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#### ▶ To cite this version:

Fatima Hassan Hellani, Nasreddine Saidi, Inès Leleu, Jacques Roland, Nathalie Martin, et al.. Role of astrocyte senescence induced by Plasmodium in the pathogenesis of Cerebral Malaria. 15th ISNI congres of International Society of Neuroimmunology, Nov 2021, Nice, France. 2020. hal-03457641

#### HAL Id: hal-03457641 https://hal.science/hal-03457641

Submitted on 30 Nov 2021

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# Role of astrocyte senescence induced by *Plasmodium* in the pathogenesis of Cerebral

# Malaria

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# Introduction

Cerebral malaria (CM) is a fatal neurological syndrom due to Plasmodium (P.) falciparum infection, which kills 405,000 people annually(1). A central event involved in the pathogenesis of CM is the sequestration of infected red blood cells in the brain capillaries and the inflammatory response of activated glial cells (2). However, molecular mechanisms involved in the induction of the pro-inflammatory response of glial cells during CM remains unknown.

In this study, we have examined if the senescence of glial cells induced by P. berghei ANKA infection participates in the neuroinflammatory response associated to CM.

Cellular senescence is a permanent state of cell cycle arrest characterized by morphologic, metabolic and molecular changes (3). The senescence phenotype is often characterized by the activation of a DNA damage response that promotes an increase of senescence-associated- $\beta$ -galactosidase (SA- $\beta$ -gal) activity and expression of cell cycle-inhibitory proteins, most frequently p53/p21<sup>WAF1/Cip1</sup> and/or p16<sup>INK4a</sup>/RB. In addition to being irreversibly arrested, senescent cells produce a battery of factors including pro-inflammatory cytokines involved in CM characterizing the senescence-associated secretory phenotype (SASP) (4).



Cell size Morphologic alterations

## **Materials and methods**



Senolytic

drugs



SR-B1

SUM

Lysosomal content SA-B-Gal activity



SASP factors (IL-6, IL-1β TNF-a, MMP-1, etc.) 🛉

Neurotrophic factor (NGF, IGF-1, etc.)

Plasmodium berghei

ANKA

C57BL/6 (CM<sup>S</sup>)

## **1- Increase of SA-β-gal activity and cell cycle inhibitors** expression in brain of CM<sup>S</sup> mice.



## 2- P. berghei ANKA infection-induced astrocytes senescence in the brain of CM<sup>S</sup> mice.





**Remerciements à Jacques-Roland, and Sophie Salome-Desnoulez** 







