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Factors influencing post-exercise plasma protein carbonyl

concentration

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Abstract

Exercise of sufficient intensity and duration can cause acute oxidative stress. Plasma protein carbonyl (PC) moieties are abundant, chemically stable and easily detectable markers of oxidative stress that are widely used for the interpretation of exercise-induced changes in redox balance. Despite many studies reporting acute increases in plasma PC concentration in response to exercise, some studies, including those from our own laboratory have shown decreases. This review will discuss the differences between studies reporting increases, decreases and no change in plasma PC concentration following exercise in humans; highlighting participant physiology (i.e. training status) and study design (i.e. intensity, duration and novelty of the exercise bout) as the main factors driving the direction of the PC response to exercise. The role of the 20S proteasome system is proposed as a possible mechanism mediating the clearance of plasma PC following exercise. Resting and exerciseinduced differences in plasma protein composition and balance between tissues are also discussed. We suggest that exercise may stimulate the clearance of plasma PC present at baseline, while simultaneously increasing reactive oxygen species production that facilitates the formation of new PC groups. The balance between these two processes likely explains why some studies have reported no change or even decreases in plasma PC level postexercise when other biomarkers of oxidative stress (e.g., markers of lipid peroxidation) were elevated. Future studies should determine factors that influence the balance between PC clearance and formation following acute exercise.

Keywords: Protein oxidation, exercise, proteasome, protein degradation, reactive oxygen species

1 Introduction

2 Exercise can induce a wide range of whole body physiological adaptations that improve metabolic health and lower oxidative stress [1-3]. Oxidative stress is a biological 3 4 state whereby reactive oxygen species (ROS) overwhelm antioxidant defences, increasing the 5 oxidation of proteins, lipids and DNA. It is widely accepted that transient increases in 6 exercise-induced ROS can initiate a diverse range of signalling pathways that lead to adaptation [4-6]. Indirect biomarkers of exercise-induced oxidative stress, such as protein 7 8 oxidation [7–9], lipid peroxidation [8–10] and antioxidant capacity [8,11,12] are routinely 9 measured to give an indication of altered redox balance. One of the most frequently examined 10 biomarkers of protein oxidation is plasma protein carbonyl (PC) concentration. Carbonylation 11 is a stable and quantifiable post-translational protein modification which is ten times more 12 abundant than other protein adducts, such as 4-Hydroxynonenal and glycooxidation end-13 products [13,14]. Most biomarkers of oxidative stress increase in plasma in response to 14 exercise [11,12,15], as would be expected following an acute bout of increased metabolic 15 activity. However some studies have reported decreases in plasma PC concentration post-16 exercise alongside increases in other biomarkers of oxidative stress [8,16,17]. The focus of this review is to explore key physiological factors that might explain these different 17 18 responses, with a primary emphasis on aerobic, steady state exercise bouts where PC groups 19 have been measured in blood plasma or serum of human participants exercising under fasted 20 conditions. It is beyond the scope of this review to include the results of studies that have 21 examined the effect of habitual diet or dietary supplementation (e.g., high dose antioxidants) 22 on resting or exercise-induced changes in PC levels.

23

24 **Protein carbonyl formation**

25 PC groups are present in all proteins (carboxylic acid (-COOH) groups), form the basis of their structural integrity, and influence their capacity to function and interact with 26 27 other molecules. Formation of additional, non-native PC groups can be a result of a variety of 28 irreversible, non-enzymatic oxidative pathways (Figure 1) that are a normal part of metabolic processes [18]. These include direct oxidation of amino acids (in conjunction with oxidising 29 30 agents such as transition metal clusters in protein structures) and secondary oxidation reactions with lipid peroxidation or glucose-protein oxidation products [19]. PC groups are 31 32 highly polar, thus increasing the proteins susceptibility to further oxidation and/or formation 33 of cross-linkages and protein aggregates (via ketone or aldehyde links) [20]. Excessive 34 introduction of new PC groups can result in altered or disrupted protein function and 35 exposure of previously embedded hydrophobic groups in the protein core, resulting in 36 targeted proteolytic degradation by 20S and 11S proteasome systems [20,21]. The degree of 37 PC formation is dependent on the presence of conjugated metals (e.g., iron [19]), the 38 orientation of specific amino acids that are more susceptible to PC formation (i.e., Proline, Arginine, Lysine, and Threonine) and importantly, the magnitude and origin of ROS 39 40 production in relation to the protein [22].

