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64th ASH Annual Meeting Abstracts

## **POSTER ABSTRACTS**

# 623.MANTLE CELL, FOLLICULAR, AND OTHER INDOLENT B CELL LYMPHOMAS: CLINICAL AND EPIDEMIOLOGICAL

**Dissecting MRD Kinetics By Automated Computational Analysis to Improve Outcome Prediction in Mantle Cell Lymphoma: A Bioinformatic Substudy from the Fondazione Italiana Linfomi (FIL) MCL0208 Clinical Trial** Francesca Cordero<sup>1,\*</sup>, Simone Ferrero, MD<sup>2,3</sup>, Simone Pernice<sup>1,\*</sup>, Elisa Genuardi<sup>3,\*</sup>, Roberta Sirovich<sup>4,\*</sup>, Beatrice Alessandria<sup>3,\*</sup>, Andrea Evangelista<sup>5,\*</sup>, Simone Ragaini<sup>2,3,\*</sup>, Maurizio Martelli, MD<sup>6,\*</sup>, Alice Di Rocco, MD<sup>6,\*</sup>, Alessandro Re, MD<sup>7,\*</sup>, Chiara Pagani, MD<sup>8,\*</sup>, Vittorio Stefoni, MD<sup>9,\*</sup>, Federica Cavallo, MD<sup>3,2,\*</sup>, Carola Boccomini<sup>10,\*</sup>, Monica Balzarotti, MD<sup>11,\*</sup>, Vittorio Ruggero Zilioli, MD<sup>12,\*</sup>, Maria Gomes da Silva<sup>13,\*</sup>, Luca Arcaini<sup>14,\*</sup>, Melania Celli<sup>15,\*</sup>, Gian Maria Zaccaria<sup>16,\*</sup>, Dora Tortarolo<sup>1,\*</sup>, Sergio Cortelazzo<sup>17,\*</sup>, Marco Ladetto<sup>18</sup>

<sup>1</sup>Department of Computer Science, University of Torino, Torino, Italy

<sup>2</sup>Division of Hematology, AOU Città della Salute e della Scienza di Torino, Torino, Italy

<sup>3</sup>Department of Molecular Biotechnologies and Health Sciences, Division of Hematology, University of Torino, Torino, Italy <sup>4</sup>Department of Mathematics G. Peano, University of Torino, Torino, Italy

<sup>5</sup>Unit of Cancer Epidemiology, CPO Piemonte, A.O.U. Città della Salute e della Scienza di Torino, Torino, Italy

<sup>6</sup>Hematology, Department of Translational and Precision Medicine, Azienda Ospedaliera Policlinico Umberto I, Sapienza University of Rome, Roma, Italy

<sup>7</sup>Hematology Division, ASST-Spedali Civili di Brescia, Brescia, Italy

<sup>8</sup>Department of Hematology, ASST Spedali Civili, Brescia, Italy

<sup>9</sup>Hematology, Department of Translational and Precision Medicine, Università di Bologna, Bologna, Italy

<sup>10</sup>Division of Hematology 2, AOU Città della Salute e della Scienza di Torino, Torino, Italy

<sup>11</sup>Unit of Hematology, Humanitas Clinical and Research Center, Rozzano, Italy

<sup>12</sup>Division of Hematology, ASST Grande Ospedale Metropolitano Niguarda, Milan, Italy

<sup>13</sup>Division of Hematology, Instituto Português de Oncologia de Lisboa, Lisbon, Portugal

<sup>14</sup>Division of Hematology, Fondazione IRCCS Policlinico San Matteo, Pavia, Italy

<sup>15</sup>U.O. di Ematologia, Ospedale degli Infermi di Rimini, Rimini, Italy

<sup>16</sup>Hematology and Cell Therapy Unit, IRCCS Istituto Tumori 'Giovanni Paolo II', Bari, Italy

<sup>17</sup>Oncology unit, Clinica Humanitas/Gavazzeni, Bergamo, Italy

<sup>18</sup>Division of Hematology, AO SS Antonio e Biagio e Cesare Arrigo, Alessandria, Italy

\*Asterisk with author names denotes non-ASH members.

**Abstract Background and Aims.** Minimal residual disease (MRD) detection is a validated outcome predictor in mantle cell lymphoma (MCL). Recently, the clinical relevance of repeated MRD monitoring has been described in a prospective clinical trial [Ferrero, Blood 2022]. Nonetheless, complex patterns of MRD kinetics over time and availability of results obtained by heterogeneous tissues (bone marrow, BM, vs peripheral blood, PB) might generate substantial interpretation issues and hamper an easy-to-use application of this predictive biomarker. To overcome the limitation of empirical analysis we tested CONNECTOR, a novel automated computational framework to facilitate the interpretation of MRD kinetics and stratify patients in risk classes based on a solid, algorithm-derived, classification.

**Patients and methods.** ASO RQ-PCR MRD data already generated from BM and PB samples in the FIL MCL0208 trial [Ferrero, Blood 2022], offering first-line high dose chemoimmunotherapy and autologous transplantation (ASCT) to youngers MCL patients, were employed. CONNECTOR (https://qbioturin.github.io/connector/) is a data-driven framework based on Functional Data Analysis for analyzing longitudinal high-dimensional data in a straightforward and revealing way. CONNECTOR allows the aggregation of time-series data through an unsupervised approach in informative clusters. Moreover, we developed a classification approach to predict the risk class for new MRD data. The new MRD curves were allocated to existing risk clusters defined on a training set of MRD data.

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Results. Without knowledge of any clinical or outcome data, CONNECTOR was run on 117 patients characterized by at least 3 MRD timepoints in BM (median 6, 3-9) and separately on 95 patients with at least 4 MRD timepoints in PB (median 7, 4-9). Starting from BM MRD data, CONNECTOR identified four patient risk Clusters (CrC), based on MRD kinetics, Figure1A. CrC\_A comprised 73 patients with an overall quick and stable MRD negativization; CrC\_B accounted for 14 patients characterized by later MRD negativization, alternating MRD results, or low-level MRD persistence; the 14 patients in CrC\_C were always characterized by early MRD negativization followed by late MRD reappearance; CrC\_D was composed of 16 patients with stable MRD positivity or only transient MRD negativization followed by early MRD reappearance. The different trends of MRD kinetics dissected by CONNECTOR fairly predicted patients' outcomes: median TTP was not reached for CrC\_A and B, 36 months for CrC\_C and 27 months for CrC\_D, respectively (p<0.0001), Figure1B. Similar data were generated when CONNECTOR was run on MRD results from PB samples: CrC allocation of BM vs PB was concordant for 89% of patients and survival analysis on PB results confirmed patients stratification in two groups: favorable MRD kinetics (CrC\_A and B, median TTP not reached) vs unfavorable MRD kinetics (CrC\_C and D, median TTP 35 months, p<0.0001), as depicted in Figure1B. Finally, we tested the performance of pooling together BM and PB results, to increase the number of evaluable patients (irrespective of eventual missing BM samples). Only the BM point at 12 months after ASCT was considered, together with the PB MRD results of all the available time points. CONNECTOR was used to verify if this mixed MRD pattern was able to correctly reclassify patients in the same CrC. Actually, 89 out of 95 patients (94%) were correctly reclassified (p\_mean=0.96, p\_sd= 0.07).

**Discussion.** CONNECTOR allowed unsupervised identification of patients clusters with distinct MRD kinetics and highly significant different clinical outcome. Such clusterization proved effective using BM, PB and mixed tissues, as well. Validation of CONNECTOR performance in large, independent prospective trials is currently ongoing.







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