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Research Article

Mast Cells in Myelodysplastic Syndromes

¹Karras George, ¹Garlemou Kallina, ³Mpakosi Alexandra, ²Karadonta Argyroula, ¹Cholevas Vasileios and ¹Kyriakou Despoina

¹Department of Haematology, Faculty of Medicine, University of Thessaly, Biopolis, 41334 Larisa, Greece

²University Hospital of Larisa, Biopolis, 41334 Larisa, Greece

³Department of Microbiology, General Hospital of Nikaia, Agios Panteleimon, West Macedonia, Greece

Abstract

Background and Objective: Myelodysplastic Syndromes (MDS) are clonal haemopoietic disorders with a high frequency of leukemic transformation. The role of mast cells (MCs) in this process is not well clarified. The aim was to study the number of mast cells, the secreted cytokines and the microvascular density (MVD) in MDS and controls to clarify the possible role of mast cells in the disease progression in these patients. **Materials and Methods:** Seventy-three patients and 60 healthy individuals were involved in the study. Thirty-seven belonged to the low/intermediate-1 risk group and 36 to the high/intermediate-2 group. Toluidine blue in bone marrow smears was used for MCs measurement. Tryptase and chymase expression was estimated by immunocytochemistry. The CD34⁺ cells were calculated by flow cytometry. Chymase and tryptase serum levels were measured by ELISA. The MVD was calculated by measuring the number of endothelial cells per 0.0625 mm² field area in paraffin sections. **Results:** An increased number of CD34⁺ cells, as well as MCs in the bone marrow of patients, was found. The MCs in patients expressed predominantly tryptase and did not present dysplastic features. Serum tryptase was higher in MDS compared to the normals. The MVD was higher in MDS compared to normals. There was a positive correlation between CD34⁺ cells and MCs as well as between MCs and MVD. High/intermediate-2 patients had a higher number of MCs, CD34⁺ cells and MVD compared to low/intermediate-1 cells and normals. **Conclusion:** The MCs in MDS do not seem to belong to the malignant clone, accumulate probably reactively and may contribute to the clone evolution by supporting angiogenesis and tumour microenvironment.

Key words: Mast cells, myelodysplastic syndromes, angiogenesis, refractory anaemia, CD34⁺ cells, tryptase, bone marrow microenvironment

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Corresponding Author: Karras George, Department of Haematology, Faculty of Medicine, University of Thessaly, Biopolis, 41334 Larisa, Greece

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Competing Interest: The authors have declared that no competing interest exists.

Data Availability: All relevant data are within the paper and its supporting information files.

INTRODUCTION

Mast cells (MC) are derived from the haematopoietic progenitor cell¹ and the committed MC was isolated in 2005². They contribute to tissue regeneration and tissue and organ homeostasis that present continuous increase and reconstruction through mediators they produce and store³⁻¹⁷. Their role in angiogenesis is also well-recognized¹⁸.

In addition, MCs have a significant role in neuroimmunological regulation, neurogenesis and behaviour^{15,16}. They also recognize and react to antigens, toxins and pathogens they have an antigen presentation function and they have a role in immune tolerance through a complicated network of receptors, mediators and enzymes¹⁹⁻⁵².

In malignancies the role of MCs is controversial. In most cases increased presence of MCs is related to poor prognosis and rapid disease progression^{53,54}. In some cases, MCs are related to better prognosis⁵⁵. The MCs are attracted to the tumour environment from factors secreted by the tumour. They contribute to tumour growth through the secretion of angiogenic and mitogenic factors⁵⁶. Their antineoplastic function is achieved through the inhibition of cell growth, induction of apoptosis, reduction of cell mobilization and enhancement of antineoplastic inflammation⁵⁷.

In haematological malignancies, MCs have been described to belong to the malignant clone or be reactive and infiltrate the microenvironment.

Myelodysplastic syndromes are a heterogeneous group of clonal hematopoietic stem cell malignancies characterized by dysplastic features, ineffective haemopoiesis and frequent evolution to acute leukaemia⁵⁸⁻⁶⁵.

There were some publications on MCs in MDS, microvascular density (MVD) and disease progression. The results are controversial and the number of patients is small^{66,67}.

Hence, this article studied the number of MCs, the tryptase and chymase production and the MVD as well as their relation to the risk status of the patient to contribute to the study of the role of these cells in these syndromes.

MATERIALS AND METHODS

Study area: The study was carried out at the Department of Hematology, University Hospital of Larisa, University of Thessaly, Larisa, Greece from 2009 to 2017.

