

different extent (average deviations: -3% , -18% , and -33%).

In conclusion, the presented study showed that our measurement procedure, based on ED and trueness-based ID-LC/tandem MS measurement of dialysate, qualifies as a method for standardization of FT₄ measurements. At the same time, our results demonstrate that samples processed according to the C37-A protocol are suitable for use in standardization. These results provide a basis for worldwide standardization of FT₄ measurements under the auspices of the International Federation of Clinical Chemistry and Laboratory Medicine (23). An inaugural meeting of the group is foreseen at the 2006 AACC Annual Meeting (24).

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References

- Ladenson PW, Singer PA, Ain KB, Bagchi N, Bigos ST, Levy EG. American Thyroid Association Guidelines for detection of thyroid dysfunction. *Arch Intern Med* 2000;160:1573–5.
- Dayan CM. Interpretation of thyroid function tests [Review]. *Lancet* 2001;357:619–24.
- Bartalena L, Robbins J. Variations in thyroid hormone transport proteins and their clinical implications [Review]. *Thyroid* 1992;2:237–45.
- Helenius T, Liewendahl K. Improved dialysis method for free thyroxine in serum compared with five commercial radioimmunoassays in nonthyroidal illness and subjects with abnormal concentrations of thyroxin-binding globulin. *Clin Chem* 1983;29:816–22.
- Hay ID, Bayer MF, Kaplan MM, Klee GG, Larsen PR, Spencer CA. American Thyroid Association assessment of current free thyroid hormone and thyrotropin measurements and guidelines for future clinical assays. The Committee on Nomenclature of the American Thyroid Association. *Clin Chem* 1991;37:2002–8.
- Nelson JC, Weiss RM, Wilcox RB. Underestimates of serum free thyroxine (T₄) concentrations by free T₄ immunoassays. *J Clin Endocrinol Metab* 1994;79:76–9.
- Nelson JC, Wilcox RB. Analytical performance of free and total thyroxine assays [Review]. *Clin Chem* 1996;42:146–54.
- Wang R, Nelson JC, Weiss RM, Wilcox RB. Accuracy of free thyroxine measurements across natural ranges of thyroxine binding to serum proteins. *Thyroid* 2000;10:31–9.
- d'Herbomez M, Forzy G, Gasser F, Massart C, Beaudonnet A, Sapin R. Clinical evaluation of nine free thyroxine assays: persistent problems in particular populations. *Clin Chem Lab Med* 2003;41:942–7.
- National Academy of Clinical Biochemistry (NACB). Laboratory Medicine Practice Guidelines: Laboratory Support for the Diagnosis and Monitoring of Thyroid Disease. NACB, Demers LM, Spencer CA (2002).
- Clinical Laboratory and Standards Institute (CLSI). Measurement of Free Thyroid Hormones; Approved Guideline. CLSI document C45-A [ISBN 1-56238-548-8]. CLSI, Wayne, PA, 2004.
- Ekins R. The science of free hormone measurement. *Proc UK NEQAS Meeting* 1998;3:35–59.
- Holm SS, Hansen SH, Faber J, Staun-Olsen P. Reference methods for the measurement of free thyroid hormones in blood: evaluation of potential reference methods for free thyroxine [Review]. *Clin Biochem* 2004;37:85–93.
- Soldin SJ, Soukhova N, Janic N, Jonklaas J, Soldin OP. The measurement of free thyroxine by isotope dilution tandem mass spectrometry. *Clin Chim Acta* 2005;358:113–8.
- Tikanoja SH, Liewendahl BK. New ultrafiltration method for free thyroxin compared with equilibrium dialysis in patients with thyroid dysfunction and nonthyroidal illness. *Clin Chem* 1990;36:800–4.
- Nelson JC, Tomei RT. Direct determination of free thyroxine in undiluted serum by equilibrium dialysis/radioimmunoassay. *Clin Chem* 1988;34:1737–44.
- De Brabandere VI, Hou P, Stöckl D, Thienpont LM, De Leenheer AP. Isotope dilution-liquid chromatography/electrospray ionization-tandem mass spectrometry for the determination of serum thyroxine as a potential reference method. *Rapid Commun Mass Spectrom* 1998;12:1099–103.
- Van Uytvanghe K, Stöckl D, Thienpont LM. Development of a simplified sample pretreatment procedure as part of an isotope dilution-liquid chromatography-tandem mass spectrometry candidate reference measurement procedure for serum total thyroxine. *Rapid Commun Mass Spectrom* 2004;18:1539–40.
- Thienpont LM, Van Uytvanghe K, Marriott J, Stokes P, Siekmann L, Kessler A, et al. Metrologic traceability of total thyroxine measurements in human serum: efforts to establish a network of reference measurement laboratories. *Clin Chem* 2005;51:161–8.
- Clinical Laboratory and Standards Institute (CLSI). Preparation and Validation of Commutable Frozen Human Serum Pools as Secondary Reference Materials for Cholesterol Measurement Procedures; Approved Guideline. CLSI document C37-A [ISBN 1-56238-392-2]. CLSI, Wayne, PA, 2004.
- Clinical Laboratory and Standards Institute (CLSI). Evaluation of Precision Performance of Quantitative Measurement Methods; Approved Guideline. CLSI document EP5-A2 (ISBN 1-56238-542-9). CLSI, Wayne, PA, 2004.
- Holm SS, Andreasen L, Hansen SH, Faber J, Staun-Olsen P. Influence of adsorption and deproteination on potential free thyroxine reference methods. *Clin Chem* 2002;48:108–14.
- IFCC. 8.3. SD Working Groups. 8.3.33 Standardization of thyroid function tests. <http://www.ifcc.org/divisions/CPD/handbook/Handbook> (accessed April 2006).
- American Association of Clinical Chemistry. 2006 Annual Meeting http://www.aacc.org/2006AM/full_brochure.pdf. p. 22 (accessed April 2006).

