



**Università Politecnica delle Marche  
Ancona  
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## **Applicazioni “interattomiche” per la comprensione delle malattie rare: il caso della Fibrosi Cistica**



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## **RARE DISEASES (facts)**

**There are nearly 8000 rare diseases (some even without a name), it turns out that these “rare” diseases affect more than 300 million people worldwide (about 5-6% of the world’s population). The economic cost of 379 rare diseases in the USA in 2019 was estimated to nearly 1 trillion USD .**

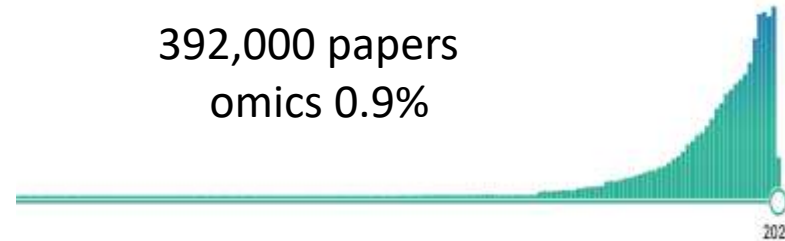
**94% of rare diseases lack an approved treatment**

**A large percentage of rare diseases are without diagnosis**

**Most of the known disease variants fall in coding regions of the genome, but much less is known about the role of noncoding region variants (90% of the eukaryotic genome) and structural variants in disease**



## RARE DISEASES (technologies)



**Transcriptomics, proteomics, phenomics, metabolomics, lipidomics and methylomics are already used, but epi-transcriptomics, metagenomics, fluxomics, glycomics and other omics are yet to be evaluated for the purposes of disease-gene prioritisation.**



## **RARE DISEASES (Challenges)**

**All the omics analyses rely on the systematic collection of well-organized large-scale datasets. Therefore, sharing patient-level data in public and controlled access repositories is highly relevant.**

**Despite these technological advancements, diagnosing rare diseases poses several challenges. Many clinicians struggle to interpret these novel findings, creating a demand for specialists who can translate results into clinical insights. Testing is often unavailable in low-resource medical settings, necessitating referrals to specialized centers. Omics, require sophisticated technologies, data repositories, bioinformatic pipelines, and age/tissue-matched controls, further complicating accessibility.**

## ***Cystic Fibrosis***

**Cystic fibrosis is the most frequent rare disease: around 4% of the population is a healthy carrier and around 200 new cases are recorded per year in Italy.**

# Cystic Fibrosis

- ▶ Autosomal recessive disease
- ▶ More than 2000 mutations in the gene coding for **CFTR**
- ▶ The deletion of Phenylalanine in position 508 ( $\Delta F$ ) present in 90% of patients worldwide



bronchopulmonary obstruction and opportunistic infections  
(pseudomonas aeruginosa, mycobacterium)

## [Tissue transglutaminase activation modulates inflammation in cystic fibrosis .](#)

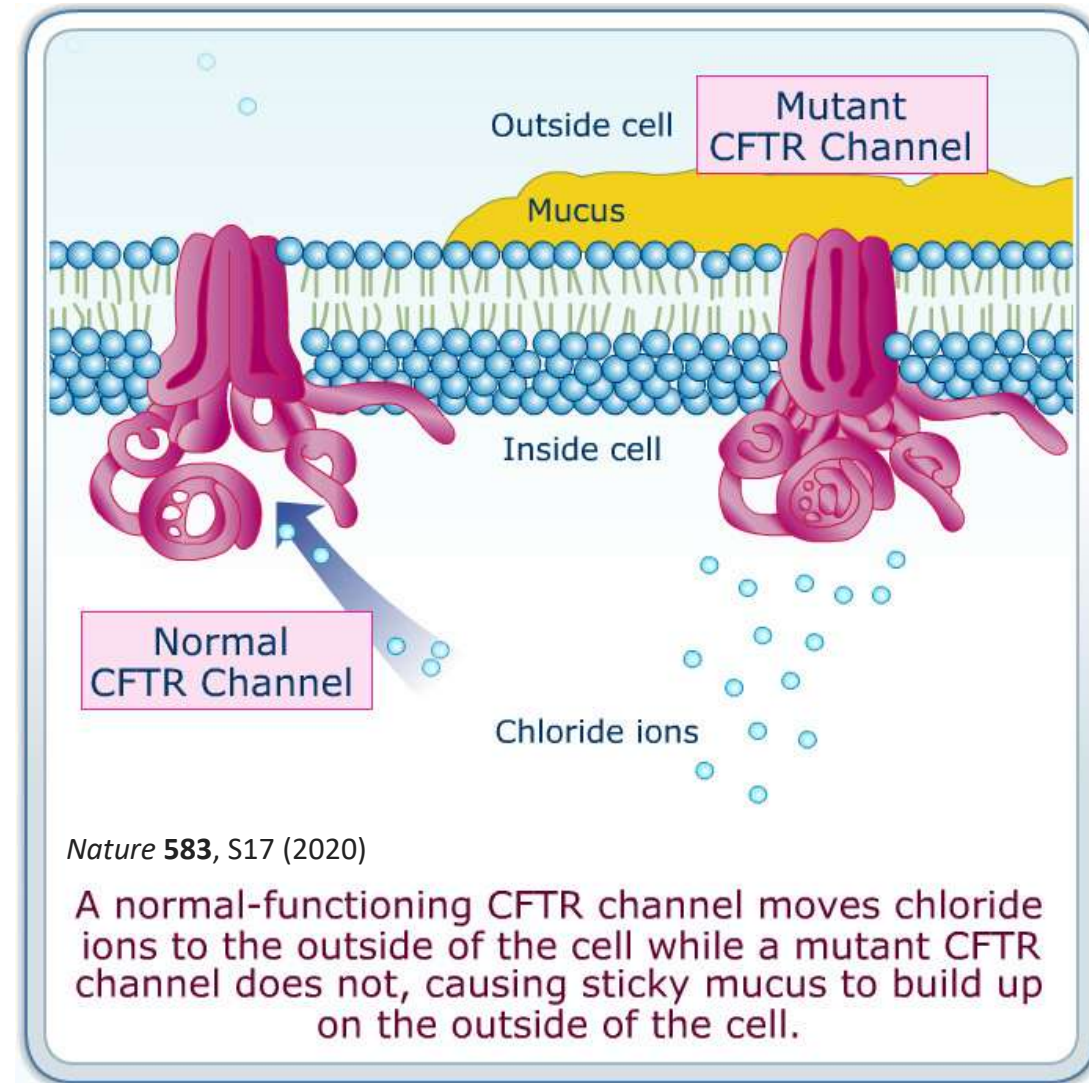
Maiuri L et al. *J Immunol.* 2008, 180(11):7697-705.

