

Supplementary Material

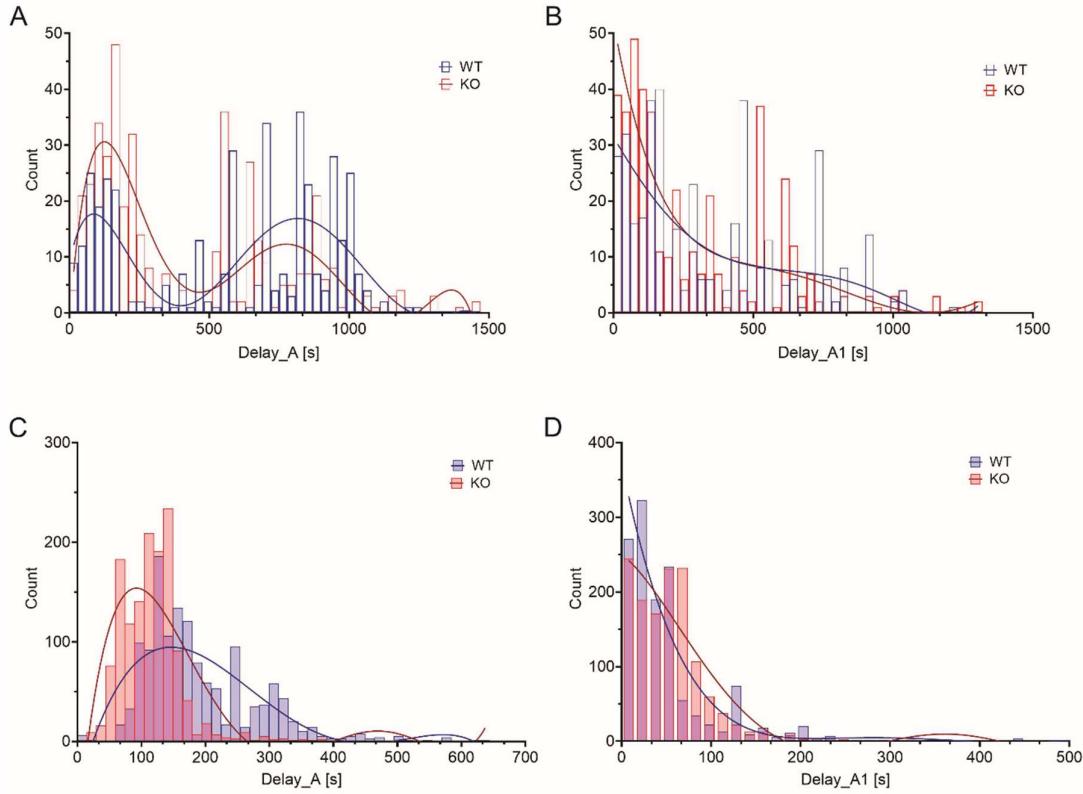


Figure S1: Distribution of activation delays after stimulation with glucose and forskolin in WT and KO mice. (A-B) Distribution of activation delays (Delay_A, panel A) and any-cell-first-responder delays (Delay_A1, panel B) after stimulation with 6 mM glucose + 10 μ M forskolin; (C-D) Distribution of activation delays (panel C) and any-cell-first-responder delays (panel D) after stimulation with 12 mM glucose.

