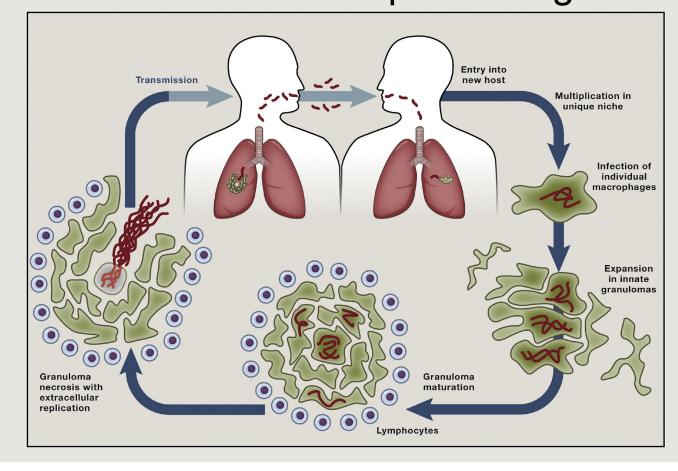


# **Role of Internal Carbon Mobilization for an Optimal Re-Growth of Dormant** Mycobacterium tuberculosis Arihant Boli, Juhyeon Lim, Hyungjin Eoh

# Abstract

Mycobacterium tuberculosis (Mtb) is the bacteria that results in the pathogenic infection of Tuberculosis, a serious pulmonary infection that is fatal if left untreated. In the presence of antibiotics. Mtb enters a persister state, in which the bacteria is phenotypically dormant and the infection subdues. However, upon the elimination of antibiotics, Mtb exits the persister state and returns back to a virulent state. We hypothesize that this reactivation process of Mtb include the internal consumption of upstream glycolytic metabolites as a source of downstream glycolytic intermediates as upstream ones are highly accumulated with reciprocal depletion of downstream ones during the dormant stage, and during re-activation these metabolites. This study examined the role of accumulated glycolytic intermediates as a source of internal carbon mobilization for an optimal re-growth of dormant Mtb. For this, we used hypoxia as this condition engendered Mtb into a dormant state, and metabolomic remodeling during re-growth was analyzed through LCMS mediated metabolomics. The outcome of this study will uncover new therapeutic targets to clear Mtb infection.



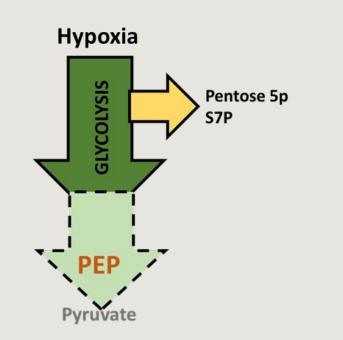


Fig. 1.1. Cycle of replication, dormancy, and re-activation in vivo of Mycobacterium tuberculosis

## Method

### 1. Mtb Culture

- a. 4 Strains of Mtb  $\rightarrow$  RV (Wildtype), RV ATC, PYK (pyk knockdown), PYK ATC,
- b. Utilized m7H9 or m7H10 growth media
- c. Created mutant strains via Cripr—dCas9
- Filter culture system to give hypoxia for re-activation Hypoxia Chamber

### 2. Checking Reactivation Growth

- e. Replicating culture:
  - Bacteria set to optical density of 0.05 (day 0)
  - Optical density checked every 3-4 days to record replication growth
- Reactivation culture:
- Re-activation culture created through disturbing Mtb filters upon 3 days of hypoxic conditions
- Optical density checked every 3-4 days to record reactivation growth

