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Exercise decreases PP2A-specific reversible thiol oxidation in human erythrocytes: Implications for redox biomarkers

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Abstract

New readily accessible systemic redox biomarkers are needed to understand the biological roles reactive oxygen species (ROS) play in humans because overtly flawed, technically fraught, and unspecific assays severely hamper translational progress. The antibody-linked oxi-state assay (ALISA) makes it possible to develop valid ROS-sensitive target-specific protein thiol redox state biomarkers in a readily accessible microplate format. Here, we used a maximal exercise bout to disrupt redox homeostasis in a physiologically meaningful way to determine whether the catalytic core of the serine/threonine protein phosphatase PP2A is a candidate systemic redox biomarker in human erythrocytes. We reasoned that: constitutive oxidative stress (e.g., haemoglobin autoxidation) would sensitise erythrocytes to disrupted ion homeostasis as manifested by increased oxidation of the ion regulatory phosphatase PP2A. Unexpectedly, an acute bout of maximal exercise lasting ~16 min decreased PP2A-specific reversible thiol oxidation (redox ratio, rest: 0.46; exercise: 0.33) without changing PP2A content (rest: 193 pg/ml; exercise: 191 pg/ml). The need for only 3-4 µl of sample to perform ALISA means PP2A-specific reversible thiol oxidation is a capillary—fingertip blood—compatible candidate redox biomarker. Consistent with biologically meaningful redox regulation, thiol reductant-inducible PP2A activity was significantly greater (+10%) at rest compared to exercise. We establish a route to developing new readily measurable protein thiol redox biomarkers for understanding the biological roles ROS play in humans.

Key words: ALISA, Protein thiol, Biomarker, Redox Signalling, Oxidative stress, Exercise, PP2A, Human.

40 **Introduction**

41

42 Unravelling regulatory roles for reactive oxygen species (ROS), a global term
43 subsuming chemically heterogeneous free radical and non-radical molecules [1], in humans
44 relies on using biomarkers to understand redox functionality: what ROS do and how they do
45 it [2]. For example, redox biomarkers (e.g., lipid peroxidation products) together with several
46 molecular surrogates (e.g., mRNA transcripts) show that manipulating ROS with small-
47 molecule antioxidants (e.g., vitamin C) impairs exercise adaptations [3–6]. ROS, therefore,
48 regulate exercise adaptations (i.e., what) by activating beneficial cell signalling cascades (i.e.,
49 how) [7–9]. Discovering regulatory roles for ROS in humans is intrinsically linked to leveraging
50 technical advances to develop new redox biomarkers [10].

51

52 The antibody-linked oxi-state assay (ALISA) is a new technical advance for measuring
53 target-specific protein thiol redox state in a microplate [11]. A protein thiol (i.e., cysteine
54 residue) can be reduced or oxidised (oxidation can be reversible or irreversible) [12,13]. The
55 vast majority of the >200,000 cysteine residues in the human proteome are highly reduced
56 (>85%) [14]. Much redox signalling is mediated by direct and/or indirect electron exchange
57 between ROS and protein thiols [15–20]. Changing protein thiol redox state can control their
58 activity, phase, lifetime, location, and interactome [21–24]. Redox signals can be catalysed
59 and reversed by the glutathione and thioredoxin dependent systems. Measuring target-
60 specific protein thiol redox state has great potential for understanding the pleiotropic roles
61 ROS play in humans [25]. Reasons to measure target-specific protein thiol redox state are
62 threefold:

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64 1. **Redox signalling.** Target-specific protein thiol redox state changes can be used to infer
65 context- and process-specific roles for ROS dependent redox signalling [26–28].
66 Measuring exercise-induced redox signalling is important because mechanistic links
67 tying antioxidants to the target-specific protein thiol redox state changes responsible
68 for regulating exercise adaptations are lacking [29].

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70 2. **Biomarkers.** A target-specific protein thiol redox state change delivers an accessible
71 and easily measurable process-specific or general candidate biomarker. For example,
72 it might be possible to find new biomarkers of systemic exercise-induced oxidative
73 stress: as chemically defined by a target-specific protein thiol redox state change.
74 Beyond expanding the available analytical arsenal [30], clearly defining oxidative
75 stress by specifying the change is essential because chemical ambiguity fosters
76 misunderstanding [31–33]. Interpretational clarity stems from target-specific protein
77 thiol redox state changes reflecting a difference in reversible thiol oxidation formation
78 and/or reduction [34,35].

