



XIX CONGRESSO NAZIONALE SIES 2026

**IMMUNE CELLS AND EXTRACELLULAR VESICLE-BASED LIQUID BIOPSY REVEAL PROGNOSTIC
AND IMMUNOMETABOLIC BIOMARKERS OF EARLY RESPONSE TO AZACITIDINE AND
VENETOCLAX IN ACUTE MYELOID LEUKEMIA**

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Firenze | 4-6 marzo 2026
Palazzo degli Affari

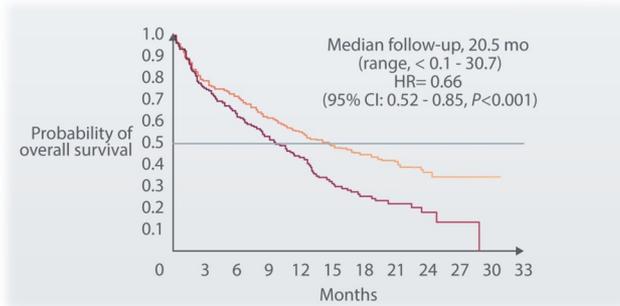


Disclosures of Federico Pasini

Company name	Research support	Employee	Consultant	Stockholder	Speakers bureau	Advisory board	Other
Nothing to declare							

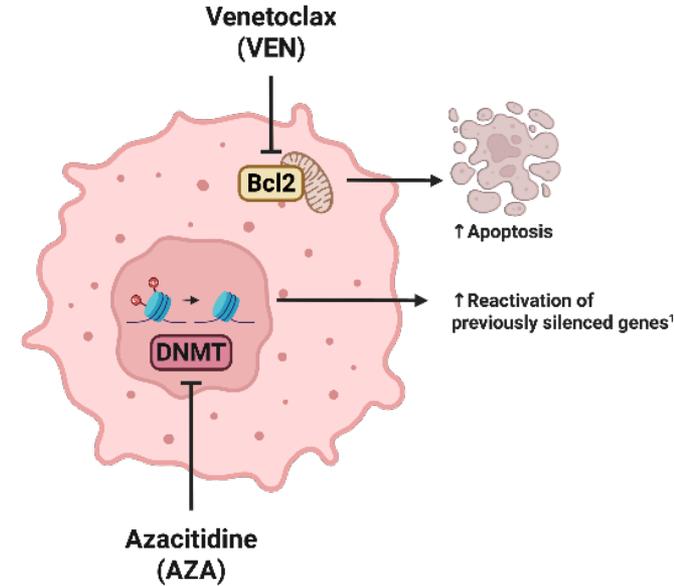
Azacitidine + Venetoclax in Acute Myeloid Leukemia: Innovative, though not without limitations

- Acute myeloid leukemia (AML) is an aggressive malignancy with high relapse rate
- Treatment with the oral BCL2 inhibitor venetoclax (VEN) represents the current standard-of-care for AML patients unfit for intensive chemotherapy, often associated to Azacitidine (AZA)

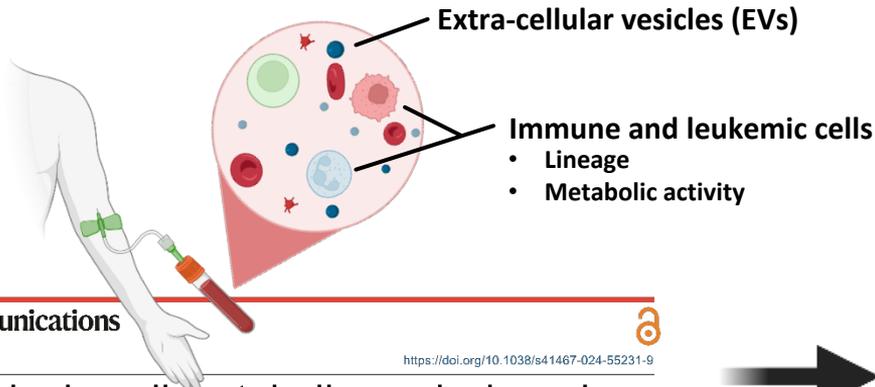


● AZA + VEN
● AZA

Forsberg et al. Haematologica 2024



- The metabolism of leukemic stem cells (LSCs) plays a pivotal role in leukemogenesis, influencing relapse and shaping therapeutic response.