41

43

44 Exercise-induced changes in protein carbonylation

A variety of cells can produce ROS in response to exercise, most notably myocytes [23], leukocytes [24] and endothelial cells [25]. Superoxide (O_2^{-}) is produced from a range enzymatic sources within these cell types during exercise, such as nicotinamide adenine dinucleotide phosphate (NADPH) oxidase, xanthine oxidase and nitric oxide synthases [26]. These cells can also release O_2^{-} into the extracellular space via enzymes expressed on the

^{42 [}Insert Figure 1 here]

50 plasma membrane such as NADPH oxidase [26-28] or via passive diffusion of uncharged 51 ROS, such as hydrogen peroxide [29]. Consequently, plasma proteins are susceptible to the 52 formation of PC groups during exercise [7,9,30,31], with evidence that new PC moieties are 53 stable in plasma for up to 4 hours, before selective degradation [32,33]. Increased formation 54 of plasma PC groups following exercise is considered to be a non-specific reflection of 55 increased systemic oxidative stress (i.e. the origin of ROS and biological impact is unknown). 56 As a result, many studies over the last 15 years have investigated associations between 57 changes in plasma PC concentration and aspects of the acute physiological stress caused by 58 exercise, in a variety of populations [7,9].

59

60 Factors influencing post-exercise changes in protein carbonylation

61 *Exercise intensity*

62 Aside from the potential for greater production of ROS from the cellular sources 63 discussed above, specific hypoxic mechanisms can also contribute during high intensity 64 exercise. Repetitive cycles of temporary occlusion (hypoxia) and re-oxygenation of the blood 65 vessels surrounding actively contracting muscle can produce large quantities of ROS via the 66 enzyme xanthine oxidase [34], that may increase PC formation. Previous studies have 67 reported increases in plasma PC concentration following high-intensity exercise to exhaustion relative to baseline values [12,31,35-37]. Lamprecht et al [30] assessed the 68 69 impact of exercise intensity on plasma PC formation by examining three 40-minute cycling bouts of different intensities (70, 75 and 80% $\dot{V}O_{2max}$) in three independent groups of 70 71 moderately active participants (Table 1). Plasma PC concentration increased in response to cycling at 80% $\dot{V}O_{2max}$ only, suggesting that exercise intensity is a determinant of the 72 73 formation of new non-native carbonyl moieties in the bloodstream. However it is likely that 74 exercise duration and other physiological factors also have an impact upon these processes.

76 *Exercise duration*

77 There is some evidence to suggest that exercise duration is also a key factor in post-78 exercise plasma PC formation. Bloomer et al [7] reported that 120 minutes of cycling at 70% $\dot{V}O_{2max}$ caused a greater increase in post-exercise plasma PC concentrations than 30 and 60-79 80 minute bouts of the same exercise intensity in male and female participants. Moreover, an 81 exercise bout of moderate intensity, but long duration (ultra-endurance running: 174 km, 30-82 44 hours) has been reported to elicit immediate and prolonged (7 days) post-exercise 83 increases in plasma PC concentration [38]. This increased protein oxidation occurred 84 simultaneously with a decline of exogenous and/or endogenous antioxidants, which may have 85 reduced the capacity to clear PC groups within this seven day period. The roles of exercise 86 intensity and duration on post-exercise PC concentrations are inevitably linked; however 87 there is evidence that other physiological factors (e.g., training status) contribute [7,15].

88

89 <u>Training status and habituation to exercise</u>

90 Exercise-induced ROS production can initiate a cascade of cellular signals which 91 result in various post-translational modifications (i.e. phosphorylation, acetylation and thiol 92 modifications), and up-regulate the expression of antioxidant and stress proteins following 93 exercise [4,39,40]. Differences in the resting expression of antioxidant proteins between 94 participants (i.e. due to training status and/or habituation to exercise) will no doubt have 95 consequences for changes in exercise-induced oxidative stress. Indeed, it has been 96 demonstrated that exercise training can stimulate an increased expression of endogenous 97 antioxidant proteins [5,41], with some direct evidence of this in humans [42]. However, it is 98 largely unclear how much resting antioxidant capacity impacts upon the acute oxidative 99 stress response to exercise. Elevated endogenous antioxidant capacity may enable a buffering 100 of ROS production subsequently reducing the magnitude of oxidative stress biomarker 101 formation during exercise in trained individuals [43]. Bloomer et al [7] reported increases in 102 plasma PC concentration immediately following cycling exercise (30 minutes at 70% $\dot{V}O_{2max}$) (*Table 1*), but in a separate study, with different research participants, reported no 103 104 change in plasma PC level following an identical cycling protocol [15]. Differences in the 105 physiology and exercise training habits of the participants featured in these studies, rather 106 than the intensity and duration of the exercise bout are likely to have influenced the 107 differential net changes in PC post-exercise. For example, there were clear differences in both the aerobic fitness ($\dot{V}O_{2max}$: 57 ± 5 ml/kg/min [7] vs. 45 ± 8 ml/kg/min [15]) and time 108 109 engaged in aerobic exercise prior to the study (10.4 \pm 1.6 hours/week [7] vs. 2.8 \pm 2.2 110 hours/week [15]). In addition, the participants in the study published by Bloomer et al, 2005 111 [15] were more resistance $(3.8 \pm 1.8 \text{ hours/week})$, than aerobically trained. This suggests that 112 the exercise stimulus implemented in both studies was more novel for the participants in the 113 2005 study [15]. This supports some previous work reporting a greater magnitude of 114 oxidative stress biomarker formation following unaccustomed exercise [44,45]. Thus, a 115 combination of factors such as the novelty of exercise, and aspects of study design (i.e., 116 exercise intensity and duration) are likely to interact to govern the magnitude of increase in 117 plasma PC formation following exercise. However, it is perhaps more challenging to explain 118 why decreases in plasma PC level have been observed.