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80 3. **Validity & specificity.** Unlike many overtly flawed or severely limited redox biomarker
81 assays [36], ALISA is valid and target-specific. In considering specificity, global protein
82 carbonylation, for example, is a general metric: it reports mean proteome-wide
83 carbonylation. Consequently, relating altered protein carbonylation to function is
84 challenging without targeted analysis. ALISA can relate functionally annotated redox
85 switches to target-specific protein thiol redox state changes. For example, reversible
86 thiol oxidation inhibits the tyrosine protein phosphatase PTP1B by disabling

87 nucleophilic catalysis [37–39]. One could infer an increase in PTP1B-specific reversible
 88 thiol oxidation as inhibitory, especially when combined with a protein activity proxy
 89 [35]. Understanding change functionality is key to interpreting oxidative stress (i.e.,
 90 the change may be good, bad, or neutral).

91
 92 To advance translational research, we used exercise to disrupt redox homeostasis in a
 93 biologically meaningful way to unearth target-specific protein thiol redox responses in human
 94 erythrocytes. Chronic oxidative stress sensitises erythrocytes to additional exercise-induced
 95 mechanical, thermal, chemical, and ionic stress [40]. We reasoned that: erythrocytes might
 96 struggle to defend ion homeostasis when exercise-induced oxidative stress is imposed, which
 97 may, in turn, increase the oxidation of the catalytic core subunit (PPP2CA, UniProt: #P67775)
 98 of the ion regulating and redox-sensitive serine/threonine protein phosphatase PP2A [41,42].
 99 PP2A was selected, therefore, because it plays a key homeostatic role in the erythrocyte and
 100 it is a redox-sensitive target protein. Here, we used ALISA to measure PP2A-specific redox
 101 state at rest and immediately after maximal exercise in human erythrocytes. Experiments
 102 were designed to unambiguously answer a clear question: does an acute bout of maximal
 103 exercise change PP2A redox state? The answer could pave the way to developing new target-
 104 specific systemic redox biomarkers for unravelling the biological roles ROS play in humans.

105 Results

106 Experimental approach

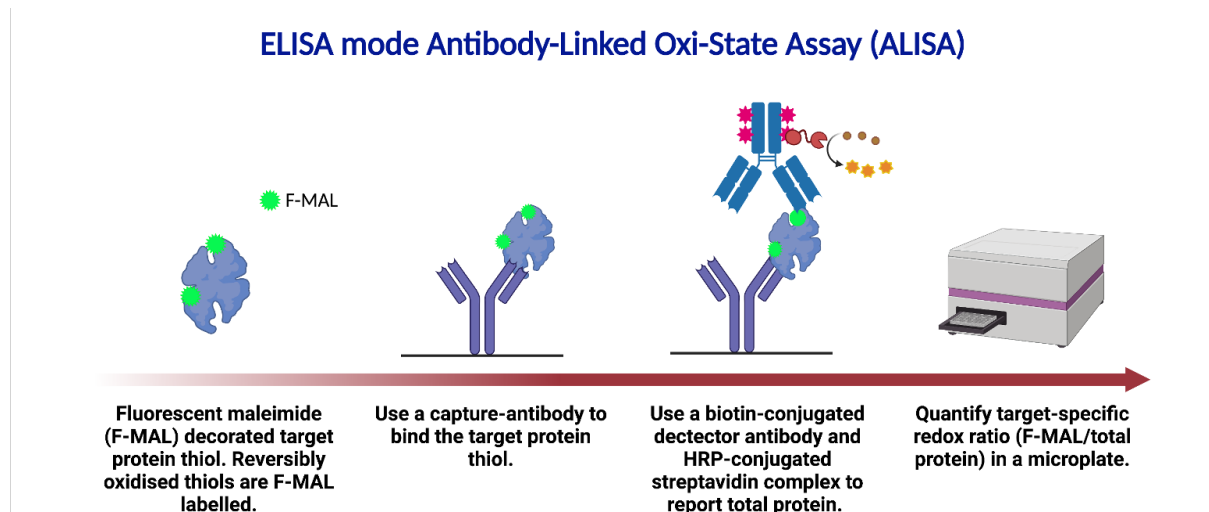
107
 108 Blood samples were collected from adult human males at rest and immediately after
 109 an acute bout of maximal exercise. Physiological markers confirmed the maximal nature of
 110 the exercise stimulus ([Table 1](#)). Blood samples were processed for ALISA. ALISA uses a thiol-
 111 reactive fluorescent-conjugated maleimide reporter (i.e., F-MAL) to label reversibly oxidised
 112 thiols and a capture antibody to bind the target protein [11]. The target-specific reversible
 113 thiol oxidation (i.e., the F-MAL signal) is normalised to the total amount of capture antibody
 114 bound target using a biotin-conjugated detector antibody and recombinant protein standard
 115 curve ([Figure 1](#)). Target-specific redox state is calculated as $(F-MAL/[target])$.

