

Health Benefits of Exercise

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Overwhelming evidence exists that lifelong exercise is associated with a longer health span, delaying the onset of 40 chronic conditions/diseases. What is beginning to be learned is the molecular mechanisms by which exercise sustains and improves quality of life. The current review begins with two short considerations. The first short presentation concerns the effects of endurance exercise training on cardiovascular fitness, and how it relates to improved health outcomes. The second short section contemplates emerging molecular connections from endurance training to mental health. Finally, approximately half of the remaining review concentrates on the relationships between type 2 diabetes, mitochondria, and endurance training. It is now clear that physical training is complex biology, invoking polygenic interactions within cells, tissues/organs, systems, with remarkable cross talk occurring among the former list.

The aim of this introduction is briefly to document facts that health benefits of physical activity predate its readers. In the 5th century BC, the ancient physician Hippocrates stated: “All parts of the body, if used in moderation and exercised in labors to which each is accustomed, become thereby healthy and well developed and age slowly; but if they are unused and left idle, they become liable to disease, defective in growth and age quickly.” However, by the 21st century, the belief in the value of exercise for health has faded so considerably, the lack of exercise now presents a major public health problem (Fig. 1) (Booth et al. 2012). Similarly, the lack of exercise was classified as an actual

cause of chronic diseases and death (Mokdad et al. 2004).

Published in 1953, Jeremy N. Morris and colleagues conducted the first rigorous epidemiological study investigating physical activity and chronic disease risk, in which coronary heart disease (CHD) rates were increased in physically inactive bus drivers versus active conductors (Morris et al. 1953). Since this pioneering report, a plethora of evidence shows that physical inactivity is associated with the development of 40 chronic diseases (Table 1), including major noncommunicable diseases such as type 2 diabetes (T2D) and CHD, and as premature mortality (Booth et al. 2012).

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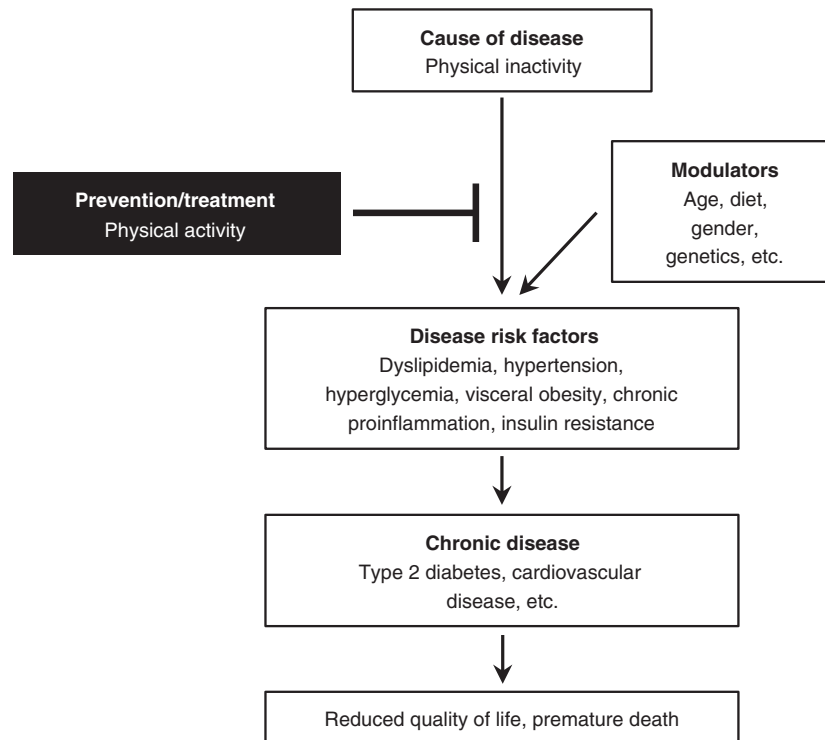


Figure 1. Simplistic overview of how physical activity can prevent the development of type 2 diabetes and one of its complications, cardiovascular disease. Physical inactivity is an actual cause of type 2 diabetes, cardiovascular disease, and tens of other chronic conditions (Table 1) via interaction with other factors (e.g., age, diet, gender, and genetics) to increase disease risk factors. This leads to chronic disease, reduced quality of life, and premature death. However, physical activity can prevent and, in some cases, treat disease progression associated with physical inactivity and other genetic and environmental factors.

In this review, we highlight the far-reaching health benefits of physical activity. However, note that the studies cited here represent only a fraction of the >100,000 studies showing positive associations between the terms “exercise” and “health.” In addition, we discuss how exercise promotes complex integrative responses that lead to multisystem responses to exercise, an underappreciated area of medical research. Finally, we consider how strategies that “mimic” parts of exercise training compare with physical exercise for their potential to combat metabolic disease.