## [Transglutaminase 2 Regulates Innate Immunity by Modulating the STING/TBK1/IRF3 Axis.](#)

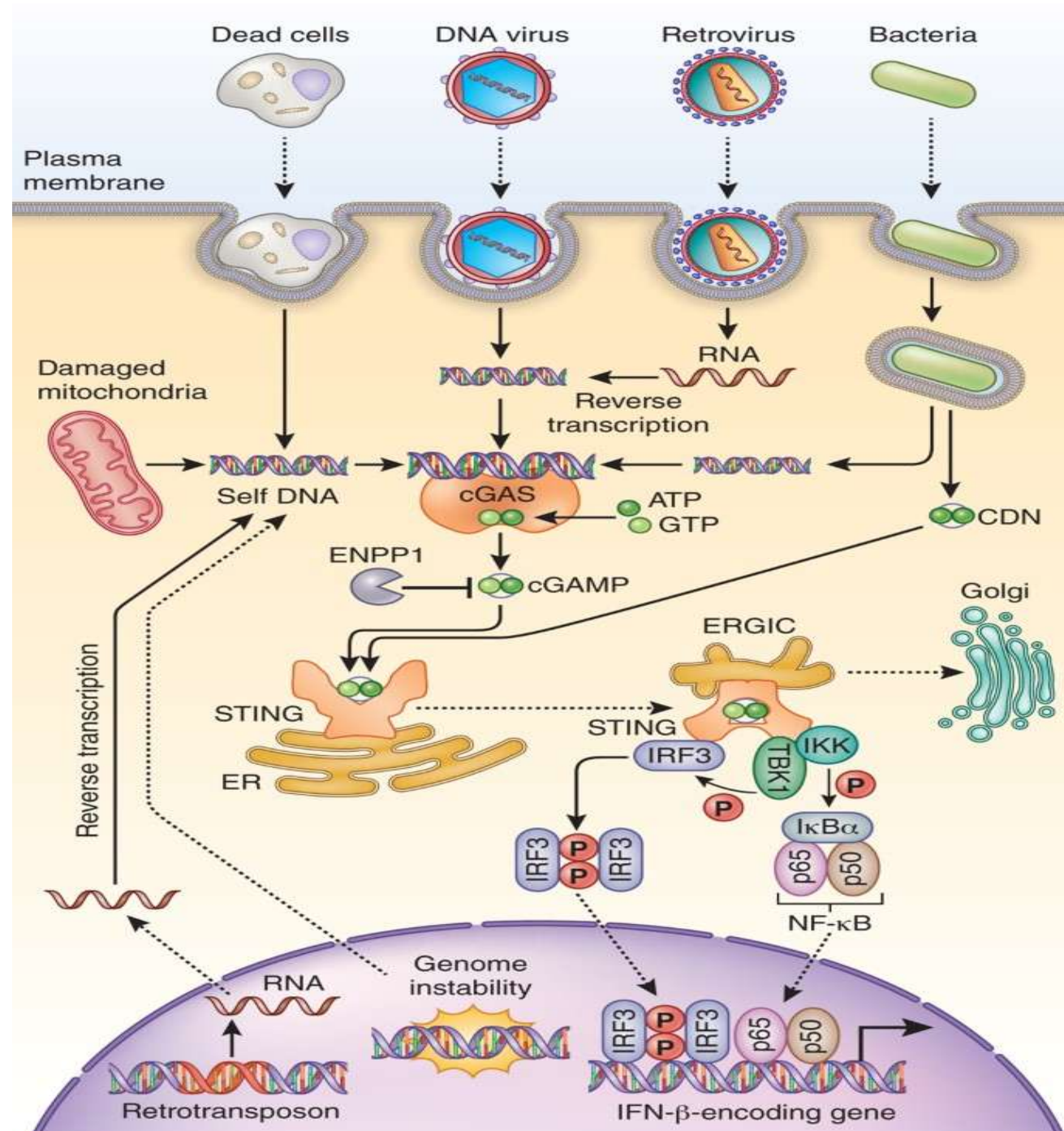
Occhigrossi L, Rossin F, D'Eletto M, Farrace MG, Ciccocanti F, Petrone L, Sacchi A, Nardacci R, Falasca L, Del Nonno F, Palucci I, Smirnov E, Barlev N, Agrati C, Goletti D, Delogu G,

Fimia GM, **Piacentini M.**

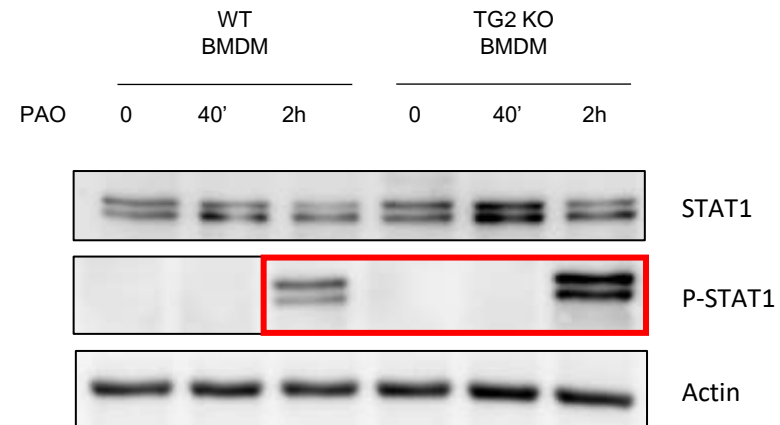
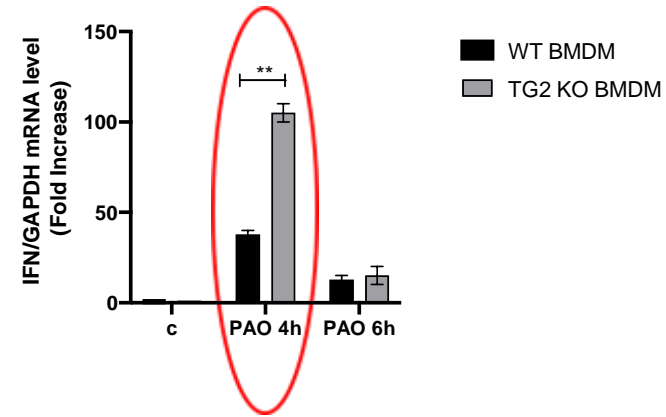
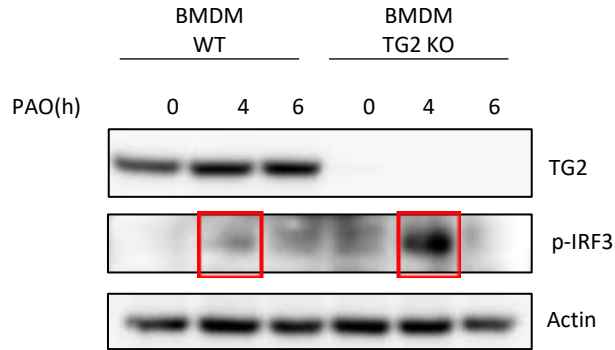
*J Immunol.* 206(10):2420-2429. 2021



# The cGAS/STING/TBK1/IRF3/IFN- $\beta$ innate immunity pathway



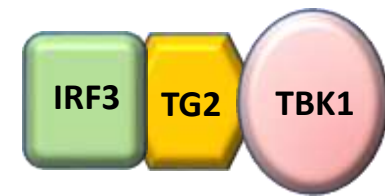
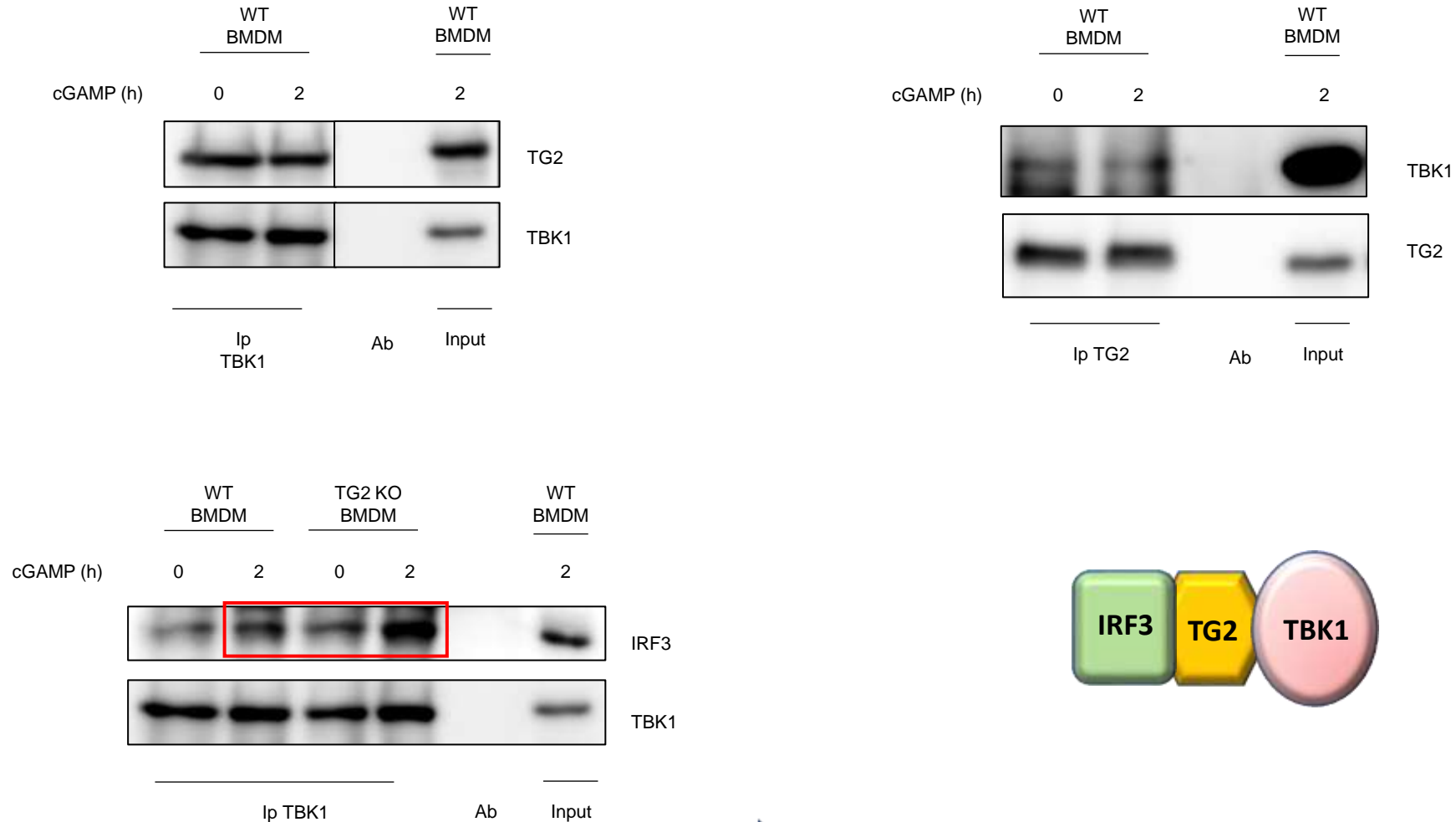
# The absence of Transglutaminase type 2 (TG2) in BMDMs infected with *Pseudomonas aeruginosa* (PAO) leads to increased levels of p-IRF3 and IFN $\beta$



