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Increased reactive oxygen species production during reductive stress: The roles of mitochondrial glutathione and thioredoxin reductases

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ABSTRACT

Both extremes of redox balance are known to cause cardiac injury, with mounting evidence revealing that the 17 injury induced by both oxidative and reductive stress is oxidative in nature. During reductive stress, when elec- 18 tron acceptors are expected to be mostly reduced, some redox proteins can donate electrons to O₂ instead, which 19 increases reactive oxygen species (ROS) production. However, the high level of reducing equivalents also con-20 comitantly enhances ROS scavenging systems involving redox couples such as NADPH/NADP⁺ and GSH/GSSG. 21 Here our objective was to explore how reductive stress paradoxically increases net mitochondrial ROS production 22 despite the concomitant enhancement of ROS scavenging systems. Using recombinant enzymes and isolated perme-23 abilized cardiac mitochondria, we show that two normally antioxidant matrix NADPH reductases, glutathione re- 24 ductase and thioredoxin reductase, generate H₂O₂ by leaking electrons from their reduced flavoprotein to O₂ 25 when electron flow is impaired by inhibitors or because of limited availability of their natural electron acceptors, 26 GSSG and oxidized thioredoxin. The spillover of H₂O₂ under these conditions depends on H₂O₂ reduction by 27 peroxiredoxin activity, which may regulate redox signaling in response to endogenous or exogenous factors. 28 These findings may explain how ROS production during reductive stress overwhelms ROS scavenging capability, 29 generating the net mitochondrial ROS spillover causing oxidative injury. These enzymes could potentially be 30 targeted to increase cancer cell death or modulate H₂O₂-induced redox signaling to protect the heart against 31 ischemia/reperfusion damage. 32

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38 1. Introduction

Both the production and removal of reactive oxygen species (ROS) in 39 the mitochondrial matrix play critical roles regulating mitochondrial 40 41 function. ROS-induced redox modifications of mitochondrial proteins are required for redox signaling to adjust metabolism to changing condi-42tions and to regulate cell death and survival pathways. Conventionally, an 43 excess of ROS and reactive nitrogen species (RNS) relative to reducing 44 45equivalents is defined as oxidative stress. Conversely, a relative shortage of ROS compared with reducing equivalents in the form of redox couples 46 (GSH/GSSG, NADPH/NADP⁺, NADH/NAD⁺, etc.) is defined as reductive 47

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http://dx.doi.org/10.1016/j.bbabio.2015.02.012 0005-2728/© 2015 Published by Elsevier B.V. stress [1]. However, the latter definition is somewhat confusing because 48 in isolated mitochondria, reductive stress, in the form of high matrix 49 NADH/NAD⁺ ratios, is known to promote excessive ROS production 50 to a level exceeding ROS scavenging capability, resulting in net 51 H_2O_2 spillover from mitochondria [2]. Specifically, experiments have 52 demonstrated that ROS production by alamethicin-permeabilized mito- 53 chondria increases steeply at high NADH/NAD ratios [3], which is tradi- 54 tionally attributed to complex I of the respiratory chain, but can also 55 involve certain tricarboxylic acid cycle (TCA) enzymes, specifically 56 those containing lipoamide dehydrogenase, the ROS-producing E3 57 component of alpha-ketoglutarate dehydrogenase and pyruvate dehy-58 drogenase [3]. 59

When the NADH/NAD⁺ pool becomes highly reduced, so does the 60 NADPH/NADP⁺ pool, which increases matrix antioxidant power to 61 compensate for increase in ROS production and limit ROS spillover. 62 NADPH is a key reducing equivalent supplying the major H_2O_2 scavenging 63 systems in the matrix, the glutathione/glutathione peroxidase3 (Gpx)/ 64 glutathione reductase (GR) and the peroxiredoxin3 (Prx)/thioredoxin2 65 (Trx)/thioreductase2 (TrxR2) systems (Fig. 1). Although catalase is also 66 present in the matrix [4,5], it has a much lower affinity for H_2O_2 [6] and 67 therefore is not likely to play an important role at micromolar concentrations of H_2O_2 . Thus, matrix antioxidant power is ultimately dependent on 69

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Abbreviations: CDNB, 1-chloro-2,4-dinitrobenzene; FCCP, trifluorocarbonylcyanide phenylhydrazone; GDH, glutamate dehydrogenase; GR, glutathione reductase; Gpx1, glutathione perioxidase; GSH, reduced glutathione; GSSG, oxidized glutathione; H₂O₂, hydrogen peroxide; ICDH, isocitrate dehydrogenase; pCMB, p-chloromer<u>c</u>uribenzoic acid; pCMPS, p-chloromercuriphenyl-sulphonate; Prx, peroxiredoxin; O₂ • , superoxide; TCA, tricarboxylic acid; TrxR1, thioredoxin reductase 1; TrxR2, thioredoxin reductase 2; Trx, thioredoxin 1 and 2

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Fig. 1. Proposed mechanism of ROS generation by GR and TrxR2 during reductive stress. Flow of electrons (e^-) as indicated from NADPH to FAD to cysteine-related sulfhydryl groups (S) and then to the natural electron acceptors GSSG for GR, and oxidized Trx for TrxR2. If oxidized GSSG or Trx is in limited supply, the flavin reduction state is increased and electron flow is shunted from FADH₂ to O₂ as an alternative electron acceptor, with one-electron reduction producing superoxide, two-electron reduction producing H₂O₂ or four-electron reduction reduction producing H₂O. Similarly, thiol reactive agents which react with SH groups in GR or TrxR2 also prevent electron transfer to the natural electron acceptors, resulting in superoxide, H₂O₂ and H₂O production instead.

an adequate supply of NADPH generated by transhydrogenase, isocitrate
dehydrogenase, malic enzyme and glutamate dehydrogenase to keep the
matrix NADPH/NADP⁺ pool reduced.