nature communications

Article

<https://doi.org/10.1038/s41467-024-55231-9>

Parallel single-cell metabolic analysis and extracellular vesicle profiling reveal vulnerabilities with prognostic significance in acute myeloid leukemia

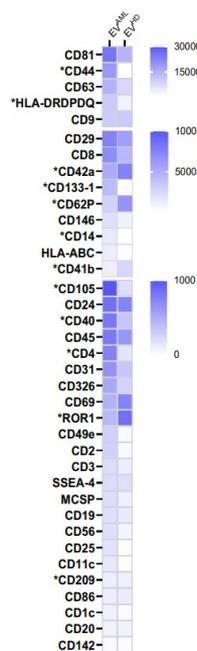
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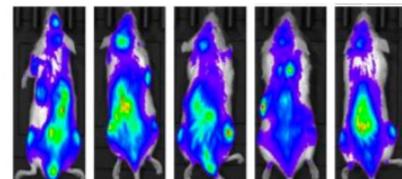
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Check for updates

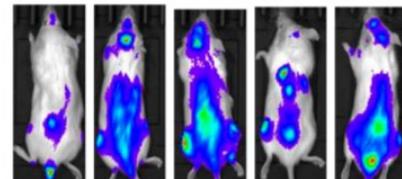
Dorian Forte ¹, Roberto Maria Pellegrino ², Paolo Falvo ^{3,4}, Paulina Garcia-Gonzalez⁵, Husam B.R. Alabed², Filippo Maltoni^{1,11}, Davide Lombardi^{3,4}, Samantha Bruno ¹, Martina Barone ⁶, Federico Pasini ¹, Francesco Fabbri⁷, Ivan Vannini⁷, Benedetta Donati⁸, Gianluca Cristiano¹, Chiara Sartor^{1,11}, Simona Ronzoni⁹, Alessia Ciarrocchi ⁶, Sandra Buratta², Lorena Urbanelli ², Carla Emiliani^{2,10}, Simona Soverini¹, Lucia Catani^{1,11}, Francesco Bertolini ^{3,4}, Rafael José Argüello ⁶, Michele Cavo^{1,11} & Antonio Curti ⁶ ✉



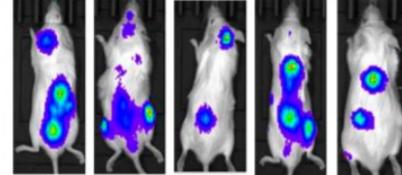
MOLM-13+ adv EV^{AML}



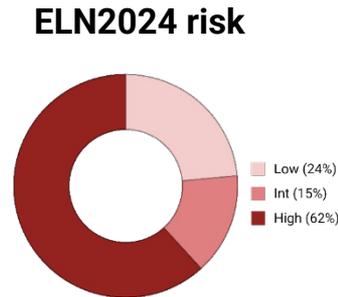
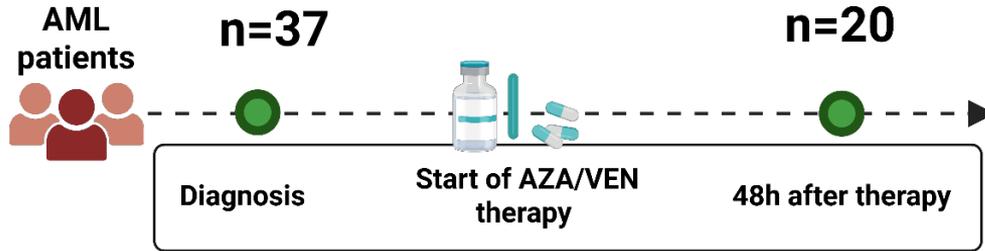
MOLM-13+ inter EV^{AML}



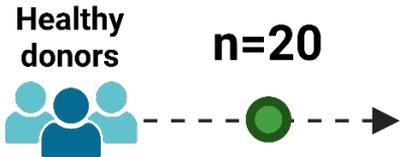
MOLM-13+ fav EV^{AML}



Can we use liquid biopsy at early timepoints to monitor patients' response to AZA/VEN in AML?