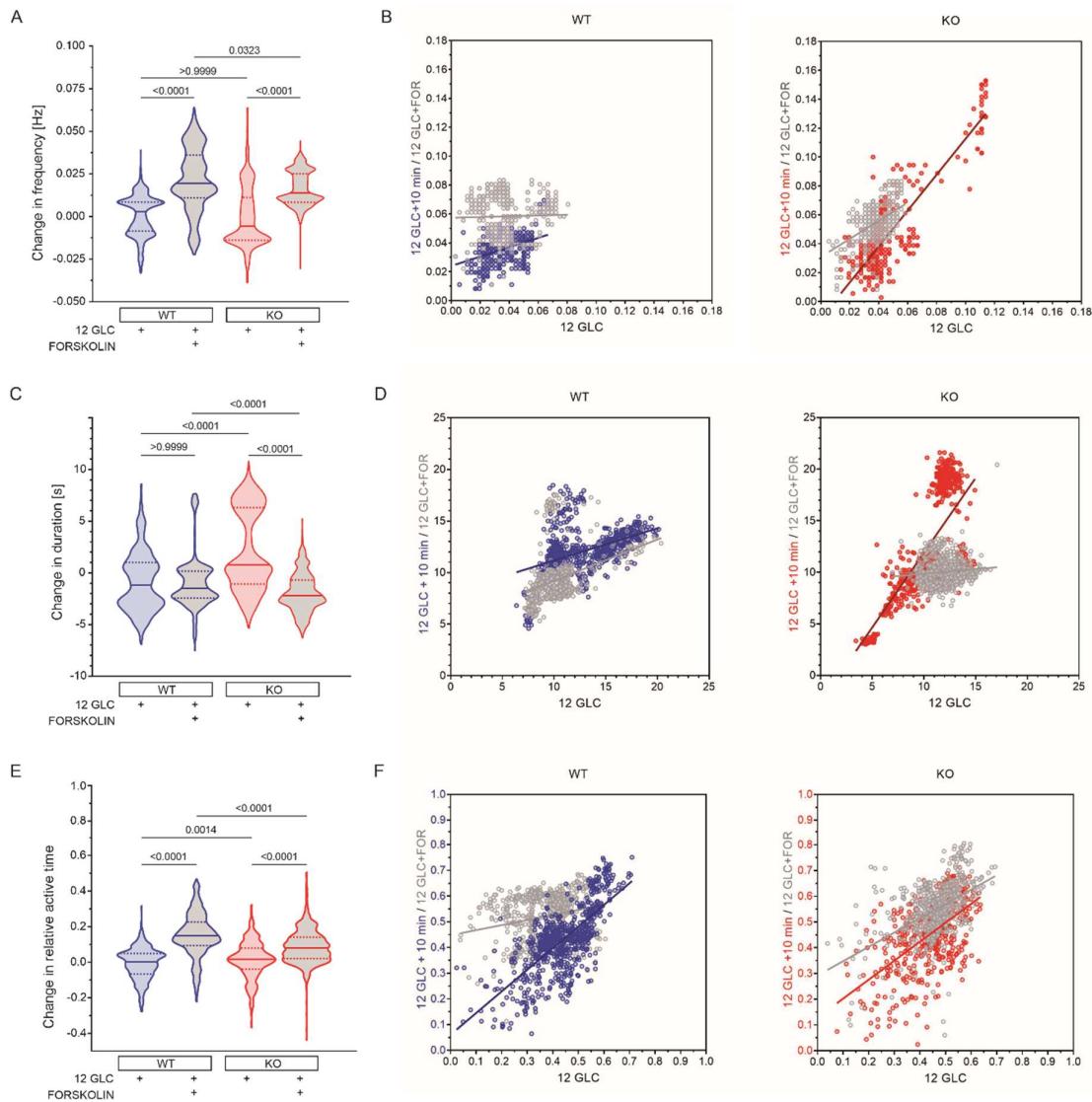


Figure S2: The effect of forskolin on frequency, duration and AT and the role of Epac2A. (A, C, E) The differences in (a) oscillation frequency, 1st quartile/median/3rd quartile (Q1/M/Q3) in Hz: -0.008/0.003/0.008 (12 mM glucose in WT), 0.011/0.020/0.036 (12 mM glucose + forskolin in WT), -0.014/0.006/0.011 (12 mM glucose in KO) and 0.08/0.014/0.025 (12 mM glucose + forskolin in KO); (C) duration Q1/M/Q3 in seconds: -3.1/-2.1 (12 mM glucose in WT), -2.4/-1.5/0.17 (12 mM glucose + 10 μ M forskolin in WT), -1.1/0.8/0.6.3 (12 mM glucose in KO) and -3.1/-2.2/-0.7 (12 mM glucose + 10 μ M forskolin in KO); and (E) relative active time Q1/M/Q3: -0.066/0.003/0.050 (12 mM glucose in WT), 0.094/0.150/0.225 (12 mM glucose + forskolin in WT), -0.039/0.017/0.078 (12 mM glucose in KO) and 0.020/0.080/0.141 (12 mM glucose + forskolin in KO) between the first and the second part of the plateau phase, separately for WT and KO mice and for both protocols; (B, D, F) Graphs represent the corresponding changes in frequency (B), duration (D) and relative active time (F) in individual cells in both WT and KO mice between the first and the second part of the plateau phase, separately for WT and KO mice and for both protocols. Data pooled from the following number of cells/islets: 687/11 (12 mM glucose in WT), 656/11 (12 mM glucose + 10 μ M forskolin in WT), 548/7 (12 mM glucose in KO), and 631/9 (12 mM glucose + 10 μ M forskolin in KO). Data were analyzed using one-way ANOVA on ranks (Kruskal-Wallis test) followed by Dunn's multiple comparisons test, p values are indicated on graphs.