Patients: Seventy-three consecutive patients have been studied 37 females and 36 males with MDS who were referred to the University Hospital of Larisa between the years 2009

and 2012. The median age of males was 72 years (range 35-86) and of females 69 years (range 59-81). For MDS classification FAB₂ and WHO 2008 classification systems were used. According to these systems, 21 patients had refractory anaemia (RA) (11 females and 10 males). The median age of males was 62 years (range 59-86) and of females 78 years (range 66-85). Eighteen patients had Refractory Anemia with Excess of Blasts (RAEB) 8 males and 10 females with the median age for males 60.5 years (range 35-75) and for females 74 years (range 58-79). Eighteen patients had Refractory Anemia with Excess of Blasts in Transformation (RAEB-T), 10 males and 8 females, with the median age for the males 66 years (range 55-77) and for the females 79.5 (range 59-80). Sixteen patients had Refractory anaemia with ringed sideroblasts (RARS) 8 males and 8 females with the median age for males 64.5 years (range 63-86) and for females 75 years (range 65-88). Patients with CMML and secondary MDS were excluded.

In addition, the patients were classified according to IPSS-1997 as low/intermediate-1 risk (37 patients) and high/intermediate-2 risk (36 patients). The median age of patients in the low/intermediate-1 risk group was 72.5 years (range 57-88) and in the high/intermediate-2 risk group was 68 years (range 35-80).

In the study, 60 normal individuals have been included 32 males and 28 females. The median age for the males was 75 years (range 56-90) and for females 74 years (range 66-90).

Methods: Bone marrow (BM) smears and paraffin sections were prepared from the posterior iliac crest. Mast cells were calculated using toluidine blue stain (mean value of positive cells per 1000 nucleated cells in 3 slides). For estimating the expression of tryptase and chymase mouse-anti-human moAbs were used (ABCAM -Discovery Drive, Cambridge Biomedical Campus Cambridge, CD2OAX, UK and MA5-11717, CC1 -Thermo Fisher Scientific, Waltham, Massachusetts, USA, respectively) and PAP kit DAKO-(Agilent, Waldbronn, Germany). Every cell was graded from 0+–4+ depending on the intensity of the stain and the score was calculated as the mean value of 200 cells as Komi and Redegeld⁶⁸ described.

Haemopoietic progenitor CD34⁺ cells were measured by flow cytometry using the anti-CD34 moAb (K567-FITC).

The levels of chymase and tryptase in serum were measured using ELISA kits (INVITROGEN-Thermo Fisher Scientific, Waltham, Massachusetts, USA) according to the manufacturer's instructions.

Microvascular density (MVD) was investigated as Kyriakou *et al.*⁶⁷ described. Microvessels were measured per 0.0625 mm² field area.

Statistical analysis: For statistical analysis, the t-test was used for the comparison of parametric values and Wilcoxon's test (Bonferroni's correction) for nonparametric values. The correlation coefficient (R^2) was used for examining possible correlations between various parameters.

RESULTS

CD34⁺ cells in bone marrow: Table 1 shows the percentage of CD34⁺ cells in each group of patients is shown. In detail, CD34⁺ cells in the normal individuals were 1.08±0.87%. In MDS patients they were 5.3±3.83%. Specifically, in various MDS categories, CD34⁺ cells were 2.52±1.2% in RA, 1.37±0.5% in RARS, 8.94±2.01% in high/intermediate-2 and 2.027±1.117% in low/intermediate-1. There was a significant difference between high/intermediate-2 and normals ($p<0.01$), between high/intermediate-2 and low/intermediate-1 ($p<0.01$), between low/intermediate-1 and normals ($p<0.01$) and between RA and normals ($p<0.01$). There was no significant difference between RARS and normals ($p>0.1$).

Mast cells in bone marrow (BMMCs): The percentage of mast cells in bone marrow in each group of patient are shown in Table 1. In normals, the percentage of mast cells was 1.82±0.93%. In MDS it was 5.767±3.45%. In RA it was 4.38±1.16%, in RARS 2.18±0.75%, in low/intermediate-1 3.43±1.48% and in high/intermediate-2 8.16±3.26%. There was a significant difference between high/intermediate-2 and low/intermediate-1 ($p<0.01$) as well as the normals ($p<0.01$). There was a significant difference between low/intermediate and normals ($p<0.01$) as well as between RA and normals ($p<0.01$). There was no difference between RARS and normals ($p>0.1$). In the total MDS group, the mast cells were higher than the normals ($p<0.01$).

Immunocytochemical detection of tryptase and chymase in the bone marrow: Results are presented in Table 1. In the total MDS group, the tryptase positivity score in bone marrow mast cells (BMMC) was 2.328±1.45 while in the normals it was 0.8±0.82 and the difference was statistically significant ($p<0.01$). In high/intermediate-2 the score was 3.694±0.467 and it was significantly higher than that of low/intermediate-1 (1±0.57, $p<0.01$) and normal ($p<0.01$). In low/intermediate-1

the score did not differ from the score in normals ($p>0.1$). In MDS-RA and MDS-RARS the scores were 1.19±0.60 and 0.75±0.447, respectively and did not differ from the normals and low/intermediate-1 ($p>0.1$), but it was significantly lower than that of high/intermediate-2 ($p<0.01$).

