specificity. However, we did not think that an Agatston Ca score > 400 significantly impaired the specificity of MDCT, as only 1 in 5 patients with a Ca score > 400 did not show acute coronary syndrome. With similar devices, in the presence of a Ca score > 400, the proportion of nonconclusive studies increases, luminal stenosis is overestimated and the specificity of the technique is severely limited; moreover, a Ca score of > 400 has been shown to be an excellent predictor for significant coronary disease. Along the same lines, Goldstein et al. recommended performing single-photon emission computed tomography (SPECT) when Ca scores were > 100, markedly lower than the 400 score used in our study.

Finally, in our opinion, the cost-effectiveness differences between the studies by Hoffman et al. and Goldstein et al. were not exclusively due to the differences in the cut-off values chosen for stenosis (50% vs 70%). Moreover, there were differences in the prevalence of acute coronary syndrome in the MDCT group (9% vs 4.4%), as well as large differences in the percentage of additional tests conducted in the control groups of the 2 studies (45% vs 100%, respectively), which contributed to the discrepancies observed.

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The Genetic Background of Left Ventricular Hypertrabeculation / Noncompaction Remains Vague

El trasfondo genético de la hipertrabeculación/miocardiopatía no compactada ventricular izquierda sigue sin estar claro

To the Editor,

We read with interest the article by Rodríguez-Serrano et al. about familial left ventricular hypertrabeculation/noncompaction (LVHT) associated with a novel alpha-cardiac actin gene (ACTC1)-mutation in 4 family members (II:4, III:4, III:6, IV:1), of whom 3 (II:4, III:4, III:6) presented with noncompaction and 1 with hypertrabeculation of the explanted heart. We have the following comments and concerns.

We do not agree with the statement that the described ACTC1-mutation “caused” LVHT. LVHT is associated with mutations in a large number of different genes but no proof has ever been provided for any of these associations that a particular mutation is truly causative of this myocardial abnormality. Reservations against a causal relation comes from the following arguments: first, in most cases of hereditary disease in which LVHT has been described, only a limited number of mutation carriers also had LVHT. Second, LVHT may be a dynamic abnormality that may not be present at birth in single patients (acquired LVHT) and may more rarely even disappear during life. Third, most of the few patients with acquired LVHT did not carry a mutated gene and did not have LVHT on previous echocardiographic or other cardiac imaging studies. Fourth, according to the authors themselves, the pathogenicity of the detected ACTC1 variant was neither confirmed nor excluded by in silico analysis.

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are the diseases most frequently associated with LVHT\(^7\) and because of the uncertainty whether the ACTCI alteration is a polymorphism or a pathogenic mutation. It would also be worthwhile to conduct a neurological examination in family members who did not show LVHT. Were creatine-kinase serum levels normal in all investigated patients?

Overall, this interesting report could benefit from clarification of some inconsistencies. It is also important to discuss the absence of LVHT on echocardiography in patient IV:1. The more details that are provided about patients or families with LVHT, the more likely the cryptogenic pathogenesis of this still enigmatic myocardial abnormality will be clarified.

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**The Genetic Background of Left Ventricular Hypertrabeculation/Noncompaction Remains Vague. Response**

**El trasfondo genético de la hipertrabeculación/miocardiopatía no compactada ventricular izquierda sigue sin estar claro. Respuesta**

**To the Editor,**

We appreciate the comments by Drs Finsterer and Zarrouk-Mahjoub.

These authors seem to question the genetic basis of left ventricular noncompaction (LVNC), contradicting the position of the European Society of Cardiology/American Heart Association (ESC/AHA).\(^1–3\) Although helpful, functional studies are not routinely performed. Instead, evidence in the literature, cosegregation, consequences in the protein and in silico studies are usually employed (as we did). Mutation carriers may not exhibit the phenotype because of an incomplete penetrance\(^2\) and diagnostic difficulties, such as different sets of criteria, suboptimal echocardiographic quality and reproducibility,\(^3\) and unavailability of magnetic resonance imaging.

Is LVNC acquired? Can it disappear? These issues are unresolved\(^2,4\) and have not been addressed.

In silico studies are not the only data to assess a mutation. Additional information supported the pathogenic effect of ACTCI\(^128907\) (third paragraph, page 859). The genetic heterogeneity of LVNC is unquestionable.\(^2,3\)

The preferred term is LVNC (PubMed) and the ESC considers “hypertrabeculation” to be incorrect.\(^2\) Even so, the above-mentioned authors prefer LVHT. We use LVNC if the criteria are fulfilled and hypertrabeculation (see the Figure in the paper by Rodríguez-Serrano et al.\(^2,5\) when these criteria cannot be assessed. Accordingly, hypertrabeculation for the heart explant (histologic criteria for LVNC are lacking) should also have been used within the text, but was changed for LVNC because of word count constraints.

Individuals II:4 and III:4 fulfilled the criteria of Chin and Stollberger whereas individual III:6 did not.

The echocardiogram of patient IV:1, thoroughly reviewed, lacked LVNC. There were no histopathologic studies or stored pictures or tissues. Image acquisition limitations at the intensive care unit (small infant heart with an LV assist device) could explain the discrepancy but it is also possible that no discrepancy was actually present, the situation being a cardiomyopathy presenting with different phenotypes, namely restrictive cardiomyopathy in an infant (which can also be caused by ACTCI mutations\(^2\) and LVNC in adults. Many circumstances may account for this phenomenon (age-dependent expression of modifier genes, additional mutations…).

Finally, neurological signs/symptoms and creatine kinase elevation were ruled out.

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