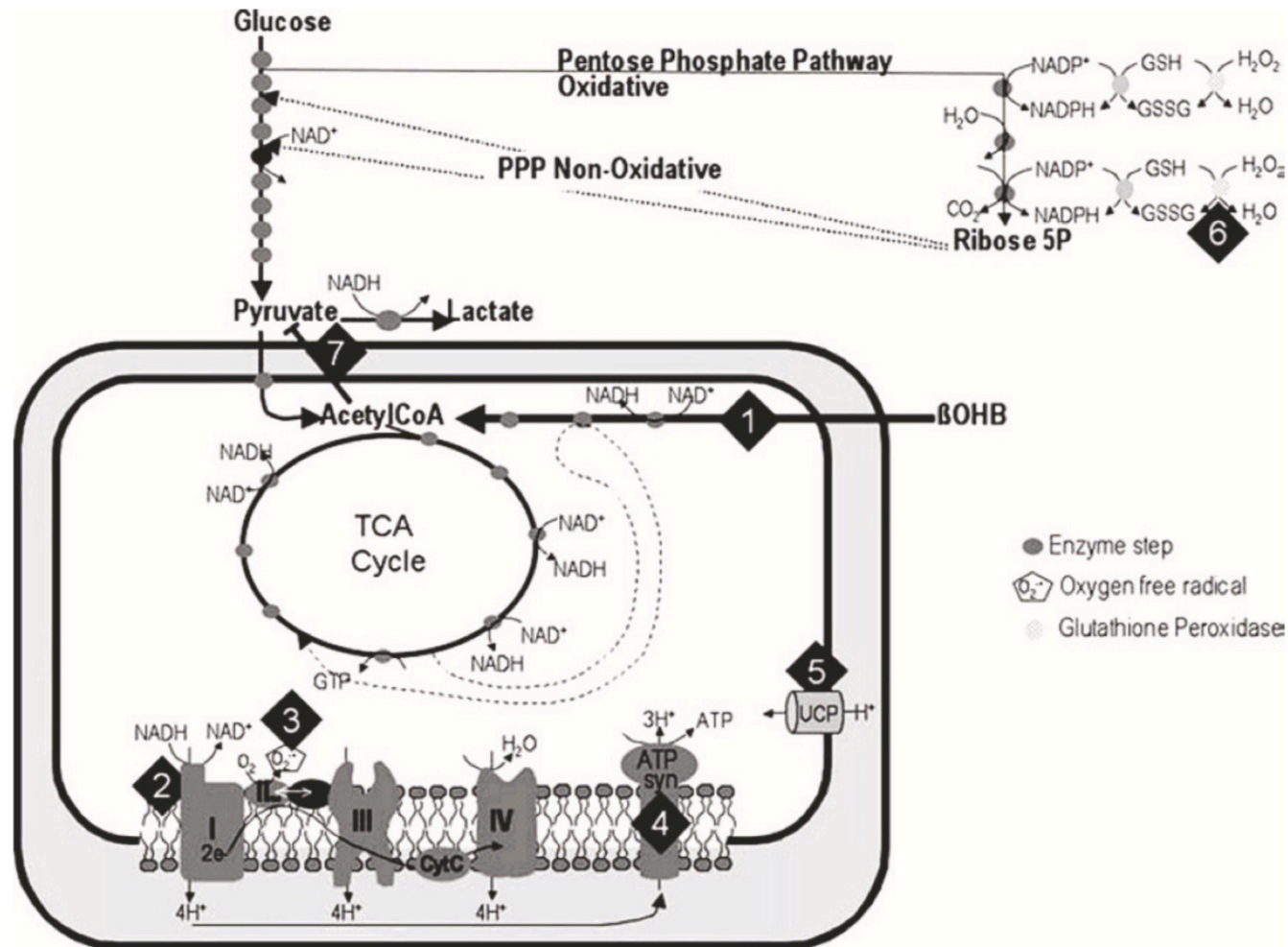


**\*\*Longevity pathways regulated by ketones and CR\*\* The crosstalk**

# Factors Increase/Decrease Ketosis

	Effect on ketogenesis	Mechanisms of action
Environmental factors		
Low carbohydrate diet	↑	Low insulin, leptin
Low calorie diet	↑	Low insulin, leptin, T3 High IGF-I, ghrelin
Physical activity	↑	Low insulin High AMPK gene expression High IGF-I
Moderate alcohol consumption	↑	High adiponectin, ghrelin Low insulin
Intrinsic factors		
AMPK activity	↑	Directly induces ketogenesis
ARNT/HIF1beta	↑	Stimulates AMPK activity

