Check for updates

OPEN ACCESS

EDITED BY Xiaofeng Yang, Temple University, United States

REVIEWED BY Katja Odening, University Hospital of Bern, Switzerland

*CORRESPONDENCE Oscar Campuzano Scar@brugada.org Georgia Sarquella-Brugada georgia.sarquella@sjd.es

[†]These authors have contributed equally to this work and share first authorship

⁺These authors have contributed equally to this work and share senior authorship

SPECIALTY SECTION

This article was submitted to Sex and Gender in Cardiovascular Medicine, a section of the journal Frontiers in Cardiovascular Medicine

RECEIVED 11 February 2023 ACCEPTED 24 March 2023 PUBLISHED 04 April 2023

CITATION

Díez-Escuté N, Arbelo E, Martínez-Barrios E, Cerralbo P, Cesar S, Cruzalegui J, Chipa F, Fiol V, Zschaeck I, Hernández C, Campuzano O and Sarquella-Brugada G (2023) Sex differences in long QT syndrome.

Front. Cardiovasc. Med. 10:1164028. doi: 10.3389/fcvm.2023.1164028

COPYRIGHT

© 2023 Diez-Escuté, Arbelo, Martinez Barrios, Cerralbo, Cesar, Cruzalegui, Chipa, Fiol, Zschaeck, Hernández, Campuzano and Sarquella-Brugada. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or

reproduction is permitted which does not comply with these terms.

Sex differences in long QT syndrome

Nuria Díez-Escuté¹¹, Elena Arbelo^{2,3,4,5†}, Estefanía Martínez-Barrios¹, Patricia Cerralbo¹, Sergi Cesar¹, José Cruzalegui¹, Freddy Chipa¹, Victoria Fiol¹, Irene Zschaeck¹, Clara Hernández¹, Oscar Campuzano^{4,6,7*†} and Georgia Sarguella-Brugada^{1,5,6*†}

¹Arrhythmia, Inherited Cardiac Diseases and Sudden Death Unit, Hospital Sant Joan de Déu, Barcelona, Spain, ²Arrhythmia Section, Cardiology Department, Hospital Clínic, Universitat de Barcelona, Barcelona, Spain, ³IDIBAPS, Institut d'Investigació August Pi I Sunyer (IDIBAPS), Barcelona, Spain, ⁴Centro de Investigación Biomédica en Red en Enfermedades Cardiovasculares, Madrid, Spain, ⁵European Reference Network for Rare, Low Prevalence and Complex Diseases of the Heart, Amsterdam, Netherlands, ⁶Medical Science Department, School of Medicine, University of Girona, Girona, Spain, ⁷Cardiovascular Genetics Center, University of Girona-IDIBGI, Girona, Spain

Long QT Syndrome (LQTS) is a rare, inherited channelopathy characterized by cardiac repolarization dysfunction, leading to a prolonged rate-corrected QT interval in patients who are at risk for malignant ventricular tachyarrhythmias, syncope, and even sudden cardiac death. A complex genetic origin, variable expressivity as well as incomplete penetrance make the diagnosis a clinical challenge. In the last 10 years, there has been a continuous improvement in diagnostic and personalized treatment options. Therefore, several factors such as sex, age diagnosis, QTc interval, and genetic background may contribute to risk stratification of patients, but it still currently remains as a main challenge in LQTS. It is widely accepted that sex is a risk factor itself for some arrhythmias. Female sex has been suggested as a risk factor in the development of malignant arrhythmias associated with LQTS. The existing differences between the sexes are only manifested after puberty, being the hormones the main inducers of arrhythmias. Despite the increased risk in females, no more than 10% of the available publications on LQTS include sex-related data concerning the risk of malignant arrhythmias in females. Therein, the relevance of our review data update concerning women and LQTS.