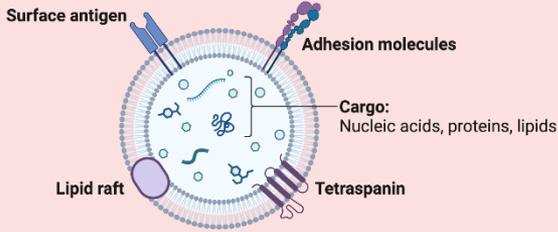


Mutational profile



Extracellular Vesicles (EVs) Represent a Promising Tool for Liquid Biopsy in AML

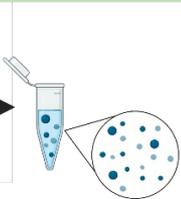
Extracellular vesicles in blood



- Actively released by viable cells
- "Cell Biopsy" ¹
- Abundant in blood (10^4 from 1 cell)
- Stable in blood



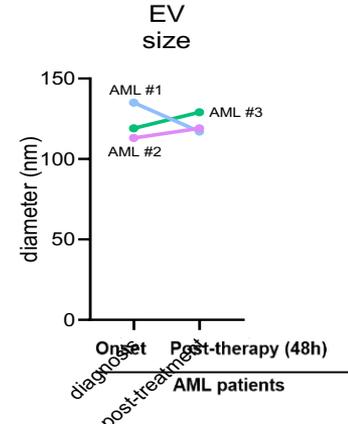
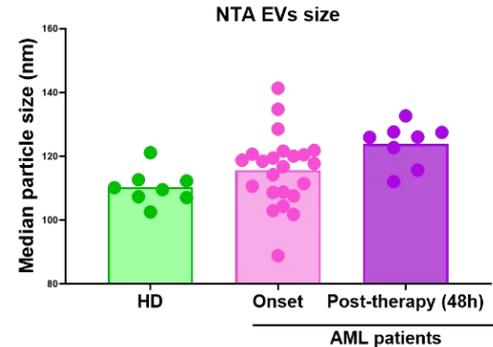
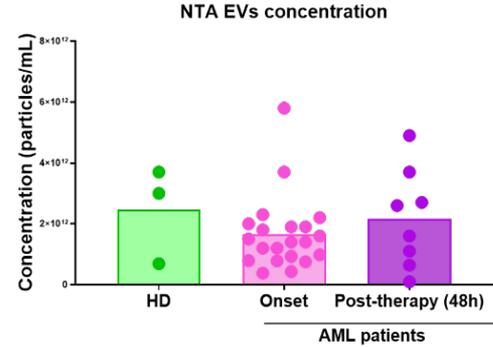
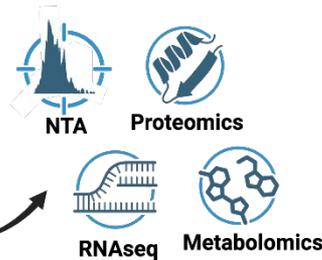
Size-exclusion chromatography



Isolated EVs

EV-TRACK ID: EV230004

EVs characterization

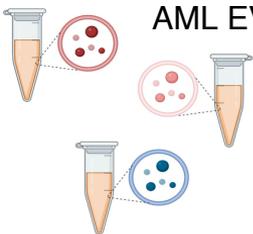




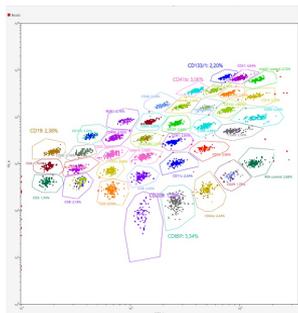
Multiplex Proteomic Profiling of AML-Derived EVs at Onset Reveals Prognostic Markers

AML EVs (T0)=18

AML EVs (48h)=18

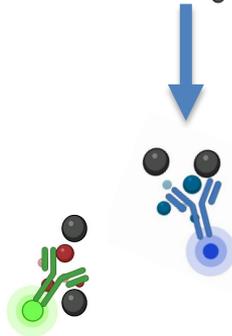
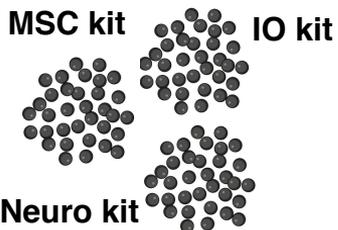


