

## Supplementary Information

### **TCR and CD28 concomitant stimulation elicits a distinctive calcium response in naive T cells**

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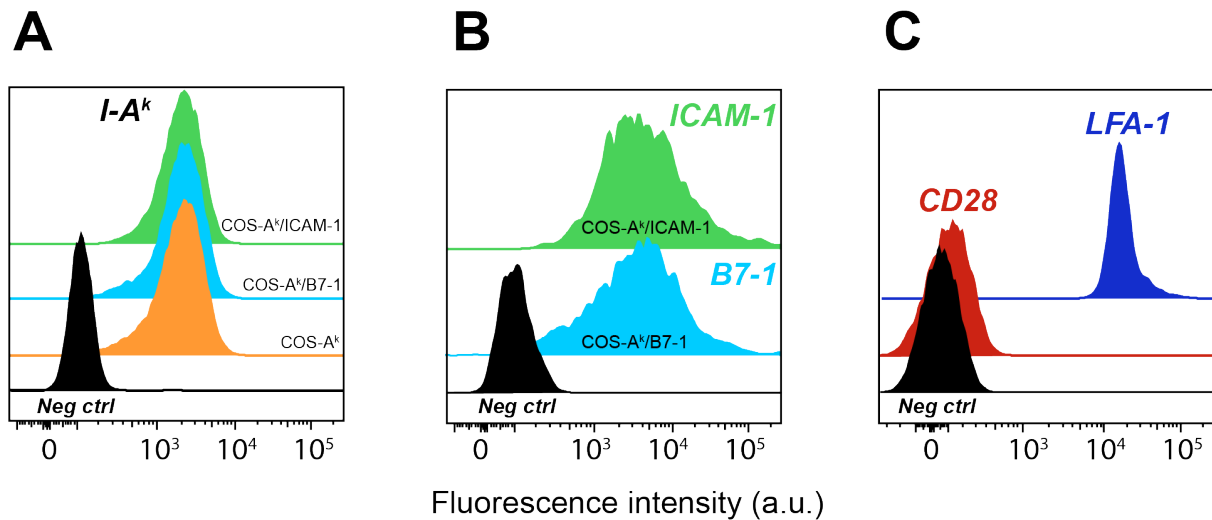
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