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POSTER ABSTRACTS

621.LYMPHOMAS: TRANSLATIONAL-MOLECULAR AND GENETIC

Prospective Evaluation of Minimal Residual Disease in Waldenström Macroglobulinemia across Different Tissues and Treatments: Results of the "BIO-WM" Trial of the Fondazione Italiana Linfomi (FIL) Martina Ferrante¹, Daniela Drandi¹, Silvia Zibellini², Luigi Marcheselli³, Chiara Varraso², Veronica Peri, MD^{4,5}, Irene Dogliotti, MD⁵, Davide Musto⁴, Emilia Cappello⁶, Federica Cavallo^{4,7}, Angela Ferrari⁸, Michele Merli⁹, Giulia Zamprogna¹⁰, Luca Laurenti, MD¹¹, Simona Tomasetti¹², Emanuele Cencini, MD¹³, Giacomo Loseto, MD¹⁴, Silvia Finotto, MD¹⁵, Monia Marchetti, MD¹⁶, Francesca Re, MD¹⁷, Antonello Sica, MD¹⁸, Jacopo Olivieri¹⁹, Giulia Vittoria Facchetti⁶, Cristina Jiménez²⁰, Noemi Puig, MDPhD²⁰, Giulia Turra⁶, Benedetto Bruno, MD PhD^{5,21}, Ramon Garcia-Sanz, MD PhD²⁰, Marzia Varettoni²², Simone Ferrero, MD^{5,4} ¹ Division of Hematology, Department of Molecular Biotechologies and Health Sciences, University of Torino, Torino, Italy ²Division of Hematology, Fondazione IRCCS Policlinico San Matteo, Pavia, Italy ³Fondazione Italiana Linfomi, Clinical Trial Office, Modena, Italy ⁴ Division of Hematology, Department of Molecular Biotechnologies and Health Sciences, University of Torino, Torino, Italy ⁵ Division of Hematology and Stem Cell Transplant Unit, University Hospital AOU Città della Salute e della Scienza, Torino, Italv ⁶Department of Molecular Medicine, University of Pavia, Pavia, Italy ⁷ Division of Hematology and Stem Cell Transplant Unit, AOU Città della Salute e della Scienza, Torino, Italy ⁸IRCCS - Arcispedale Santa Maria Nuova, Hematology, ITA, Reggio Emilia, Italy ⁹Ospedale di Circolo e Fondazione Macchi, Varese, Italy ¹⁰Department of Hematology, Niguarda Cancer Center, ASST Grande Ospedale Metropolitano Niguarda, Milano, Italy ¹¹S. Ematologia, Dipartimento Scienze Radiologiche Radioterapiche ed Ematologiche, Fondazione Policlinico Universitario A. Gemelli IRCCS, Rome, Italy ¹²Ematologia, Ospedale degli Infermi di Rimini, Rimini, Italy ¹³Hematology, Azienda Ospedaliera Universitaria Senese & University of Siena, Siena, Italy ¹⁴IRCCS Istituto Tumori "Giovanni Paolo II", Hematology Unit, Bari, Italy ¹⁵Oncologia1, Dipartimento di Oncologia, Istituto Oncologico Veneto IRCCS, Padova, Italy ¹⁶Ematologia, Ospedale Civile SS Antonio e Biagio e Cesare Arrigo, Alessandria, Italy ¹⁷UO Ematologia e CTMO, Azienda Ospedaliera Universitaria di Parma, Parma, Italy ¹⁸Oncologia Medica ed Ematologia, AOU Università degli Studi della Campania Luigi Vanvitelli, Naples, Italy ¹⁹Clinica Ematologica, Centro Trapianti e Terapie Cellulari "Carlo Melzi", Azienda Sanitaria Universitaria Integrata di Udine, Udine, Italy ²⁰ Hospital Universitario de Salamanca (HUSAL), IBSAL, IBMCC (USAL-CSIC), CIBERONC, Salamanca, Spain ²¹ Division of Hematology, Department of Molecular Biotechologies and Health Sciences, University of Torino, Division of Hematology and Stem cell transplant unit, AOU Città della Salute e della Scienza, Torino, Italy ²²Division of Hematology, Fondazione IRCCS Policlinico San Matteo, Pavia, Italy Introduction. MYD88 ^{L265P} is the hallmark mutation in Waldenström Macroglobulinemia (WM) and is becoming increasingly important in the management of IgM-gammopathies, due to its role as prognostic and predictive biomarker. Recently, MYD88

