

The response to Convalescent-Plasma in COVID-19 induced ARDS does not differ across immunological sub-phenotypes

Immunological sub-phenotypes and response to Convalescent-Plasma (CP) in COVID-19 induced ARDS (C-ARDS): a hierarchical cluster analysis

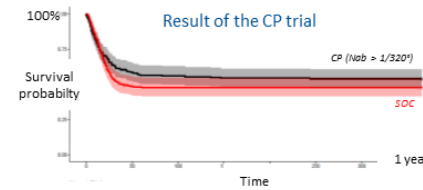
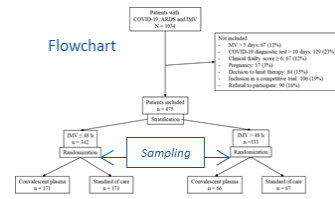
INTRODUCTION

The CONFIDENT trial tested the effect of passive immunization with plasma collected from convalescents and high titers of neutralizing antibodies may reduce mortality in patients with COVID-19 associated ARDS when administered early after the introduction of IMV. It demonstrated a 9.7% reduction in day-28 mortality (p=0.03) (Misset, *New Engl J Med*, 2023 in press).

As the immunological response to COVID-19 is likely heterogeneous across patients, we hypothesized that immunologically similar COVID-19 clusters might differ in their responses to convalescent-plasma treatment.

METHODS

- Secondary analysis of the CONFIDENT trial
- **Inclusion criteria** : Adult, Frailty scale < 6, SARS-CoV-2 pneumonia, IMV <5 days (stratification at 48h)
- **Intervention** : 2 x 250 mL bags of CP with **NAb > 1/320°**
- Control : standard care (SC)



- **Sampling at the time of inclusion** (in 13/17 centers having accepted the secondary part of the trial).
- **Multiplex Luminex** technique (ELISA combined with flow cytometry)
- **20 cytokines, chemokines and cell adhesion markers.**
- Descriptive statistics (median [IQR]), **hierarchical cluster analysis** and search for association between the identified clusters and CP effect on day-28 mortality.

CONCLUSIONS

In a cohort of patients with COVID-associated ARDS included in a convalescent plasma trial, we identified 4 sub-phenotypes based on their immune response associated with different clinical profiles. The response to convalescent plasma as assessed by the day-28 mortality was not different between the 4 sub-phenotypes.

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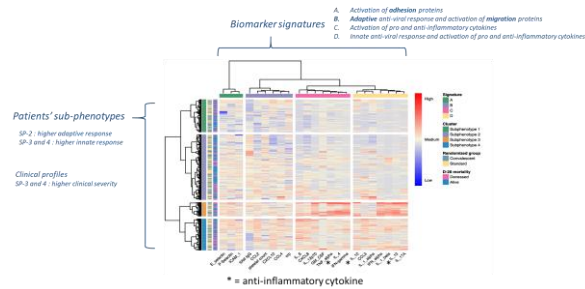
RESULTS

Population

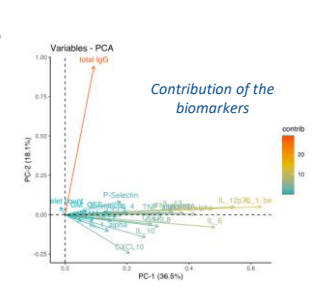
Similar to the total trial population (n=475)

Patients, n	391
Age, y	64 [56-72]
BMI, kg/m ²	30,0 [26,4-34,5]
APACHE II	13 [9-17]
SOFA	6 [4-8]
Pao2/FIO2	120 [90-160]
Steroids	375 (95,9%)
Group (CP/SC)	196 / 195
Inclusion IMV < 48h	299 (76,5%)

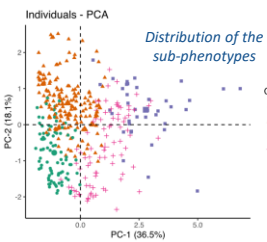
Hierarchical clustering



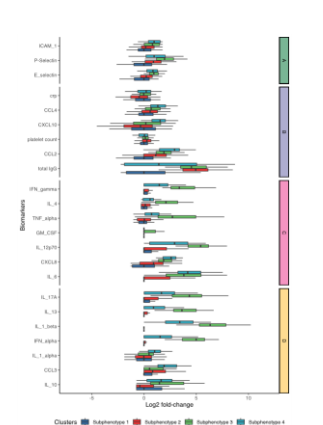
Principal component analysis 1



Principal component analysis 2



Biomarkers by sub-phenotypes



Effect of CP/SC by sub-phenotypes