HD EVs=10



Spectral flow cytometry analysis

EV capture beads



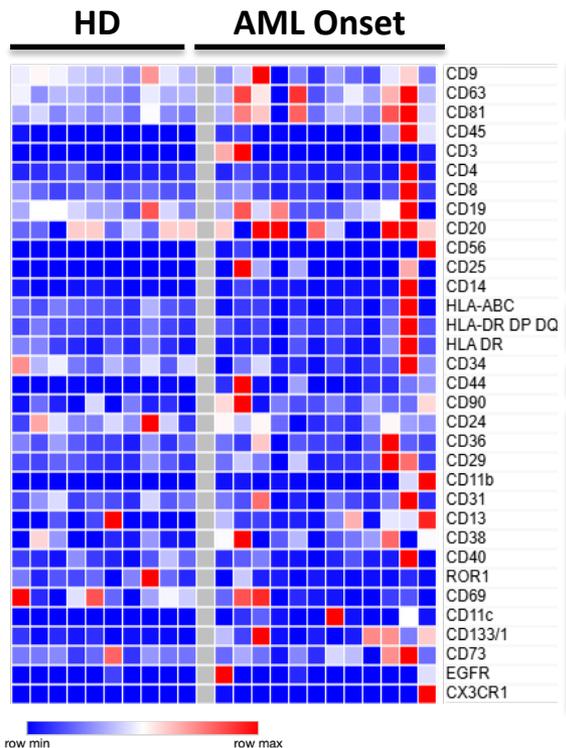
AML patients according to ELN2024

Adverse	Intermediate	Favorable	id
Blue	Red	Red	CD54
Blue	Red	White	CD44
Blue	White	Red	CD81
Blue	White	Red	CD63
Blue	White	Red	HLA-DR DP DQ
Blue	White	Red	HLA DR
Blue	White	Red	HLA-ABC
Blue	White	Red	CD29
Blue	White	Red	CD8
Blue	White	Red	CD36

row min row max



Multiplex Proteomic Profiling of EVs show a specific profile in AML compared to HD



**Tetraspanins
"EV Markers"**

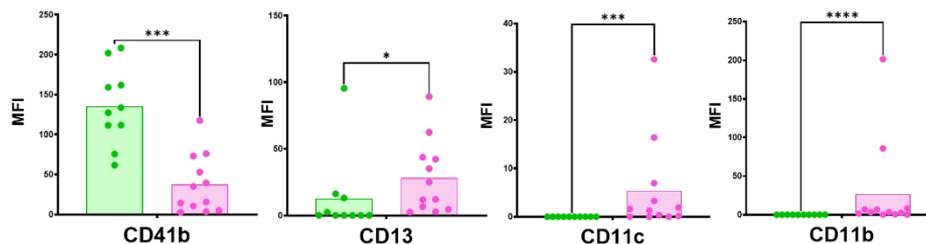
**Immune
lineages**

HLA markers

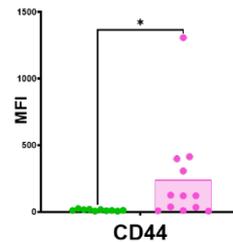
Blasts markers

**Immuno-metabolic
markers**

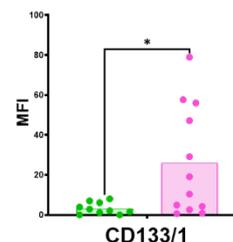
Differentiation and lineage markers



LSCs



Stemness

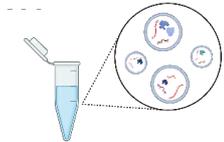


● HD

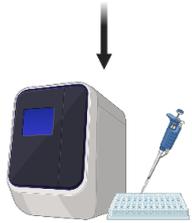
● AML Onset



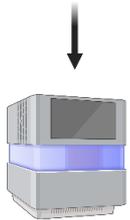
EV RNA sequencing highlights a T cell and metabolic involvement