119

120 Studies reporting *decreases* in plasma PC concentration after exercise

121 The available evidence suggests exercise intensity [30] and duration [7] influence the 122 magnitude of ROS production and the associated increase in biomarkers of oxidative stress. It 123 is therefore not surprising that many studies have reported no change in plasma PC 124 concentration following sub-maximal exercise [15,46–50]. However, counter-intuitively, many studies report a *decrease* in plasma PC concentration following both sub-maximal and maximal exercise [8,16,17,51] (*Table 1*). Importantly, these changes have sometimes been reported alongside increases in other biomarkers of oxidative stress [8,17]. This is unexpected and implies that exercise stimulates clearance processes alongside exerciseinduced ROS production.

We have shown that steady state submaximal cycling (60% and 80% $\dot{V}O_{2max}$ for 27 130 131 [moderate intensity] and 20 minutes [high intensity] respectively) undertaken by untrained 132 males ($\dot{V}O_{2max}$; 42.7 ± 5.0 ml/kg/min) elicits a 9% (moderate intensity, p<.0001) and 4% (high intensity, p<.0001) mean decrease in plasma PC concentration (see figure 2: A and B) 133 [8]. Importantly, these changes occurred in parallel with increases in plasma lipid 134 135 hydroperoxides (LOOH), total antioxidant capacity (TAC) and cellular markers of oxidative 136 stress [52]. Interestingly, the participants in this study engaged in less than 3 hours of generic 137 aerobic exercise per week, indicating that these bouts were relatively unaccustomed. These 138 findings are not limited to just moderately trained individuals. We have also shown a 7% 139 decrease (p=.016) in plasma PC concentration immediately after a bout of cycling exercise (70% $\dot{V}O_{2max}$, 75 min) in highly trained cyclists ($\dot{V}O_{2max}$; 63.7 ± 5.3 ml/kg/min) (Wadley et 140 141 al, 2015; In Preparation; see figure 2D). Finally, our findings are not limited to sub-maximal 142 exercise: we have also shown a 10% mean decrease (p=.002) in plasma PC concentration immediately after a graded exercise test to volitional exhaustion (i.e., 100% \dot{V} O_{2max}, 143 approximately 15-minutes) in very active young men ($\dot{V}O_{2max}$; 61.9 ± 4.7 ml/kg/min; see 144 145 Figure 2C) (Turner *et al*; unpublished data) and a 13% mean decrease (p<.0001) following a bout of low volume high intensity interval exercise (10×1 minute stages at $90\% \dot{V}O_{2max}$, with 146 9×1 minute rest intervals at 40% $\dot{V}O_{2max}$) in moderately active participants ($\dot{V}O_{2max}$; 42.7 ± 147 5.0 ml/kg/min) [8]. 148

149 There are a number of other studies that report (but do not always comment upon) 150 decreases in plasma PC levels following exercise (Table 1). For example, Chevion et al, [16] reported decreases in PC concentration following a study involving two walks (50km and 151 152 80km) of unreported intensity. The magnitude of this decrease in PC concentration was 153 greater following the first walk compared to the second, highlighting again that the novelty of 154 the exercise stimulus may be a key factor modulating post-exercise PC concentration. Furthermore, reductions in plasma PC level have been reported immediately [17,51] and up 155 156 to four hours [17] following submaximal cycling exercise in moderately trained participants.