# Ketones Increase AcetylCoA Pool



**Figure 1. Potential neuroprotective mechanisms of ketones.** Ketones (1) require only three enzymatic steps to enter the tricarboxylic acid (TCA) cycle, (2) reduce the NAD couple, (3) decrease free radical formation, (4) increase production of ATP, (5) increase mitochondrial uncoupling, (6) increase glutathione peroxidase activity, and (7) inhibit pyruvate entry into the TCA cycle. CoA, coenzyme A; GSH, reduced glutathione; GSSG, oxidized glutathione; NAD<sup>+</sup>, nicotinamide adenine dinucleotide; NADH, nicotinamide adenine dinucleotide hydrogenase; NADP, nicotinamide adenine dinucleotide phosphate; NADPH, nicotinamide adenine dinucleotide phosphate hydrogenase; OHB, hydroxybutyrate; PPP, pentose phosphate pathway; UCP, uncoupling protein. Reprinted with permission from [43].

# T cell metabolism drives immunity

Michael D. Buck, David O'Sullivan, and Erika L. Pearce

Department of Pathology and Immunology, Washington University School of Medicine, St. Louis, MO 63110

Lymphocytes must adapt to a wide array of environmental conditions during their development, during which they undergo a dramatic search in this area has yielded surprising findings on ways and metabolites, which have been found to regulate differentiation, function and fate. In this review, we provide an up-to-date resource on lymphocyte metabolism.

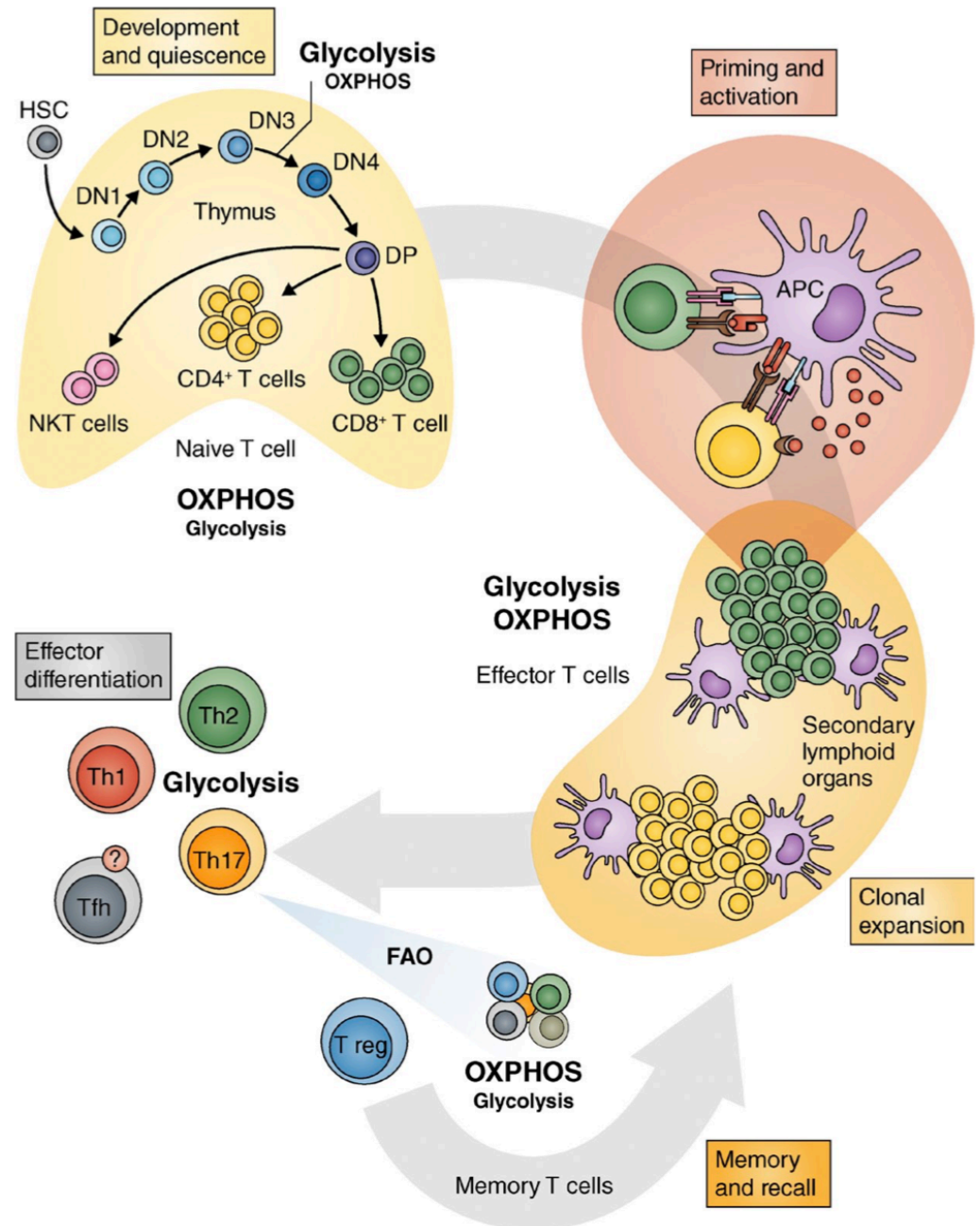
*"Part of the secret of success in life is to eat what you like and let the food fight it out inside."*  
— Mark Twain