EXERCISE IMPROVES CARDIORESPIRATORY FITNESS

There is arguably no measure more important for health than cardiorespiratory fitness (CRF)

(commonly measured by maximal oxygen uptake, VO_{2max}) (Blair et al. 1989). For example, Myers et al. (2002) showed that each 1 metabolic equivalent (1 MET) increase in exercise-test performance conferred a 12% improvement in survival, stating that “ VO_{2max} is a more powerful predictor of mortality among men than other established risk factors for cardiovascular disease (CVD).” Low CRF is also well established as an independent risk factor of T2D (Booth et al. 2002) and CVD morbidity and mortality (Kodama et al. 2009; Gupta et al. 2011). Similarly, Kokkinos et al. (2010) reported that men who transitioned from having low to high CRF decreased their mortality risk by ~50% over an 8-yr period, whereas men who transitioned from having high to low CRF increased their mortality risk by ~50%.



Table 1. Worsening of 40 conditions caused by the lack of physical activity with growth, maturation, and aging throughout life span

1. Accelerated biological aging/premature death
2. Aerobic (cardiorespiratory) fitness (VO_{2max})
3. Arterial dyslipidemia
4. Balance
5. Bone fracture/falls
6. Breast cancer
7. Cognitive dysfunction
8. Colon cancer
9. Congestive heart failure
10. Constipation
11. Coronary (ischemic) heart disease
12. Deep vein thrombosis
13. Depression and anxiety
14. Diverticulitis
15. Endometrial cancer
16. Endothelial dysfunction
17. Erectile dysfunction
18. Gallbladder diseases
19. Gestational diabetes
20. Hemostasis
21. Hypertension
22. Immunity
23. Insulin resistance
24. Large arteries lose more compliance with aging
25. Metabolic syndrome
26. Nonalcoholic fatty liver disease
27. Obesity
28. Osteoarthritis
29. Osteoporosis
30. Ovarian cancer
31. Pain
32. Peripheral artery disease
33. Preeclampsia
34. Polycystic ovary syndrome
35. Prediabetes
36. Rheumatoid arthritis
37. Sarcopenia
38. Stroke
39. Tendons being less stiff
40. Type 2 diabetes

The breadth of the list implies that a single molecular target will not substitute for appropriate daily physical activity to prevent the loss of all listed items.

Importantly then, from the above paragraph, physical activity and inactivity are major environmental modulators of CRF; increasing and decreasing it, respectively, often through independent pathways. Findings from rats se-

lectively bred for high or low intrinsic aerobic capacity show that rats bred for high capacity, which are also more physically active, have 28%–42% increases in life span compared to low-capacity rats (Koch et al. 2011). Endurance exercise is well recognized to improve CRF and cardiometabolic risk factors. Exercise improves numerous factors speculated to limit VO_{2max} including, but not restricted to, the capacity to transport oxygen (e.g., cardiac output), oxygen diffusion to working muscles (e.g., capillary density, membrane permeability, muscle myoglobin content), and adenosine triphosphate (ATP) generation (e.g., mitochondrial density, protein concentrations).

Data from the HERITAGE Family Study has provided some of the first knowledge of genes associated with VO_{2max} plasticity because of endurance-exercise training. Following 6 wk of cycling training at 70% of pretraining VO_{2max} , Timmons et al. (2010) performed messenger RNA (mRNA) expression microarray profiling to identify molecules potentially predicting VO_{2max} training responses, and then assessed these molecular predictors to determine whether DNA variants in these genes correlated with VO_{2max} training responses. This approach identified 29 mRNAs in skeletal muscle and 11 single-nucleotide polymorphisms (SNPs) that predicted ~50% and ~23%, respectively, of the variability in VO_{2max} plasticity following aerobic training (Timmons et al. 2010). Intriguingly, pretraining levels of these mRNAs were greater in subjects that achieved greater increases in VO_{2max} following aerobic training, and of the 29 mRNAs, >90% were unchanged with aerobic training, suggesting that alternative exercise intervention paradigms or pharmacological strategies may be needed to improve VO_{2max} in individuals with a low responder profile for the identified predictor genes (Timmons et al. 2010). Keller et al. (2011) found that, in response to endurance training, improvements in VO_{2max} were associated with effectively up-regulating proangiogenic gene networks and miRNAs influencing the transcription factor-directed networks for runt-related transcription factor 1 (RUNX1), paired box gene 3 (PAC3), and sex-determining region Y box 9 (SOX9).

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Collectively, these results led the investigators to speculate that improvements in skeletal muscle oxygen sensing and angiogenesis are primary determinates in training responses in VO_{2max} (Keller et al. 2011).

Clinically important concepts have emerged from the pioneering HERITAGE Family Study. One new clinical concept is that a threshold dose–response relationship influences the percentage of subjects responding with an increase in VO_{2max} to endurance training volumes (with volume being defined here as the product of intensity \times duration), as previously published (Slentz et al. 2005, 2007). Ross et al. (2015) later extended the aforementioned Slentz et al. studies. After a 24-wk-long endurance training study (Ross et al. 2015), percentages of women and men identified as nonresponders to the training (i.e., defined as not increasing their VO_{2peak}) progressively fell inversely to a two stepwise progressive increase in endurance-exercise training volume, as described next. Thirty-nine percent (15 of 39) of training subjects did not increase their VO_{2peak} in response to the low-amount, low-intensity training; 18% (9 of 51) had no increase in VO_{2peak} in the group having high-amount, low-intensity training; and 0% (0 of 31) who underwent high-amount, high-intensity training did not increase their VO_{2peak} . A biological basis for the dose–response relationship in the previous sentence could be made from an analysis of interval training (IT) and IT/continuous-training studies published from 1965 to 2012 (Bacon et al. 2013). A second older concept is being reinvigorated; Bacon et al. (2013) indicate that different endurance-exercise intensities and durations are needed for different systems in the body. They suggest that very short periods of high-intensity endurance-type exercise may be needed to reach a threshold for peripheral metabolic adaptations, but that longer training durations at lower intensities are required to see large changes in maximal cardiac output and VO_{2max} .