KEYWORDS

long QT syndrome, gender, arrhythmias, sudden cardiac death, woman

1. Introduction

Congenital long QT syndrome (LQTS) is a cardiac channelopathy, characterized by ventricular repolarization and polymorphic ventricular tachycardia (*torsades de pointes*, TdP), leading to malignant arrhythmias, syncope and sudden cardiac death (SCD) at a young age. It is clinically recognized by a prolonged QT interval in the surface electrocardiogram (ECG). LQTS is one of the most common inherited arrhythmia conditions (1:2000/1:2500). LQTS is caused by rare genetic alterations in cardiac ion channels or accessory ion channel subunits, mainly following an autosomal-dominant pattern of inheritance. To date, there are 17 genes potentially associated with LQTS, but definite deleterious alterations have been identified in three genes (*KCNQ1, KCNH2* and *SCN5A*) that account for about 90% of all LQTS cases (1). Due to the potential risk of malignant arrhythmias, it is crucial to accurately identify and manage patients. Continuous advances in diagnosis and personalized treatment, as well as prevention, have been achieved in the last few years. However, risk stratification remains the main,

10.3389/fcvm.2023.1164028

challenge in clinical practice at present. Variable expressivity and incomplete penetrance are hallmarks of LQTS, impeding a conclusive risk stratification. Currently, the existence of differences in sex in LQTS is widely accepted, showing females with an increased risk of developing polymorphic ventricular arrhythmia or SCD than men after the onset of adolescence (2). In addition, women experience a decreased risk during pregnancy, but increased in postpartum period and perimenopause. These differences exist due to hormone levels, which vary depending on the menstrual cycle, gestation, and the postnatal period. Despite being widely accepted as definite risk factor, no more than 10% of studies focused on the pathophysiological mechanism occurring in females suffering from LQTS have been published to date (Figure 1). Herein, we will review evidence from basic and clinical studies involving female-susceptibility to LQTS. A better understanding of the role of sex-related differences in LQTS will lead to improvement in risk stratification.

2. Clinical findings

Congenital LQTS is a primary cardiac electric disease characterized by the prolongation of the QT interval, usually associated with T-wave abnormalities. To make a definite diagnosis, it is essential to exclude secondary causes of the prolonged QT interval, such as QT-prolonging drugs or electrolyte imbalances (3). High risk of life-threatening arrhythmias includes T wave alternans and functional 2:1 atrioventricular block in the ECG. In addition to the baseline ECG, QTc behavior can also be assessed during stress testing and 24 h Holter recording, preferably 12-lead. Therefore, the "Schwartz score" was developed as a diagnostic criterion to support the diagnosis of the disease if a score of >=3,5 is attained. Adrenaline testing, supine-to-standing ECG or mental stress tests are of lower diagnosis yield (4). It is widely accepted that QT intervals are generally longer in healthy women; therefore, sex-specific cut-off values for prolonged QTc should be applied in order to accurately diagnose LQTS -470 ms in men and 480 ms in women- (5). In addition, women are more susceptible of developing a QT prolongation at slower heart rates, making QTc duration at rest and during sleep critical markers of arrhythmic risk (6). Therefore, adult women with LQTS are at higher risk of malignant arrhythmias due to the influence of a hormone, requiring pharmacological therapy (7). Taking into account all these points, a novel risk score for patients with long QT syndrome has been developed, offering accurate prognostic information to guide clinicians in identifying the patients at the highest risk of life-threatening arrhythmias (8).

The use of β -blockers (preferably nadolol) is the most effective therapy in both sexes (especially in LQTS type 1 and 2), despite the fact that response varies by sex and underlying genotype. Patients with LQTS type3 may benefit from mexiletine or even flecainide; left cardiac sympathetic denervation may be also offered despite rarely and only for special cases. Finally, a reduced number of LQTS patients are suitable for ICD, such as primarily survivors of cardiac arrest and patients at a high-risk for SCD, with recurrent syncope despite adequate pharmacological therapy.