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 119 **Table 1. Adult male participant ($n = 8$) data by physiological parameter. SD denotes the standard deviation.**

Parameter	Mean	SD	Range
Age (years)	38	9	21-47
Height (cm)	179.3	5.8	174-190
Body mass (kg)	79.9	9.5	60.8-88.8
Body fat (%)	12.3	3.8	4.4-16.5
VO _{2max} (ml/kg/min)	47.4	5.7	41-59
Max work rate (Watts)	365	26	342-408
Exercise time (seconds)	946	78	858-1074
Max Heart rate (bpm)	176	9	158-188

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Figure 1. ELISA mode ALISA workflow. Left to right. Reversibly oxidised target protein thiols are decorated with a fluorescent-conjugated thiol-reactive maleimide (F-MAL) reporter using standard labelling protocols (see methods). A microplate immobilised capture antibody is used to bind the F-MAL decorated target protein. After washing (omitted for clarity, see methods), a biotin-conjugated (red groups) detector antibody is used to bind a distinct target protein epitope. Streptavidin-conjugated horseradish peroxidase (HRP) and a turn-on fluorescent HRP substrate are used to quantify total protein relative to a recombinant protein standard curve (omitted for clarity) in a microplate. To terminate the HRP reaction and unmask the F-MAL groups for measurement the antibody-target complex is denatured. Target-specific protein thiol redox state is calculated as: F-MAL/total target.

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No maximal exercise-induced change in pan-erythrocyte reversible protein thiol oxidation in humans

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Consistent with the presence of many F-MAL decorated proteins, qualitative SDS-PAGE analysis identified several distinct F-MAL positive bands corresponding to proteins containing at least one reversibly oxidised thiol in resting and exercised erythrocyte lysates. (Figure 2A). We quantified their redox state relative to haemoglobin (Hb) protein content in a microplate (i.e., total F-MAL/Hb). No exercise-induced change in pan-erythrocyte reversible thiol oxidation was observed (Figure 2B). Given the trend towards statistical significance, it might well be that the global analysis was underpowered.

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Maximal exercise decreases PP2A-specific reversible thiol oxidation in human erythrocytes

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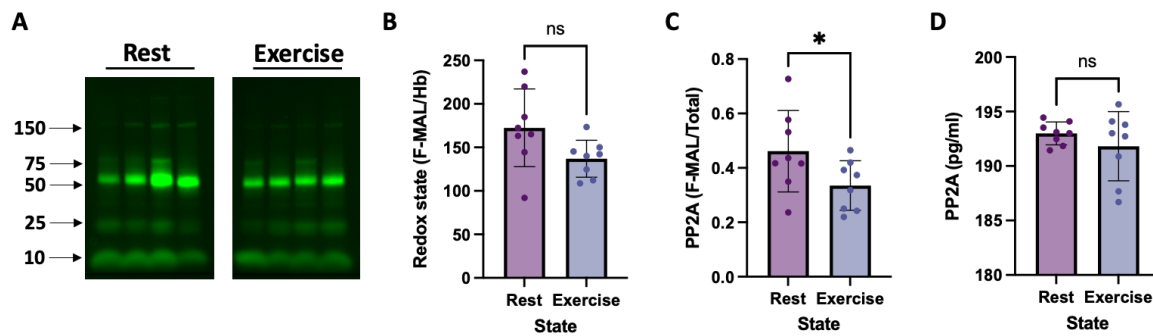
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To measure PP2A-specific reversible thiol oxidation, we used ALISA [11]. Exercise significantly decreased PP2A-specific reversible thiol oxidation without changing PP2A content in human erythrocytes (Figure 2C-D). ALISA required only 3-4 μ l of sample to measure PP2A-specific redox state and content. The coefficient of variation (CV) of resting and exercise technical replicates for PP2A redox state was 8 and 4.5%, respectively. Note that the CV for ALISA will vary on a target-by-target basis.