A comparable example exists for resistance training. Maximal resistance loads require a minimum of 2 min/per wk for each muscle group recruited by a specific maneuver to obtain a strength training adaptation [(8 contrac-

tions/set \times 2 sec/contraction \times 3 sets/day) \times 2 days/wk] = 96 sec]. As of 2016, one opinion from Sarzynski et al. (2016) for the molecular mechanisms by which endurance exercise drives VO_{2max} include, but are not limited to, calcium signaling, energy sensing and partitioning, mitochondrial biogenesis, angiogenesis, immune functions, and regulation of autophagy and apoptosis.

Perhaps more importantly, lifelong aerobic exercise training preserves VO_{2max} into old age. CRF generally increases until early adulthood, then declines the remainder of life in sedentary humans (Astrand 1956). The age-related decline in VO_{2max} is not trivial, as Schneider (2013) reported a \sim 40% decline in healthy males and females spanning from 20 to 70 yr of age. However, cross-sectional data show that with lifelong aerobic exercise training, trained individuals often have the same VO_{2max} as a sedentary individual four decades younger (Booth et al. 2012). Myers et al. (2002) found that low estimated VO_{2max} increases mortality 4.5-fold compared to high estimated VO_{2max} . They concluded, “Exercise capacity is a more powerful predictor of mortality among men than other established risk factors for cardiovascular disease.” Given the strong association between CRF, chronic disease, and mortality, we feel identifying the molecular transducers that cause age-related reductions in CRF may have profound implications for improving health span and delaying the onset of chronic disease. In two of our recent papers, transcriptomics was performed on the triceps muscle (Toedebusch et al. 2016) and on the cardiac left ventricle (Ruegsegger et al. 2017). We were addressing the question of what molecule initiates the beginning of the lifelong decline in aerobic capacity with aging. Aerobic capacity (VO_{2max}) involves, at a minimum, the next systems/tissues, as oxygen travels through the mouth, airways, pulmonary membrane, pulmonary circulation, left heart, aorta/arteries/capillaries, and sarcoplasm/myoglobin to mitochondria. We allowed female rats access, or no access, to running wheels from 5 to 27 wk of age. Surprisingly, voluntary running had no effect on the delay in the beginning of the lifetime decrease in



VO_{2max}. Our skeletal muscle transcriptomics elicited no molecular targets, whereas gene networks suggestive of influencing maximal stroke volume were identified in the left ventricle transcriptomics (Rueggsegger et al. 2017).

Publications concerning the effects of exercise on the brain (from 54 to 216 papers listed on PubMed from 2007 to 2016) have increased 400%. In addition, a 2016 study (Schuch et al. 2016) of three previous papers reported that humans with low- and moderate-CRF had 76% and 23%, respectively, increased risk of developing depression compared to high CRF in three publications. With this forming trend, the next section will consider exercise and brain health.

EXERCISE IMPROVES MENTAL HEALTH

Many studies support physical activity as a non-invasive therapy for mental health improvements in cognition (Beier et al. 2014; Bielak et al. 2014; Tian et al. 2014), depression (Kratz et al. 2014; McKercher et al. 2014; Mura et al. 2014), anxiety (Greenwood et al. 2012; Nishijima et al. 2013; Schoenfeld et al. 2013), neurodegenerative diseases (i.e., Alzheimer's and Parkinson's disease) (Bjerring and Arendt-Nielsen 1990; Mattson 2014), and drug addiction (Zlebnik et al. 2012; Lynch et al. 2013; Peterson et al. 2014). In 1999, van Praag et al. (1999) showed the survival of newborn cells in the adult mouse dentate gyrus, a hippocampal region important for spatial recognition, is enhanced by voluntary wheel running. Similarly, spatial pattern separation and neurogenesis in the dentate gyrus are strongly correlated in 3-mo-old mice following 10 wk of voluntary wheel running (Creer et al. 2010), and the development of new neurons in the dentate gyrus is coupled with the formation of new blood vessels (Pereira et al. 2007). Many exercise-related improvements in cognitive function have been associated with local and systemic expression of growth factors in the hippocampus, notably, brain-derived neurotrophic factor (BDNF) (Neeper et al. 1995; Cotman and Berchtold 2002). BDNF promotes many developmental functions in the brain, including neuronal cell survival, differen-

tiation, migration, dendritic arborization, and synaptic plasticity (Park and Poo 2013). In rat hippocampus, regular exercise promotes a progressive increase in BDNF protein for up to at least 3 mo (Berchtold et al. 2005). In an opposite manner, BDNF mRNA in the hippocampus is rapidly decreased by the cessation of wheel running, suggesting BDNF expression is tightly related to exercise volume (Widenfalk et al. 1999).