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Cell-Free Plasma DNA as a Novel Marker of Aseptic Inflammation Severity Related to Exercise Overtraining

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Background: Circulating free plasma DNA is implicated in conditions associated with tissue injury, including exercise-induced inflammation, and thus is a potential marker for athletic overtraining.

Methods: We measured free plasma DNA along with C-reactive protein (CRP), creatine kinase (CK), and uric acid (UA) in 17 recreationally trained men participating

in a 12-week resistance training regimen (8 resistance multi-joint exercises selected to stress the entire musculature: bench press, squat, leg press, snatch, hang clean, dead lifts, barbell arm curls, and rowing), consisting of 4 training periods (t1, t2, t3, and t4).

Results: Plasma DNA concentrations increased markedly after t1, t2, and t3 and returned to baseline after t4. There were substantial differences between t2 and t1 and between t3 and t2 plasma DNA concentrations. CRP increased by 300% after t2 and by 400% after t3 (there was no difference between t2 and t3 CRP values) compared with baseline (t0). CK increased only after t3. UA increased after t2 and t3, with a greater increase after t3.

Conclusions: This study demonstrates that, after chronic excessive resistance exercise, plasma DNA concentrations increase in proportion to training load, suggesting that plasma DNA may be a sensitive marker for overtraining-induced inflammation.

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Strenuous exercise may lead to transient muscle fiber damage, with indications such as soreness, edema, performance deterioration, and protein release into plasma (1–6). Muscle injury has been attributed to local ATP depletion, calcium homeostasis disturbance, and oxidative stress induced by generation of free radicals (7, 8). These adverse effects are seen in athletes affected by overtraining, which has been defined as an accumulation of exercise training and nontraining stress, leading to long-term decrement in athletic performance (despite continued training) with or without related physiologic and psychological signs. Overtraining can lead to an acute breakdown with subsequent repair of skeletal muscle and is associated with increased susceptibility to infections attributable to changes in the functional status of immune cells (9).