EVs RNA extraction



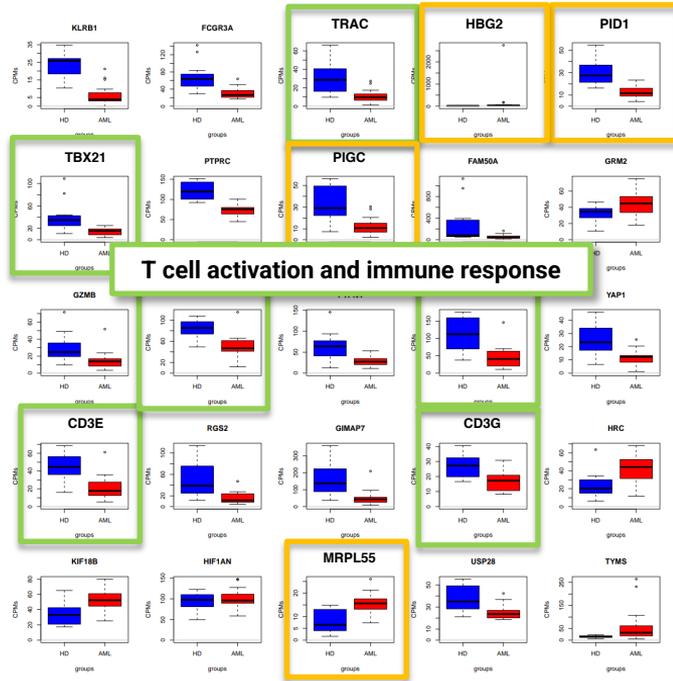
cDNA library preparation



Sequencing

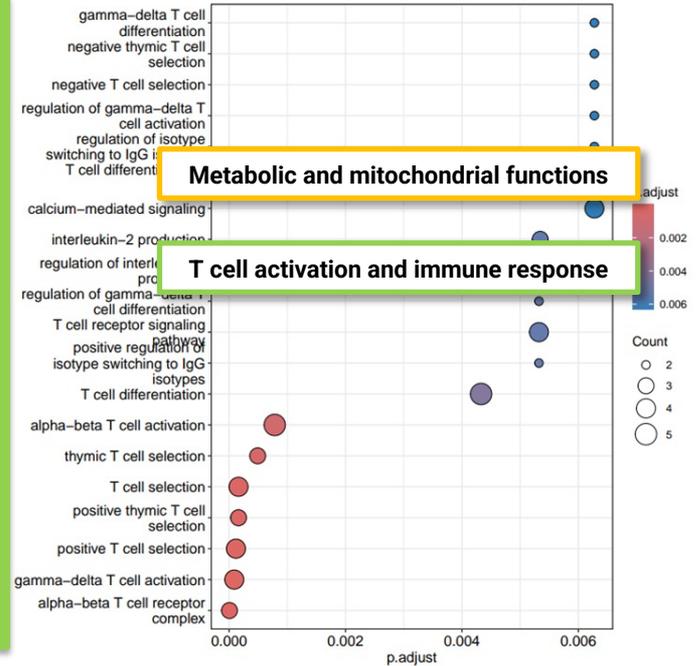
N = 12 HD vs 25 AML ONSET

Gene expression



T cell activation and immune response

Over-Representation Analysis (ORA)



