

Illuminating the fundamental structure of elementary Ca<sup>2+</sup> signals with a combination of realtime nanoscopy and CaCLEAN in living cardiac myocytes

Peter Lipp<sup>1</sup>, Marcel Lauterbach<sup>2</sup>, Qinghai Tian<sup>1</sup>

<sup>1</sup>Molecular Cell Biology, Center for Molecular Signaling (PZMS) & <sup>2</sup>Molecular Imaging, Center for Integrative Physiology and Molecular Medicine (CIPMM), Medical Faculty, Saarland University, Germany

E-Mail: [peter.lipp@uks.eu](mailto:peter.lipp@uks.eu), [marcel.lauterbach@uks.eu](mailto:marcel.lauterbach@uks.eu), [qinghai.tian@uks.eu](mailto:qinghai.tian@uks.eu)

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In cardiac myocytes, the cellular Ca<sup>2+</sup> transient determines the performance of the entire heart and is established by the recruitment of elementary Ca<sup>2+</sup> release signals, the so-called Ca<sup>2+</sup> sparks [1]. For more than 25 years sparks are believed to originate from clusters of the principal Ca<sup>2+</sup> release channel, the ryanodine receptor (RyR) on the sarcoplasmic reticulum, [2]. Surprisingly, super-resolution imaging of immunolabelled fixed cardiac myocytes indicated an average density of RyR clusters of  $\sim 2.2/\mu\text{m}^3$  [3] suggesting cluster distances well below the spread of an individual Ca<sup>2+</sup> spark (FWHM  $\sim 2.8 \mu\text{m}$ ). A novel instant SIM (iSIM, VisiTech Ltd, Sunderland, UK) microscope allows super resolution imaging ( $<130 \text{ nm}$  lateral and  $<270 \text{ nm}$  axial) at unprecedented acquisition rates ( $\sim 175 \text{ frames/s}$ , @1840x1024 pixels with a Hamamatsu ORCA Flash4 sCMOS camera). This realtime nanoscopy together with our computational methodology recently introduced [4] combining astronomical imaging processing with our knowledge of release and diffusion of Ca<sup>2+</sup> enabled us – for the first time – to identify fundamental nanoscopic Ca<sup>2+</sup> release bursts (CRB) that build up the microscopic Ca<sup>2+</sup> spark. On average, around 6 of such CRBs fire during a normal Ca<sup>2+</sup> spark. Physiological up or down regulation of the spark's amplitude by e.g. dissociation of the RyR accessory protein FKBP506 or phosphorylation of the RyR results in reduced ( $\sim 4.7$ ) and increased ( $\sim 9$ ) numbers of contributing CRBs, respectively. STED microscopy of fixed and immunolabelled ventricular myocytes clearly supported our functional findings in that microscopic (i.e. confocal) RyR clusters clearly comprised many (up to  $>10$ -15) RyR sub clusters. Contributing CRBs fire in succession and spatial proximity appears to determine the particular activation order. Registration of a longer time series of spontaneous Ca<sup>2+</sup> sparks and their associated CRBs reveals a well-ordered distribution of release sites reflecting the RyR distribution. In this study we combined innovative realtime nanoscopy with the novel algorithmic approach CaCLEAN to unravel the fine-structure of elementary Ca<sup>2+</sup> release signals, Ca<sup>2+</sup> sparks. We established – in contrast to the current view – that sparks indeed comprise a multitude of CRBs that build up the microscopic Ca<sup>2+</sup> transient. We believe that individual RyR clusters represent the morphological correlate to these CRBs. Physiological and pathological modulation of RyR results in a graded recruitment of CRBs to the Ca<sup>2+</sup> spark and results in its altered properties.

- [1] M. J. Berridge, P. Lipp, and M. D. Bootman, "The versatility and universality of calcium signalling," *Nat Rev Mol Cell Biol*, vol. 1, no. 1, pp. 11–21, Oct. 2000.
- [2] H. Cheng, W. J. Lederer, and M. B. Cannell, "Calcium sparks: elementary events underlying excitation-contraction coupling in heart muscle," *Science*, vol. 262, no. 5134, pp. 740–744, Oct. 1993.
- [3] Y. Hou, I. Jayasinghe, D. J. Crossman, D. Baddeley, and C. Soeller, "Nanoscale analysis of ryanodine receptor clusters in dyadic couplings of rat cardiac myocytes.," *Journal of molecular and cellular cardiology*, vol. 80, pp. 45–55, Mar. 2015.
- [4] Q. Tian, L. Kaestner, L. Schröder, J. Guo, and P. Lipp, "An adaptation of astronomical image processing enables characterization and functional 3D mapping of individual sites of excitation-contraction coupling in rat cardiac muscle.," *Elife*, vol. 6, p. 665, Nov. 2017.