Findings by Wrann et al. (2013) highlight one mechanism by which endurance exercise may up-regulate BDNF expression. To summarize, Wrann et al. (2013) noted that exercise increases the activity of the estrogen-related receptor α (ERR α)/peroxisome proliferator-activated receptor γ coactivator 1 α (PGC-1 α) complex, in turn increasing levels of the exercise-secreted factor FNDC5 in skeletal muscle and the hippocampus, whose cleavage products provide beneficial effects in the hippocampus by increasing BDNF gene expression. While future research should determine whether the FNDC5 cleavage-product was produced locally in hippocampal neurons or was secreted into the circulation, this finding eloquently displays one mechanism responsible for brain health benefits following exercise. Similarly, work by van Praag and colleagues suggests that exercise or pharmacological activation of AMP-activated protein kinase (AMPK) in skeletal muscle enhances indices of learning and memory, neurogenesis, and gene expression related to mitochondrial function in the hippocampus (Kobilo et al. 2011, 2014; Guerrieri and van Praag 2015).

Insulin-like growth factor 1 (IGF-1), is central to many exercise-induced adaptations in the brain. Like BDNF, physical activity increases circulatory IGF-1 levels and both exercise and infusion of IGF-1 increase BrdU⁺ cell number and survivability in the hippocampus (Trejo et al. 2001). Similarly, the protective effects of exercise on various brain lesions are nullified by anti-IGF-1 antibody (Carro et al. 2001).

In 1979, Greist et al. (1979) provided evidence that running reduced depression symptoms similarly to psychotherapy. However, the precise mechanisms by which exercise prevents and/or treats depression remain largely unknown. Of the proposed mechanisms, increases

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in the availability of brain neurotransmitters and neurotrophic factors (e.g., BDNF, dopamine, glutamate, norepinephrine, serotonin) are perhaps the best studied. For example, tyrosine hydroxylase (TH) activity, the rate-limiting enzyme in dopamine formation, in the striatum, an area of the brain's reward system, is increased following 7 days of treadmill running in an intensity-dependent manner (Hattori et al. 1994). Voluntary wheel running is also highly rewarding in rats, and voluntary wheel running in rats lowers the motivation to self-administer cocaine, suggesting exercise may be a viable strategy in the fight against drug addiction (Larson and Carroll 2005).

Similar to the above examples, secreted factors from skeletal muscle have been linked to the regulation of depression. Agudelo et al. (2014) showed that exercise training in mice and humans, and overexpression of skeletal muscle PGC-1 α 1, leads to robust increases in kynurenine amino transferase (KAT) expression in skeletal muscle, an enzyme whose activity protects from stress-induced increases in depression in the brain by converting kynurenine into kynurenic acid. Additionally, overexpression of PGC-1 α 1 in skeletal muscle left mice resistant to stress, as evaluated by various behavioral assays indicative of depression (Agudelo et al. 2014). Simultaneously, they report gene expression related to synaptic plasticity in the hippocampus, such as BDNF and CamkII, were unaffected by chronic mild stress compared to wild-type mice. Collectively, these findings suggest exercise-induced increases in skeletal muscle PGC-1 α 1 may be an important regulator of KAT expression in skeletal muscle, which, via modulation in plasma kynurenine levels, may alleviate stress-induced depression and promote hippocampal neuronal plasticity.

TYPE 2 DIABETES, MITOCHONDRIA, AND EXERCISE

T2D Predictions Show a Pandemic

In a 2001 *Diabetes Care* article (Boyle et al. 2001), investigators at the U.S. Centers for Disease Control (CDC) predicted 29 million U.S.

cases of T2D would be present in 2050. Unfortunately, the 2001 prediction of 29 million was reached in 2012! For 2012, the American Diabetes Association reported that 29 million Americans had diagnosed and undiagnosed T2D, which was 9% of the American population (Dwyer-Lindgren et al. 2016). More rapid increases in T2D are now predicted by the CDC than in the previous estimate. The CDC now predicts a doubling or tripling in T2D in 2050. The tripling would mean that one out of three U.S. adults would have T2D in their lifetime by 2050 (Boyle et al. 2010), which would be >100 million U.S. cases. The International Diabetes Federation (IDF) reports T2D cases worldwide. In 2015, the IDF reported that 344 and 416 million North American (including Caribbean) and worldwide adults, respectively, had T2D. Furthermore, the IDF predicts for 2040 that 413 and 642 million, respectively, will have T2D. In sum, T2D is now pandemic, and the pandemic will increase in numbers without current apparent action within the general public.

Type 2 Diabetes Prevalence Is Based on a Strong Genetic Predisposition

The Framingham study found that T2D risk in offspring was 3.5-fold and sixfold higher for a single and two diabetic parent(s), respectively, as compared to nondiabetic offspring (Meigs et al. 2000). Thus, T2D is gene-based.

Noncoding regions of the human genome contain >90% of the >100 variants associated with both T2D and related traits that were observed in genome-wide association studies (Scott et al. 2016). Another 2016 paper (Kwak and Park 2016) lists at least 75 independent genetic loci that are associated with T2D. Taken together, T2D is a complex genetic disease (Scott et al. 2016).