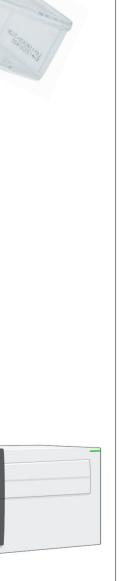
### 3. LC-MS Metabolomics

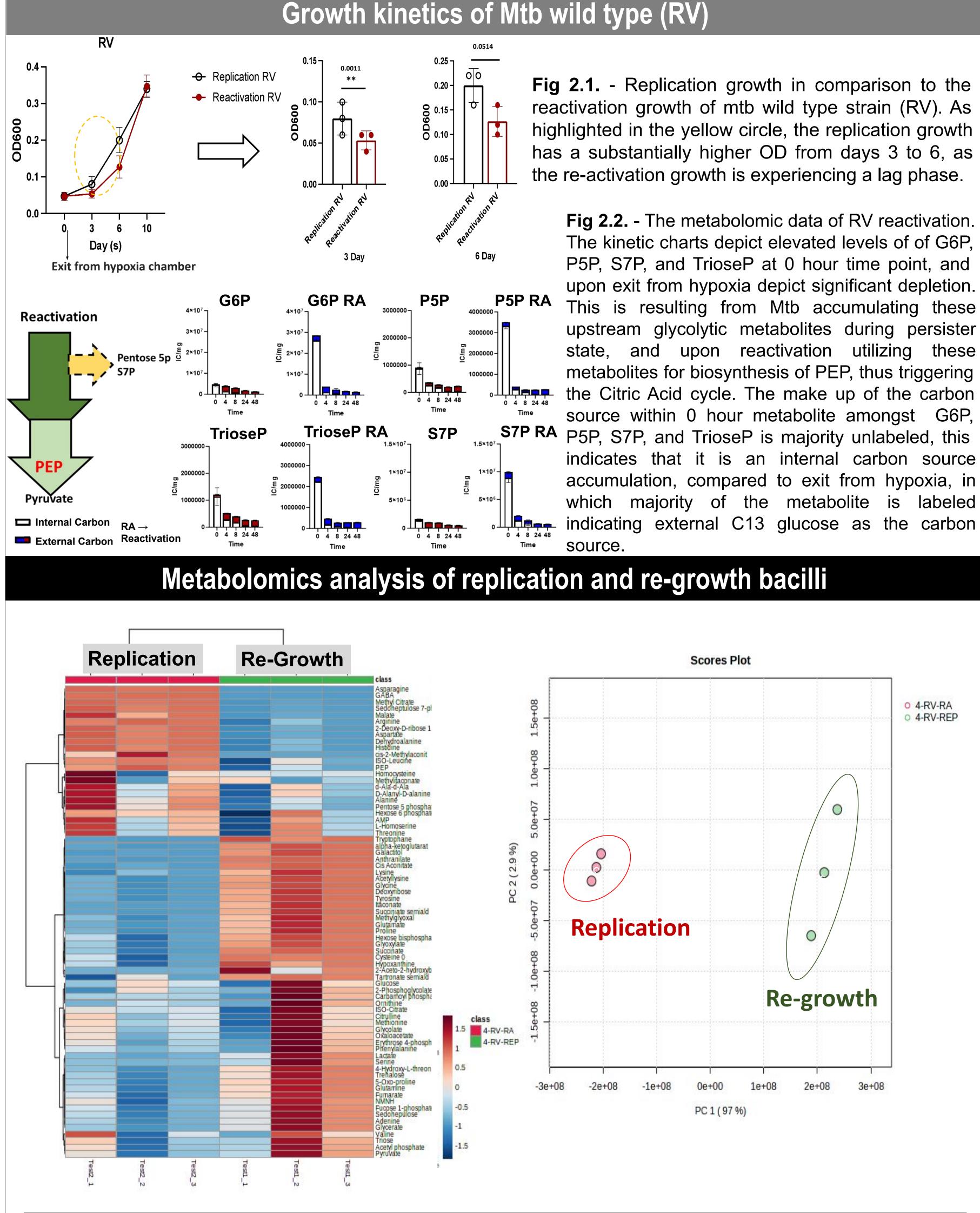
- Samples for metabolomics transferred to C13 isotope plates for detection of Internal vs External carbon
- Liquid Chromatography Mass Spectrometry used to detect metabolites present in replication and re-growth samples
- Principal component analysis and Heat Map analysis performed via Metaboanalyst V. 5.0