**➔ the cGAS/STING pathway is upregulated in TG2 KO BMDMs**



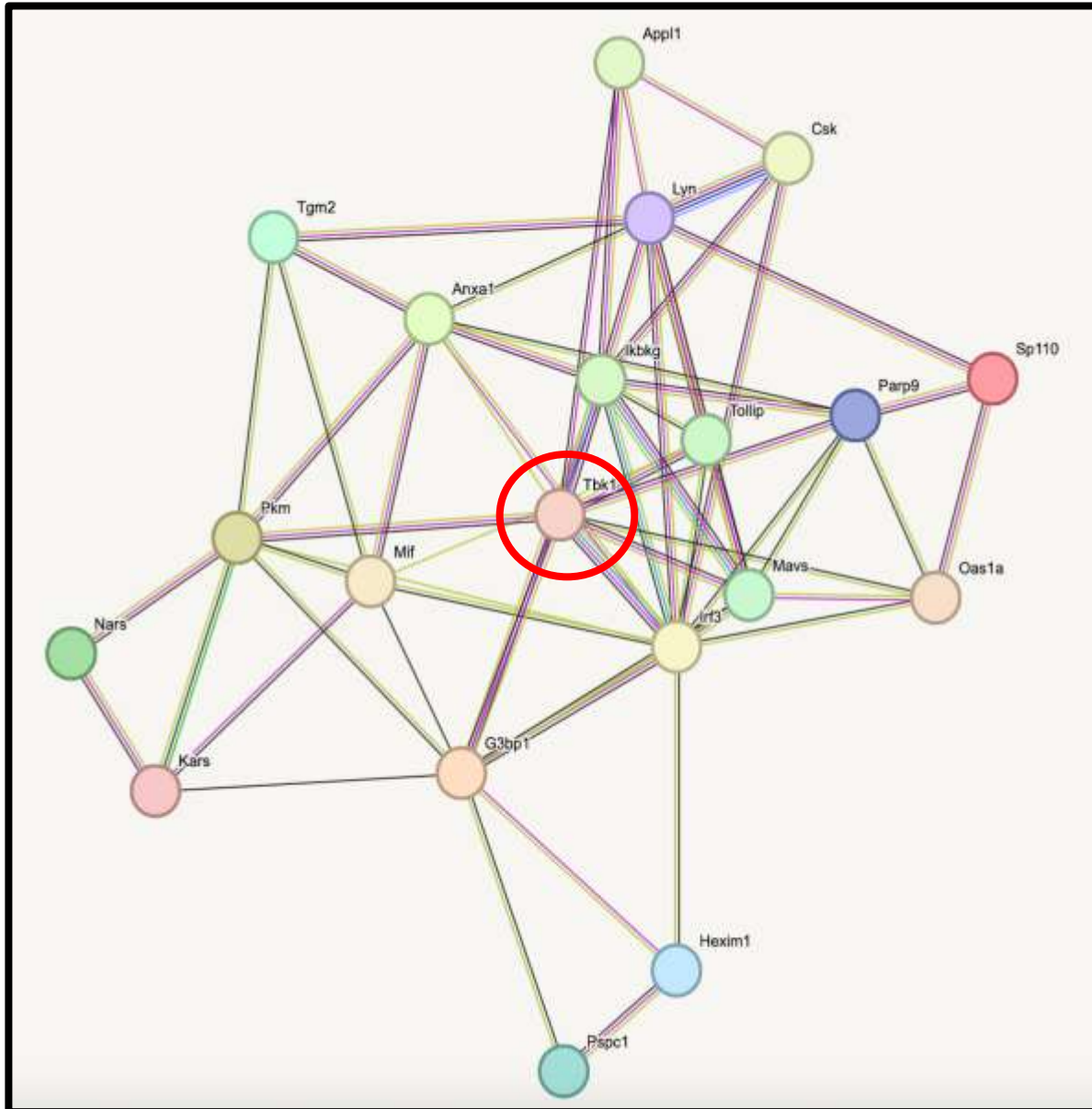
## TG2 interacts with TBK1 and this interaction limits the interaction between IRF3 and TBK1 in BMDMs



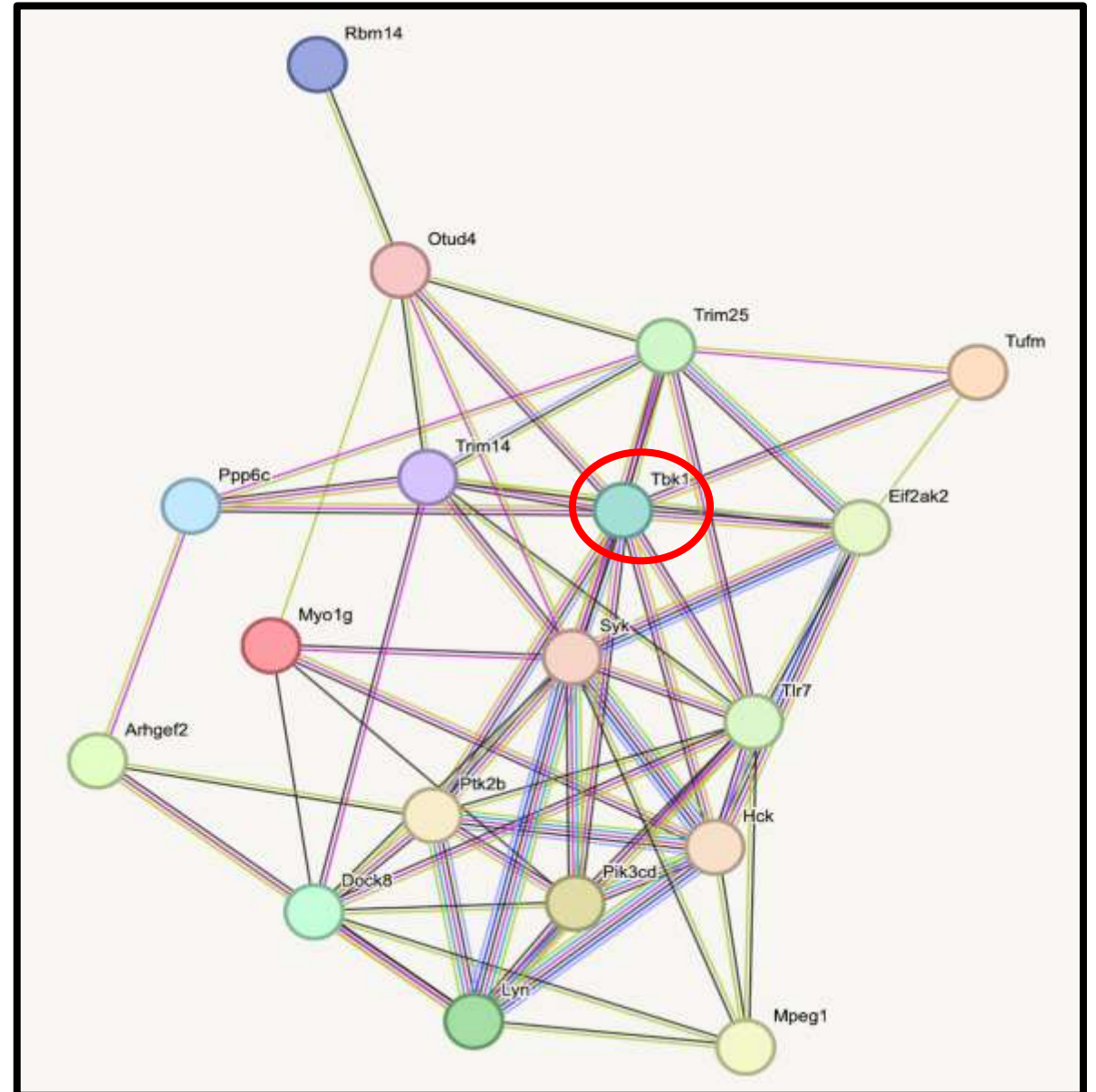
TG2 reduces the interaction of TBK1 with IRF3