Type 2 Diabetes Is Modulated by Lifestyle, with Exercise as the More Powerful Lifestyle Factor

Three large-scale epidemiological studies have been performed on prediabetics, each in a different geographical location. The first study,



and only study to have separate study arms for diet and exercise, was in China. The pure exercise intervention group had a 46% reduction in the onset of T2D, relative to the nontreated group, after 6 yr of the study (Pan et al. 1997). Diet alone reduced T2D by 31% in the Chinese study. The second study on T2D was the Finnish Diabetes Prevention Study. It found a 58% reduction in T2D in the lifestyle intervention (combined diet and exercise) in its 522 prediabetic subjects after a mean study duration of 3.2 yr (Tuomilehto et al. 2001). The latest of the three studies was in the U.S. Diabetes Prevention Program. The large randomized trial ($n = 3150$ prediabetics) was stopped after 2.8 yr, because of harm to the control group. T2D prevalence in the high-risk adults was reduced by 58% with intensive lifestyle (diet and exercise) intervention, whereas the drug arm (metformin) of the study only reduced T2D by 31%, both compared to the nonintervention group (Knowler et al. 2002). Thus, if differences in genetics in the above three differing ethnicities are not a factor, combined exercise and diet remain more effective in T2D prevention than the drug metformin two decades ago.

Exercise Increases Glucose by Signaling Independent of the Insulin Receptor

A single exercise bout increases glucose uptake by skeletal muscle, sidestepping the insulin receptor and thus insulin resistance in T2D patients (Holloszy and Narahara 1965; Goodyear and Kahn 1998; Holloszy 2005). After insulin binding to its receptor, insulin initiates a downstream signaling cascade of tyrosine autophosphorylation of insulin receptor, insulin receptor substrate 1 (IRS-1) binding and phosphorylation, activation of a PI3K-dependent pathway, including key downstream regulators protein kinase B (Akt) and the Akt substrate of 160 kDa (AS160), ultimately promoting glucose transporter 4 (GLUT4) translocation to the plasma membrane (Rockl et al. 2008; Stanford and Goodyear 2014). Despite normal GLUT4 levels, insulin fails to induce GLUT4 translocation in T2D (Zierath et al. 2000). However, exercise activates a downstream insulin-

signaling pathway at AS160 and TBC1 domain family member 1 (TBC1D1) (Deshmukh et al. 2006; Maarbjerg et al. 2011), facilitating GLUT4 expression translocation to the plasma membrane independent of the insulin receptor. We contend that exercise could be considered as a very powerful tool to primarily attenuate the T2D pandemic.

Complex Biology of T2D Interactions with the Complex Biology of Exercise

An important consideration from the above is that T2D is such a genetically complex disease that a single gene has not been proven to be sufficiently causal to be effective, at this stage in time, to be a successful target for pharmacological treatment. The expectation for a single molecule target has been met for infectious diseases, which are often monogenic diseases. For example, a vaccine against smallpox was highly successful. Edward Jenner in 1796 produced the first successful vaccine. An important fact is that exercise is genetically complex. The literature allows us to speculate that exercise is at least as genetically complex as the approximately 75 genes associated with T2D (Kwak and Park 2016). An example indicating that exercise is complex biology follows. RNA sequencing analysis of all 119 vastus lateralis muscle biopsies found that endurance training for 4 days/wk for 12 wk produced the differential expression of 3404 putative isoforms, belonging to 2624 different genes, many associated with oxidative ATP production in 23 women and men aged 29 yr old (Lindholm et al. 2016). Our notion is that over 2600 genes suggests complex biology.

A “Case-Type” Study of the Molecular Underpinnings of Exercise, Mitochondria, and T2D Interactions

A PubMed search for the terms “diabetes mitochondria exercise molecular” elicited 74 papers. We arbitrarily selected some of the most recent 50 (spanning from mid-2014 into January 2017), with the assumption they would be representative of any other papers that we did not

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find in our search. Papers fell into our two arbitrary categories of single gene studies versus “omic”-type studies. First, subcategories of studies that develop themes will be arbitrarily presented.

Recent Studies Show Single Gene Manipulation Alters Mitochondrial Level and Running Performance

Numerous reports in the past couple of years observed that single gene manipulations increase mitochondrial gene expression and activity, which was also associated with increased exercise performance/capacity. A few of these are presented below:

1. Irisin was shown to increase oxidative metabolism in myocytes and increase PGC-1 α mRNA and protein (Vaughan et al. 2014), which extends the first observation made earlier in adipose tissue by Spiegelman (Bostrom et al. 2012).
 2. Patients with impaired glucose tolerance underwent low-intensity exercise training. Patients whose mitochondrial markers increased to levels that were measured in a separate cohort of nonexercised healthy individuals recovered normal glucose tolerance (Osler et al. 2015). In opposition, those patients whose mitochondria markers did not improve, remained with impaired glucose tolerance.
 3. In 2003, muscle PGC-1 α mRNA was shown to be induced by endurance-exercise training in human skeletal muscle (Short et al. 2003). PGC-1 α was shown to have multiple isoforms (Lin et al. 2002). After a 60-min cycling bout, human vastus lateralis biopsies were taken from both sexes in their mid-20s. Additional biopsies were taken 30 min, and at 2, 6, and 24 hr postexercise. At 30 min postexercise, PGC-1 α -ex1b mRNA and PGC-1 α mRNA increased 468- and 2.4-fold, respectively, whereas PGC-1 α -ex1b protein and PGC-1 α protein increased 3.1-fold and no change, respectively. Gidlund et al. (2015) interprets the above data as implying PGC-1 α -ex1b could be responsible
- for other changes that have previously been recorded before the increase in total PGC-1 α postexercise.
4. Mice with knockout of the kinin B1 receptor gene had higher mitochondrial DNA quantification and of mRNA levels of genes related to mitochondrial biogenesis in soleus and gastrocnemius muscles and had higher exercise times to exhaustion, but did not have higher VO_{2max} (Reis et al. 2015).
 5. Mice do not normally express cholesteryl ester transfer protein (CETP), which is a lipid transfer protein that shuttles lipids between serum lipoproteins and tissues. Overexpression of CETP in mice after 6 wk on a high-fat diet increased treadmill running duration and distance, mitochondrial oxidation of glutamate/malate, but not palmitoyl-carnitine oxidation, and doubled PGC-1 α mRNA concentration (Cappel et al. 2015).
 6. The myokine musclin is a peptide secreted from exercising muscle during treadmill running. Removal of musclin release during running results in lowered VO_{2max}, lower skeletal muscle mitochondrial content and respiratory complex protein expression, and reduced exercise tolerance (Subbotina et al. 2015).
 7. Lactate dehydrogenase B (LDHB), which produces pyruvate from lactate, was overexpressed in mouse skeletal muscle. Increases in markers of skeletal muscle mitochondria were associated with increased running distance in a progressive speed test, and increased peak VO₂ (Liang et al. 2016).
 8. Another example of endurance-type exercise adaptations is the 2016 paper that transcription factor EB (TFEB) regulates metabolic flexibility in skeletal muscle independent of PGC-1 α during endurance-type exercise (Mansueto et al. 2017). Lack of metabolic flexibility, termed “metabolic inflexibility,” is important because it is common in T2D. One definition of metabolic inflexibility is its inability to rapidly switch between glucose and fatty acid substrates for ATP production



when nutrient availability changes from high blood glucose levels immediately after a meal to decreasing below 100 mg/dl when not eating for hours after a meal. A clinical consequence of T2D-induced metabolic inflexibility is prolonged periods of hyperglycemia, because skeletal muscle is more insulin insensitive in T2D. In contrast, after sufficient endurance exercise, skeletal muscle increases its insulin sensitivity by a second pathway that is independent of proximal postreceptor insulin signaling (see Stephenson et al. 2014 for further discussion).

Studies Showing that Manipulation of One Signaling Molecule Does Not Alter Expression of All Genes with Mitochondrial Functions Found in Skeletal Muscles of Wild-Type Animals to Exercise Training

A 2010 review article (Lira et al. 2010) concludes from gene-deletion studies that p38 γ MAPK/PGC-1 α signaling controls mitochondrial biogenesis' adaptation to endurance exercise in skeletal muscle. Two studies do not completely agree with the conclusion in the review article. The Pilegaard laboratory published a 2008 study (Leick et al. 2008) that did not confirm their hypothesis that PGC-1 α was required for every metabolic protein adaptive increase after endurance-exercise training by skeletal muscle. They reported that PGC-1 α was not required for endurance-training-induced increases in ALAS1, COXI, and cytochrome *c* expression (Leick et al. 2008). Their interpretation, at that time, was that molecules other than PGC-1 α can exert exercise-induced mitochondrial adaptations. A second study published in 2012 rendered a similar verdict. A 12-day program of endurance training led to the middle portion of the gastrocnemius muscle demonstrating a similar 60% increase in mitochondrial density in both wild-type and PGC-1 α muscle-specific knockout mice (Myo-PGC-1 α KO) (Rowe et al. 2012). The paper concludes that PGC-1 α is dispensable for endurance-exercise's induction of skeletal muscle mitochondrial adaptations.

Exercise signaling targets have actions that are independent of PGC-1 α , which is specific

to endurance exercise. In 2002, two groups identified PGC-1 β , a transcriptional coactivator closely related to PGC-1 α (Kressler et al. 2002; Lin et al. 2002). Later in 2012, the PGC-1 α 4 variant of PGC-1 α was found to induce skeletal muscle hypertrophy and strength (Ruas et al. 2012). The importance of the finding of a PGC-1 α variant is that it partially explains the phenotypic variation for differing types of exercise. Since the 1970s (Hollloszy and Booth 1976), it has been appreciated that the biochemical and anatomical observations between endurance and resistance differed. For example, Hollloszy and Booth (1976) noted in 1976 that, whereas endurance-type exercise markedly increased skeletal muscle mitochondrial density with very minor increases in muscle fiber diameter, strength-type exercise, in contrast, increased muscle fiber diameter without increases in skeletal muscle mitochondrial density. Taken together, a drug specific for PGC-1 α will not likely mimic separate physical training for endurance, strength/resistance, and coordination types of exercise in the same subject. Thus, the common usage of the term exercise capacity is a misnomer because endurance training and resistance training were shown to have different exercise capacity phenotypes very long ago.

