

The Translational Machinery of Human CD4⁺ T Cells Is Poised for Activation and Controls the Switch from Quiescence to Metabolic Remodeling

Sara Ricciardi, Nicola Manfrini, Roberta Alfieri, Piera Calamita, Maria Cristina Crosti, Simone Gallo, Rolf Müller, Massimiliano Pagani, Sergio Abrignani, and Stefano Biffo*

*Correspondence: biffo@ingm.org

<https://doi.org/10.1016/j.cmet.2018.09.010>

(Cell Metabolism 28, 895–906.e1–e5; December 4, 2018)

In the originally published version of this manuscript, the size of Figures 4 and 6 was mistakenly reduced and enlarged, respectively, and the asterisks in Figure 5I were mistakenly removed. Furthermore, as a result of an author oversight, the name and surname of Simone Gallo appeared in the reversed order. These errors have now been corrected in the article online and in print. The authors apologize for the error and any inconvenience that may have resulted.

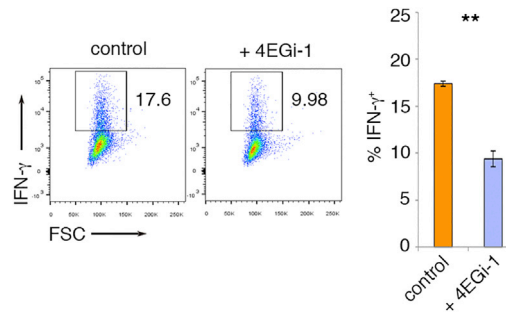


Figure 5I. Translational Activation of ACC1 via eIF4E Sustains a Metabolic Feedforward Loop that Completes the Metabolic Reprogramming to an Effector Phenotype (corrected)

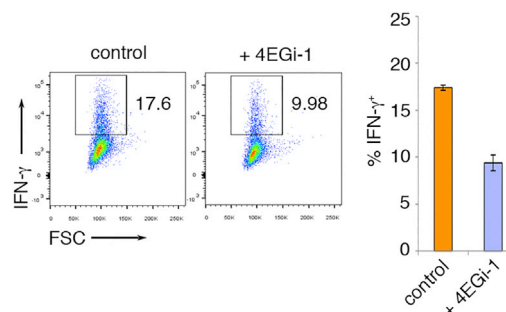


Figure 5I. Translational Activation of ACC1 via eIF4E Sustains a Metabolic Feedforward Loop that Completes the Metabolic Reprogramming to an Effector Phenotype (original)