

# Lung Hybrid Neutrophils and Extracellular Traps Are Protective in COVID-19-Associated Pulmonary Aspergillosis

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## Background

COVID-19-associated pulmonary aspergillosis (CAPA)

- *Aspergillus* superinfection in critically ill COVID-19 patients
- Increased mortality, diagnosis is difficult
- Pathophysiological insight is lacking.

## Methods

### Single-cell RNA sequencing

- 37 BAL samples of 37 critically ill COVID-19 patients
  - 22 COVID-19-only patients (no aspergillosis)
  - 6 early CAPA patients (sampling <5 days after diagnosis)
  - 9 late CAPA patients (sampling 5-11 days after diagnosis)

### Neutrophil extracellular trap (NET) levels

- 57 BAL samples of 57 critically ill COVID-19 patients
  - 33 COVID-19-only patients
  - 24 early CAPA patients

1B

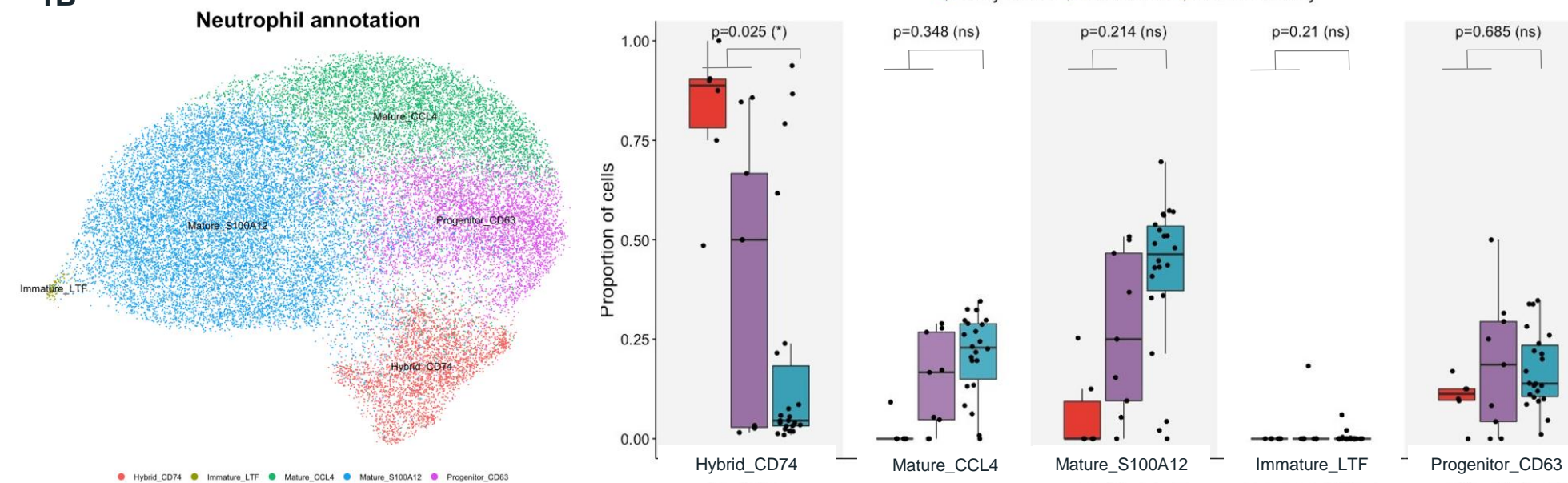


Figure 1 panel (B): UMAP of the neutrophil fraction, and box plots depicting the proportions of neutrophil subtypes in BAL fluid. P-values shown for differences between the pooled CAPA patients and the COVID-19-only patients. P-values were calculated using a generalized linear model correcting for age, Charlson Comorbidity Index at hospital admission, and administration of corticosteroids (prednisone equivalent dose 20 mg or higher) within 48 hours of BALF sampling.