Metabolic and mitochondrial functions

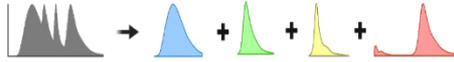
T cell activation and immune response

RNA analysis performed by F. Maltoni

Multi-parametric spectral flow cytometry allows to detect rare cell subsets

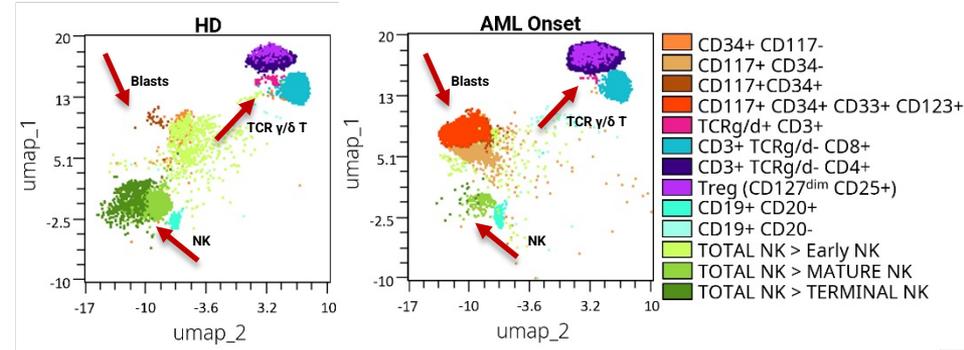


SONY ID7000 spectral flow cytometry



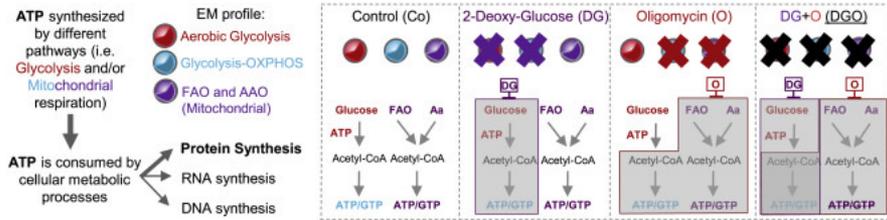
36 colors in a single tube (spectral unmixing)

Immunophenotype

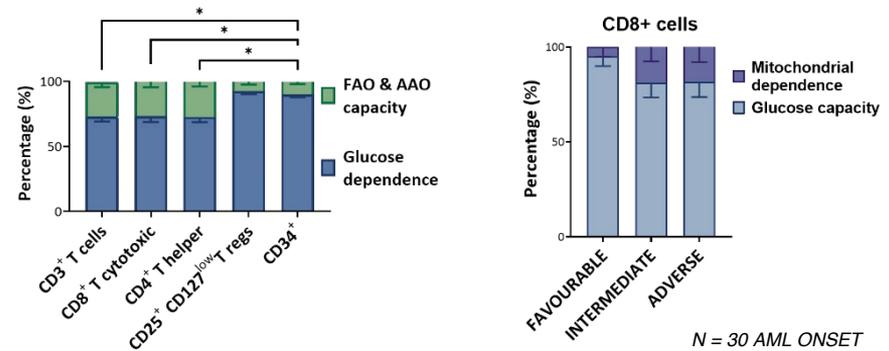


SCENITH¹

(Single Cell Energetic metabolism by profiling Translation inhibition)



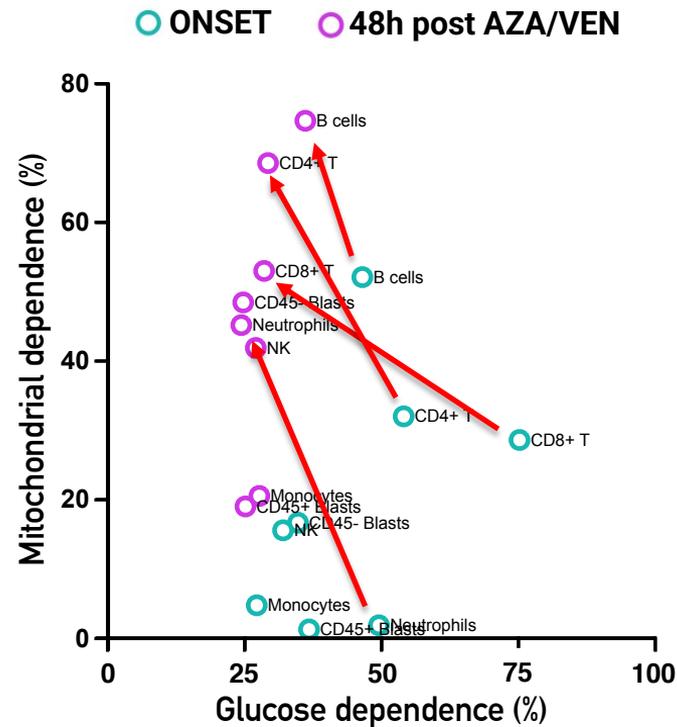
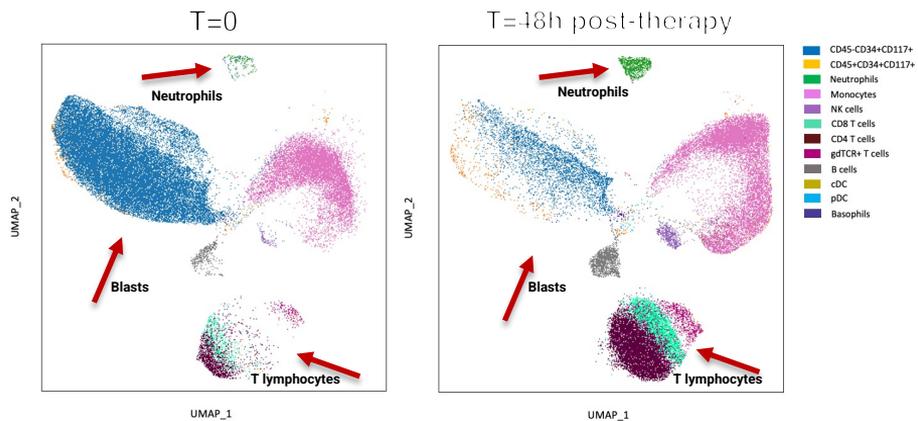
SCENITH metabolic analysis by cell type or risk