3. Genetic and cellular basis

The normal QT interval in the ECG represents the time from the beginning to the end of ventricular depolarization. LQTS is characterized by a prolonged QT interval due to cardiac repolarization dysfunction leading to risk for ventricular tachyarrhythmias, syncope, and even SCD (3). Indeed, QT prolongation is related to a combination of modifiable and



unmodifiable risk factors. Electrolyte anomalies are the most common risk factors associated with prolonged QT. Among those, hypokalemia has the main arrhythmogenic effect, as well as hypocalcemia and hypomagnesemia. They may be responsible for prolonging the QT interval, but may also be a major risk factor for drug-induced LQTS, one of the most frequent reasons for QT prolongation (9). Genetic background, as well as older age and female sex, are the most important unmodifiable risk factors. Focused on genetics, rare deleterious alterations located in genes encoding ion channels or associated proteins have been reported as a cause of LQTS. Diagnosed families follow an autosomal dominant pattern of inheritance, with characteristic incomplete penetrance and variable expressivity. Nowadays hundreds of rare alterations have been reported in more than 15 genes (10), despite only a limited number of variants that are definitively classified as deleterious following current ACMG recommendations (11). These causative variants are located mainly in three genes (KCNQ1, KCNH2, and SCN5A), and are responsible up to 75% of all patients with LQTS (3, 10). Loss-offunction deleterious variants in the potassium channels KCNQ1 and KCNH2 are responsible for LQT1 and LQTS2, respectively. They cause decreasing activity of the slow delayed rectifier current (IKs) and rapid delayed rectifier current (IKr) (phase 3 of the action potential), respectively. In contrast, gain-of-function deleterious variants in the SCN5A gene (sodium channel, phase 0 of an action potential) are responsible for LQTS3. They cause persistent sodium influx that extends through the plateau phase. A loss of IKs or IKr function, or gain of INa function predisposes ventricular myocytes to early afterdepolarizations, then triggering malignant arrhythmias. Other rare alterations have been reported in minor genes, accounting for 5% of LQTS (*CACNA1C, CALM1, CALM2, CALM3,* and *TRDN*), whereas about 20% of all diagnosed patients do not have an identifiable deleterious variant in any of the current known genes related to LQTS (3). We must also remark that co-inheritance of a second rare variant that affects ventricular repolarization is described as a "second hit". These variants are classified as deleterious in the same gene (compound heterozygosity) or in a different gene (digenic heterozygosity). Compound deleterious variants are present in 5%–10% of LQTS patients and it is well-accepted as cause of a more severe phenotype (12). Concerning genetic risk after the onset of adolescence, women with pathogenic variants in *KCNQ1* (LQTS1) and *KCNH2* (LQTS2) are at increased risk of malignant arrhythmias, especially LQT2 due to a pore loop pathogenic variant (13) (Figure 2).

4. Pregnancy, post-partum and perimenopausal periods

It is widely accepted that there exists a slight female phenotypic predominance in LQTS, with the female sex being a risk factor for malignant arrhythmias. This increased risk occurs after adolescence due to sex hormone differences, with testosterone being a main cause of QT-interval duration in men (2). Low risk of arrhythmias is widely accepted during pregnancy but this risk increases during postpartum, menopausal, and perimenopausal periods (Figure 2). The complex interaction of sex hormones and cardiac ion currents/action potential is widely accepted but it still remains to be clarified (14). What is clear is that sex differences in the electrical substrate are not the result of a simple change in the expression of a single or even a few ionic



currents (7). As a preventive measure, a multidisciplinary approach to women with LQTS ensures comprehensive risk assessment and optimal patient care.

4.1. Pregnancy

Fluctuations in sex hormone levels during pregnancy could potentially provoke cardiac events. In addition to sex hormone levels, other internal or external factors may alter cardiac electrophysiology during pregnancy and postpartum (alterations in adrenergic activity, disrupted sleep pattern), that can contribute to changes in arrhythmic risk (14). It is widely reported that pregnancy decreases risk of malignant arrhythmias in females diagnosed with LQTS, especially in LQTS type 1 (15). The choice of b-blockers has not been established due to the limited evidence available nowadays. One thing that is for certain is that the use of b-blockers such as propranolol is effective in reducing the risk of arrhythmic events (15, 16). However, lower fetal birth weight has been reported, leading to non-selective bblocker use, mainly metoprolol (17). Because of the risk in mothers with LQTS for stillbirth and the higher risk of miscarriages, a more stringent follow-up of these patients during pregnancy might be necessary. Finally, concerning contraceptives, no conclusive studies have been published to date, with the risk of arrhythmias remaining unknown (18).