157

158 [Insert Figure 2 here]

159

160 **Possible mechanisms mediating protein carbonyl clearance in response to exercise**

161 <u>The 20S proteasome system</u>

162 The proteasome system is an organised assembly of proteins present in all cell types 163 that functions to degrade irreversibly modified proteins, such as those containing carbonyl 164 groups. The ubiquitin-independent 20S proteasome is the primary system in place to degrade oxidatively damaged proteins [21,32,53], with recent evidence also suggesting the 11S 165 166 proteasome facilitates this process under conditions of heightened oxidative stress [20]. 167 Increased exposure of carbonyl-mediated hydrophobic groups within the protein core can 168 increase targeted degradation by these proteasome systems within cells. The 20S proteasome 169 is also excreted into extracellular fluids such as plasma [54-56] and is known to be 170 enzymatically functional [57], suggesting that it could cleave PC groups in plasma directly. 171 Studies in humans have found that exercise can acutely increase ubiquitin-dependent 172 proteasome gene expression in response to resistance [58–60] and ultra-endurance exercise 173 [61]. However, the impact of acute shorter-duration aerobic-type exercise on the ubiquitin174 independent 20S and 11S proteasome subunits (either intra- or extra-cellular; expression or 175 activity) and the relationship with plasma PC has not been investigated in humans. Data from 176 rats has indicated that chymotrypsin-like activity of the ubiquitin-independent 20S 177 proteasome subunit increases in brain tissue following 8 weeks of exercise overload (1 hour 178 swimming/day, 5 days a week for 6 weeks) [62].

179 Exercise bouts below a certain intensity and/or duration, in combination with the other physiological parameters discussed above may activate intracellular and extracellular 180 181 proteasome pathways to clear modified proteins and thus lower the concentration of PC 182 groups in plasma (Figure 3). The net increase in PC level observed in response to high 183 intensity or prolonged duration exercise may result from the formation of new PC groups 184 outnumbering proteolytic clearance of those present at baseline. Furthermore, inferences 185 from cell culture data [53] led Radak et al [63] to propose that proteasome activity may follow a hormetic-type response with increasing exercise-induced oxidative stress. 186 187 Proteasome activity may be reduced at higher exercise intensities or after prolonged duration 188 exercise, due to ROS-induced inactivation of the functional 20S or 11S proteasome. Future 189 work is needed to explore this mechanism.

190

191 <u>Resting or exercise-induced differences in total plasma protein composition and balance</u>
 192 <u>between tissues</u>

193 Studies examining plasma PC concentration typically express results relative to the 194 total protein concentration of plasma (i.e., nM of carbonyls per mg of protein) [64]. 195 Measuring protein carbonylation in this way does not account for possible exercise-induced 196 shifts in protein composition. This is relevant because certain plasma proteins, such as 197 serotransferrin and fibrinogen (approximately 2-4% of total plasma proteins) are more 198 susceptible to exercise-induced oxidation than other plasma proteins [18]. Therefore, individual differences in baseline plasma protein composition may influence post-exercise
changes in PC. Furthermore, exercise could induce shifts in the proportion of certain proteins
in plasma, which could result in exaggerated or suppressed protein carbonylation.

202 Although not as susceptible to oxidation as fibrinogen, albumin, the most abundant 203 plasma protein, has also been shown to exhibit a high concentration of PC groups [65]. The 204 extent to which albumin becomes oxidised in response to exercise is dependent on intensity, 205 and albumin carbonylation is marked at intensities of 80% VO_{2max} or more [30]. Inter-206 individual variation in exercise-induced proteinurea or albuminuria has been documented 207 [66,67], and might affect the amount and type of plasma proteins that can be oxidised during 208 exercise. Furthermore, it has been shown that during and following both aerobic and 209 resistance exercise, there is an increase in the uptake and turnover of proteins such as 210 albumin and fibrinogen by muscle [68,69]. Thus, differences in PC excretion and balance 211 between tissues might explain to some degree, alterations in the composition of plasma proteins, which may increase or decrease plasma PC level. 212

213

214 Experimental approaches for monitoring changes in plasma protein carbonyl level

The evidence presented in this review highlights the uncertainty with regards to the mechanistic underpinnings of post-exercise decreases in plasma protein carbonylation. Despite the factors discussed, experimental design and the analytical techniques used to quantify protein carbonyl concentration in the cited studies warrants discussion.

