



sense, it would have been very interesting to analyse whether PRS combined with results of SCB and other risk factors⁶ could better define the arrhythmic risk in BrS. In other words, an important study, an important first step that needs further scientific follow-up; an important analysis that, like all significant analyses, gives us some answers but opens up multiple new questions (Figure 1).

Conflicts of interest: none declared.

References

1. Tadros R, Tan HL, for the ESCAPE-NET Investigators, el Mathari S, Kors JA, Postema PG, Lahrouchi N, Beekman L, Radivojkov-Bлагоjevic M, Amin AS, Meitinger T, Tanck MW, Wilde AA, Bezzina CR. Predicting cardiac electrical response to sodium-channel blockade and Brugada syndrome using polygenic risk scores. *Eur Heart J* 2019;**40**:3097–3107.
2. Strauss DG, Vicente J, Johannesen L, Blinova K, Mason JW, Weeke P, Behr ER, Roden DM, Woosley R, Kosova G, Rosenberg MA, Newton-Cheh C. Common genetic variant risk score is associated with drug-induced QT prolongation and torsade de pointes risk. *Circulation* 2017;**135**:1300–1310.
3. Collins FS, Varmus H. A new initiative on precision medicine. *N Engl J Med* 2015; **372**:793–795.
4. Bezzina CR, Barc J, Mizusawa Y, Remme CA, Gourraud JB, Simonet F, Verkerk AO, Schwartz PJ, Crotti L, Dagradi F, Guicheney P, Fressart V, Leenhardt A, Antzelevitch C, Bartkowiak S, Borggrefe M, Schimpf R, Schulze-Bahr E, Zumhagen S, Behr ER, Bastiaenen R, Tfelt-Hansen J, Olesen MS, Kääh S, Beckmann BM, Weeke P, Watanabe H, Endo N, Minamino T, Horie M, Ohno S, Hasegawa K, Makita N, Nogami A, Shimizu W, Aiba T, Froguel P, Balkau B, Lantieri O, Torchio M, Wiese C, Weber D, Wolswinkel R, Coronel R, Boukens BJ, Béziau S, Charpentier E, Chatel S, Despres A, Gros F, Kyndt F, Lecointe S, Lindenbaum P, Portero V, Violleau J, Gessler M, Tan HL, Roden DM, Christoffels VM, Le Marec H, Wilde AA, Probst V, Schott JJ, Dina C, Redon R. Common variants at SCN5A–SCN10A and HEY2 are associated with Brugada syndrome, a rare disease with high risk of sudden cardiac death. *Nat Genet* 2013;**45**:1044–1049.
5. Dooijes D, Siemlink M, Baas AF. Evaluation of gene panels for inherited cardiac disease—is less more? *Neth Heart J* 2019;**27**:297–298.
6. Sieira J, Conti G, Ciconte G, Chierchia GB, Casado-Arroyo R, Baltogiannis G, Di Giovanni G, Saitoh Y, Juliá J, Mugnai G, La Meir M, Wellens F, Czaplá J, Pappaert G, de Asmundis C, Brugada P. A score model to predict risk of events in patients with Brugada syndrome. *Eur Heart J* 2017;**38**:1756–1763.

Corrigendum

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In the original version of this article, a small number of the MMM Investigators were missing or recorded incorrectly, and an author's affiliation was incorrect. This has now been corrected online and in print.

The authors apologise for the error.

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