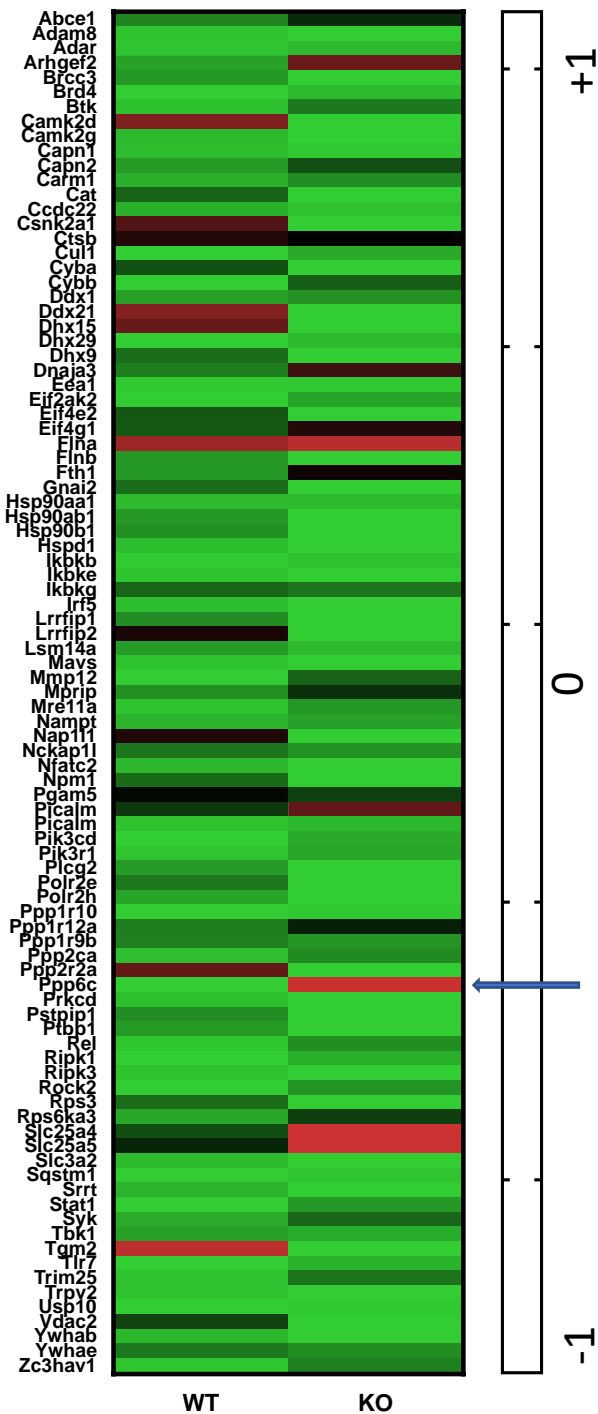
# TBK1 interactome

plus TG2

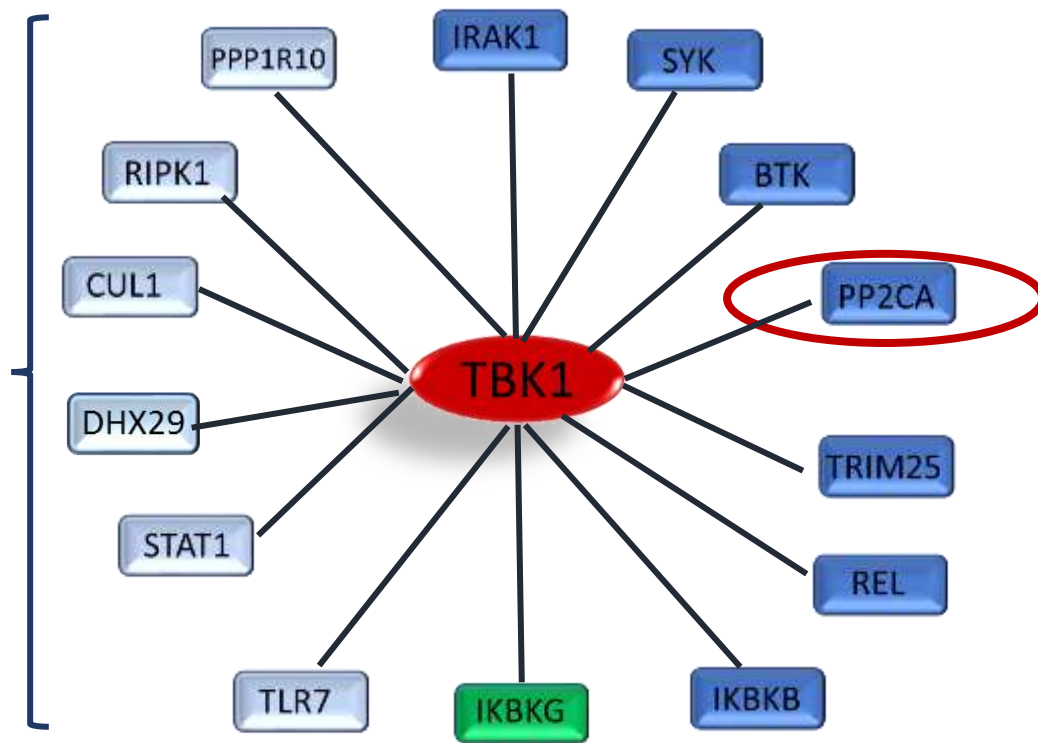


minus TG2





## The absence of TG2 influences the TBK1 interactome



Only in KO

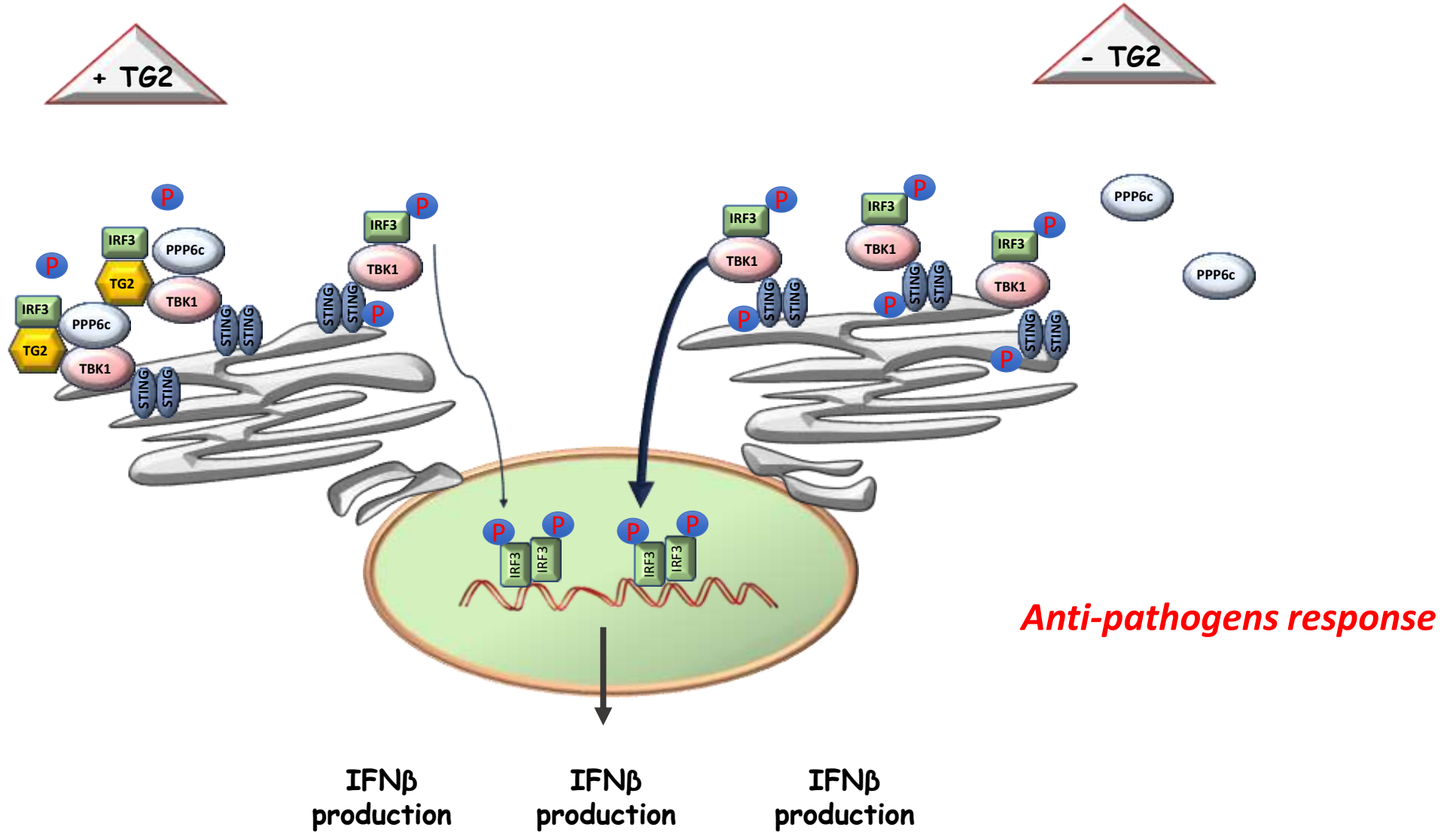


Particularly interesting is the drastic decrease of the phosphatase