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160 **Figure 2. Maximal exercise decreases PP2A-specific reversible thiol oxidation in human erythrocytes. A.**
161 **Representative qualitative SDS-PAGE gel image showing the presence of several distinct F-MAL positive bands**
162 **in resting and exercised erythrocyte lysates. Arrows mark various molecular weights (MW) in kilodaltons**
163 **(kDa). B. Quantifying the F-MAL signal relative to Hb in a microplate revealed no significant (paired t-test, $P =$**
164 **0.0832) impact of exercise on global proteome-wide erythrocyte redox state in humans ($n = 8$). C. ALISA**
165 **revealed a significant (paired t-test, $P = 0.0367$) exercise-induced decrease in PP2A-specific reversible thiol**
166 **oxidation in human erythrocytes ($n = 8$). D. No significant (paired t-test, $P = 0.3099$) exercise-induced change**
167 **in PP2A protein content was observed in human erythrocytes ($n = 8$). Panels B to D report the mean and**
168 **standard deviation.**

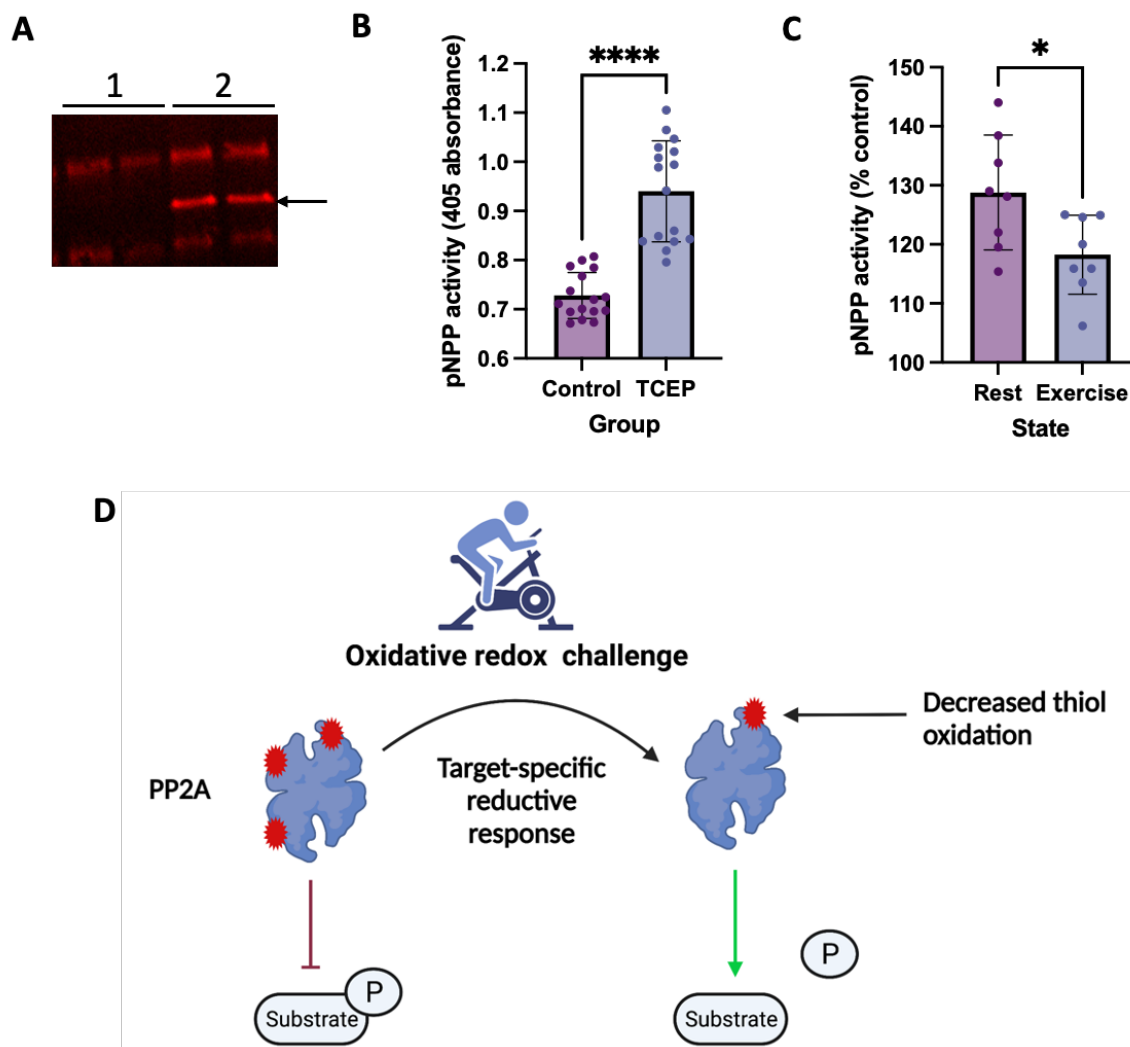
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170 **Chemically reversing protein thiol oxidation increases PP2A activity**