Circulating plasma DNA is altered both quantitatively and qualitatively under a variety of conditions, including tissue injury, pregnancy, cancer, and trauma (10–13). Plasma DNA concentration has been correlated with injury severity and thus is a potential marker for risk stratification (11). We investigated the use of the plasma DNA response as an inflammation marker during a chronic resistance exercise (RE) protocol of progressively increased training volume. We then compared this marker with conventional indices such as C-reactive protein (CRP), creatine kinase activity (CK), and uric acid (UA), which are associated with exercise-induced muscle damage and acute-phase response.

Participants were recruited from a volunteer database, by word of mouth and by fliers posted within the University and the local community. A written informed consent was signed by all participants. Procedures were in accordance with the Helsinki Declaration for the Ethical Treatment of Human Subjects. Ethics approval was given by the institutional review board.

Our study participants were healthy volunteers: age, 21.56 (2.6) years; body weight, 77 (7.1) kg; body height,

1.77 (0.11) m; and body fat, 12.2 (2.1)%. They were 17 recreationally trained men who participated in a 12-week resistance training regimen (8 resistance multi-joint exercises were selected to stress the entire musculature: bench press, squat, leg press, snatch, hang clean, dead lifts, barbell arm curls, and rowing) consisting of four 3-week training periods (t1, t2, t3, and t4). The first and fourth training periods (t1 and t4) included low-volume training: 2 training days per week, 2 sets per exercise, 10–12 reps per set at 70% of their maximal strength (1 repetition maximum, 1RM); t2 included high-volume training, 4 training days per week, 4 sets per exercise, 6–10 reps per set at 75%–85% of 1RM; and t3 included very-high-volume training, 6 training days per week, 6 sets per exercise, and 1–6 reps per set at 85%–100% of 1RM. Consecutive training periods were separated by a 5-day rest period. The mean training volume (tonnage) lifted per training period is presented in Table 1.

Blood samples were drawn at baseline (participants abstained from resistance training for at least 8 weeks before the study) and at 96 h after the last training session of each period.

We isolated plasma DNA from 400 μ L plasma after 10 min centrifugation at 800g, followed by 10 min centrifugation at 16 000g. DNA was extracted with the QIAamp DNA Blood MiniKit (Qiagen) and eluted into 50 μ L of RNase/DNase-free H₂O. We measured free-DNA by quantitative real-time PCR with the LightCycler (Roche Diagnostics) with the following primers: forward, 5'-AGG TGA ACG TGG ATG AAG TT-3' and reverse, 5'-AGG GTA GAC CAC CAG CAG CC-3', for the amplification of a 189-bp fragment of the β -globin gene. PCR amplification reactions were performed with the LightCycler Fast Start DNA Master SYBR Green I (Roche Diagnostics). The PCR conditions included a first denaturation step of 95 °C (10 min), followed by 40 cycles of 95 °C (10 s), 95 °C (5 s), and 95 °C (20 s) with a temperature ramp of 95 °C/s (14). All samples were analyzed in duplicate. Each reaction included a calibration curve of 6.6 ng to 6.6 pg genomic DNA made from 4 serial dilutions of purchased human genomic DNA (0.2 g/L in 10 mmol/L Tris-HCl, 1 mmol/L EDTA, pH 8.0), (Roche Diagnostics GmbH, Cat. No. 11691112001) plus 2 known dilutions (50 and 500 pg DNA) used as controls in all experiments. DNA values were expressed as genome equivalents per milliliter (GenEq/mL), corresponding to 6.6 pg DNA.

We measured CK, UA, and CRP with the COBAS INTEGRA 800 Clinical Chemistry System (Roche Diagnostics).

We analyzed data with SPSS-PC and calculated means (SE). We evaluated time differences with ANOVA repeated measures and used a Bonferroni test for post hoc comparisons. Data normality was tested with the 1-sample Kolmogorov-Smirnoff test; therefore a nonparametric test was not necessary. Statistical significance was set at $P < 0.05$.