The chymase score in BMMC in the total MDS group was 0.78±0.71, while in the normals it was 0.9±0.3. In high/intermediate-2 MDS the score was 0.804±0.749 and the difference between the 3 groups was not significant ($p>0.1$). In the low/intermediate-1 group the score was 0.75±0.68, in MDS-RA 0.66±0.58 and in MDS-RARS it was 0.875±0.718. There was no statistically significant difference among all groups ($p>0.1$).

Serum tryptase and chymase levels: The levels of serum tryptase in the normals were found 7.8±3.21 ng mL⁻¹, in total MDS group were 40.96±35.4, in high/intermediate-2 were 73.31±20.87, in low/intermediate-1 were 9.48±4.217, in MDS-RA 10.75±4.83, in MDS-RARS 7.806±2.503. There was a significant difference between total MDS and high/intermediate-2 compared to the low/intermediate-1 and normals ($p<0.1$). The levels of serum chymase (pg mL⁻¹) were not detected in any category of MDS and normals (Table 1).

Correlation of tryptase levels and BM CD34⁺ and mast cells: There was a strong positive correlation between serum tryptase levels and BMMC ($R^2 = 0.6837$). There was also a positive correlation between BMMCs and BM CD34⁺ cells ($R^2 = 0.4369$) (Fig. 1 and 2).

Microvascular density in bone marrow (MVD): The MVD (microvessel number/0.0625 mm²) in the entire group of MDS was 6.3±3.3 and was significantly higher than the normals (2.23±0.38, $p<0.01$). In high/intermediate-2 the MVD was 7.89±2.8, in low/intermediate-1 it was 4.8±2.2, in MDS-RA it was 4.98±1.8 and in MDS-RARS it was 5±2.0. There was a significantly higher MVD in high/intermediate-2 compared to low/intermediate-1, MDS-RA, MDS-RARS and normal ($p<0.01$). There was also a significant difference between low/intermediate-1, MDS-RA, MDS-RARS and normals ($p<0.01$). There was a positive correlation between BMMC and MVD (Fig. 3).

Table 1: Studied parameters in each group of patients and controls (CD34⁺ cells, BMMCs, MVD, tryptase score+, chymase score and serum tryptase)

Patients' group	CD34 ⁺ cells (%)	BMMCs (%)	MVD microvessels/0.0625 mm ²	Tryptase score	Chymase score	Serum tryptase (ng mL ⁻¹)
MDS	5.3±3.83	5.767±3.45	6.3±3.3	2.328±1.45	0.78±0.71	40.26±35.4
MDS-RA	2.52±1.2	4.38±1.16	4.98±1.8	1.19±0.60	0.66±0.58	10.75±4.83
MDS-RARS	1.37±0.5	2.18±0.75	5.0±2.0	0.75±0.447	0.875±0.718	7.806±2.503
MDS low/int-1	2.027±1.117	3.43±1.48	4.8±2.2	1.0±0.57	0.75±0.68	9.48±4.217
MDS high/int2	8.94±2.01	8.16±3.26	7.89±2.8	3.694±0.467	0.804±0.749	73.31±20.87
Normals	1.08±0.87	1.82±0.93	2.23±0.38	0.8±0.82	0.9±0.447	7.8±3.21

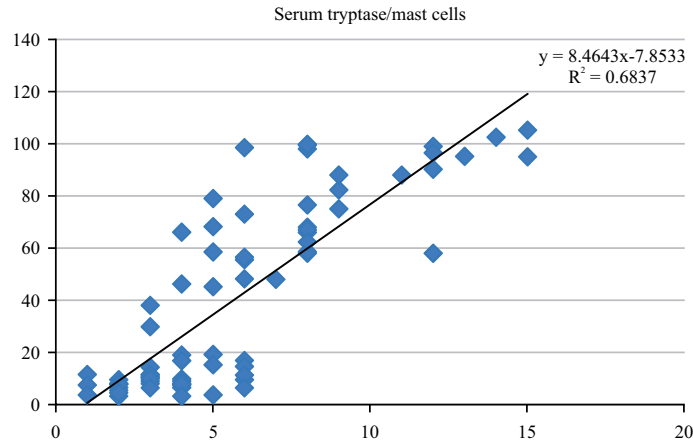


Fig. 1: Correlation between bone marrow mast cells (x-axis) and serum tryptase levels (y-axis) in MDS patients
There is a strong positive correlation between the 2 parameters ($R^2 = 0.6837$)