In a 2015 *Diabetes* paper (Wong et al. 2015), Muoio's laboratory concluded that changes in glucose tolerance and total body fat depended upon how much energy is expended in contracting muscle rather than muscle mitochondrial content or substrate selection. A finding to support the previous sentence was the glucose tolerance tests (GTTs). MCK-PGC-1 α mice and their nontransgenic (NT) littermates were not different in GTT, with both being the most glucose intolerant after 10 wk of high-fat feeding. Adding 10 wk of voluntary wheel running to the two high-fat-feed groups during the next 10-wk period (weeks 11–20 of the experiment) lowered the glucose intolerance, and then during weeks 21–30 of the experiment, glucose intolerance was further lowered by adding 25% caloric restriction with the high-fat food and running during the final 10 wk. The percentage weight lost after 30 wk of high-fat feeding was positively related to greater running distances.

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No single front-runner gene candidate could be identified by principle component analysis. Taken together, the paper suggests “doubts” that pharmacological exercise mimetics that increase muscle oxidative capacity will be effective antiobesity and/or antidiabetic agents. Rather, Muoio and investigators suggest energy expenditure by muscle contraction induces localized shifts in energy balance inside the muscle fiber, which then initiates a broad network of metabolic intermediates regulating nutrient sensing and insulin action. A further discussion of complex biology produced by polygenicity continues next.

POLYGENICITY OF EXERCISE LEADS TO COMPLEX MULTISYSTEM RESPONSES TO IMPROVE HEALTH OUTCOMES

Multiple tissues, organs, and systems are influenced by physical activity, or the lack thereof (Table 2).

Table 2. Worsening of maximal functioning in selected major organ/tissue/systems that are caused by the lack of physical activity with growth, maturation, and aging

| Site | Organ/tissue/system loses designated function, most with aging after maturation |
|------------------------|---|
| Brain | Specific types of cognition |
| Brain | Motor coordination and balance |
| Heart | Maximal pumping volume/minute |
| Peripheral circulation | Maximal capacity to supply blood to working muscles |
| Peripheral circulation | Prevention of capillary rarefaction in feet |
| Skeletal muscle | Less mass and strength |
| Skeletal muscle | Lower insulin sensitivity |
| Pancreas | Loss of β cells |
| Aerobic capacity | Lower |
| Bone | Less mass and strength |
| Liver | Excessive fat storage |

The higher their maximal function is before the end of each item's maturation, the longer chances are that the quality of life will remain optimal. The breadth of the list implies that a single molecular target will not substitute for appropriate daily physical activity to prevent the loss of all listed items.

To present one extreme, that most will agree, one molecule will not describe the 1000s of molecules adapting to aerobic, resistance, and coordination exercise training. On the opposite extreme, many could likely agree that usage of the various “omics” underlying all adaptations to physical activity will differ (i.e., not be identical in most aspects) among the next list: various cell types within a tissue/organ, tissues/organs, and various intensities of physical activity (i.e., the thresholds among gene responses for health benefits will differ because of the presence of responders and non-responders, or protein isoform type); during various types cycling (circadian or menstrual); postprandial versus fasting between meals; male and female; child, adult, and the elderly; trained and untrained; aerobic- and resistance-exercise types; and so forth. Others have repetitively written that only ~59% of the risk reduction for all forms of CVD have been shown to be caused by effects through traditional factors (Mora et al. 2007; Joyner and Green 2009). Thus, we pose the next question: what is the identity of all molecules in the yet-to-be-discovered gap between our knowledge of single gene functions and the totality of personalized prescription of physical activity to maximize the period of life free of any chronic disease, termed health span?

While approaches using single-gene manipulations are valuable tools, research must also focus on integrating exercise-responsive molecules into networks that maintain or improve health. This process will reveal complex, multi-system, polygenic networking essential for the advancement of many goals pertaining to exercise physiology, such as tailoring exercise prescriptions and implementing personalized medicine. One example is the developing myokine network with auto-, para-, and endocrine molecules. The first myokine interleukin (IL)-6 began to be described as early as 1994 by the Pedersen laboratory (Ullum et al. 1994), with a history of its development as the first exercise myokine recounted in 2007 (Pedersen et al. 2007). Since their discovery, myokine action within and at a distance from their origins in skeletal muscle have been increasingly studied,