Plasma DNA increased significantly after t1, t2, and t3, and returned to baseline concentrations after t4 (Table 1, Fig. 1). Plasma DNA concentrations differed significantly

for t2 vs t1 and for t3 vs t2. CRP increased compared with baseline by 300% after t2, and by 400% after t3 (there was no difference between t2 and t3 values). CK increased only after t3 (Table 1). UA (Table 1) increased after t2 and t3, with a greater increase after t3. Performance increased after t2 and declined thereafter (Table 1).

An overtraining response to the exercise protocol was demonstrated by a decreased performance level, despite increased training and subsequent lack of recovery within 3 weeks, and despite decreased training volume in t4 (9). Tissue damage and inflammation were confirmed by increased plasma DNA and CRP concentrations. After each training period, Pearson correlation coefficients were calculated between plasma DNA, CRP, CK, UA, and training load. The results showed substantial correlations between plasma DNA and CK at t2 ($r = 0.793$, $P < 0.01$) and at t3 ($r = 0.744$, $P < 0.01$). We observed no statistically significant correlations between plasma DNA, CRP, CK, and UA. Plasma DNA concentrations were significantly related to training load (average tons lifted in all exercises per week in each training period) only in t2 ($r = 0.755$, $P < 0.01$) and t3 ($r = 0.786$, $P < 0.01$).

Circulating DNA has been shown to increase 9- to 17.5-fold after long-distance running (21–246 km) (7, 14), but changes in concentration in response to chronic overload training have not been investigated. We measured plasma DNA 96 h after the last training session to look for changes reflecting chronic exercise-induced inflammation. Indeed, plasma DNA remained increased for 96 h postexercise after the t1, t2, and t3 periods. Increased plasma DNA concentrations have persisted up to 3 h after major injury in trauma patients (15), 2 or more days (as in this study) after exhaustive exercise (14), and several days in patients with multiple organ dysfunction syndrome (10).

Increased plasma DNA concentrations correlated with injury severity in trauma and stroke patients (11, 16)

indicate that circulating DNA concentration is a potential risk stratification marker. We observed a significant correlation of plasma DNA with CK (in t2 and t3) but not with CRP. Demonstration of a similar response of plasma DNA to progressive training that does not lead to overtraining would indicate that plasma DNA can be used as an overtraining marker.

Muscular overtraining impairs athletic performance, increases susceptibility to infections, and induces muscle tissue injury as well as psychological, immunological, and biochemical dysfunctions (9). Although there is a strong demand for early diagnosis of overtraining, specific sensitive markers have not been identified. CRP is a sensitive marker for inflammation regardless of etiology (17) and increases in response to intense exercise (14, 18, 19). Our results provide evidence of a delayed (4 days) CRP response to RE, which was also seen (3–6 days) after intense running (20, 21). In our study, basal CRP concentrations were within the reference interval (0–10 mg/L), but after t3, mean CRP (4.8 mg/L) approached the reference interval for acute infections (5 mg/L for adults) (22). Unlike plasma DNA, however, CRP concentrations did not reflect the difference in magnitude of exercise overload between t3 and t4. Similarly, CK increased only after t3. The response of plasma DNA, but not of CRP or CK, seemed to be proportional to the exercise training overload.

The mechanisms underlying the presence of circulating DNA are generally unknown. Apoptosis, necrosis, active release, and impaired clearance have been implicated (23–25), but information regarding their role during exercise is scarce. RE has been reported to generate reactive oxygen species (ROS) and oxidative stress (26). ROS have been associated with postexercise inflammation and muscle damage propagation (27). An inflammatory response during the repair of overtraining-induced muscle damage (28) promotes neutrophil and macrophage infiltration of

Table 1. Training volume and performance as well as CK, CRP, and UA concentration changes at rest (Baseline), and after low- (t1), high- (t2), very high- (t3), and low-volume (t4) resistance training