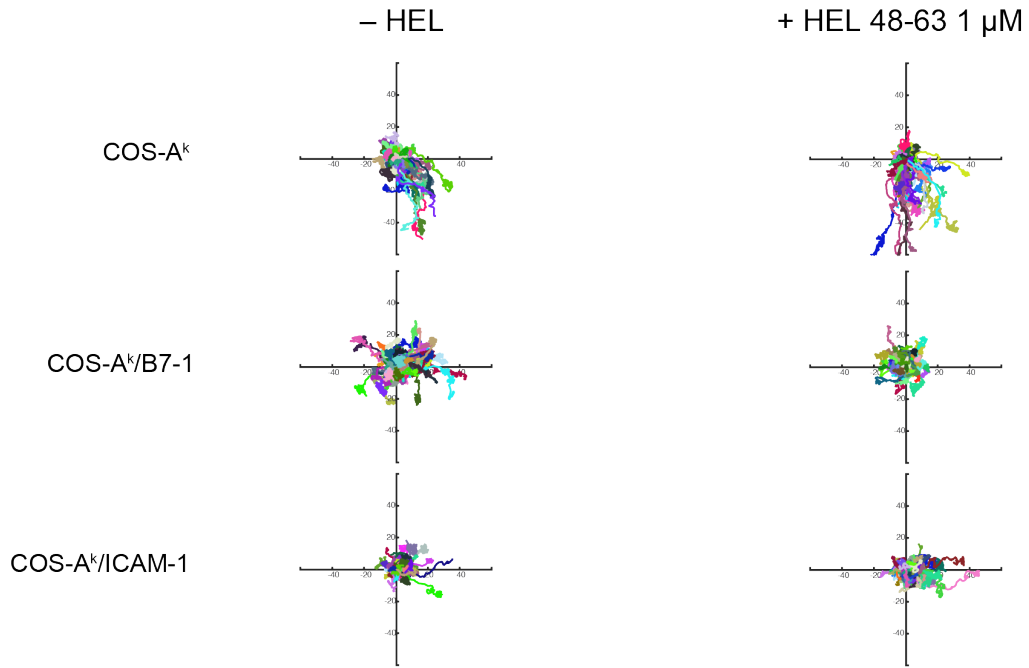
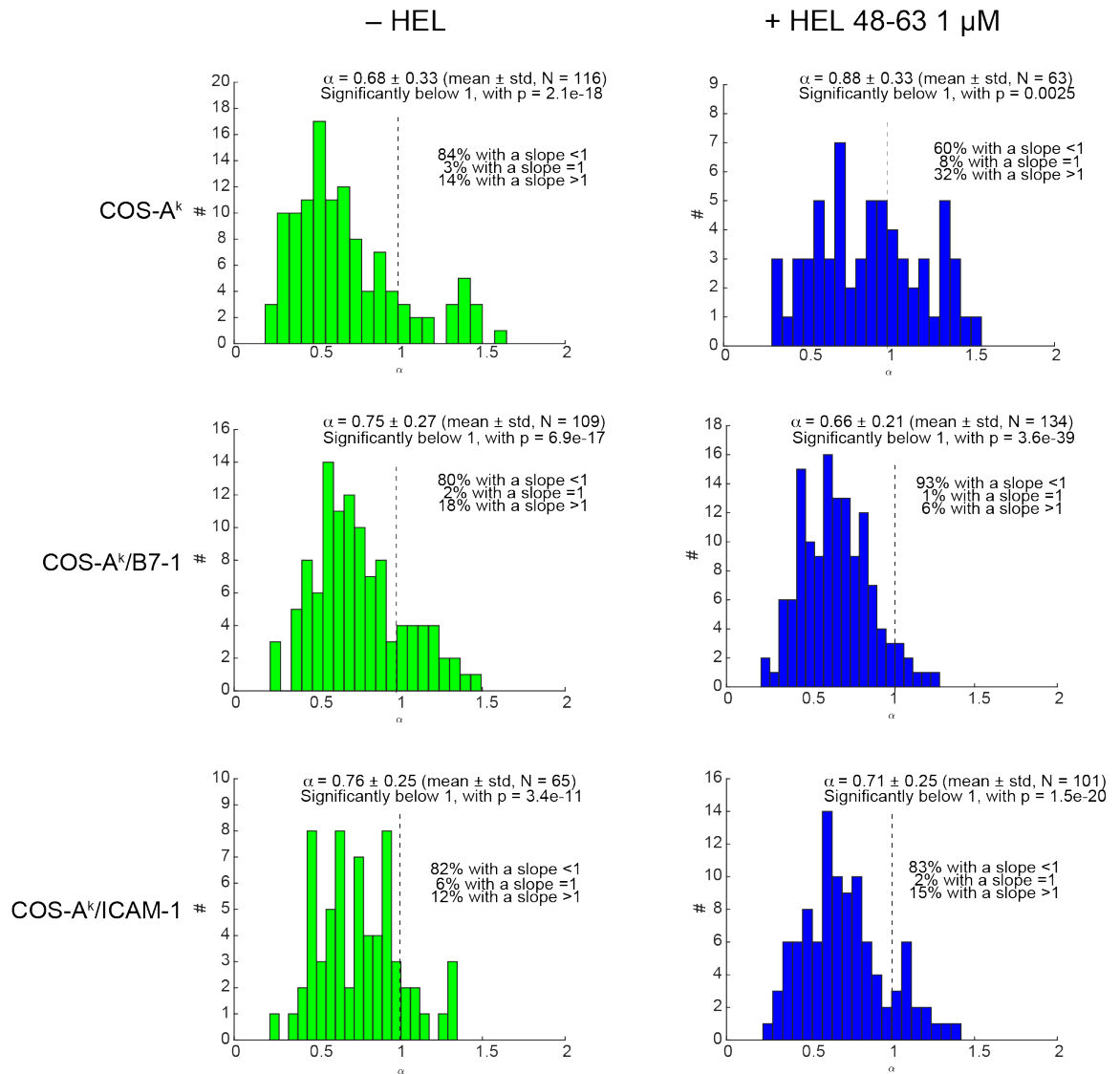
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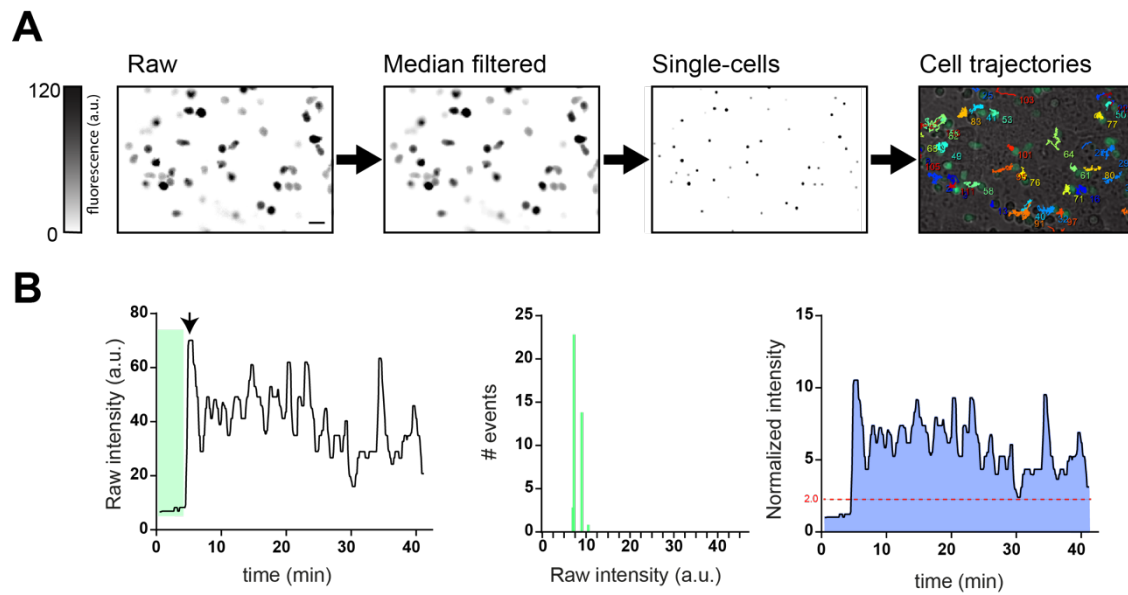
The Supplementary Information includes four supplementary figures (Supplementary Figures S1 – S4)



