

# 11P - Senescent Immune Phenotype (SIP) status that predicts resistance to Immune Checkpoint Blockers (ICB) among CMV+ advanced Non-Small Cell Lung Cancer (aNSCLC) patients is not associated with chronic type I IFN signature

M. Naigeon<sup>1</sup>, C. de Oliveira<sup>1</sup>, F.-X. Danlos<sup>2</sup>, B. Duchemann<sup>1</sup>, P. Saulnier<sup>3</sup>, L. Boselli<sup>1</sup>, J-M Jouriaux<sup>1</sup>, F. Griscelli<sup>1</sup>, A. Marabelle<sup>2</sup>, L. Lacroix<sup>3</sup>, L. Cassard<sup>1</sup>, B. Besse<sup>5</sup>, N. Chaput<sup>1</sup>

1.Laboratory of Immunomonitoring in Oncology, CNRS-UMS 3655 and INSERM-US23, Paris Saclay University, Gustave Roussy, Villejuif, Cedex, France, 2.Drug Development Department, INSERM U1015-LRTI, Paris Saclay University, Gustave Roussy, Villejuif, France, 3.Inserm, CNRS, Analyse moléculaire, modélisation et imagerie de la maladie cancéreuse, Genomic platform Molecular Biopathology unit and Biological Resource Center, Paris Saclay University, Gustave Roussy, Villejuif, Cedex, France, 4.Department of Biopathology, Gustave Roussy, Villejuif, France, 5.Département de Cancer Medicine, Paris Saclay University, Gustave Roussy, Villejuif, France. Study sponsored by Malakoff Médéric.

## BACKGROUND

- Immunosenescence is a progressive remodeling of immune functions with a multifactorial etiology (aging, chronic inflammation, persistent infections, cancer).
- CMV has been shown to act as chronic antigenic stressor and to accelerate immune ageing by affecting peripheral blood T cell phenotypes, including loss of CD28 or overexpression of CD57.
- Latent viral infections were shown to be associated with chronic type I IFN signature that might promote lymphocyte senescence.
- We defined SIP as the proportion of CD28<sup>-</sup>CD57<sup>+</sup>KLRG1<sup>+</sup>CD8<sup>+</sup> circulating T cells. We showed that a high pretreatment SIP (>39.5%, SIP+) was associated with resistance to ICB in patients with aNSCLC<sup>3</sup>.

## OBJECTIVE

We aimed to assess the role of SIP combined to CMV status on outcomes in aNSCLC patients and the association between SIP and chronic type I IFN signature.

## PATIENTS AND METHODS

Baseline SIP status was assessed by flow cytometry on fresh blood samples from ICB-treated and polychemotherapy-treated (PCT) aNSCLC patients. Type I IFN score was calculated by the sum of relative expression of 5 IFN-stimulated genes (*IFI44*, *IFIT1*, *IFITM1*, *LY6E*, *MX1*), determined by RT-qPCR. Soluble factors associated to interferons (IFN $\alpha$ , IFN $\beta$ , IFN $\gamma$ , IP-10, PD-L1) were quantified using the MSD assay.