	Baseline	t1	t2	t3	t4
Mean training volume, tonnage/wk (SE) ^a	N/A	2.2 (0.2)	7.8 (0.6) ^d	14.8 (1.4) ^{d,c}	1.7 (0.3) ^f
Performance (maximal strength), kg (SE) ^b	96.6 (17.1)	99.2 (11.7)	109.6 (14.8) ^{c,d}	104.5 (14.9) ^c	107.2 (12.5) ^{c,d}
Cell-free plasma DNA, 8GenEq/mL (SE)	31.4 (13.8)	143.5 (22.9) ^c	289.9 (41.1) ^{c,d,e}	605.7 (116.4) ^{c,f}	74.8 (29.6) ^{c,d,f}
	(14.0–65.9)	(76.6–180.6)	(196.9–383.0)	(251.8–959.5)	(27.7–141.8)
CK activity, units/L (SE)	102.6 (19.1)	134.5 (20.1)	296.6 (49.6)	368.1 (44.0) ^{c,d}	161.7 (23.8) ^f
	(59.4–149.8)	(86.7–82.2)	(185.5–407.6)	(264.4–513.9)	(105.7–217.7)
CRP, mg/L (SE)	0.84 (0.17)	0.86 (0.19)	3.364 (1.07) ^{c,d}	4.8 (1.04) ^{c,d}	1.42 (0.30) ^f
	(0.45–1.24)	(0.42–1.31)	(1.00–5.77)	(1.89–6.64)	(0.73–2.11)
UA, mg/L (SE)	39.1 (4.0)	42.4 (4.0)	51.2 (3.7) ^c	66.6 (5.8) ^{c,d}	43.8 (4.3) ^f
	(30–48.2)	(33.3–51.5)	(42.8–59.6)	(53.4–79.5)	(34.1–53.5)

^a Mean number of tons lifted per week for all resistance exercises employed in the protocol. NA, not applicable.

^b Mean maximal strength in 1 representative exercise (squat).

^c Significant difference with baseline (T0).

^d Significant difference with t1.

^e Difference between t3 and t4.

^f Significant difference between t4 and t3.

^g Significant difference with t4. Minimum and maximum values are shown in parentheses.

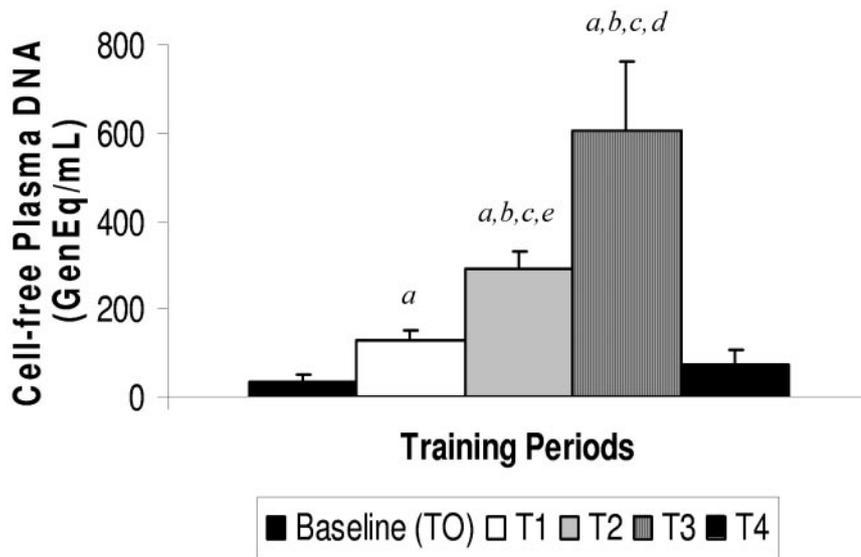


Fig. 1. Cell-free plasma DNA concentration in 17 study participants measured sequentially at rest (t0) and after low- (t1), high- (t2), very high- (t3), and low-volume (t4) resistance training.