Immuno-metabolic shift after 48h of AZA/VEN therapy

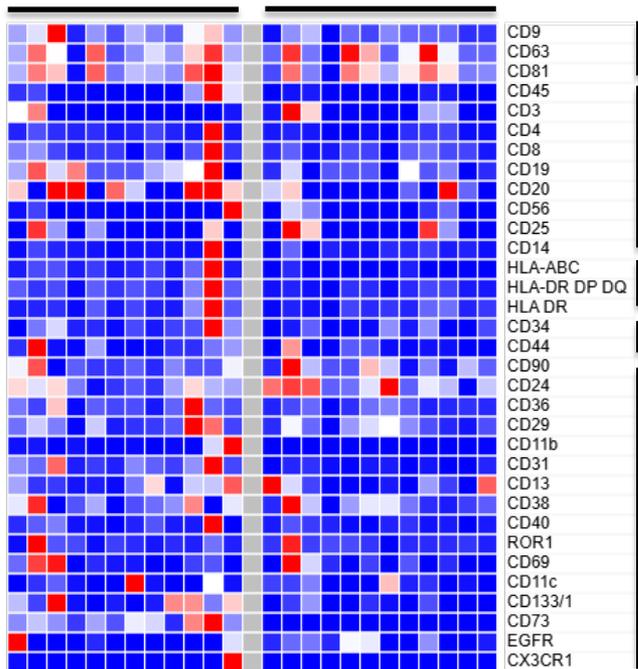
SCENITH analysis performed on whole peripheral blood
 -> Analysis in real-time



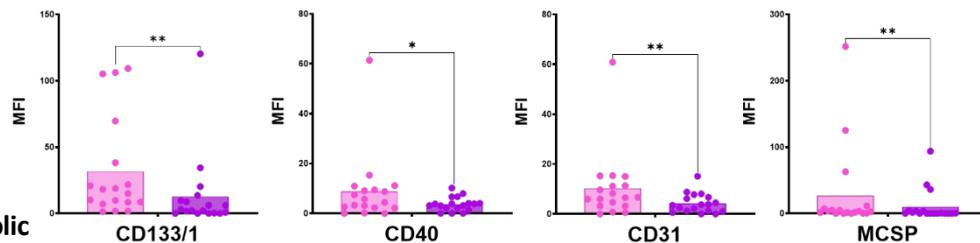
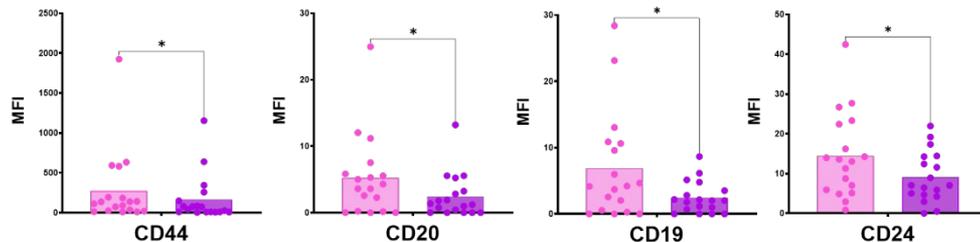