## RESULTS

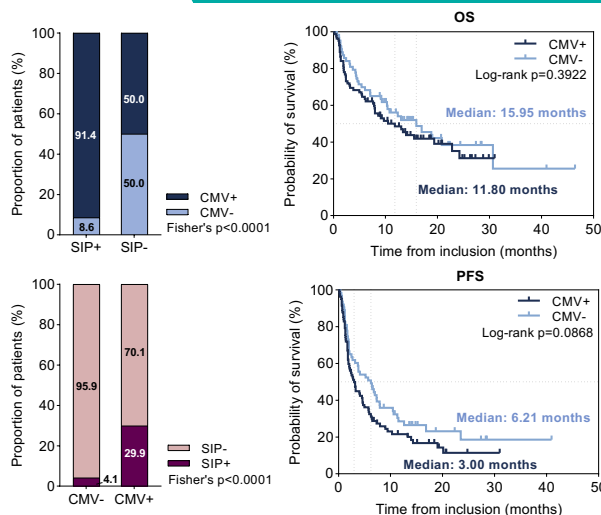


Fig. 1: SIP and CMV status

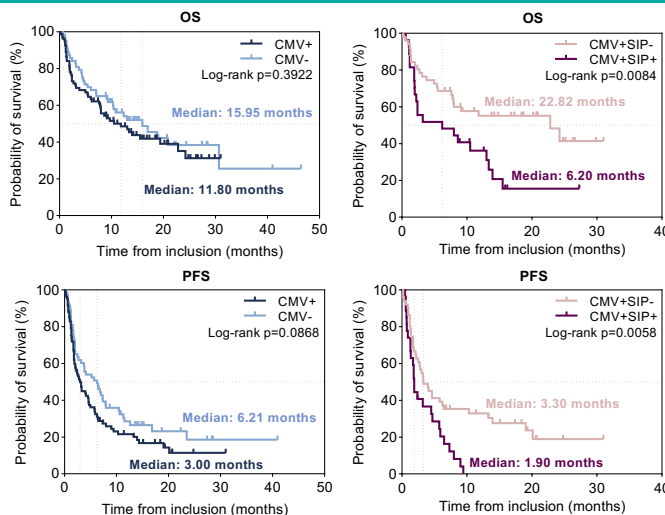


Fig. 2: PFS and OS according to CMV status

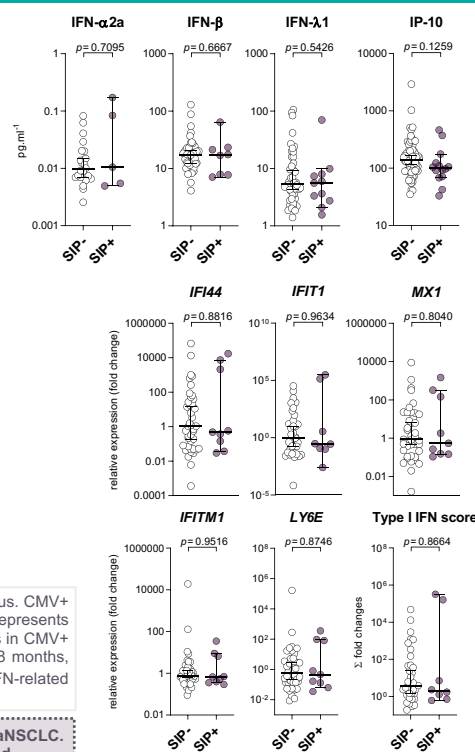


Fig. 3: PFS and OS in CMV+ patients according to SIP status

203 aNSCLC patients (142 ICB-treated, 61 PCT-treated) were evaluable for SIP (19.7% SIP+) and 180 patients (89%) for CMV status. CMV+ patient's rate was significantly higher in SIP+ compared to SIP- patients (91.4% vs 50%,  $p < 0.0001$ ). Among CMV+ patients, SIP+ represents only 30% (Fig.1). In ICB-treated patients, CMV status was not sufficient to predict outcomes (median PFS 6.21 in CMV- vs 3 months in CMV+ patients,  $p = 0.087$ ) (Fig.2). Among CMV+ patients, SIP+ patients had lower PFS (1.9 vs 3.3 months,  $p = 0.006$ ) and OS (6.2 vs 22.8 months,  $p = 0.008$ ) than SIP- (Fig.3). Otherwise, at baseline, no association between SIP and type I IFN signature nor plasmatic levels of IFN-related factors was found (Fig.4).

## CONCLUSION

SIP+ patients are CMV+, SIP+ identifies patients with poorer outcomes in CMV+ ICB-treated aNSCLC. No association between type I IFN signature or IFN-related plasmatic factors and SIP was found.

Fig. 4: IFN-related plasmatic factors and type I IFN signature according to SIP status