- ^a Significant difference with baseline (t0)
- ^b Significant difference with t1.
- ^c Significant difference between t3 and t2.
- ^d Significant difference between t4 and t3.
- ^e Significant difference with t4.

muscles (2), most likely initiated by ROS (29). Thus, peroxidative muscle damage is exacerbated during post-exercise inflammation. Neutrophils and macrophages generate superoxide (30), which may be converted to hydrogen peroxide, which then reacts with superoxide in the presence of a transition metal to form hydroxyl radical (31). After exercise, neutrophil counts in muscle are increased for 5 days, and peak macrophage counts occur at 2 to 7 days (1–2). UA, which we used as an oxidative stress marker because of its free-radical scavenging properties, increased after t2 and t3, indicating that chronic RE induced oxidative stress. ROS generation leads to DNA damage in the form of DNA cross-links, oxidized nucleosides, and strand breaks (32). More work is needed to elucidate the origin of plasma DNA during exercise.

This study demonstrates that plasma DNA concentrations increase in proportion to training load after chronic excessive RE, suggesting that plasma DNA may be a sensitive marker for monitoring and quantification of overtraining in athletes. In particular, our finding that plasma DNA was substantially correlated with mean training load during periods of high-volume and very-high-volume training (t2 and t3) supports this theory. Plasma DNA measurement is noninvasive and requires limited time and personnel. Further investigation of the origin of plasma DNA in the overtrained state is warranted.

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References

1. Fielding R, Manfredi T, Ding W, Fiatarone M, Evans W, Cannon J. Acute phase response to exercise. III: neutrophil and IL-1 β accumulation in skeletal muscle. *Am J Physiol* 1993;265:R166–72.
2. Evans W, Cannon J. Metabolic effects of exercise-induced muscle damage. *Exerc Sports Sci Rev* 1991;18:99–125.
3. Smith LL. Acute inflammation: the underlying mechanism in delayed onset muscle soreness? *Med Sci Sports Exerc* 1991;23:542–51.
4. Belcastro A, Arthur G, Albisser T, Raj D. Heart, liver, and skeletal muscle myeloperoxidase activity during exercise. *J Appl Physiol* 1996;80:1331–5.
5. Cannon J, Orencole S, Fielding R, Meydani M, Meydani S, Fiatarone M, et al. Acute phase response in exercise: interaction of age and vitamin E on neutrophils and muscle enzyme release. *Am J Physiol* 1990;259:R1214–9.
6. Malm C, Sjodin LB, Sjoberg B, Lenkei R, Renstrom P, Lundberg IE, et al. Leukocytes, cytokines, growth factors, and hormones in human skeletal muscle and blood after uphill or downhill running. *J Physiol* 2004;556:983–1000.
7. Atamaniuk J, Vidotto C, Tschan H, Bachl N, Stuhlmeier KM, Miller MM. Increased concentrations of cell-free plasma DNA after exhaustive exercise. *Clin Chem* 2004;50:1668–70.
8. Mars M, Govender S, Weston A, Naicker V, Chuturgoon A. High intensity exercise: a cause of lymphocyte apoptosis? *Biochem Biophys Res Commun* 1998;249:366–70.
9. Kreider R, Fry AC, O'Toole M. Over-training in sport: terms, definitions, and prevalence. In: Kreider R, Fry AC, O'Toole M, eds. *Over-training in Sport*. Champaign, IL: Human Kinetics, 1998:47–66.
10. Lam NY, Rainer TH, Chan LY, Joynt GM, Lo YM. Time course of early and late changes in plasma DNA in trauma patients. *Clin Chem* 2003;49:1286–91.
11. Rainer TH, Wong LKG, Lam W, Yuen E, Lam NYL, Metrewell C, et al. Prognostic use of circulating plasma nucleic acid concentrations in patients with acute stroke. *Clin Chem* 2003;49:562–9.
12. Lo YMD, Tein MSC, Pang CCP, Yeung CK, Tong KL, Hjeim NM. Presence of donor-specific DNA in plasma of kidney and liver-transplant recipients. *Lancet* 1998;351:1329–30.
13. Chen XQ, Stroun M, Magnenat JL, Nicod LP, Kurt AM, Lyautey J, et al. Microsatellite alterations in plasma DNA of small cell lung cancer patients. *Nat Med* 1996;2:1033–5.
14. Margeli A, Skenderi K, Tsironi M, Hantzi E, Matalas AL, Vrettou C, et al. Dramatic elevations of interleukin-6 and acute phase reactants in athletes participating in the ultradistance foot race Spartathlon: severe systemic inflammation and lipid and lipoprotein changes in protracted exercise. *J Clin Endocrinol Metab* 2005;90:3914–8.
15. Rainer TH, Lo YM, Chan LY, Lit LC, Cooks RA. Derivation of a prediction rule for posttraumatic organ failure using plasma DNA and other variables. *Ann N Y Acad Sci* 2001;945:211–20.
16. Lo YM, Rainer TH, Chan LY, Hjeim NM, Cooks RA. Plasma DNA as a prognostic marker in trauma patients. *Clin Chem* 2000;46:319–23.
17. Kushner I, Broder ML, Karp D. Serum C-reactive protein: kinetics after acute myocardial infarction. *J Clin Invest* 1978;61:235–42.
18. Weight LM, Alexander D, Jacobs P. Strenuous exercise: analogous to the acute phase response? *Clin Sci (Lond)* 1991;81:677–83.
19. Strachan AF, Noakes TD, Kotzenberg G, Nel AE, de Beer FC. C-reactive protein concentrations during long-distance running. *Br Med J* 1984;289:1249–51.
20. Fallon KE. The acute phase response and exercise: the ultramarathon as prototype exercise. *Clin J Sports Med* 2001;11:38–43.
21. Dufaux B, Order U. Complement activation after prolonged exercise. *Clin Chim Acta* 1989;179:45–50.
22. Thomas L. *Labor und Diagnose (Laboratory and Diagnosis)*. Marburg: Medizinische Verlagsgesellschaft, 1995:781.

