

EDITORIAL COMMENT

Further Evidence of Harm From Exercise in ARVD/C*



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Exercise is loathsome. And it cannot be any benefit when you are tired, and I was always tired.

—Mark Twain (1)

The evidence implicating exercise as a pathogenic factor in arrhythmogenic right ventricular dysplasia/cardiomyopathy (ARVD/C) continues to grow. Early reports of this condition recognized its occurrence with athletic activity, and later studies strengthened this association (2,3). Retrospective quantification of exercise for 87 people with desmosome mutations demonstrated that those with the highest levels of exercise developed earlier and more severe manifestations of this inherited cardiomyopathy (3). In a complementary analysis of 82 patients with ARVD/C, those without desmosomal mutations or family history of ARVD/C had performed more intense exercise (4). These findings support an emerging paradigm in which both genetic predisposition and environmental factors interact around a threshold for phenotypic expression of ARVD/C (5).

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Many questions remain regarding the pathogenesis of ARVD/C, and in this issue of the *Journal*, Cruz et al. (6) highlight an alternative approach to its investigation. Using an adeno-associated virus (AAV)-mediated gene transfer protocol, they overexpressed a common human *PKP2* mutation in mice. A single

intravenous injection of an AAV vector, harboring the *PKP2* p.Arg735Ter mutation, led to cardiac production of truncated plakophilin-2 for 10 months. Unfortunately, this was insufficient for recapitulating ARVD/C. However, on the basis of prior human, murine, and cellular studies implicating exercise as harmful in the context of desmosome mutation (3,7,8), Cruz et al. (6) subjected these AAV-treated mice to forced swim exercise. Indeed, exercised *PKP2* mutant mice demonstrated reduced global right ventricular (RV) ejection fraction compared with exercised mice with wild-type *PKP2* transgene or sedentary mutant mice.

This work is notable for some important findings. The authors demonstrate that AAV achieves robust cardiac expression, with recapitulation of RV dysfunction induced by the addition of mutant human *PKP2*. We can explain more than 50% of this disease on the basis of a desmosome mutation, yet rare or novel variation in these genes commonly occurs among asymptomatic control subjects (9,10). The use of this technology would allow uncertain desmosome variants to be assessed similarly, which is certainly easier than generating mice with knock-in mutations. Somewhat surprisingly, the investigators were able to induce cardiac dysfunction despite a normal content of murine plakophilin-2 by overexpressing the truncated human protein. The majority of *PKP2* mutations led to premature transcript termination, which suggests it is caused by insufficient protein. Furthermore, *Pkp2* haploinsufficiency led to sodium current deficits in murine hearts, and *PKP2* knockdown in HL-1 myocytes produced features of ARVD/C (11,12). However, the initial report of *PKP2* mutations causing ARVD/C demonstrated that some truncating mutations have normal mutant *PKP2* messenger ribonucleic acid content, without degradation by nonsense-mediated messenger ribonucleic acid decay (13). The findings by Cruz et al. (6) add further support for a dominant-negative

*Editorials published in the *Journal of the American College of Cardiology* reflect the views of the authors and do not necessarily represent the views of JACC or the American College of Cardiology.

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pathogenesis imparted by *PKP2* mutations, in addition to prior evidence regarding haploinsufficiency. This work also contributes to the growing body of evidence that indicates that hearts with desmosome mutations are more susceptible to exercise-induced dysfunction than those without a desmosome mutation.

Although this AAV model is efficient and relatively inexpensive compared with other methods of investigation, it falls short in a few ways. First, the pathological hallmarks of disease are largely absent, including a lack of fibrofatty scar, plakoglobin redistribution, or even ultrastructural changes in the cardiac desmosome by transmission electron microscopy. Still, Cruz et al. (6) demonstrated punctate junctional distribution of the principal gap junction protein connexin-43 (Cx43) in exercised mutant mice compared with continuous distribution of Cx43 at myocyte junctions in sedentary mutant mice. Abnormal Cx43 distribution is an important, yet nonspecific, histological feature of ARVD/C (14). Although the authors reported a small increase in the QRS duration in mutant mice compared with controls, they did not show significant arrhythmias or decreased survival as a consequence of mutant *PKP2* overexpression.

Aerobic exercise has remarkable cardiovascular benefits, decreasing the risk of atherosclerosis, reducing death from heart failure, lowering blood pressure, decreasing abdominal fat, lowering glycosylated hemoglobin in diabetes mellitus, and improving sarcopenia (15-18); however, endurance athletics is not without some risk. La Gerche et al. (19) investigated 40 subjects who participated in endurance races (3 to 11 h in duration) and showed that

this level of exercise causes acute RV, but not left ventricular, dysfunction. The mechanism of RV dysfunction in response to exercise is not known. One simple explanation is that the thin-walled RV is stretched by the larger volumes and cardiac output required by aerobic activity, and desmosome mutations cause a loss of myocyte cell adhesion, which worsens the response to exercise. Although this explanation is appealing, it does not adequately account for some of the other mechanistic data published for ARVD/C. Several investigations have highlighted an important role for diminished Wnt/ β -catenin signaling in ARVD/C, although exercise appears to stimulate the canonical Wnt pathway (20,21). Peroxisome proliferator-activated receptor (PPAR)- γ and PPAR- α stimulation of induced pluripotent stem cell-derived cardiomyocytes from patients with ARVD/C leads to exaggerated lipogenesis and apoptosis, which implicates this pathway in ARVD/C pathogenesis (22). Because exercise upregulates the PPAR- γ pathway, this may contribute to the harmful effects of exercise for patients with ARVD/C (23).

As several groups continue to address these tantalizing questions regarding the pathogenic effects of exercise, improved understanding of the interaction of desmosome mutations, aerobic activity, and cardiac metabolism should lead to better estimation of risk and more effective therapies to prevent ARVD/C and sudden cardiac death.

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KEY WORDS arrhythmia, cardiomyopathy, exercise, sudden cardiac death