PPP6C that is the regulatory subunit of the PP2CA which is well known to dephosphorylate IRF3

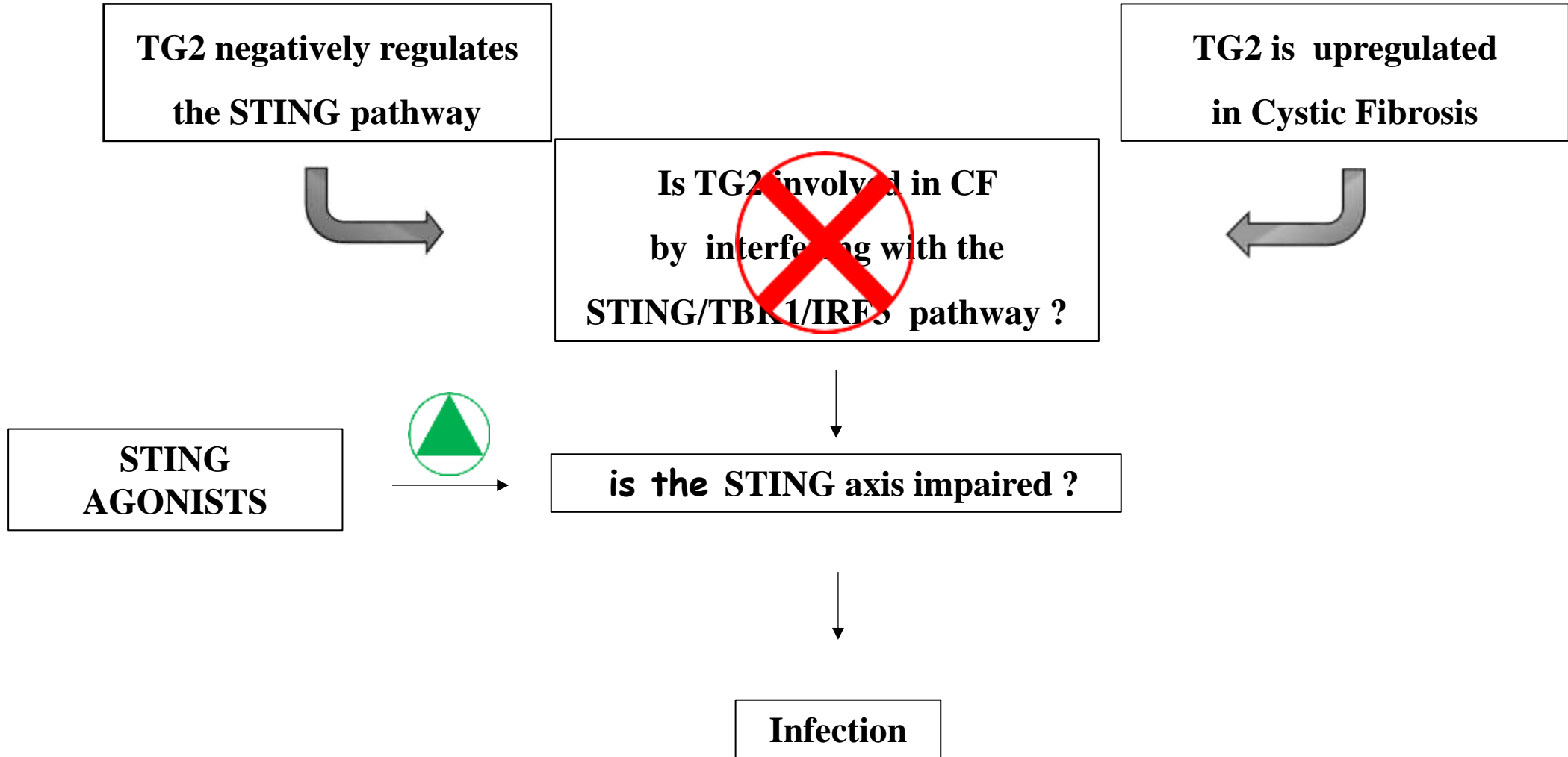
[PPP6C Negatively Regulates STING-Dependent Innate Immune Responses.](#)

Ni G, Ma Z, Wong JP, Zhang Z, Cousins E, Major MB, Damania B. *mBio*. 2020 Aug 4;11(4):e01728-20.

