

Office of Student Affairs and Admissions SGS at New Jersey Medical School Rutgers, The State University of New Jersey 185 South Orange Avenue, MSB C-696 Newark, NJ 07103 http://njms.rutgers.edu/gsbs/ p. 973-972-4511 f. 973-972-7148

YOU ARE INVITED TO ATTEND THE DEFENSE OF THE DOCTORAL DISSERTATION

"Long-chain Fatty Acid Sensing by GPR132 Regulates CD8⁺ T Cell Responses to Infection"

By

Giuseppina Marchesini Tovar Molecular Biology, Genetics, and Cancer Program B.Sc. Biotechnology, 2017, Universidad Politécnica de Madrid, Madrid, Spain

> Thesis Advisor: Tessa Bergsbaken, Ph.D. Assistant Professor Rutgers New Jersey Medical School Department of Pathology, Immunology and Laboratory Medicine

> > Friday, April 11th, 2025 11:00 A.M. Cancer Center, G1196

Join Zoom presentation

https://rutgers.zoom.us/j/92660071853?pwd=bPSDVpG6iylIgi98gawTRyQRqbhIX5.1&from=addon

Meeting ID: 926 6007 1853 Password: 589436

ABSTRACT

CD8⁺ T cells undergo robust expansion upon priming to control pathogen replication and provide host immunity; however, T cell function must be tightly regulated to ensure effective immunosurveillance without immunopathology. G protein coupled receptors sense environmental signals and are essential for modulating T cell function, and the lipid sensing receptor GPR132 is rapidly upregulated by T cells upon activation and maintained after infection is resolved. Ligands for this receptor include oxidized long-chain fatty acids derived from dietary linoleic acid, and these are increased under proinflammatory conditions. Additionally, mimics of these ligands are synthesized by commensal and pathogenic microorganisms. GPR132-deficient (KO) mice develop an autoimmune syndrome accompanied by expansion of the T cell compartment, suggesting GPR132 as a key target for T cell regulation. We examined the cell-intrinsic role of GPR132 in CD8⁺ T cell responses during infection and found that KO T cells displayed significantly enhanced expansion over wild-type (WT) cells. However, WT and KO T cells had similar proliferation and gene expression profiles upon TCRengagement in vitro. There were also no GPR132-dependent changes in T cell expansion observed during lymphopenia-driven proliferation in vivo, or when GPR132KO mice were aged in a SPF facility, indicating GPR132 inhibits T cell expansion specifically in the context of inflammation. An in-depth inquiry into the mechanism of action revealed that GPR132 limits T cell proliferation at peak of infection, while subsequently escalating cell death during contraction. Moreover, when T cells lacked GPR132 signaling, we observed an increase in the maintenance of CX3CR1⁺ effectors. Transcriptomic analysis of splenic WT and KO antigenspecific CD8⁺ T cells supported this finding, as GPR132 deficiency results in enlarged frequencies of cytotoxic effector cells at the expense of the memory compartment after infection. Additionally, formation of tissueresident memory pools was diminished. GPR132 also regulated effector function, as granzyme production was reduced while cytokine production was increased in KO T cells. Lastly, despite increased numbers of KO memory T cells, they displayed similar expansion potential as WT memory cells, yet retained enhanced proinflammatory cytokine production during rechallenge. Altogether, these studies reveal a role for GPR132 in detecting self and commensal lipids to regulate T cell responses against pathogens and indicate that GPR132 can be targeted to modulate T cell number and function without induction of autoimmunity.