The majority of studies included in *Table one* controlled for factors such as age, training status, dietary habits (i.e. fasted exercise trials), and the time of day that exercise was undertaken. All of these variables have been shown to alter protein balance through changes in protein uptake to skeletal muscle during exercise [68,69], highlighting the rigorous experimental approach taken in these studies. However, an important consideration is the 224 timing of blood sampling. Many of the studies presented assessed protein carbonyl 225 concentration immediately following cessation of the exercise bout. It is conceivable that changes within the minutes and hours after exercise might give more insight into the 226 227 'conundrum' of post-exercise changes in protein carbonyl level. Indeed, independent studies 228 have shown increases [70] and decreases [17] in protein carbonyl concentration up to 4 hours 229 post-exercise. In this regard, it is clear that extensive timecourse analysis of protein carbonyl 230 concentration is needed to validate these findings and importantly, to elucidate the 231 mechanisms influencing decreases following exercise.

232 It is important to note that a variety of analytical techniques are used to quantify 233 protein carbonylation in the literature. These have primarily included spectrophotometric and 234 ELISA-based methods, which have recently come under scrutiny with regards to their 235 sensitivity [71] and reproducibility [72]. However, while these studies provide an example of 236 an important limitation of quantifying protein carbonyl concentration, they do not explain 237 why protein carbonyl concentration might decrease following exercise. Furthermore, with 238 independent laboratories now reporting decreases in protein carbonyl level following exercise 239 [8,16,17,51], this reduces the chances of experimental variability causing this effect.

240

241 [Insert Figure 3 here]

242

243 Conclusions

Plasma protein carbonyl concentration is a marker of oxidative stress routinely used to assess exercise-induced redox regulation. The evidence presented in this review suggests that certain exercise conditions can result in a net *decrease* in plasma PC concentration following exercise, which occurs in parallel with increases in other biomarkers of oxidative stress. Exercise intensity (>70% $\dot{V}O_{2max}$) and prolonged duration (>60 minutes) appear to be the main contributing factors in the observed post-exercise increases in PC concentration. The factors influencing *decreases* in protein carbonyl level are more difficult to interpret, but likely involve the clearance of oxidised proteins from plasma, potentially by plasma proteasomes, excretion, or uptake into active tissues. Studies wishing to assess markers of oxidative stress in response to exercise should assess PC concentration together with other biomarkers over an extended time course (immediately after and up to 4 hours post-exercise) for a true representative assessment [31]).

256

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260 article and revising it critically for important intellectual content.

261

262 **Declaration of Interest**

263 None of the authors declare a conflict of interest and have no financial interest in the study.

264

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Figure Legends

Figure 1: Formation of non-native carbonyl groups in proteins. Non-native carbonyl groups can be introduced into proteins by; (1) direct oxidation of amino acids, (2) via secondary oxidation products of lipid peroxidation, and (3) the oxidation products of reducing sugars. An example carbonyl modification is included for each pathway; (A) (2-amino-3-ketobutyric acid, (B) malondialdehyde-Lysine adduct, and (C) carboxymethyl lysine (3)). The non-native carbonyl group in each example is indicated by a dashed circle around the C=O bond of the amino acid or amino acid side chain (adduct).

Figure 2: Published and unpublished data indicating decreases in protein carbonyl concentration (nM/mg protein) in response to exercise. PC level decreases during the last minute (End-Exercise) (A) 27 minutes of cycling at 60% $\dot{V}O_{2max}$ and (B) 20 minutes of cycling at 80% VO_{2max} in untrained young men (n=10; mean ± SD: age 22 ± 3 yrs; $\dot{V}O_{2max}$ 42.7 ± 5.0 ml/kg/min) [8]; C) PC level decreases immediately following a $\dot{V}O_{2max}$ test to exhaustion in trained young men (n=10; mean ± SD: age 23 ± 3 yrs; $\dot{V}O_{2max}$ 61.9 ± 4.7 ml/kg/min) (Turner & Aldred; unpublished work); D) PC level decreases immediately following 75 minutes of cycling at 70% $\dot{V}O_{2max}$, and is elevated above baseline levels 2 hours post-exercise in trained young men (n=12; mean ± SD: age: 28 ± 4 years, $\dot{V}O_{2max}$ 63.7 ± 5.3 ml/kg/min) (Wadley et al, 2015; *in preparation*). Individual (grey bars) and mean (black line) PC concentration changes are reported. Mean percentage change and statistical significance are indicated above each graph. Paired samples T-tests (A-C) and repeated measures ANOVA (D) were performed using SPSS (PASW Statistics, 22.0).

Figure 3: Proposed formation/ clearance of protein carbonyl groups in response to exercise in humans. Exercise can cause an increase in plasma protein carbonylation when the exercise bout is of sufficient intensity and/or duration. Activation of clearance mechanisms may drive a decrease in PC groups present at baseline when certain exercise conditions are met.