4.2. Postpartum

Concerning post-partum follow-up, women with LQTS, especially LQTS type 2, show a higher risk of VT and SCD in comparison to the relatively low risk during pregnancy. After postpartum, the risk of cardiac events returns to basal levels before pregnancy. However, there are no general recommendations and approved schemes on how women with LQTS should be supervised after delivery. Limited data suggest increased risk in the early post-partum period, particularly in patients with LQTS2 (15). In addition, the use of b-blockers is mostly well tolerated during the postpartum period to prevent life-threatening manifestations. It is also important to note that b-blockers are secreted in breast milk, but hypoglycemia and bradycardia may occur in breastfed infants, albeit rarely (19). Therefore, dose adjustment of beta-blockers may be needed and postpartum care remains the same as in routine cases (20).

4.3. Menopausal and perimenopausal periods

Nowadays, only one study has been published focusing on females after the onset of menopause who have been diagnosed with LQTS. Available data suggest a higher risk of malignant arrhythmias in LQTS-type2 women (21). In contrast, there are studies in post-menopausal period showing that estrogen replacement therapy (ERT) prolongs the QTc interval more, compared to those postmenopausal women taking no hormones or taking combined estrogen-progestin replacement therapies (22, 23). It suggests that progesterone has a similar protective effect to testosterone (24).

5. Role of hormones in arrhythmogenic risk

The underlying molecular mechanisms that cause patients with LQTS to have sex-dependent variability in arrhythmogenic risk remain to be elucidated. However, several studies suggest that it results from the effect of certain hormones on cardiac ion channels. Animal experiments have shown that estradiol could act as a pro-arrhythmic agent in LQTS2, whereas progesterone would have a protective role (25). This effect of estradiol is attributed to its interaction with certain potassium ion currents (26), and due to the effect on the transcription of some genes like KCNE2 and RyR2, being mediated by an estrogen receptordependent process (27, 28). However, such effects have not been supported in human studies, in which a more complex interaction between estrogen and progesterone appears to exist (29). For example, a recent study in women with LQTS found an inverse relationship of RR interval with estradiol levels during the menstrual cycle (30). In addition, progesterone was found to show an inverse association with the corrected QT interval and with the ratio of progesterone to estradiol in women with LQTS2. This shortening of the QT interval was observed during the luteal phase, and was mainly attributed to increased progesterone levels in this phase (30). Such associations were not maintained in women with LQTS1, supporting that the observed differences in LQTS subtypes are due to different effects of hormones on the channels, with different sensitivity between genotypes. Likewise, it has been shown that progesterone may have anti-arrhythmic effects, which may help to reduce the risk of arrhythmias (31). These effects of progesterone on the QT interval are related to an increase in intracellular calcium reuptake from the sarcoplasmic reticulum mediated by SERCA2a (32). Similar to progesterone, testosterone decreases I_{CaL} current and increases potassium channel currents (I $_{\rm Kr}$, I $_{\rm Ks}$, I $_{\rm K1}$), reducing the QTc interval in animals and humans (33-35).

During pregnancy, the combination of hormones is complex. Apart from the interaction of estrogen and progesterone, oxytocin released towards the end of pregnancy, during labor and lactation, would have direct effects on sodium channels and cause a direct acute inhibition on the potassium I_{Ks} channel (36), which could explain the increased risk in post-partum women with LQTS2. Data on the potential impact of prolactin on cardiac electrophysiology are limited, but some animal studies suggest that it would act similarly to oxytocin (36).