## Results

### scRNA-seq

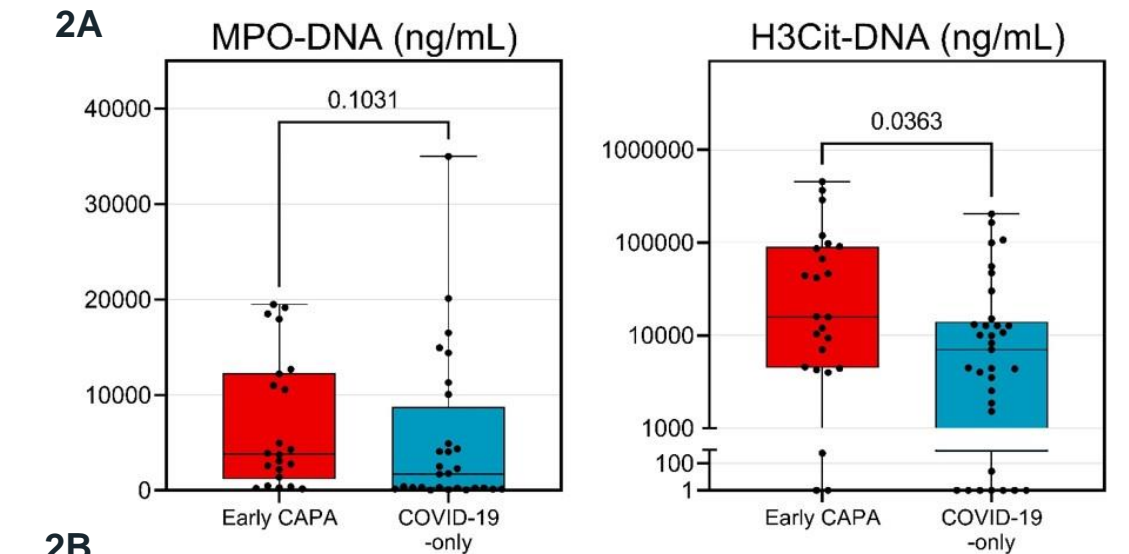
- 69008 cells passed quality filtering
- Lower neutrophil proportions in CAPA vs. COVID-19-only (Fig. 1A)
- Neutrophil subclustering (Fig. 1B)
  - Immature & mature clusters
  - “Hybrid” neutrophil cluster
    - Genes ~ antigen-presenting functions
    - Dominant in CAPA

### NETosis

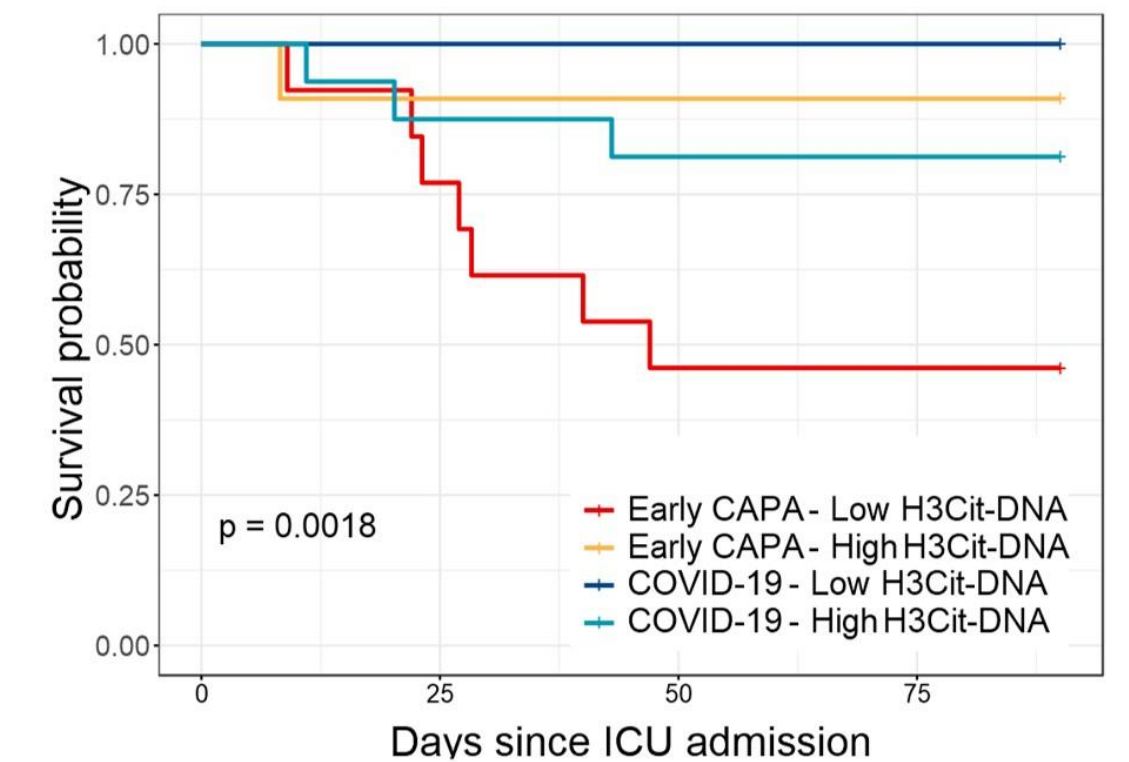
- Significantly higher citrullinated histone H3 DNA complexes levels (H3Cit-DNA) in CAPA patients (Fig. 2A)
  - Explains low neutrophil proportions
- Significantly lower survival in CAPA patients with the lowest H3Cit-DNA levels (Fig. 2B).