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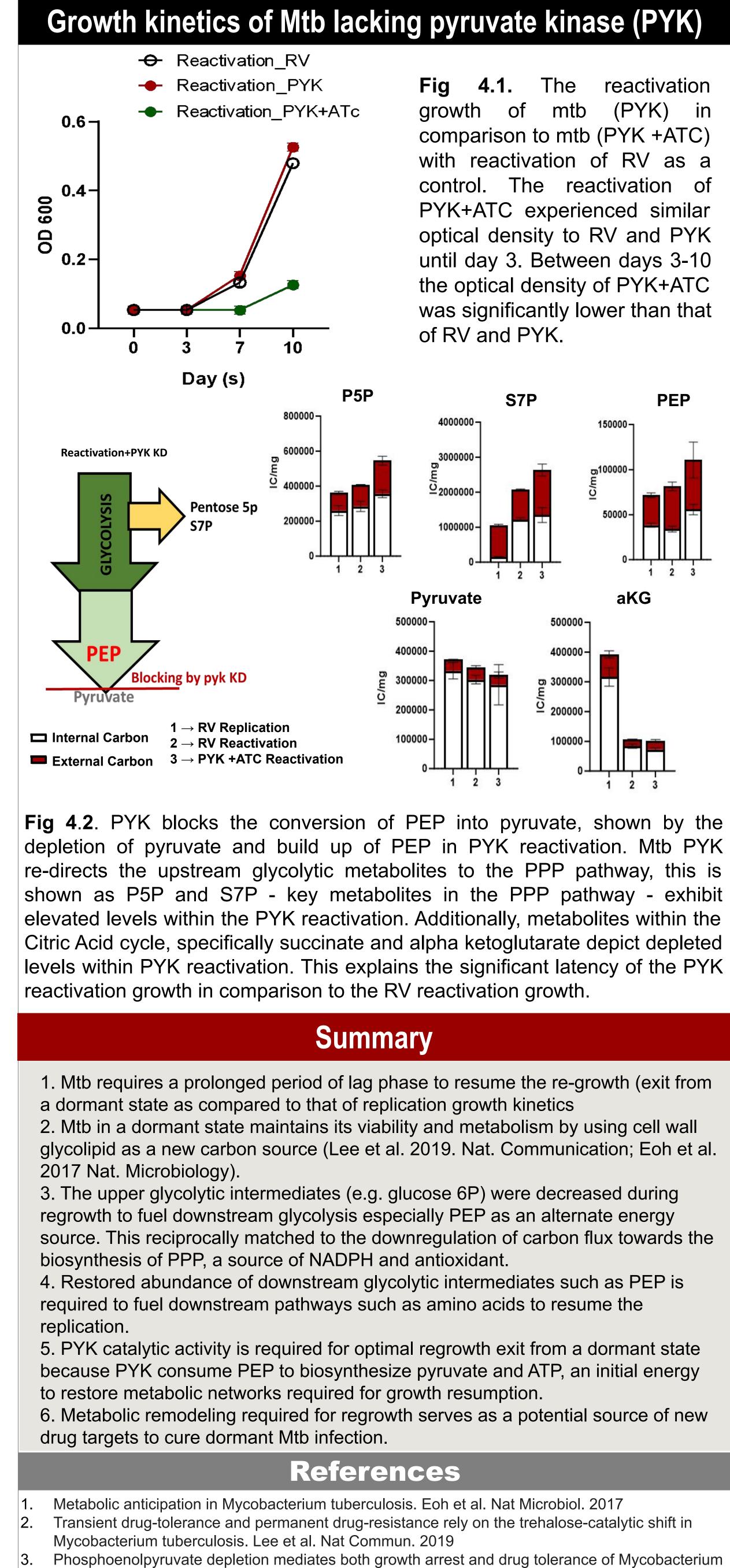


# **Fig 3.1**

Left – Central carbon metabolism (e.g., glycolysis and pentose phosphate pathway) and cell wall glycolipid precursor metabolism (e.g., trehalose or alpha glucan) belong to high ranked pathways that were different between the two conditions.

Right – PCA analysis showed the Mtb metabolic networks in a replication and a re-growth condition were clearly different.

Fig 2.2. - The metabolomic data of RV reactivation. The kinetic charts depict elevated levels of G6P, P5P, S7P, and TrioseP at 0 hour time point, and upon exit from hypoxia depict significant depletion. his is resulting from Mtb accumulating these metabolites for biosynthesis of PEP, thus triggering the Citric Acid cycle. The make up of the carbon source within 0 hour metabolite amongst G6P, indicates that it is an internal carbon source accumulation, compared to exit from hypoxia, in which majority of the metabolite is labeled indicating external C13 glucose as the carbon







The reactivation growth of mtb (PYK) in with reactivation of RV as a control. The reactivation of PYK+ATC experienced similar

tuberculosis in hypoxia. Lim J et al. Proc Natl Acad Sci U S A. 2021