However, experimental evidence indicates that the enhancement of 73antioxidant power by the reduced NADPH/NADP⁺ pool during reductive 74 stress is insufficient to counterbalance the increased ROS production 7576 resulting from the reduced NADH/NAD⁺ pool, leading to net ROS spillover, matrix oxidation, cytotoxicity and protein aggregation causing 77 cardiomyopathy [7–10]. These observations and others support the 78 79 "redox-optimized ROS balance" hypothesis recently put forth by O'Rourke and colleagues [11,12]. According to this hypothesis, redox balance is 80 81 lost and ROS production increases at both extremes of oxidation and 82 reduction of redox couples involved in respiratory chain activity (NADH/NAD⁺) or ROS scavenging (NADPH/NADP⁺, GSH/GSSG). In 83 84 other words, the extent of ROS production is defined by the overall change of matrix redox environment in either direction. In a highly re-85 86 duced environment, ROS production is accelerated to the point that it overwhelms ROS scavenging capacity, even though the latter should 87 be maximally potentiated [11,12]. 88

To better understand the factors that lead to net ROS production and 89 90 oxidative cytotoxicity during reductive stress, we explored the possibility 91 that two of the NADPH-dependent reductases that normally enhance antioxidant function, namely GR and TrxR2, begin to directly generate 92significant ROS when the NADPH/NADP⁺ pool becomes highly reduced. 93 This hypothesis is based on the premise that these enzymes belong to 94the same family of disulfide reductase flavoenzymes as lipoamide dehy-95drogenase, the E3 component of α -ketoglutarate dehydrogenase and 96 pyruvate dehydrogenase. Lipoamide dehydrogenase has been shown 97 to produce ROS with a rate equal to or even higher than complex I 98 when the NADH/NAD⁺ pool is highly reduced [3,13]. It is structurally 99 similar and catalyzes the transfer of electrons between pyridine nucle-100 otides and disulfides with very similar reaction chemistry as GR and 101 TrxR's. Indeed, the cytoplasmic isoform of thioredoxin reductase 102 (TrxR1) has been previously shown to generate ROS robustly in the 103 presence of NADPH when its preferred endogenous substrate, oxidized 104 105 thioredoxin 1 (Trx), is in limited supply [14]. Under these conditions, it is thought that the reduced flavoprotein redox site of TrxR1 directly 106 donates electrons obtained from NADPH to molecular O₂, to generate 107 superoxide and H₂O₂ by one or two electron reduction of O₂, respectively. 108

In this study, we tested the hypothesis that, analogous to lipoamide 109 dehydrogenase and TrxR1, GR and TrxR2 generate ROS when their 110 natural electron acceptors (GSSG and oxidized Trx, respectively) are in 111 limited supply, as illustrated schematically in Fig. 1. Using recombinant 112 GR and TrxR1, we show that both enzymes generate robust amounts of 113 H₂O₂ when provided with NADPH in the absence of their preferred 114 electron acceptors, or in the presence of inhibitors which interfere 115 with electron transfer to their preferred electron acceptors (Fig. 1). 116 Moreover, we present evidence that these same reactions occur with 117 GR and TrxR2 in cardiac mitochondria exposed to reductive stress, and 118 that the amounts of ROS generated under these conditions are quantita- 119 tively comparable to those generated by respiratory complexes and TCA 120 cycle dehydrogenases during reductive stress. Finally, we show that ROS 121 spillover during reductive stress is very sensitive to Prx inhibition by 122 Zn²⁺. These findings help to explain how respiratory complexes and 123 NAD⁺-related matrix dehydrogenases, together with GR and TrxR2 124 directly generating ROS, combine to overwhelm NADPH-dependent 125 antioxidant mechanisms, resulting in ROS efflux from the matrix during 126 reductive stress. 127

2. Material and methods

This study was approved by the UCLA Chancellor's Animal Research 129 Committee (ARC 2003-063-23B) and performed in accordance with the 130 Guide for the Care and Use of Laboratory Animals published by the 131 United States National Institutes of Health (NIH Publication No. 85-23, 132 revised 1996) and with UCLA Policy 990 on the Use of Laboratory Animal Subjects in Research (revised 2010). 134

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All measurements were carried out using customized Fiber Optic 135 Spectrofluorometer (Ocean Optics) in a continuously stirred cuvette at 136 room temperature (22 to 24 °C). The cuvette was partially open to air 137 during isolated mitochondria experiments, but was tightly closed 138 when O_2 consumption by recombinant GR was measured. Mitochondria 139

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