## Conclusion

- Extremely high NET levels in CAPA lower respiratory tract, which are likely to be protective in CAPA patients specifically
- Hybrid neutrophil formation in CAPA, probably upon encountering the fungus



2B



Number at risk

Strata	0	25	50	75
Early CAPA - Low H3Cit-DNA	13	10	6	6
Early CAPA - High H3Cit-DNA	11	10	10	10
COVID-19 - Low H3Cit-DNA	17	17	17	17
COVID-19 - High H3Cit-DNA	16	14	13	13

Figure 2 panel (A): Myeloperoxidase (MPO) DNA complexes were analyzed as biomarkers for general NET-formation, while citrullinated histone H3 DNA (H3Cit-DNA) levels were analyzed for PAD4-dependent NET-formation, in BALF samples from early CAPA and COVID-19-only patients. Similar MPO-DNA levels were found in early CAPA patients, while H3Cit-DNA levels were significantly higher in early CAPA compared to COVID-19-only patients. P-values calculated using Mann-Whitney U test.

Figure 2 panel (B): Kaplan-Meier analysis of patients with NETosis analyses, divided in early CAPA and COVID-19-only patients and subdivided according to H3Cit-DNA levels (cut-off at 20000 ng/mL for early CAPA and at 8000 ng/mL for COVID-19-only). Log-rank test was used to compare survival distributions. For the comparison early CAPA (low H3Cit-DNA) versus early CAPA (high H3Cit-DNA), the p-value was 0.033.

1A

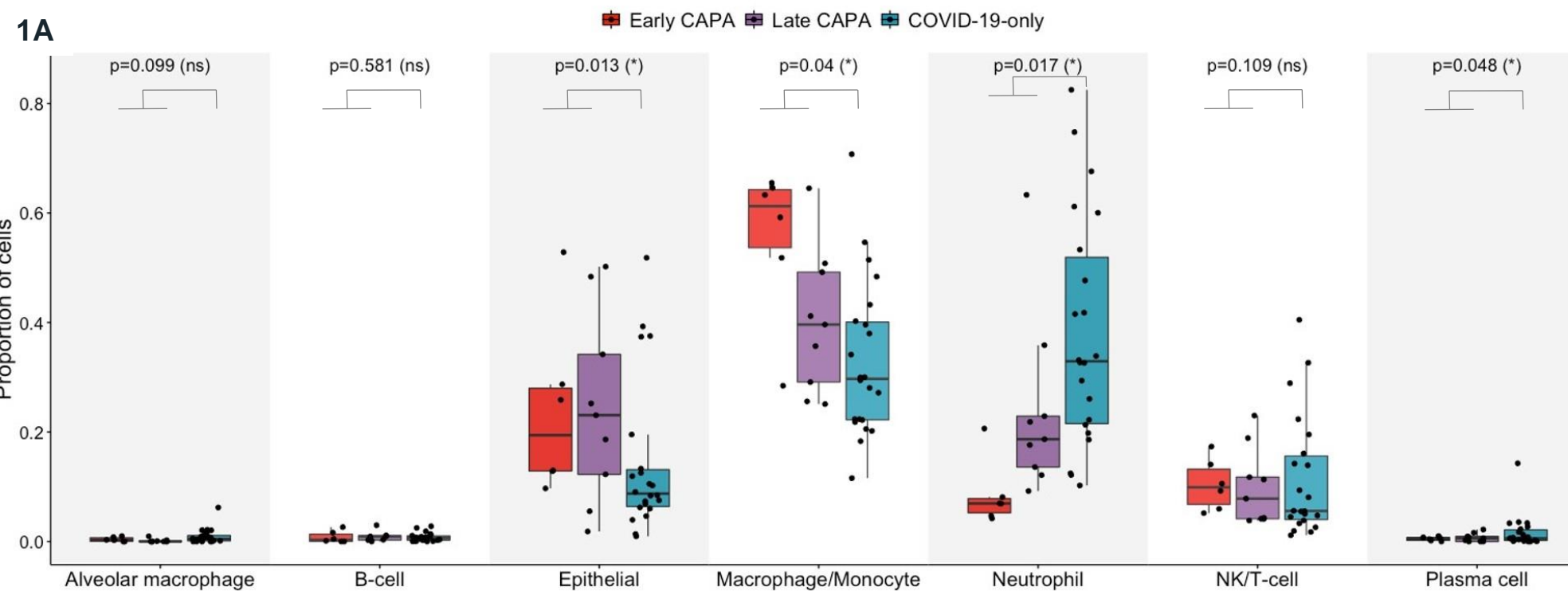


Figure 1 panel (A): BALF neutrophil proportions as analyzed by single-cell RNA sequencing using the Seurat R package are significantly lower in CAPA patients compared to COVID-19-only patients. Patients with early CAPA have significantly lower BALF neutrophil proportions than patients with late CAPA. Macrophage/monocyte and epithelial cell proportions are reciprocally increased in CAPA patients compared to COVID-19-only patients. P-values shown for differences between the pooled CAPA patients and the COVID-19-only patients. P-values were calculated using a generalized linear model correcting for age, Charlson Comorbidity Index at hospital admission, and administration of corticosteroids (prednisone equivalent dose 20 mg or higher) within 48 hours of BALF sampling.

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