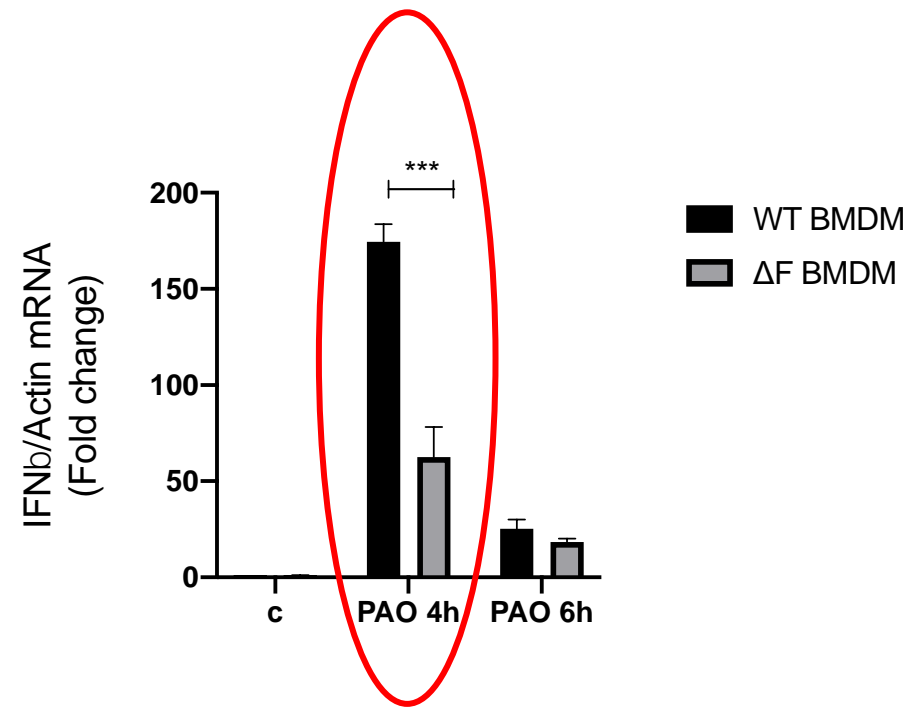
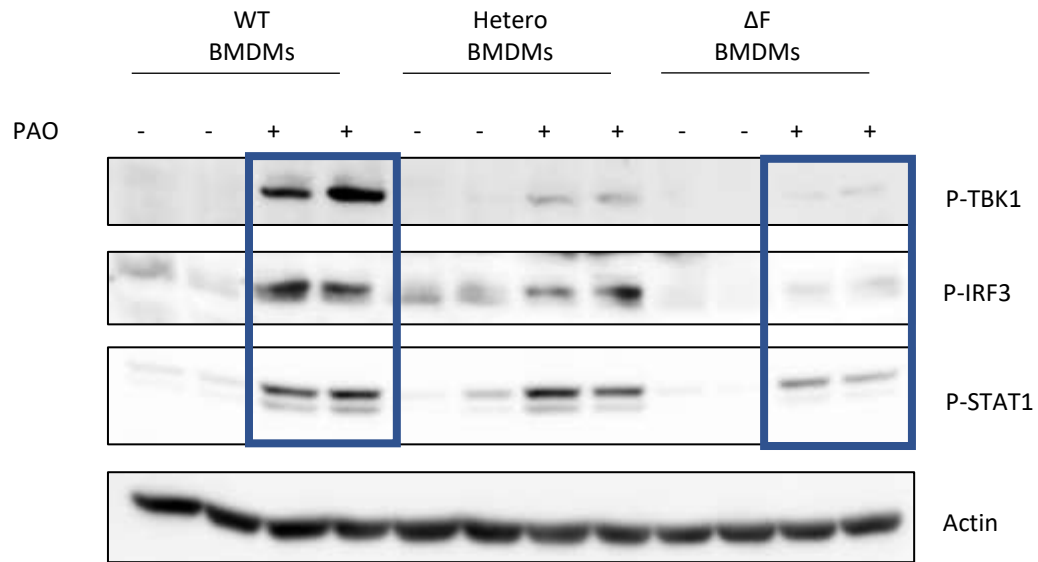
# Working model



# Work Hypothesis

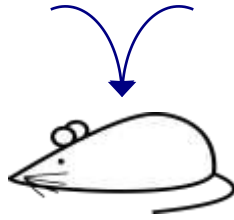


**Primary BMDMs from  $\Delta F508del$ -CFTR mice infected with *Pseudomonas aeruginosa* do not activate the STING/TBK1/IRF3 pathway**



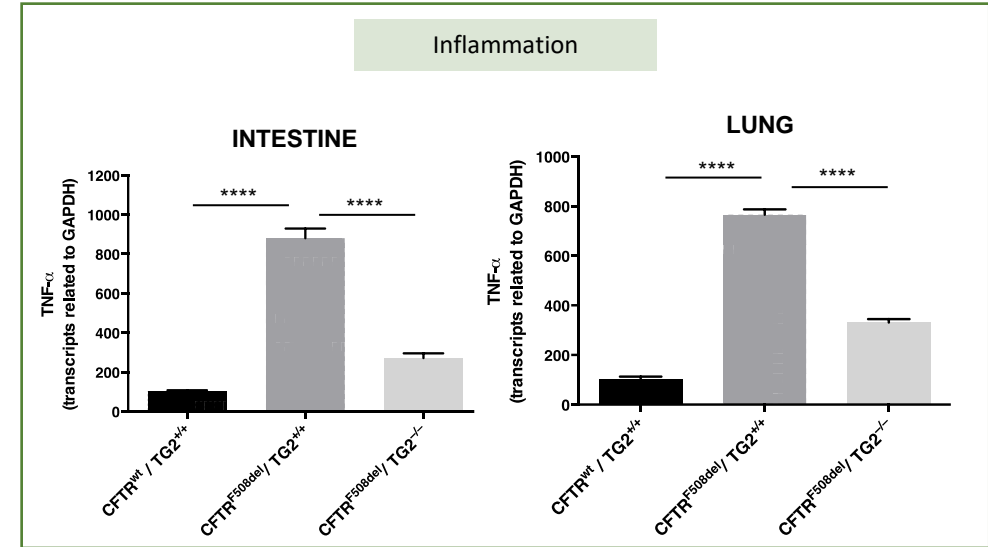
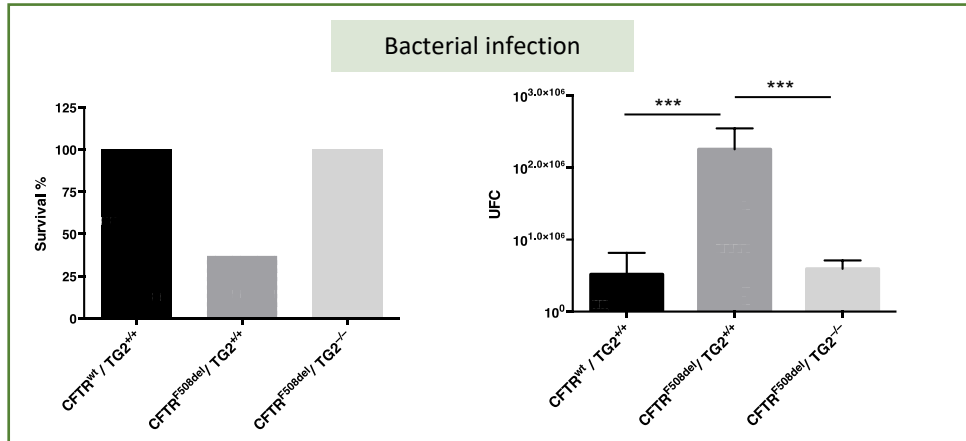
CFTR<sup>wt</sup>/F508del/TG2<sup>+/+</sup>

CFTR<sup>wt</sup>/wt/TG2<sup>-/-</sup>



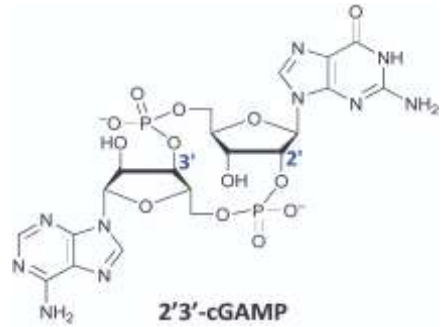
CFTR<sup>F508del</sup>/TG2<sup>-/-</sup>