23. Jahr S, Hentze H, Englisch S, Hardt D, Fackelmayer FO, Hesch RD, et al. DNA fragments in the blood plasma of cancer patients: quantitations and evidence for their origin from apoptotic and necrotic cells. *Cancer Res* 2001;61:1659–65.
 24. Stroun M, Lyautey J, Lederrey C, Olson-Sand A, Anker P. About the possible origin and mechanism of circulating DNA apoptosis and active DNA release. *Clin Chim Acta* 2001;313:139–42.
 25. Atamaniuk J, Ruzicka K, Stuhlmeier KM, Karimi A, Eigner M, Mueller MM. Cell-free plasma DNA: a marker for apoptosis during hemodialysis. *Clin Chem* 2006;52:523–6.
 26. Liu JF, Chang WY, Chan KH, Tsai WY, Lin CL, Hsu MC. Blood lipid peroxides and muscle damage increased following intensive resistance training of female weightlifters. *Ann N Y Acad Sci* 2005;1042:255–61.
 27. McCord J. Superoxide radical: controversies, contradictions, and paradoxes. *Proc Soc Exp Biol Med* 1995;209:112–7.
 28. Tiidus PM. Radical species in inflammation and over-training. *Can J Physiol Pharmacol* 1998;76:533–8.
 29. Atalay M, Marnila P, Lilius EM, Hanninen O, Sen CK. Glutathione-dependent modulation of exhausting exercise-induced changes in neutrophil function in rats. *Eur J Appl Physiol* 1996;74:342–7.
 30. Pyne D. Regulation of neutrophil function during exercise. *Sports Med* 1994;17:245–58.
 31. Jaeschke H. Mechanisms of oxidant stress-induced acute tissue injury. *Proc Soc Exp Biol Med* 1995;209:104–11.
 32. Ogonovszky H, Sasvari M, Dosek A, Berkes I, Kaneko T, Tahara S, et al. The effects of moderate, strenuous, and over-training on oxidative stress markers and DNA repair in rat liver. *Can J Appl Physiol* 2005;30:186–95.
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