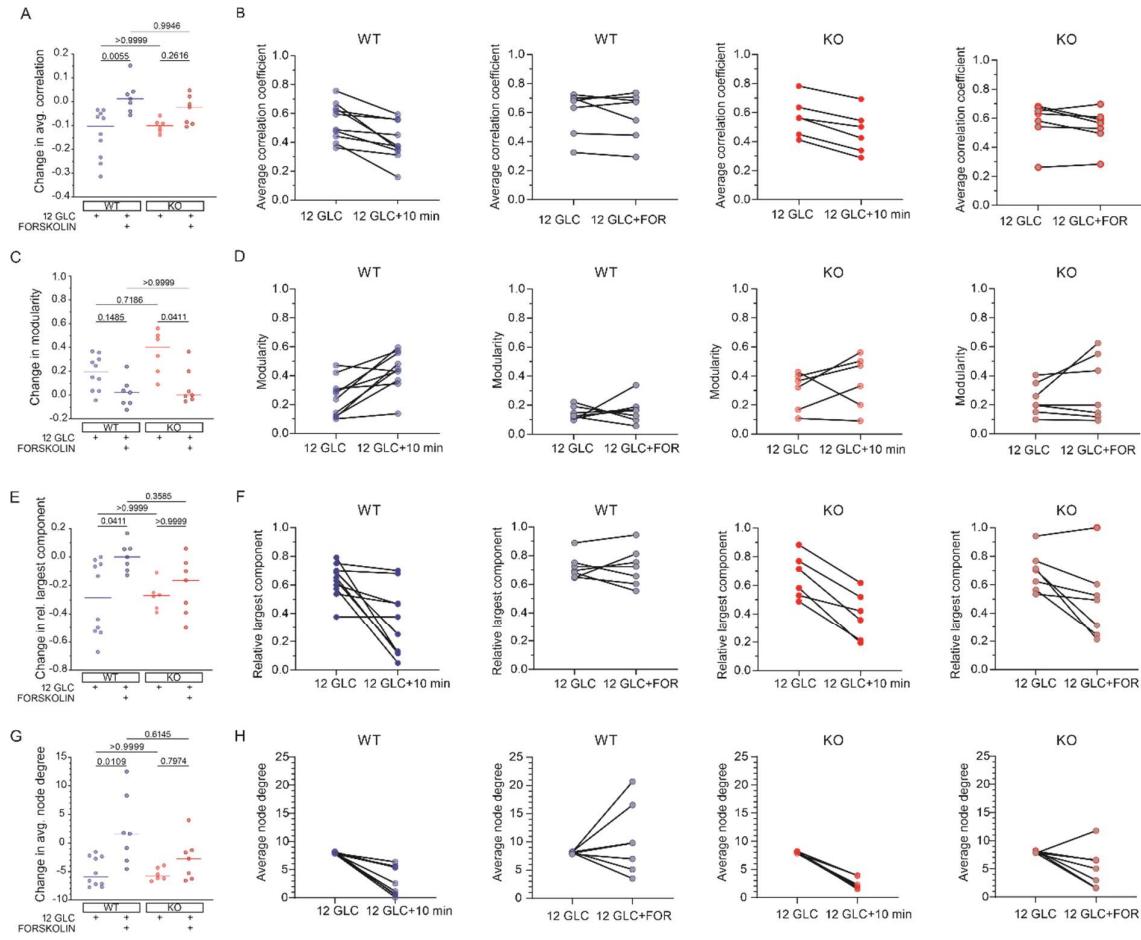


Figure S3: Changes in beta cell synchronicity and functional network parameters. Panels (A, C, E, G) denote differences in parameter values between the first and second part of the plateau phase, separately for both types of mice and for both protocols (stimulation with prolonged 12 mM glucose and stimulation with 12 mM glucose + 10 μ M forskolin in the second part of the stimulation). Individual dots represent the average values in individual islets, whereas the horizontal line denotes the median value. Panels (B, D, F, H) represent the corresponding changes in individual islets in different groups.