AML EVs 48h post AZA/VEN show a decreased level of immune-associated markers

Onset Post-therapy



row min row max

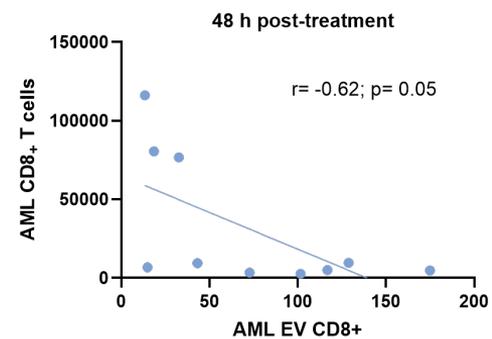
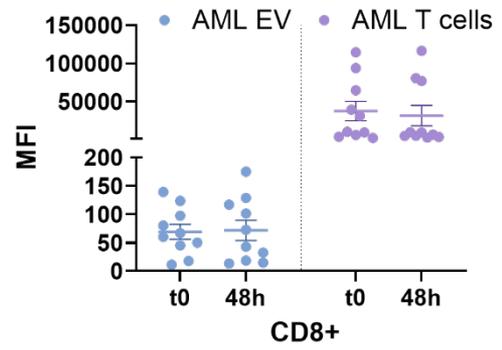
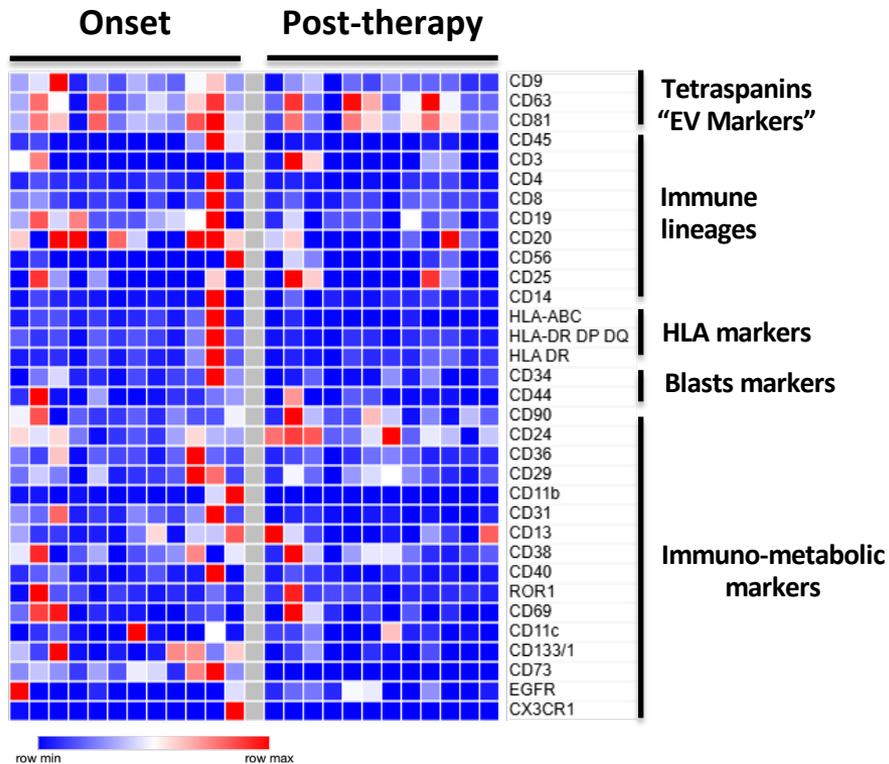


● ONSET

● 48h post AZA/VEN



AML EVs 48h post AZA/VEN show a decreased level of immune-associated markers



Conclusions

- Liquid biopsy provides useful insights in the monitoring of therapy response at early timepoints
- Immune cells metabolism at diagnosis shows prognostic value and after therapy shows a different metabolic profile
- AML EV surface markers correlate with ELN risk classification and show a different profile after therapy at early timepoints, suggesting an immune involvement

Clinical potential: Integrating EV signatures with immune-metabolic crosstalk may reveal novel prognostic biomarkers and allow to monitor therapy response

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Read our paper on EVS
from AML at Onset

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IRCCS Istituto di ricovero e cura a carattere scientifico