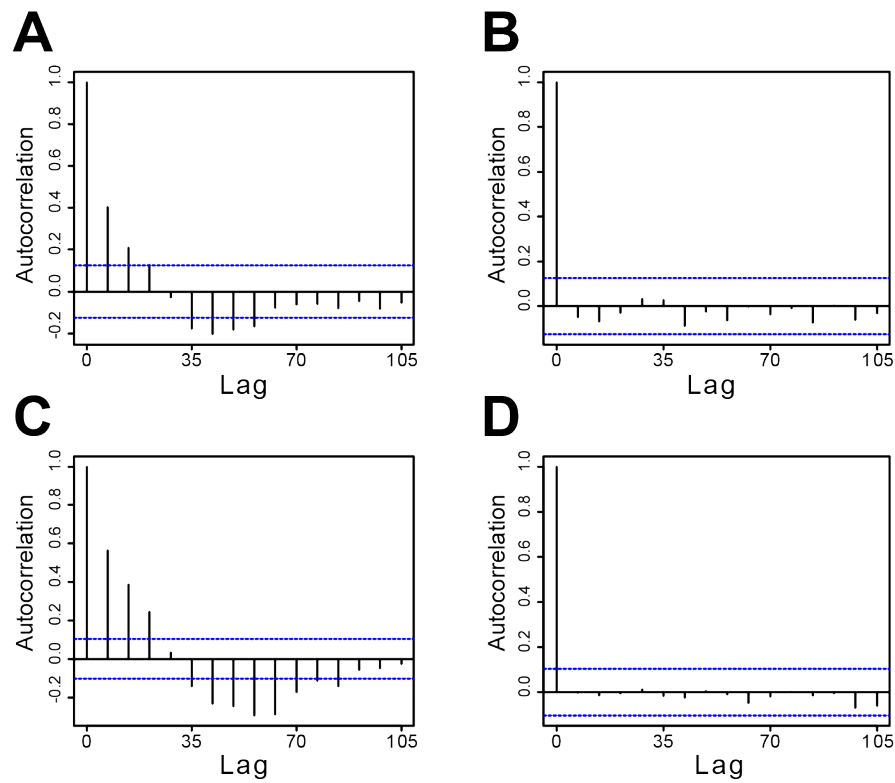
**Figure S1. Flow cytometry analysis of COS APCs and 3A9 TCRtg CD4<sup>+</sup> T cells. (A)** Expression of I-A<sup>k</sup> on COS-A<sup>k</sup>, COS-A<sup>k</sup>/B7-1, COS-A<sup>k</sup>/ICAM-1 cells, respectively. **(B)** Expression of B7-1 on COS-A<sup>k</sup>/B7-1 and ICAM-1 on COS-A<sup>k</sup>/ICAM-1 cells, respectively. **(C)** Expression of CD28 and LFA-1 on 3A9 TCRtg CD4<sup>+</sup> T cells. In these experiments, we used PE- (for I-A<sup>k</sup>, CD28 and LFA-1) and APC- (for ICAM-1 and B7-1) conjugated mAbs that all display the protein/dye ratio at about 1:1 according to the suppliers' communications. In **(B)** and **(C)** we used the labeling mAbs at concentrations close to the saturating binding conditions in order to better compare the relative expression levels of different surface antigens. All data are representatives of three independent experiments.