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172 To determine whether the exercise-induced decrease in PP2A-specific reversible thiol
173 oxidation is biologically meaningful, we quantified Tris-(2carboxyethyl)phosphine (TCEP)-
174 inducible para-Nitrophenylphosphate (pNPP)-phosphatase activity. To measure TCEP-
175 inducible pNPP activity, we used immunoprecipitation (IP) to “pull-down” PP2A. PP2A pull-
176 down was confirmed by immunoblot (Figure 3A). To prevent artificial oxidation during the IP,
177 PP2A was pulled-down in the presence of *N*-ethylmaleimide (NEM) [43]. IP eluents were split
178 into two aliquots: (1) solvent control and (2) TCEP. Consistent with the ability of NEM to inhibit
179 PP2A [44] and reversible thiol oxidation, TCEP significantly increased PP2A activity compared
180 to the solvent control regardless of state (Figure 3B). Comparing the thiol-reductant induced
181 increase in resting compared to exercise eluents revealed that TCEP-induced a significantly
182 greater increase in PP2A activity at rest compared to exercise (Figure 3C). Chemically
183 decreasing PP2A-specific reversible thiol oxidation increased pNPP-specific PP2A activity.
184 Data are consistent with a canonically “oxidative” stimulus—exercise—activating a selective
185 target-specific and functionally relevant “reductive” response (Figure 3D).

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189 **Figure 3. Chemically reversing thiol oxidation increases PP2A activity.** A. Immunoblot result showing the
190 presence of a PP2A-specific band at ~37 kDa (marked by an arrow) in the PP2A capture antibody eluent (2),
191 but not the rabbit “sham” isotype control eluent (1). The image shows both technical replicates. B. A
192 significant (paired t-test, $P < 0.0001$) TCEP-inducible increase in PP2A activity compared to the solvent control,
193 as assessed by the monitoring pNPP dephosphorylation at 405 nm, following PP2A IP ($n = 16$). For this
194 experiment, IP eluents ($n = 16$, 8 rest + 8 exercise) were divided into two and treated with either a solvent
195 control ($n = 16$) or TCEP ($n = 16$). Hence the number of data-points are greater than 8 for this panel. C. The
196 TCEP-induced increase in PP2A activity was significantly (paired t-test, $P = 0.0164$) greater in IP eluents from
197 rested ($n = 8$) compared to exercised human erythrocytes ($n = 8$). This experiment refers to the comparison
198 between the rest and exercise +TCEP eluent for each participant. D. Reverse redox regulation model. Exercise
199 a canonical “oxidative” redox challenge induced a PP2A-specific reductive response manifested by decreased
200 reversible thiol oxidation in human erythrocytes. Decreased exercise-induced thiol oxidation increased PP2A
201 activity—depicted by substrate dephosphorylation. Panels B and C report the mean and standard deviation.