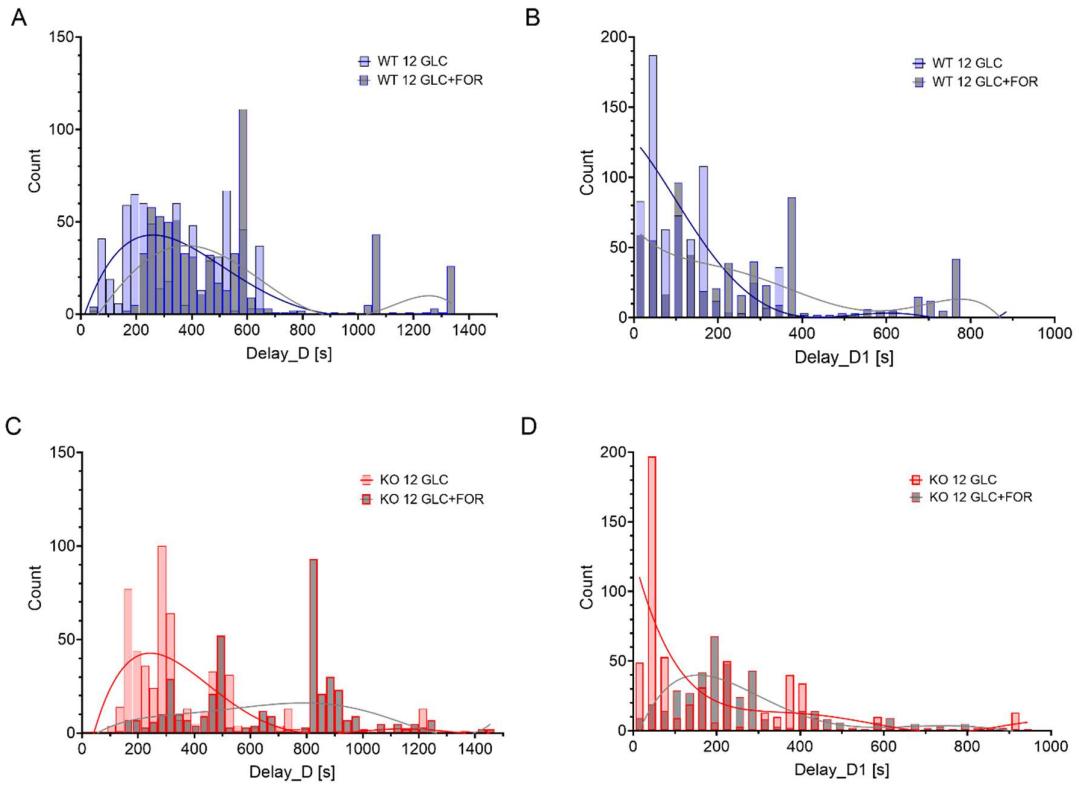


Figure S4: Distribution of deactivation delays after the end of stimulation with glucose and forskolin in WT and KO mice. (A) Distribution of deactivation delays after cessation of stimulation with 12 mM glucose (blue) and 12 mM glucose + 10 μ M forskolin (grey) in WT; (B) Distribution of the deactivation delays of cells after 1st responded cell after cessation of stimulation with 12 mM glucose (blue) and 12 mM glucose + 10 μ M forskolin (grey) in WT; (C) Distribution of deactivation delays after cessation of stimulation with 12 mM glucose (red) and 12 mM glucose + 10 μ M forskolin (grey) in KO; (D) Distribution of the deactivation delays of cells after 1st responded cell after cessation of stimulation with 12 mM glucose (red) and 12 mM glucose + 10 μ M forskolin (grey) in KO.