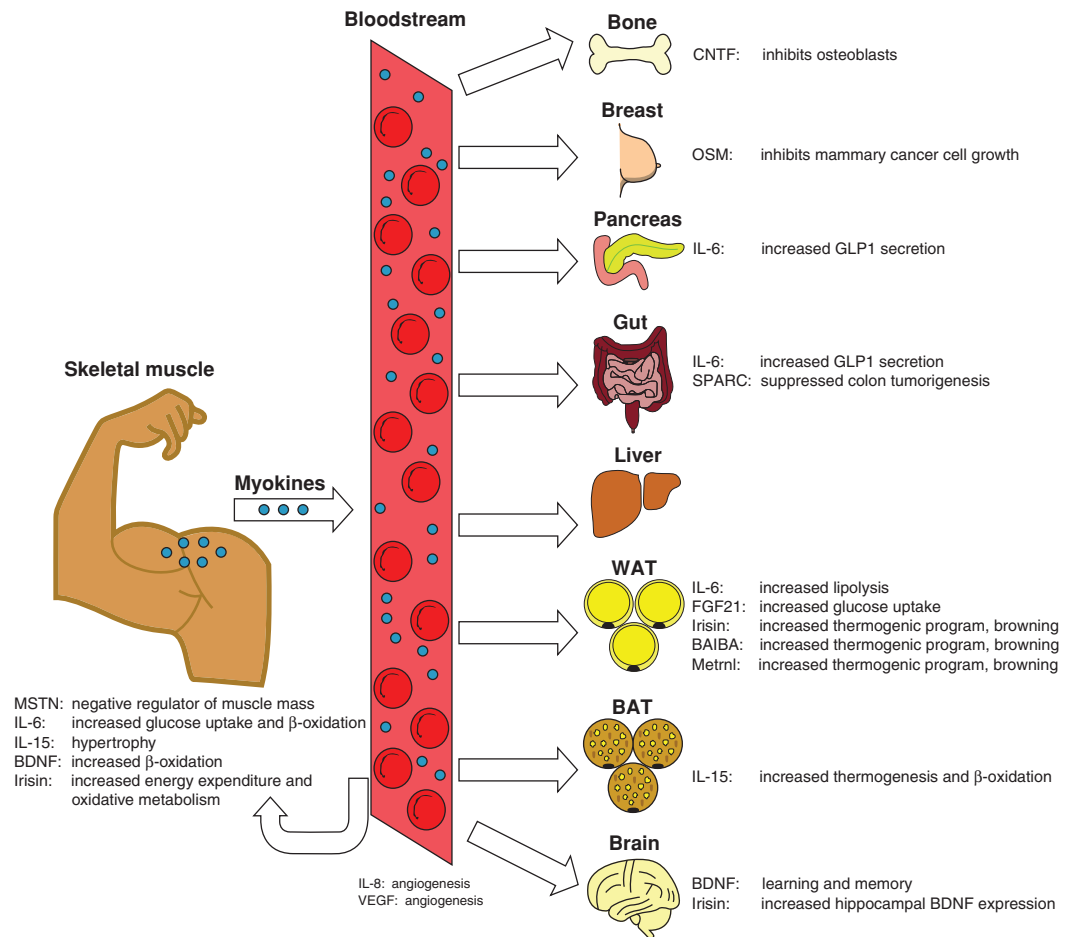


Figure 2. Figure provides an illustration of myokine production by skeletal muscle for actions within or at a distance. Myokine release promotes a high degree of intertissue cross talk. CNTF, Ciliary neurotrophic factor; OSM, oncostatin M; IL, interleukin; BDNF, brain-derived neurotrophic factor; VEGF, vascular endothelial growth factor. (From Schnyder and Handschin 2015; reprinted, with permission, courtesy of PMC Open Access.)

as schematically illustrated by Schnyder and Handschin (2015) (Fig. 2).

Similarly, maximal aerobic exercise is accompanied by tremendous stress on many systems, yet whole-body homeostasis is remarkably maintained. For example, world-class endurance athletes can increase whole-body energy production well over 20-fold (Joyner and Coyle 2008), whereas maintaining blood glucose concentrations at resting levels (Wasserman 2009). Intuitively, such effort would require sophisticated interorgan cross talk and polygenic integration of numerous functions.

Exercise Provides Too Many Benefits to “Fit into a Single Pill”

Despite the well-known benefits of exercise, most adults and many children lead relatively sedentary lifestyles and are not active enough to achieve the health benefits of exercise (Warburton et al. 2006; Fried 2016). Accelerometry measurements suggest that >90% of U.S. individuals >12 yr of age and ~50% of children aged 6–11 yr old fail to meet U.S. Federal physical activity guidelines (Troiano et al. 2008). Given this incredibly low compliance, the iden-

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tification of genetic and/or orally active agents that mimic the effects of endurance exercise might have high appeal for a majority of sedentary individuals. This high appeal has led to recent identification/development of exercise “mimetics.” In 2009, we set criteria for proper usage of the term “exercise mimetic,” based upon its common usage (Booth and Laye 2009). We gave the Oxford English Dictionary’s definition of mimetic, “A synthetic compound that produces the same (or a very similar) effect as another (especially a naturally occurring) compound.” While many exercise “mimetics” activate signaling pathways commonly associated with muscle endurance, these agents have not completely mimicked all effects for all types of exercise. For example, the AMPK activator 5-aminoimidazole-4-carboxamide ribonucleotide (AICAR), when given daily to rats over a 5-wk-period, did not increase maximal oxygen consumption (VO_{2peak}) in the sedentary group of rats that were forced to run to VO_{2peak} on treadmills, as compared to sedentary rats receiving the vehicle (Toedebusch et al. 2016). Thus, in our opinion, the published claim (Narkar et al. 2008) that AICAR is an exercise mimetic is invalidated because it did not increase VO_{2peak} . While these agents may undoubtedly have specific health benefits, it is currently impractical to assume that all of the benefits of exercise can be replaced by “exercise mimetics.”

CONCLUDING REMARKS

Exercise is a powerful tool in the fight to prevent and treat numerous chronic diseases (Table 1). Given its whole-body, health-promoting nature, the integrative responses to exercise should surely attract a great deal of interest as the notion of “exercise is medicine” continues to its integration into clinical settings.

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