





Blood 142 (2023) 2581-2582

The 65th ASH Annual Meeting Abstracts

POSTER ABSTRACTS

311.DISORDERS OF PLATELET NUMBER OR FUNCTION: CLINICAL AND EPIDEMIOLOGICAL

Clonal Hematopoiesis in Patients with Immune Thrombocytopenia: Preliminary Results of a Cross-Sectional Study Alessandro Bosi¹, Marta Canzi², Marco Capecchi, MD³, Juri Alessandro Giannotta, MD⁴, Andrea Artoni⁵, Wilma Barcellini, MD⁶, Bruno Fattizzo⁷

- ¹ Department of Oncology and Hemato-Oncology, University of Milan, Milan, Italy, Milan, Italy
- ²Department of Oncology and Hemato-Oncology, University of Milan, Milan, Italy, Milan, ITA
- ³Clinica Moncucco, Division of Hematology, Lugano, Switzerland
- ⁴Angelo Bianchi Bonomi Hemophilia and Thrombosis Center, Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico, Milano, ITA
- ⁵ Angelo Bianchi Bonomi Hemophilia and Thrombosis Center, Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico, Milan, ITA
- ⁶UOC Ematologia, UOS Fisiopatologia delle Anemie, Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico di Milano, Milan, Italy
- ⁷ Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico, Milano, ITA

Immune thrombocytopenia (ITP) is an acquired disorder marked by immune-mediated attack to platelets. However, the borders between ITP and clonal cytopenias, such as low risk myelodysplastic syndromes, may be difficult to establish. Detection of somatic mutations on peripheral blood leukocytes (PBL) through next generation sequencing (NGS) is gaining diagnostic and prognostic power in myeloid disorders and may be found also in individuals without blood malignancies, namely clonal hematopoiesis with indeterminate potential (CHIP). The prevalence and significance of CHIP in immune-mediated hematological diseases, such as ITP, is not clearly known.

Here we present the preliminary results of a cross-sectional study on adult ITP patients followed at our clinics from 2019 until March 2023. History of any hematological malignancy was an exclusion criterion. An NGS study on PBL was performed by Ion Torrents5, Ion Reporters software 5.2, evaluating mutations in 69 potentially oncogenic genes present in the Oncomine Myeloid Research Assay diagnostic panel. Only mutations with a variant allele frequency (VAF) of 2% were considered, while mutations with VAF > 40% were regarded as potentially germline and discarded. To correlate mutational status with clinical features, a two-tailed Fisher's exact test was used for analysis of dichotomic variables, while a two-sided student's t was used for continuous ones.

We included 37 ITP patients (Table 1), 14 of which (38%) had at least one mutation. After exclusion of 4 mutations with VAF > 40%, a total of 17 mutations were considered, affecting 11/37 (30%) patients (NGS-positive). Mutated genes were TET2 (7/17, 41%), DNMT3A (4/17, 18%), SRSF2 (2/17, 12%), and SF3B1, MPL, ASXL1 and NF1 in 1/17 (6%) case each. All patients, except two, had received at least one treatment line (steroids +/- intravenous immunoglobulins), 32 (86%) and 14 (38%) subjects 2 or 3 lines, respectively.

Median age of NGS-positive was significantly higher than NGS-negative patients: 73 years, interquartile range (IQR) 68 - 76 vs 62 years, IQR 52 - 70 (p = 0.02). NGS-positive had a slightly lower rate of complete response (CR; PLT > 100×10^9 /L) after steroid treatment, compared with NGS-negative patients (40% vs 60%, respectively). Among the 9 NGS-positive patients who received thrombopoietin receptor agonists (TPO-RA), only 2 were evaluated before, and the remaining after a median of 2.6 years from TPO-RA start. Considering the 9 patients treated with TPO-RA, NGS-positive had a slightly lower rate of CR than NGS-negative patients (44% vs 79%, respectively). In this respect, we did not find substantial disparities in distribution or VAF of gene mutations between patients who fully responded to TPO-RA and those failing to achieve CR.

With the caveat of a cross-sectional analysis in a limited number of cases, these data show the presence of CHIP in 1/3 of multi-treated ITP patients, a frequency similar to that of a much more elderly Italian population >80-year-old (Rossi et al, Blood. 2021), suggesting a possible role of autoimmunity/inflammation and treatment in clonal selection.

Disclosures Barcellini: Alexion, AstraZeneca Rare Disease: Consultancy, Membership on an entity's Board of Directors or advisory committees, Research Funding; Novartis: Consultancy, Honoraria, Speakers Bureau.

Table 1: Clinical features and response to treatment of ITP patients with or without CHIP

	All (n = 37)	NGS-positive (n = 11)	NGS-negative (n = 26)
Male sex	20/37 (54%)	8/11 (73%)	12/26 (46%)
Median age (IQR) at NGS evaluation, years	67 (54 - 73)	73 (68 - 76)	62 (52 - 70)
Evans syndrome	5/37 (14%)	2/11 (18%)	3/26 (16%)
Secondary ITP	5/37 (14%)	2/11 (18%)	3/26 (16%)
Median PLT at diagnosis (IQR), x 10^9/L	20 (5 - 56)	29 (26 - 57)	19 (5 - 53)
n° of treatment lines (IQR)	2 (2 - 3)	2 (2 - 4)	2 (2 - 3)
Steroid treatment	35/37 (95%)	10/11 (91%)	25/26 (96%)
Steroid response PLT>50 x 10^9/L	28/35 (80%)	8/10 (80%)	20/25 (80%)
Steroid response PLT>100 x 10^9/L	19/35 (54%)	4/10 (40%)	15/25 (60%)
Splenectomy	6/37 (16%)	2/11 (18%)	4/26 (17%)
Any TPO-RA treatment	28/37 (76%)	9/11 (82%)	19/26 (73%)
Eltrombopag	26/37 (70%)	8/11 (73%)	18/26 (69%)
Romiplostim	5/37 (14%)	2/11 (18%)	3/26 (12%)
TPO-RA response PLT>50 x 10^9/L	25/28 (89%)	8/9 (89%)	17/19 (89%)
TPO-RA response PLT> 100 x 10^9/L	19/28 (68%)	4/9 (44%)	15/19 (79%)
Fostamatinib treatment	4/37 (11%)	2/11 (18%)	2/26 (8%)
Fostamatinib response PLT>50 x 10^9/L	3/4 (75%)	2/2 (100%)	1/2 (50%)
Fostamatinib response PLT>100 x 10^9/L	2/4 (50%)	1/2 (50%)	1/2 (50%)
thrombosis	5/37 (14%)	2/11 (18%)	3/26 (12%)

IQR: interquartile range; ITP: immune thrombocytopenia, NGS: next generation sequencing, PLT: platelet, TPO-RA: thrombopoietin receptor agonist

Figure 1

https://doi.org/10.1182/blood-2023-185087