## CF mouse model knock out for TG2

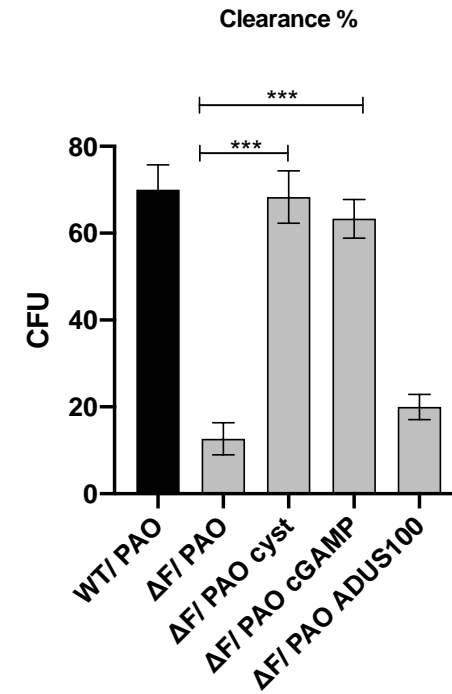
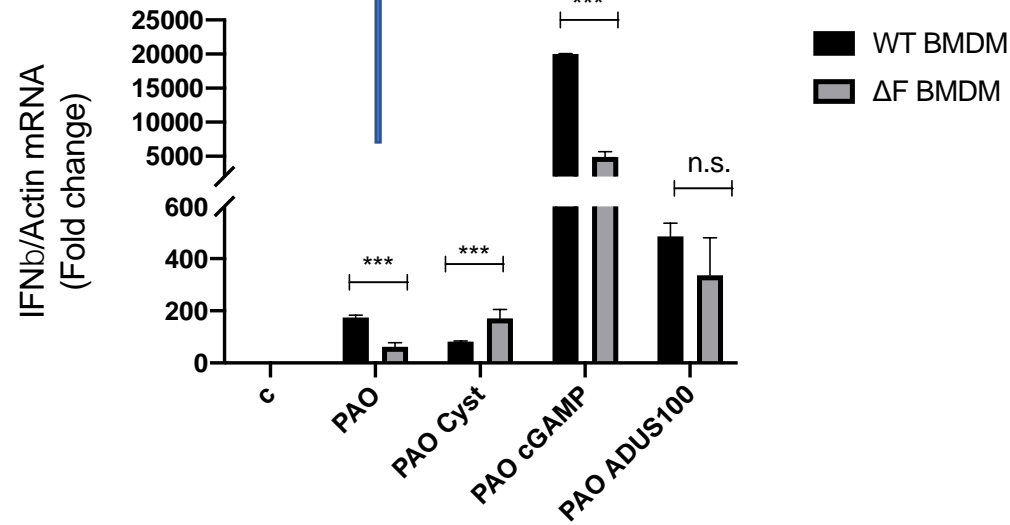
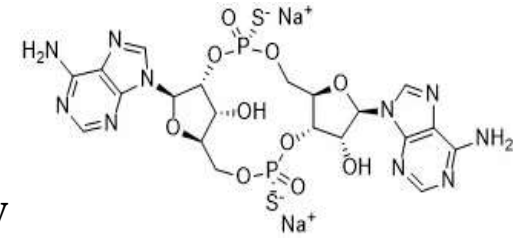


**TG2 ablation in CFTR<sup>F508del</sup> mice re-establishes the immune response to the infection with *Pseudomonas Aeruginosa*.**

The **STING agonists cGAMP** and **ADUS 100** rescue the **INF-beta** production and limit the infection in **F508del-CFTR BMDMs** infected with **Pseudomonas aeruginosa**



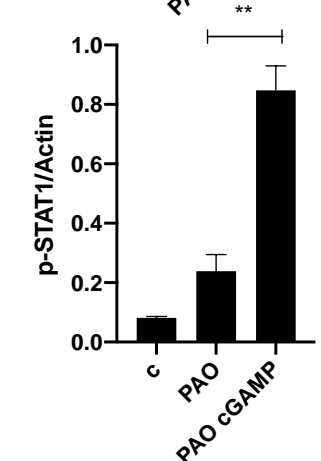
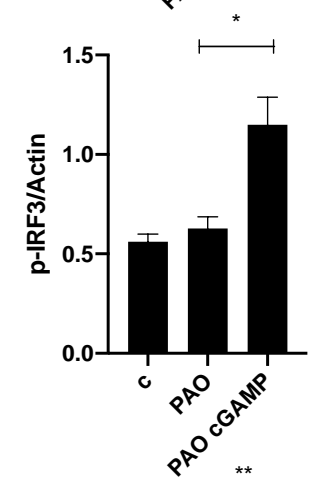
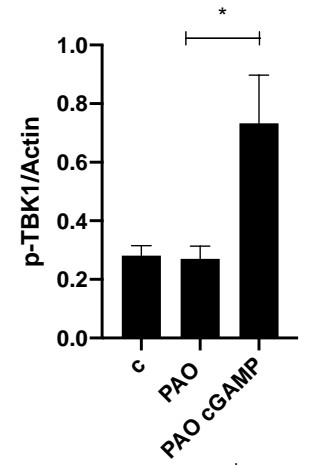
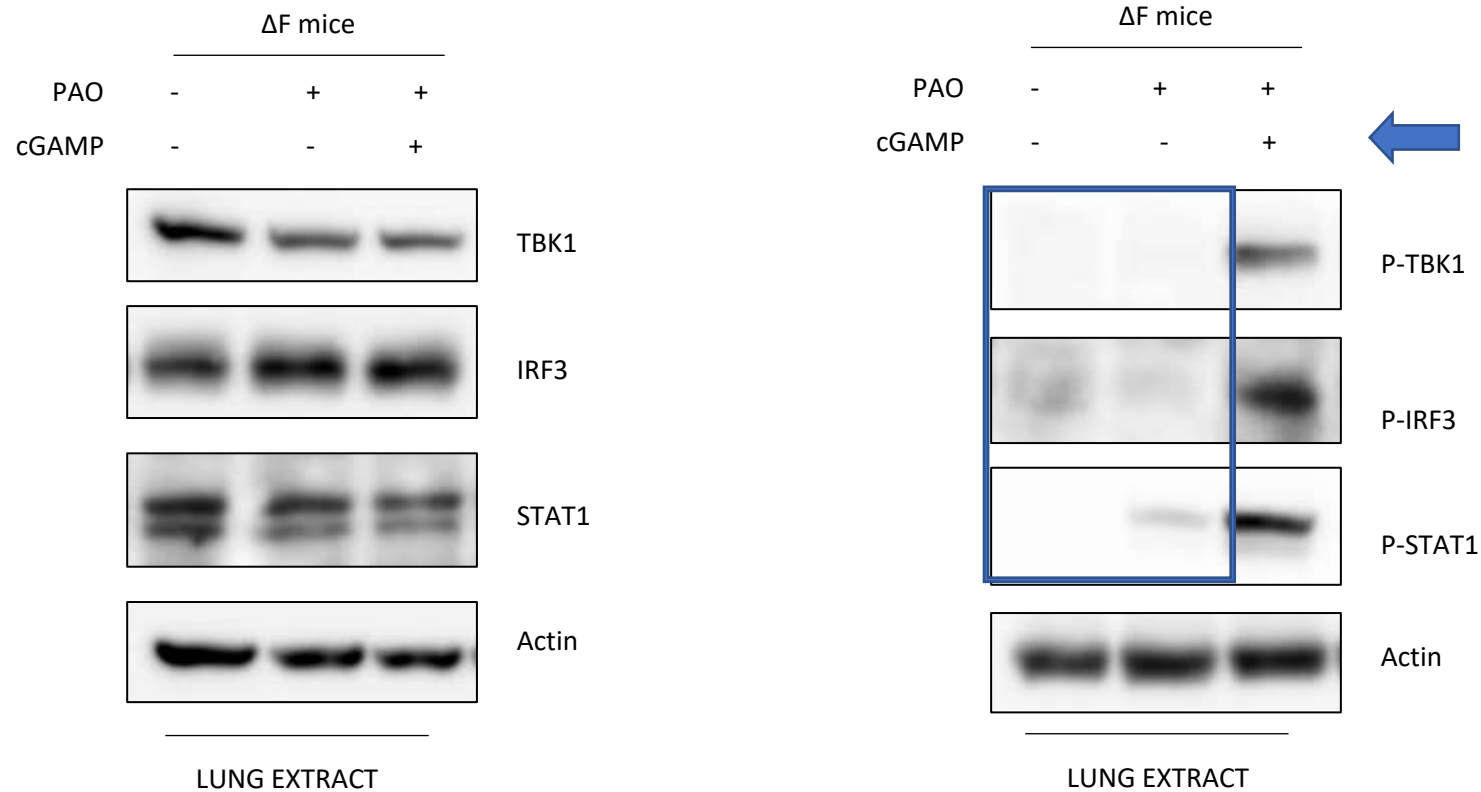
**cGAMP favours PAO internalization and clearance by BMDMs from  $\Delta F508$  mice BMDMs**



