

Unravelling the Clinical and Biological Features of Accelerated Chronic Lymphocytic Leukemia

Francesco Angotzi¹, Raffaella Pasquale², Massimo Moratti², Luca Laurenti³, Tommaso Quaranta³, Idanna Innocenti³, Gioachino Catania⁴, Massimo Gentile⁵, Marta Coscia⁶, Giulia Zamprogna⁶, Enrico Lista⁷, Paolo Sportoletti⁸, Alberto Fresa⁹, Andrea Galitzia¹⁰, Valerio Guarente¹¹, Alessandro Cellini¹, Arianna Bevilacqua¹, Chiara Adele Cavarretta¹, Andrea Serafin¹, Marco Pizzi¹², Enrico Gaffo¹³, Stefania Bortoluzzi¹⁴, Livio Trentin¹, Andrea Visentin¹

¹Hematology Unit, University of Padova, Padova, Italy; ²Hematology Unit, Azienda Sanitaria Universitaria Friuli Centrale (ASUFC), Udine, Italy; ³Fondazione Policlinico Universitario A Gemelli Roma IRCCS, Roma, Italy; ⁴Division of Hematology, Hospital Saints (A.O.SS) Antonio e Biagio and Cesare Arrigo, Alessandria, Italy; ⁵Haematology Unit of Cosenza, cosenza, Italy; ⁶Department of Medicine and Surgery, University of Insubria and Department of Oncology, ASST Sette Laghi, Ospedale di Circolo, Varese, Italy; ⁷Santa Chiara Hospital, APSS Trento, Trento, Italy; ⁸University of Perugia, Perugia, Italy; ⁹o Nazionale Tumori IRCCS - Fondazione "G. Pascale", Naples, Italy; ¹⁰Struttura complessa di Ematologia, Ospedale S Francesco, ASL Nuoro Nuoro Italy; ¹¹Hematology Unit, Ospedale "S.Bassiano" AULSS7 Pedemontana, Bassano del Grappa, Italy; ¹²Pathology Unit, Department of Medicine, University of Padova, Padova, Italy; ¹³Computational Genomics Group, Department of Surgery, Oncology and Gastroenterology, University of Padova, Padova, Italy; ¹⁴Department of Molecular Medicine, University of Padova, Padova, Italy

Background

Accelerated chronic lymphocytic leukemia (aCLL) is a rare (~1%) and understudied istological variant of CLL with aggressive histological features, considered to fall in a biological continuum between CLL and Richter Transformation (RT).

Aim

To investigate the clinical and biological features of aCLL

Methods & study design

We conducted a multicentric retrospective study enrolling **73 patients with an histologically proven aCLL diagnosis** according to the criteria defined by Giné et al. (1). across 12 Italian institutions.

Two cohorts of patients with histologically proven diagnoses of **CLL (n=155)** and **RT (n=15)** treated at the Hematology Unit of the University of Padova were used as **control groups**.

We also performed:

- **NGS analysis** (108-lymphoid gene panel) on FFPE lymph-node tissue form 12 patients (n=7 aCLL, n=2 RT, n=3 CLL)
- **scRNAseq analysis** on FFPE lymph-node tissue from 16 patients ((n=10 aCLL, n=3 RT, n=3 CLL) through the Illumina NovaSeq platform

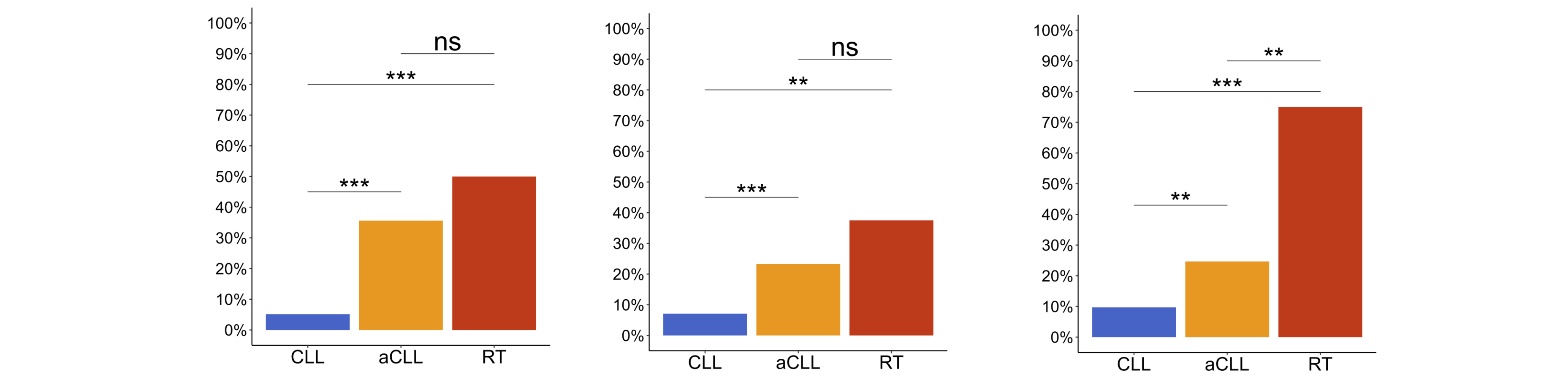
The three groups were compared for clinical and biological variables, including time-to event outcomes such as overall survival (OS) and time to next treatment (TTNT).