Simply stated, we are what we eat. Our genetics, coupled with environmental influences, dictate how we metabolize the nutrients that we consume and how this shapes our growth, function, and overall health. The same principles hold true at the cellular level. Just as a track runner quickly engages their muscles to propel themselves from rest to sprint in response to a starting gun, pathogen-derived or inflammatory signals drive T cells out of quiescence, resulting in rapid modulation of gene expression and the acquisition of new functions. These changes range from increased production of cytokines and cytolytic molecules to the ability to undergo cell division and migration. Intimately integrated into this program of activation is the regulation of cellular metabolism.

The engagement of specific metabolic pathways profoundly affects cell differentiation and function. Metabolic reprogramming is controlled by key receptor signaling events and growth factor cytokines, as well as availability of nutrients. In addition, metabolic products provide substrates that can alter the functional fate of a cell through posttranslational modifications (PTMs) or epigenetic remodeling. Several recent articles have covered these and other emerging topics in T cell metabolism (Chapman and Chi, 2014; Bird, 2015; Lochner et al., 2015; O'Sullivan and Pearce, 2015; Palmer et al., 2015; Ramsay and Cantrell, 2015; Ron-Harel et al., 2015). In this

review, we overview with a focus on the metabolic changes that occur during T cell development and activation.

**T cell development and activation**  
Although the T cell cycle is a complex process, it can be divided into several stages: development and quiescence, priming and activation, effector differentiation, clonal expansion, and memory and recall. Each stage is characterized by distinct metabolic changes that drive the progression of the cycle.



M.D. Buck, D. O'Sullivan, and E.L. Pearce's present address is Dept. of Immunometabolism, Max Planck Institute of Immunobiology and Epigenetics, 79108 Freiburg, Germany.

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## CORRESPONDENCE

Erika L. Pearce:  
pearce@ie-freiburg.mpg.de

Abbreviations used: ACC1, acetyl-CoA carboxylase 1; AMPK, adenosine monophosphate-activated protein kinase; ASS1, argininosuccinate 1; CPT1a, carnitine palmitoyl-transferase 1a; DN, double negative; DP, double positive; ERR $\alpha$ , estrogen-related receptor  $\alpha$ ; ETC, electron transport chain; FAS, fatty acid synthase; HAT, histone acetyltransferase; HDAC, histone deacetylase; IDO, indoleamine-2,3-dioxygenase; PDK1, pyruvate dehydrogenase kinase; PDPK1, 3-phosphoinositide-dependent protein kinase-1; PI3K, phosphatidylinositol 3-kinase; PKA, protein kinase A; PTM, post-translational modification; ROS, reactive oxygen species; TCA, tricarboxylic acid; TPPII, tripeptidyl peptidase II; TSC1, tuberous sclerosis complex 1.