^{1265P} detection by allele-specific quantitative PCR was proposed as reliable minimal residual disease (MRD) marker in bone marrow (BM) samples (Varettoni, Hematol Oncol 2022). Novel, more sensitive, techniques as droplet digital PCR (ddPCR) might extend the feasibility of MRD detection in WM also in non-invasive tissues, as peripheral blood (PB) or plasmatic cellfree (cf) DNA. This was a secondary endpoint of the multicenter, observational "BIOWM" (NCT03521596) trial, sponsored by the Fondazione Italiana Linfomi (FIL) and the International WM Foundation/Leukemia and Lymphoma Society. From 2018 to 2020 this trial enrolled 300 consecutive patients with primary diagnosis of WM or IgM-MGUS and a systematic biobanking was performed. Here are presented the first results of the MRD study on the patient subset who received frontline treatment, with the aim of driving correlations with clinical response, type of therapy received and outcome prediction. **Methods.** Paired

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BM, PB and plasma samples were collected for each patient at baseline (TO) and follow-up and MYD88 L265P detection was performed by ddPCR in all specimens (Drandi, Haematologica 2018). As of today, 59/300 patients (58 WM and 1 IgM-MGUS) received frontline treatment and MRD levels after therapy (T2) were detected by ddPCR: actually, T0-T2 paired samples were available in 49 (BM), 56 (PB) and 48 (cfDNA) cases, respectively. Moreover, MRD was evaluated also by 8-color multiparametric flow cytometry (MFC) in 23 BM and 24 PB samples with a minimum of 10 million cells acquired. Results. Median age of the 59 treated patients was 68 years (24-85), 32% were female, median IqM value was 2420 mg/dL (101-8840), median Hb 10.5 g/dL (8 -17) and IPSS-WM was high in 35% and intermediate in 46%. MYD88 ^{L265P} mutation rate was 94% (46/49) in BM, 80% (45/56) in PB and 90% (43/48) in cfDNA samples, respectively. Treatment was started because of anemia in 32% cases, lymphadenopathy or splenomegaly in 34%, hyperviscosity in 20%, other reasons in 13%, and the only MGUS patient required treatment for antimyelin-associated glycoprotein polyneuropathy. Thirty-one out of 59 patients received bendamustine-rituximab (BR), 23/59 dexamethasone, rituximab and cyclophosphamide (DRC) or DRC-like regimen and 5/59 a single agent therapy (namely, 3 rituximab, 1 cyclophosphamide, 1 ibrutinib). Overall response rate was 90% (87% for BR and 96% for DRC), with 19% CR, 29% VGPR and 35% PR in BR and 4% CR, 0% VGPR and 91% PR in DRC, respectively. Overall, MRD negativity after treatment was 30% (14/46) in BM, 89% (40/45) in PB and 54% (23/43) in cfDNA, respectively. Moreover, among patients still MRD positive after treatment, median MYD88 ^{L265P} tumor burden shrinkage was about 2 Logs in BM (5.5E-03 vs 2.1E-01) and around 1 Log in cfDNA (5.9E-03 vs 3.7E-02) and in PB samples (3.6E-03 vs 2.5E-02), respectively (Figure 1). Interestingly, the MRD shrinkage in BM was deeper among patients receiving BR (48% MRD negative with a median MYD88 ^{L265P} decrease of more than 2 Logs among patients still MRD positive) vs patients receiving DRC (10% MRD negative with a median decrease of 1 Log), Figure 2. Similar trends in favor of BR in inducing MRD negativity were reported in PB (96% vs 88%) and cfDNA (70% vs 35%), respectively. MRD results by MFC were overall in line with ddPCR results: namely 27% of patients reached MRD negativity in BM and 69% in PB. The median follow-up for the whole series was 41 months, resulting in a 3-years PFS of 71% and 3-years OS of 89%, respectively. Interestingly, despite the current limited follow-up for such an indolent disease, MRD positivity by ddPCR in PB predicted a dismal clinical outcome if compared with MRD negative patients (3-years PFS 40% vs 73%, p=0.038). Conclusions. To the best of our knowledge, this is the first report of prospective MRD evaluation in a WM clinical trial. Our preliminary results suggest that MYD88 ^{1265P} monitoringby ddPCR is a suitable target for MRD analysis, in the contest of common immunochemotherapeutic schedules. Moreover, different levels of disease persistence were described across BM, PB and cfDNA, with MRD monitoring in non-invasive tissues showing promising predictive value for outcome discrimination.

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Figure 1. MRD shrinkage across different tissues. *MYD88*^{L265P} values in bone marrow (BM), peripheral blood (PB) and plasma cfDNA (PL) at baseline (T0) and after treatment (T2) are depicted. MRD, minimal residual disease; DRC, dexamethasone, rituximab and cyclophosphamide.



Figure 2. Waterfall plot describing MRD shrinkage in BM according to treatment received. Decrease in *MYD88*^{L265P} values in bone marrow (BM) after therapy is depicted. Best response rates from R-Bendamustine and DRC are indicated for each patient. MRD, minimal residual disease; DRC, dexamethasone, rituximab and cyclophosphamide; CR, complete response; VGPR, very good partial response; PR, partial response; MR, minor response; SD, stable disease; PD, progressive disease.



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