Results

Median age was 68 years (range: 39-88), 94 patients (39%) were female and 150 (61%) were male, with no significant differences in terms of age or sex distribution between the three groups.

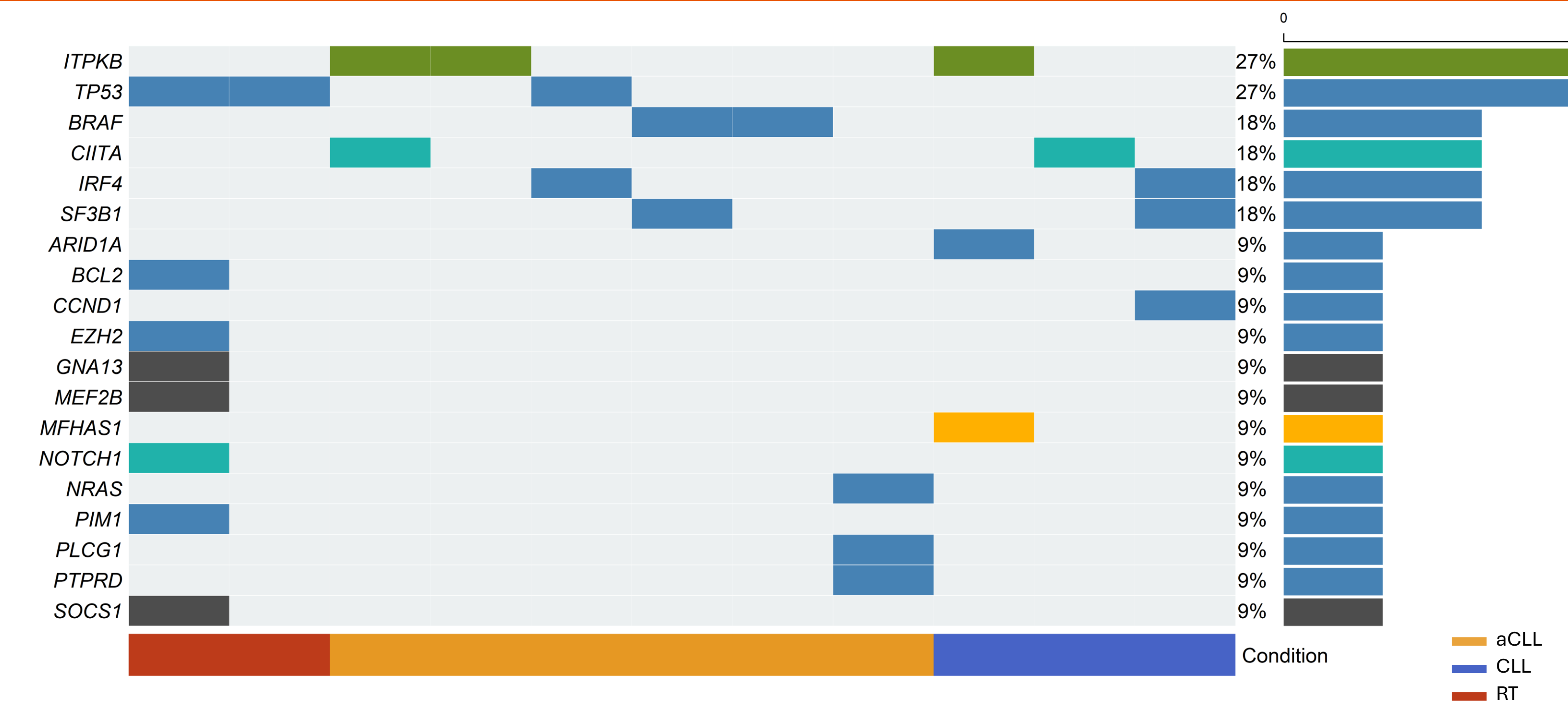
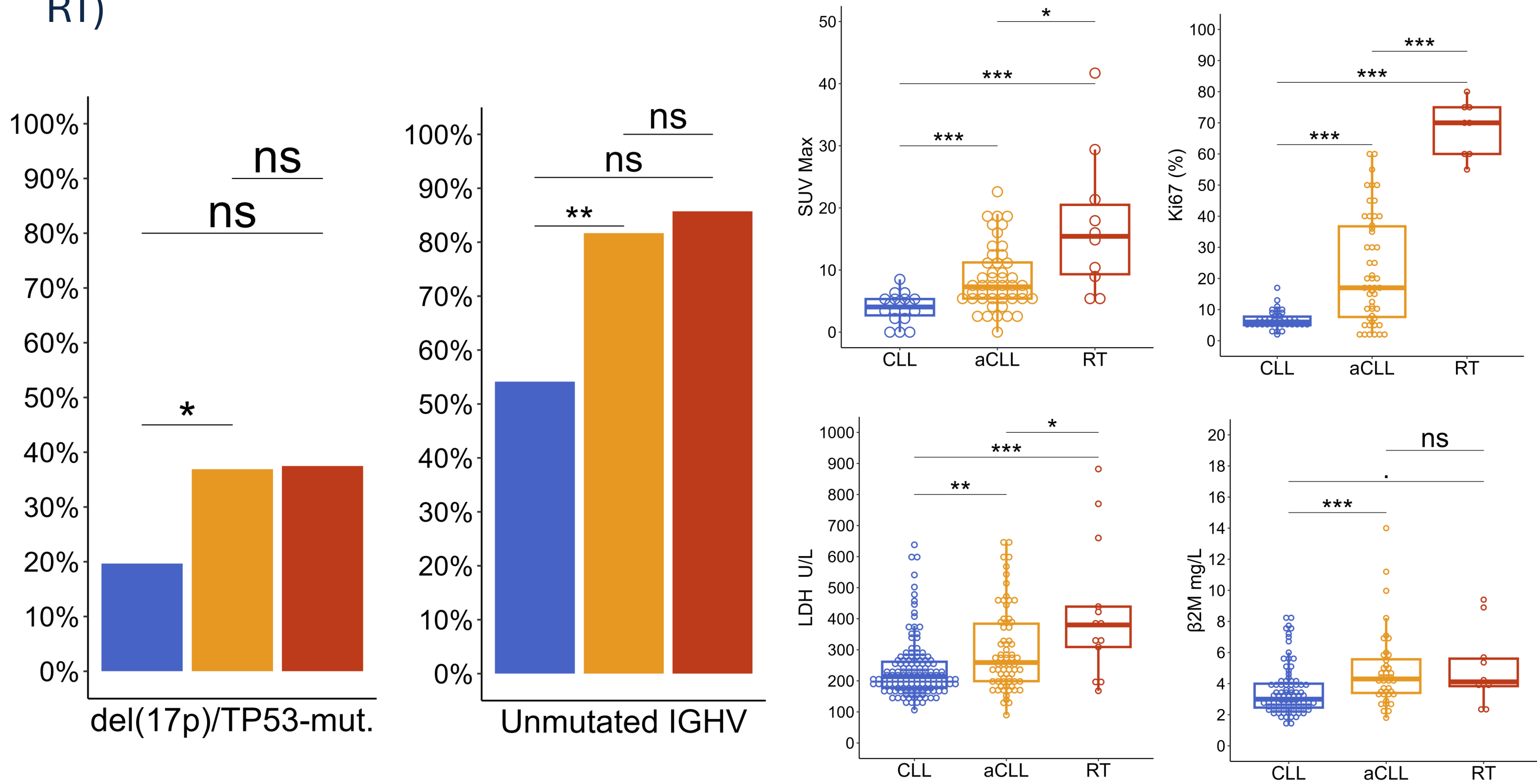
Among aCLL patients, 20 (27%) were diagnosed de-novo with aCLL, while 53 (73%) evolved from CLL after a **median time of 79 months** (range: 1.4 – 267).