6. Conclusions

LQTS is a rare heterogenous group of arrhythmogenic entities, characterized by a prolonged QT interval. In recent years, a

continuous improvement in diagnosis as well as genetic/ pathophysiological mechanism has been performed, however, risk stratification remains a current challenge. It is widely accepted that sex is an independent risk factor due to females having a high risk of malignant arrhythmias associated with LQTS. Hormones seem to be the main reason for reported sex differences, despite the fact that the link between the sex hormones and susceptibility to malignant arrhythmias is still a matter of debate. Additional studies will help to unravel the pathophysiological mechanism involved in sex differences, helping to define a proper risk stratification in LQTS patients.

Author contributions

GS-B, OC, and EA developed the concept. ND-E, EM-B, EA, PC, SC, JC, FC, VF, CH, and GS-B acquired, pre-processed, and analyzed the data. ND-E, EA, OC, and GS-B prepared the manuscript. EA, OC, and GS-B supervised the study. All authors contributed to the article and approved the submitted version.

Funding

This work was supported by Obra Social "La Caixa Foundation" (LCF/PR/GN19/50320002), Instituto de Salud

References

1. Wilde AAM, Amin AS, Postema PG. Diagnosis, management and therapeutic strategies for congenital long QT syndrome. *Heart*. (2022) 108:332-8. doi: 10.1136/ heartjnl-2020-318259

2. Costa S, Saguner AM, Gasperetti A, Akdis D, Brunckhorst C, Duru F. The link between sex hormones and susceptibility to cardiac arrhythmias: from molecular basis to clinical implications. *Front Cardiovasc Med.* (2021) 8:644279. doi: 10.3389/ fcvm.2021.644279

3. Wilde AAM, Amin AS. Clinical Spectrum of SCN5A mutations: long QT syndrome. Brugada Syndrome, and Cardiomyopathy. *JACC Clin Electrophysiol.* (2018) 4:569–79. doi: 10.1016/j.jacep.2018.03.006

4. Lankaputhra M, Voskoboinik A. Congenital long QT syndrome: a clinician's Guide. Intern Med J. (2021) 51:1999–2011. doi: 10.1111/imj.15437

5. Vink AS, Neumann B, Lieve KVV, Sinner MF, Hofman N, El Kadi S, et al. Determination and interpretation of the QT interval. *Circulation*. (2018) 138:2345–58. doi: 10.1161/CIRCULATIONAHA.118.033943

6. Wilde AAM, Semsarian C, Marquez MF, Sepehri Shamloo A, Ackerman MJ, Ashley EA, et al. European Heart rhythm association (EHRA)/heart rhythm society (HRS)/Asia pacific heart rhythm society (APHRS)/latin American heart rhythm society (LAHRS) expert consensus statement on the state of genetic testing for cardiac diseases. *Heart Rhythm.* (2022) 19(7):e1–60. doi: 10.1016/j.hrthm.2022.03.1225

7. Salama G, Bett GC. Sex differences in the mechanisms underlying long QT syndrome. *Am J Physiol Heart Circ Physiol.* (2014) 307:H640-8. doi: 10.1152/ajpheart.00864.2013

8. Mazzanti A, Trancuccio A, Kukavica D, Pagan E, Wang M, Mohsin M, et al. Independent validation and clinical implications of the risk prediction model for long QT syndrome (1-2-3-LQTS-risk). *Europace*. (2022) 24:614–9. doi: 10.1093/ europace/euab238

9. Li Z, Guo X, Guo L, Zheng L, Yu S, Yang H, et al. Sex differences in association between decreased glomerular filtration rate and prolongation of corrected QT interval in general Chinese population. *Eur J Intern Med.* (2017) 43:e33–5. doi: 10.1016/j.ejim. 2017.05.011

10. Adler A, Novelli V, Amin AS, Abiusi E, Care M, Nannenberg EA, et al. An international, multicentered, evidence-based reappraisal of genes reported to cause

Carlos III (FIS PI21/00094), and Fundació La Marató TV3 (245/ U/2020). CIBERCV is an initiative of the ISCIII, Spanish Ministry of Economy and Competitiveness. Funders had no role in study design, data collection, data analysis, interpretation, or writing of the report.

Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

congenital long QT syndrome. Circulation. (2020) 141:418-28. doi: 10.1161/ CIRCULATIONAHA.119.043132

11. Richards S, Aziz N, Bale S, Bick D, Das S, Gastier-Foster J, et al. Standards and guidelines for the interpretation of sequence variants: a joint consensus recommendation of the American college of medical genetics and genomics and the association for molecular pathology. *Genet Med.* (2015) 17:405–24. doi: 10.1038/gim.2015.30

12. Etheridge SP, Asaki SY, Niu MC. A personalized approach to long QT syndrome. Curr Opin Cardiol. (2019) 34:46–56. doi: 10.1097/HCO.000000000000587

13. Goldenberg I, Bos JM, Yoruk A, Chen AY, Lopes C, Huang DT, et al. Risk prediction in women with congenital long QT syndrome. *J Am Heart Assoc.* (2021) 10:e021088. doi: 10.1161/JAHA.121.021088

14. Asatryan B, Yee L, Ben-Haim Y, Dobner S, Servatius H, Roten L, et al. Sex-Related differences in cardiac channelopathies: implications for clinical practice. *Circulation*. (2021) 143:739–52. doi: 10.1161/CIRCULATIONAHA.120.048250

15. Seth R, Moss AJ, McNitt S, Zareba W, Andrews ML, Qi M, et al. Long QT syndrome and pregnancy. *J Am Coll Cardiol.* (2007) 49:1092–8. doi: 10.1016/j.jacc. 2006.09.054

16. Ishibashi K, Aiba T, Kamiya C, Miyazaki A, Sakaguchi H, Wada M, et al. Arrhythmia risk and beta-blocker therapy in pregnant women with long QT syndrome. *Heart.* (2017) 103:1374–9. doi: 10.1136/heartjnl-2016-310617

17. Regitz-Zagrosek V, Roos-Hesselink JW, Bauersachs J, Blomstrom-Lundqvist C, Cifkova R, De Bonis M, et al. 2018 ESC guidelines for the management of cardiovascular diseases during pregnancy. *Eur Heart J.* (2018) 39:3165–241. doi: 10. 1093/eurheartj/ehy340

18. Goldenberg I, Younis A, Huang DT, Yoruk A, Rosero SZ, Cutter K, et al. Use of oral contraceptives in women with congenital long QT syndrome. *Heart Rhythm.* (2022) 19:41–8. doi: 10.1016/j.hrthm.2021.07.058

19. de Bruin R, van Dalen SL, Franx SJ, Simons SH, Flint RB, van den Bosch GE. Risk for neonatal hypoglycaemia and bradycardia after beta-blocker use during pregnancy or lactation: a systematic review and meta-analysis protocol. *BMJ open.* (2022) 12:e055292. doi: 10.1136/bmjopen-2021-055292

20. Marcinkeviciene A, Rinkuniene D, Puodziukynas A. Long QT syndrome management during and after pregnancy. *Medicina (Kaunas).* (2022) 58(11):1694. doi: 10.3390/medicina58111694

21. Buber J, Mathew J, Moss AJ, Hall WJ, Barsheshet A, McNitt S, et al. Risk of recurrent cardiac events after onset of menopause in women with congenital long-QT syndrome types 1 and 2. *Circulation.* (2011) 123:2784–91. doi: 10.1161/CIRCULATIONAHA.110.000620

22. Haseroth K, Seyffart K, Wehling M, Christ M. Effects of progestin-estrogen replacement therapy on QT-dispersion in postmenopausal women. *Int J Cardiol.* (2000) 75:161–5; discussion 165–6. doi: 10.1016/S0167-5273(00)00317-X

23. Kadish AH, Greenland P, Limacher MC, Frishman WH, Daugherty SA, Schwartz JB. Estrogen and progestin use and the QT interval in postmenopausal women. *Ann Noninvasive Electrocardiol*. (2004) 9:366–74. doi: 10.1111/j.1542-474X. 2004.94580.x