202

203 Discussion

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205 Superimposing exercise-induced oxidative stress in erythrocyte cells already burdened
206 by chronic Hb autoxidation might be expected to comprise their ability to defend thiol redox
207 homeostasis as manifested by increased reversible thiol oxidation [40,43,45]. Unexpectedly,
208 brief maximal exercise decreased PP2A-specific reversible thiol oxidation in human
209 erythrocytes, which is associated with increased PP2A catalysed pNPP phosphatase activity.

210 This counterintuitive finding supports a “reversed” redox regulation model wherein PP2A-
211 specific reversible thiol oxidation constrains PP2A activity at rest. Exercise releases a redox
212 break on PP2A activity by decreasing reversible thiol oxidation, which may enable
213 erythrocytes to post-translationally maintain a vital function (e.g., ion homeostasis [42]).
214 Mechanistically, the target-specific “reductive” response is compatible with several
215 possibilities converging on a change in the rate of reversible thiol oxidation formation and/or
216 reduction. Accordingly, we use the word reductive in the sense that the net outcome of
217 several complex processes was a decrease in PP2A-specific reversible thiol oxidation. An
218 interpretationally essential point: The finding does not necessarily mean the enzymes
219 responsible for reversing PP2A oxidation were more active or ROS levels were decreased.
220 Further work is needed to unravel the underlying mechanism. Moreover, studies to decipher
221 the relevant PP2A-substrates and to extend our finding to different exercise types in other
222 cohorts are warranted. A persistent degree of reversible thiol oxidation in the exercised state
223 suggests fine-tuned granular redox control. Data point to a general (i.e., beta subunit
224 independent) and pervasive PP2A regulatory mechanism defined by reversible core catalytic
225 subunit thiol oxidation [46]. Given the potent PP2A inhibitor okadaic acid binds to Cys269
226 [47], it is entirely possible that oxidising Cys269 is an endogenous “natural” control
227 mechanism, and its redox state may govern the efficacy and IC₅₀ of okadaic acid. Targeted
228 redox proteomics is needed to pinpoint the oxidised residues [24,48,49] and future studies
229 should explore the chemotypes involved and their regulation. In particular, ALISA is
230 compatible with new sulfenic acid reactive turn-on fluorescent probes [50]. Our work is
231 consistent with redox regulated PP2A activity [41,51] and the wider value of the erythrocyte
232 for studying protein thiol redox biology [52].

233

234 Translational human redox biology research is beset by overtly flawed and/or severely
235 limited assays [36]. Despite hampering current progress, it has been difficult to persuade
236 many to abandon the offending assays because they are easy to use and implement—
237 convenient commercial kits are available. ALISA offers all of their advantages—easy to
238 implement, interpret, and access—without their troubling and rate-limiting drawbacks. ALISA
239 is valid and target-specific. Selected drawbacks of ALISA include (1) the inability to reveal the
240 residues oxidised, (2) fold-change analysis, and (3) the need for a suitable capture antibody,
241 and in this case, an appropriate well-validated ELISA (see [11] for an involved discussion).
242 Nonetheless, the advantages of ALISA allowed us to discover PP2A as a first-in-class systemic
243 protein thiol redox biomarker. Our work and the advent of the state-specific redox array [53]
244 sets the stage to develop protein thiol redox biomarkers for unravelling the biological roles
245 ROS play in humans. At present, PP2A should be treated as a candidate biomarker only
246 because further validation work (e.g., across different cohorts) is clearly required.

247

248 In summary, we have discovered a new readily accessible and easily measurable
249 systemic redox state change in PP2A—a key phosphatase—that is associated with an
250 unexpected redox regulation model wherein a physiological “oxidative” stimulus that is well
251 known to induce oxidative stress triggers a biologically meaningful target-specific “reductive”
252 response as defined by a decrease in target-specific reversible thiol oxidation.

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254 **Methods**

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256 **Materials**

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Supplementary Table 1 details the materials and reagents used.

Participants

Following ethical approval (#OL-ETHSHE-1436) and after obtaining written informed consent, 8 adult male participants provided a venous blood sample before and immediately after a maximal exercise test.