Compared to CLL, aCLL showed more frequent lymph nodes >5 cm, extranodal disease, systemic symptoms and higher LDH and β 2-microglobulin leves. aCLL was also enriched in patients with *TP53* disruption and U-IGHV status

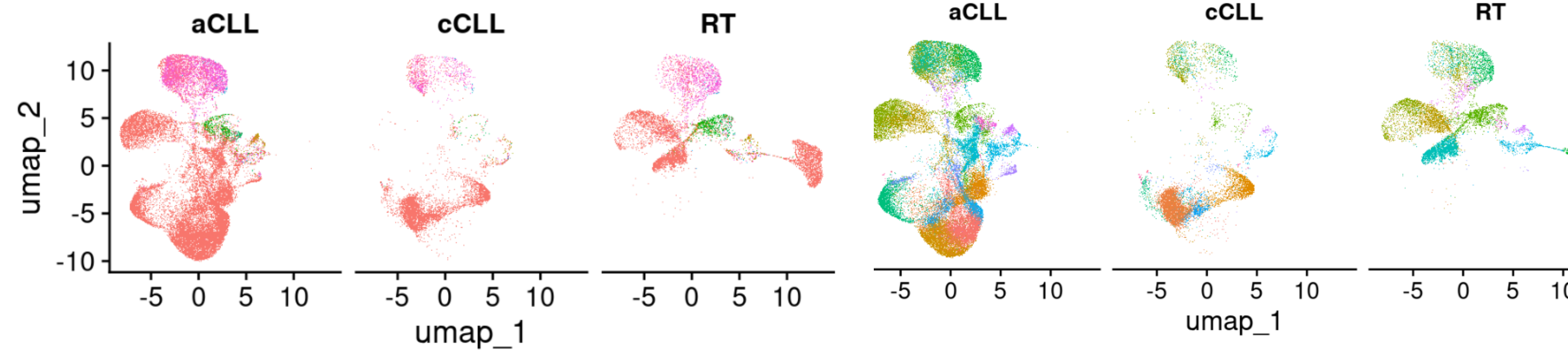


aCLL also showed:

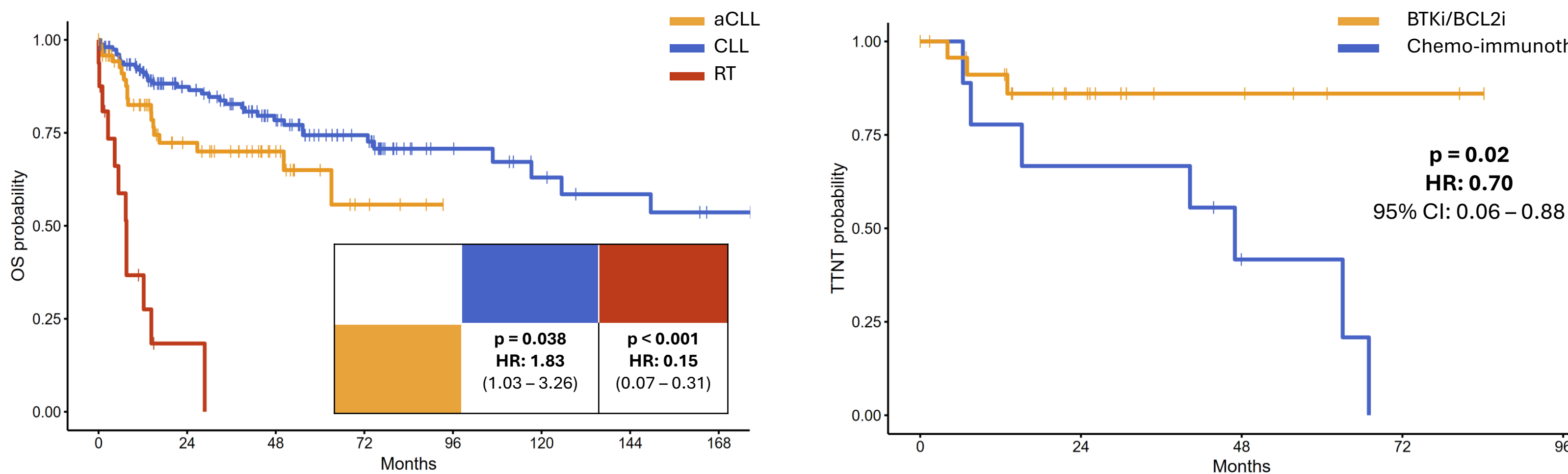
- Intermediate Ki67 expression levels (median 14% vs. 6% in CLL and 70% in RT)
- Intermediate 18-FDG PET/CT SUV-max (median 7.3 vs. 4.0 in CLL and 15.4 in RT)



aCLL tissue was comprised of multiple different clusters of B-cells copared to CLL



aCLL showed intermediate OS between CLL and RT and better outcomes when treated with BTKi/BCL2i



Conclusions

aCLL could represent a distinct biological entity enriched in adverse molecular alterations and proliferative subclones driving worse survival outcomes