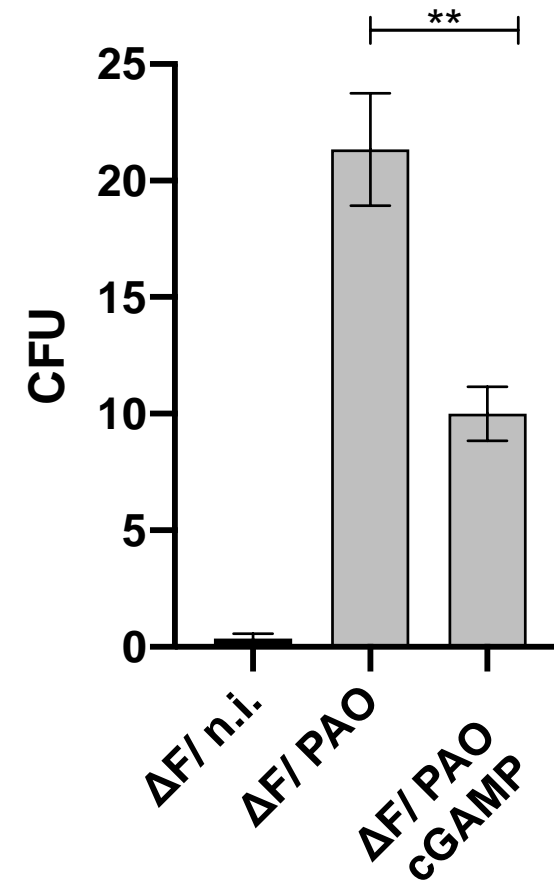
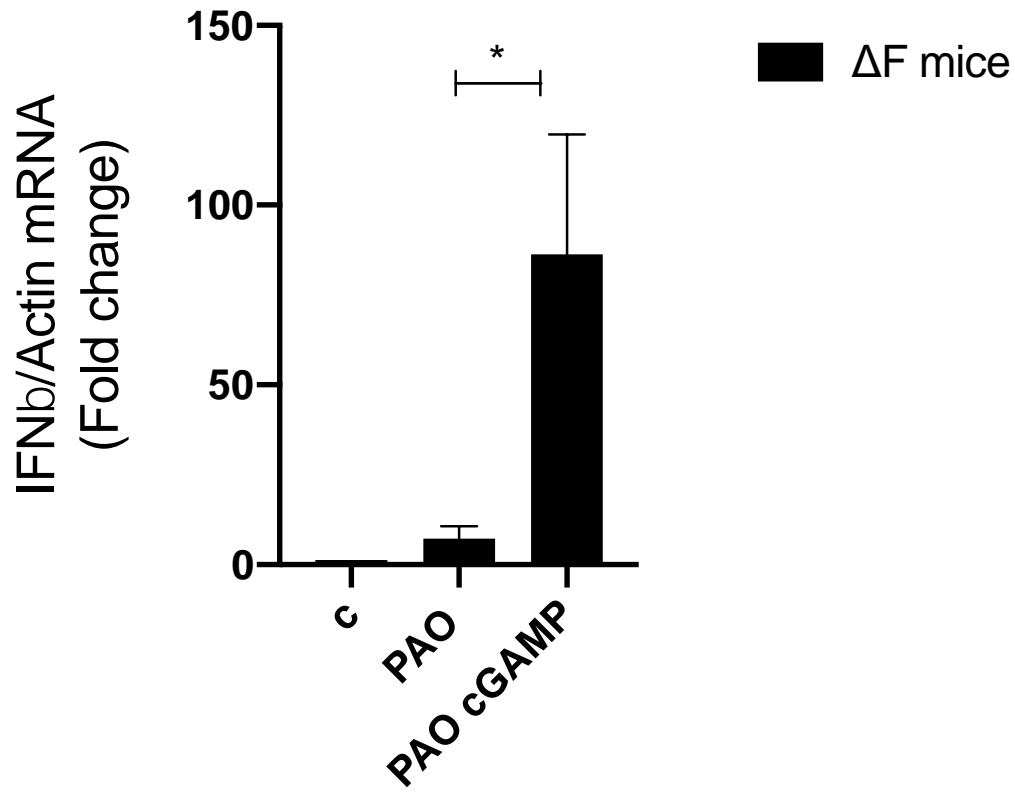


**What happens in vivo in infected  
 $\Delta F$  508del-CFTR mice?**

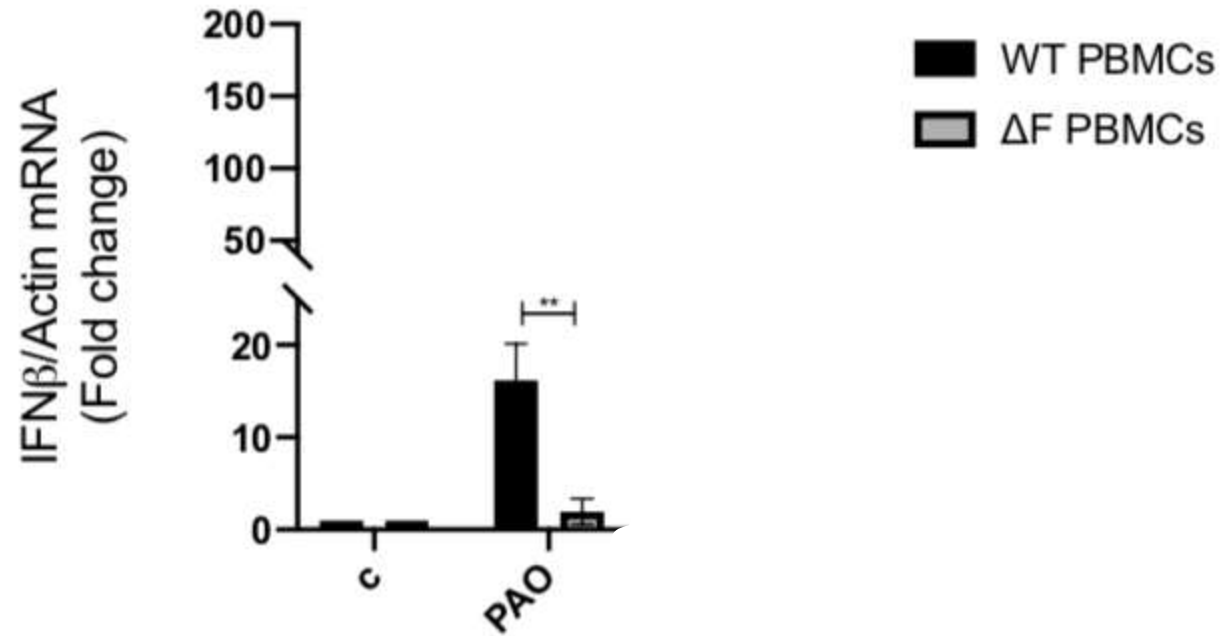
# The STING agonist cGAMP rescues the INF-beta production in $\Delta F508del$ -CFTR mice infected with *Pseudomonas aeruginosa* via the STING/TBK1/IRF3-dependent pathway



**The STING agonist cGAMP rescues the INF-beta production and increases the clearance of *Pseudomonas aeruginosa* in the lung of F508del-CFTR mice**



**Is the STING agonist cGAMP able to rescue the INF-beta production in primary macrophages from Cystic Fibrosis F508del-patients?**



# CONCLUSIONS

- The innate immunity STING/TBK1/IRF3/IFN $\beta$  pathway is defective in Cystic Fibrosis.
- STING agonists and TG2 ablation can re-establish the production of IFN $\beta$  and the clearance of *Pseudomonas aeruginosa in vivo in  $\Delta F508del$  mice.*
- The STING agonists can re-establish the production of IFN $\beta$  *ex vivo* in the PBMCs of Cystic Fibrosis  $\Delta F508del$  patients.

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