Exercise

Participants completed an incremental exercise test to exhaustion on a bicycle ergometer. Their heart rate and oxygen uptake were recorded throughout (see extended methods).

Thiol labelling

NEM (100 mM) supplemented erythrocyte aliquots were lysed in an equal volume of IP lysis buffer (25 mM Tris-HCl, 150 mM NaCl, 1 mM EDTA, 1% NP-40, 5% glycerol, pH 7.1) supplemented with a protease inhibitor tablet. Samples were centrifuged (14,000 *g* for 2 min), soluble supernatants were passed through a 6-kDa spin column to remove excess NEM. Flow throughs were treated with 5 mM neutral-TCEP for 30 min on ice. After removing excess TCEP with a spin column, samples were treated with 1 mM F-MAL (i.e., fluorescein-5-maleimide) for 30 min on ice. Excess (i.e., unreacted) F-MAL was removed with a spin column.

SDS-PAGE

Samples (50 μ M Hb) were supplemented with 2X loading buffer and resolved by SDS-PAGE on a precast 4-15% gradient gel. F-MAL signals were measured on a gel scanner.

Global-ALISA

Samples were dispensed in triplicate into microplate wells containing distilled H₂O. F-MAL was measured for 100 ms at 494 and 519 nm in a plate reader. Spectrometric Hb absorbance was measured for 100 ms at 577 nm [54]. After subtracting the background, global reversible thiol oxidation was calculated as: F-MAL/Hb.

PP2A-ALISA

Black MaxiSorp microplates were incubated with 2 μ g/ml capture PP2A antibody overnight at 4°C in binding buffer (35 mM NaHCO₃, 15 mM Na₂CO₃, pH 9.6) on a plate shaker at 350 rpm. Wells were blocked (50% PBST, 50% Superblock) for 2 h at RT at 350 rpm and washed (3 x 2 min PBST washes at 400 rpm). The recombinant PP2A standard and samples (diluted to 100 μ M Hb in PBS) were added in duplicate and incubated for 2 h at RT at 350 rpm. Excess sample was removed, wells were washed (3 x 2 min PBST washes at 400 rpm), and 0.5 μ g/ml biotin-conjugated PP2A detector antibody was added for 1 h at RT at 350 rpm. After washing, 0.05 μ g/ml of HRP-conjugated streptavidin was added for 1 h at RT at 350 rpm. After

304 a final wash, wells were incubated with QuantaBlu™ for 10 min at RT at 400 rpm. The
305 QuantaBlu™ signal was measured at 325 and 425 nm for 100 ms. Denaturing buffer (4% SDS)
306 was added to stop the HRP reaction and unmask the F-MAL1 signal. After subtracting the
307 background, PP2A-specific redox state was calculated as: F-MAL/total PP2A. All assay steps
308 were performed under light protected conditions.

309

310 **PP2A activity assay**

311

312 Lysates were incubated with anti-PP2A functionalised magnetic protein A beads
313 overnight at 4°C. After washing (3 x 2 min distilled H₂O), PP2A was eluted in gentle IP elution
314 buffer. To normalise protein content, eluents were divided into two aliquots and their pNPP
315 activity was assessed with (aliquot 1) and without (aliquot 2) 1 mM neutral-TCEP at 405 nm
316 in a microplate.

317

318 **PP2A-Immunoblot**

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320 IP eluent PP2A content was checked by immunoblotting as previously
321 described[11,26]. To avoid interference from coeluting capture antibody light and heavy
322 chains, membranes were probed with a fluorescent-conjugated PP2A antibody.

323

324 **Statistics**

325

326 After assessing normality, data were assessed using paired t-tests with alpha <0.05
327 on GraphPad Prism version 9.

328

329 **Author contributions:** Conceptualisation: D.M., D.C., G.W. and J.N.C. Methodology and Resources: J.N.C. and
330 A.T.; Investigation: D.C. and D.M.; Formal data analysis: D.M., A.T., and D.C., Writing original draft: J.N.C., Writing
331 review and editing: All authors, Funding acquisition: J.N.C., I.L.M. and D.C.

332

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336 (www.Biorender.com) and exported with the appropriate publication license.

337

338 **Conflict of interest:**

339

340 There are no conflicts of interest.

341

342 **References**

343

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