24. Schwartz JB, Volterrani M, Caminiti G, Marazzi G, Fini M, Rosano GM, et al. Effects of testosterone on the Q-T interval in older men and older women with chronic heart failure. *Int J Androl.* (2011) 34:e415–21. doi: 10.1111/j.1365-2605. 2011.01163.x

25. Odening KE, Choi BR, Liu GX, Hartmann K, Ziv O, Chaves L, et al. Estradiol promotes sudden cardiac death in transgenic long QT type 2 rabbits while progesterone is protective. *Heart Rhythm.* (2012) 9:823–32. doi: 10.1016/j.hrthm. 2012.01.009

26. Kurokawa J, Kodama M, Furukawa T, Clancy CE. Sex and gender aspects in antiarrhythmic therapy. *Handb Exp Pharmacol.* (2012):237–63. doi: 10.3389/fcvm. 2021.644279

27. Kundu P, Ciobotaru A, Foroughi S, Toro L, Stefani E, Eghbali M. Hormonal regulation of cardiac KCNE2 gene expression. *Mol Cell Endocrinol.* (2008) 292:50–62. doi: 10.1016/j.mce.2008.06.003

28. Long V, Fiset C. Contribution of estrogen to the pregnancy-induced increase in cardiac automaticity. J Mol Cell Cardiol. (2020) 147:27-34. doi: 10.1016/j.yjmcc.2020.08.005

29. Vink AS, Clur SB, Wilde AAM, Blom NA. Effect of age and gender on the QTcinterval in healthy individuals and patients with long-QT syndrome. *Trends Cardiovasc Med.* (2018) 28:64–75. doi: 10.1016/j.tcm.2017.07.012

30. Bjelic M, Zareba W, Peterson DR, Younis A, Aktas MK, Huang DT, et al. Sex hormones and repolarization dynamics during the menstrual cycle in women with congenital long QT syndrome. *Heart Rhythm.* (2022) 19:1532–40. doi: 10.1016/j. hrthm.2022.04.029

31. Asatryan B, Rieder M, Castiglione A, Odening KE. Arrhythmic risk during pregnancy and postpartum in patients with long QT syndrome. *Herzschrittmacherther Elektrophysiol.* (2021) 32:180–5. doi: 10.1007/s00399-021-00757-4

32. Moshal KS, Zhang Z, Roder K, Kim TY, Cooper L, Patedakis Litvinov B, et al. Progesterone modulates SERCA2a expression and function in rabbit cardiomyocytes. *Am J Physiol Cell Physiol.* (2014) 307:C1050-7. doi: 10.1152/ ajpcell.00127.2014

33. Montano LM, Calixto E, Figueroa A, Flores-Soto E, Carbajal V, Perusquia M. Relaxation of androgens on rat thoracic aorta: testosterone concentration dependent agonist/antagonist L-type Ca2 + channel activity, and 5beta-dihydrotestosterone restricted to L-type Ca2 + channel blockade. *Endocrinology.* (2008) 149:2517–26. doi: 10.1210/en.2007-1288

34. Masuda K, Takanari H, Morishima M, Ma F, Wang Y, Takahashi N, et al. Testosterone-mediated upregulation of delayed rectifier potassium channel in cardiomyocytes causes abbreviation of QT intervals in rats. *J Physiol Sci.* (2018) 68:759–67. doi: 10.1007/s12576-017-0590-4

35. Gutierrez G, Wamboldt R, Baranchuk A. The impact of testosterone on the QT interval: a systematic review. *Curr Probl Cardiol.* (2022) 47:100882. doi: 10.1016/j. cpcardiol.2021.100882

36. Bodi I, Sorge J, Castiglione A, Glatz SM, Wuelfers EM, Franke G, et al. Postpartum hormones oxytocin and prolactin cause pro-arrhythmic prolongation of cardiac repolarization in long QT syndrome type 2. *Europace*. (2019) 21:1126–38. doi: 10.1093/europace/euz037