**A****B** $\alpha$  value distribution

**Figure S2. T cell movements on the COS APCs.** (A) Trajectories of naive CD4<sup>+</sup> T cells upon contacting with APCs loaded or not with HEL 48-63 peptides. Naive CD4<sup>+</sup> T cells from 3A9 TCRtg mice were loaded onto monolayers of COS-A<sup>k</sup>, COS-A<sup>k</sup>/B7-1, COS-A<sup>k</sup>/ICAM-1 APCs grown to confluency on Lab-Tek chamber slides, loaded or not with 1 μM HEL 48-63. T cell trajectories were recorded for 45 min at 37 °C using a confocal video-microscope. (B) Distribution of  $\alpha$  values from the MSD analyses.



**Figure S3. The mean amplitude (MA) and response fraction (RF) of the calcium response magnitude as determined with MAAACS. (A)** Automatic tracking of fluorescent cells. Experimental raw fluorescence images (color coded with a white to black palette) were transformed into Gaussian intensity peaks as described in Materials and Methods. Scale bar, 10  $\mu\text{m}$ . **(B)** MA and RF determination. The construction of the raw fluorescence intensity distribution histogram (green area in the left panel) prior to the maximal amplitude (arrow) allows to compute a median (6.69 in this example) that is defined as the individual baseline value to which raw intensities are normalized (right panel). The mean amplitude (MA) of fluorescence intensities corresponds to the average normalized intensity on the whole trace (blue area). Based on the analysis of the average MA of the cell population under stimulating and non-stimulating conditions, we could establish an activation threshold (red dotted line; in our study, this value was set 2, as described in Materials and Methods). The response fraction (RF) could then be determined as the ratio between the time when the intensity is above the threshold and the total time during which the intensity is detected. The experimental data were taken from the same experiment shown in Figure 2.



**Figure S4. The examples of autocorrelation function analysis of the  $[Ca^{2+}]_i$  time series data.** 3A9 TCRtg CD4<sup>+</sup> T cells stained with Fluo-4 PBX were loaded respectively onto COS-A<sup>k</sup> (A & B) and COS-A<sup>k</sup>/B7-1(C & D), both pulsed with 1  $\mu$ M HEL 48-63, and observed for 45 min at 37 °C. The autocorrelation function plots of  $[Ca^{2+}]_i$  time series data are shown in (A) and (C), and the cumulative periodograms of the residual obtained by using AR fitting are shown in (B) and (D).