Supplementary figure legend. A more detailed description of figure 2 in the main text

In figure 2 a single neuronal compartment was simulated with a buffer composition as expected in CA1 pyramidal cells. In this compartment with 100 μ M CaM and 30 μ M CB the [Ca²⁺] was very rapidly (τ =10 μ s) increased by 50 μ M (Δ [Ca²⁺]_{total}). At 20 μ s (the maximum time resolution in the figure is 20 μ s) Δ [Ca²⁺]_{free} peaks at 8.8 μ M. As the total [Ca²⁺] increases, the bulk of the Δ [Ca²⁺]_{total} is rapidly bound to the N-terminus of CaM. At ~40 μ s, 36.4 μ M Ca²⁺ (73% of Δ [Ca²⁺]_{total}) is bound to the N-terminus (Δ [Ca²⁺]_N), while at that moment 5.2 μ M (10% of Δ [Ca²⁺]_{total}) and 1.7 μ M (3.3% of Δ [Ca²⁺]_{total}) are bound to the C-terminus (Δ [Ca²⁺]_C) and CB (Δ [Ca²⁺]_{CB}) respectively. After this, the Δ [Ca²⁺]_N drops with two τ 's of 0.38 ms (64%) and 2.1 ms (36%) to 0.54 μ M (1.1% of Δ [Ca²⁺]_{total}) at 30 ms. Around 0.9 ms the Δ [Ca²⁺]_C reaches its maximum of 21.4 μ M (43% of Δ [Ca²⁺]_{total}) while at that moment there is 11.0 μ M Δ [Ca²⁺]_N (22% C and 14.9 μ M Δ [Ca²⁺]_{CB} (30% of (Δ [Ca²⁺]_{total}). After this peak the Δ [Ca²⁺]_C drops with a τ of 3.7 ms to 9.8 μ M (19.7% of Δ [Ca²⁺]_{total}) at 30 ms. Over the whole experimental period, the [Ca²⁺]_{CB} steadily increases with τ 's of 0.63 ms (27%) and 3.1 ms (73%) to 39.3 μ M (78.6% of Δ [Ca²⁺]_{total}) at 30 ms. With 99.3% Δ [Ca²⁺]_{total} bound to either the N-terminus, C-terminus or CB 0.35 μ M of Δ [Ca²⁺]_{total} (i.e., 0.7%) remains unbound after 30 ms.

b) To better understand how Ca^{2+} moves through the system composed of 3 buffers, the amount of Ca^{2+} flowing between the four states, free ($\Delta[Ca^{2+}]_{\text{free}}$), bound to the N-terminus ($\Delta[Ca^{2+}]_N$), bound to the C-terminus ($\Delta[Ca^{2+}]_C$) and bound to CB (Δ [Ca²⁺]_{CB}) were calculated for 3 epochs. The concentrations of Ca²⁺ in the different states is represented by the area covered by the different circles where the grey area represents the concentration at the beginning of the epoch, while the colored circles represent the concentration at the end of the epoch. The numbers in the figure indicate percent of $\Delta [Ca^{2+}]_{total}$. During the first epoch (red), which runs from the start of the simulation (t=0 s) to when $[Ca^{2+}]_N$ peaks (t=40 µs) 86.6% of the 50 µM $\Delta [Ca^{2+}]_{total}$ (top grey circle) is directly bound by the 3 buffers (78.7% $[Ca^{2+}]_N$, 6.0% $[Ca^{2+}]_C$ and 1.9% $[Ca^{2+}]_{CB}$). Small amounts of buffered Ca^{2+} are already redistributed from $[Ca^{2+}]_N$ to $[Ca^{2+}]_N$ (4.4%) and $[Ca^{2+}]_{CB}$ (1.5%). During the second epoch (green), which runs from 40 μ s to when $[Ca^{2+}]_{C}$ peaks (t=900 μ s), and the last epoch (green, 0.9–30 ms) a further total of 11.7% of free Ca^{2+} is directly bound by the N-terminus whereas the amounts of free Ca^{2+} directly binding to either Cterminus (0.8%) or CB (0.2%) are less than 1% and for clarity are not shown. After the first 40 µs the main 'flow' of Ca^{2+} is the redistribution between the 3 buffering components. During the second epoch 33.0% moves from $[Ca^{2+}]_N$ to $[Ca^{2+}]_C$, 25.3% from $[Ca^{2+}]_N$ to $[Ca^{2+}]_{CB}$ and 1.0% from $[Ca^{2+}]_C$ to $[Ca^{2+}]_{CB}$ Remarkably, during the third epoch 19.3% moves back from $[Ca^{2+}]_C$ to $[Ca^{2+}]_N$ which is then directly redistributed to the C-terminus as part of the 44.4% that moves from $[Ca^{2+}]_N$ to $[Ca^{2+}]_{CB}$. Also a further 4.2% moves from $[Ca^{2+}]_C$ to $[Ca^{2+}]_{CB}$. All the net redistributions that take place during the 30 ms of the simulation are shown in the diagram with the black arrows where the black circles correlate with the end concentration and the white circles indicate the maxima reached during the experiment. Over the whole period more than 89% of the $\Delta [Ca^{2+}]_{total}$ is first bound by the fast

N-terminus which is later bound to the slower C-terminus and CB which eventually outcompete the N-terminus based on their higher affinity for Ca^{2+} .

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