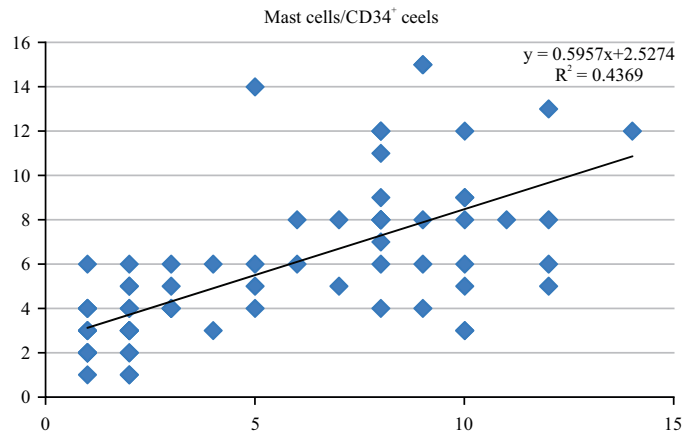


Fig. 2: Correlation between BMMCs (y-axis) and BMCD34+ cells (x-axis) in MDS patients
There was a positive correlation between the 2 parameters ($R^2 = 0.4369$)

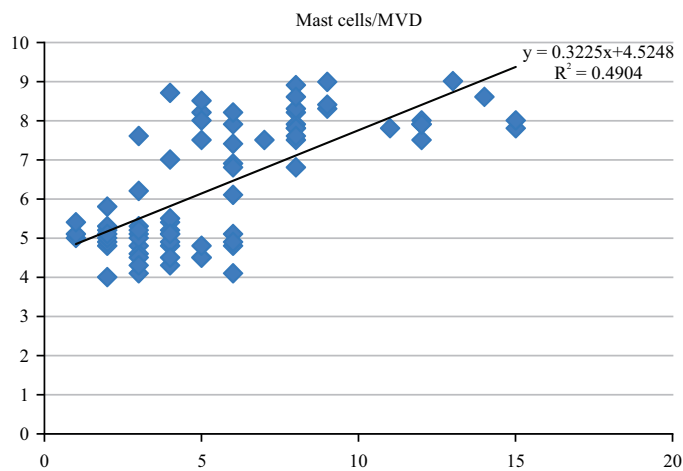


Fig. 3: Correlation between bone marrow mast cells (x-axis) and bone marrow microvessel density-MVD (y-axis) in MDS patients
There is a positive correlation between the 2 parameters ($R^2 = 0.4904$)

DISCUSSION

An increased number of MBMCs and MVD in MDS possible pathogenetic connection between them was found in our study. There were several publications regarding the presence of mast cells in malignant diseases, including haematological malignancies and their role in disease progression is controversial. The mechanisms by which they contribute are not completely clarified^{53,54,57,69,70}.

A higher number of mast cells in higher-risk MDS as well as higher numbers of BMMC in the entire MDS group compared to normals was found. This was in agreement with what has been reported earlier by Alexandrakis *et al.*⁶⁶. This also found in agreement with earlier reports by Pejler *et al.*⁷¹ that, these cells secrete tryptase predominantly that was detected by immunocytochemistry in BM as well as in the serum of patients. In reactive mastocytosis such as in allergic conditions, it has been reported that the mast cells are tryptase and chymase-producing⁷² and are peripheral and not BM mast cells. Although in a previous study there was no difference in BMMC numbers in the various MDS categories⁶⁶, an increase in BMMC in higher-risk MDS was found. The fact that was detected increased numbers of BMMC were in higher risk MDS which is considered a more advanced stage of the disease probably means that the number of BMMC increases with disease progression. Whether they contribute to the disease progression or they accumulate reactively is not clear. In earlier reports, it has been found that in some cases, the mast cells belong to the malignant clone and in these cases, they present dysplastic features^{66,73,74}. In this study dysplastic features (elongated shape, abnormal granulation and abnormal accumulation) were not found in BMMC but this does not exclude the neoplastic origin of these cells. Studies during disease progression in each case, genetic studies and MC cultures could help to clarify the origin of these cells. Probably the majority of BMMCs accumulate in response to tumour-secreted factors in BM but it is not known if the cytokines that BMMCs secrete contribute to disease progression.

It has been reported that MC is involved in angiogenesis^{16-18,56,57,75} and neoangiogenesis is important for tumour growth⁷⁶⁻⁷⁸.

We found increased MVD in MDS compared to the normals. We also found higher MVD in high/intermediate-2 risk MDS compared to the low/intermediate-1, in contrast to a previous study by Kyriakou *et al.*⁶⁷. The MVD is positively correlated with the number of BMMC. This raises the suggestion that BMMCs may contribute to tumour progression by enhancing angiogenesis. The heterogeneity of the MDS group may be responsible for the contradictive results in the literature⁶⁷.

Many previous studies suggested that MC involvement in fibrosis in many situations like liver cirrhosis, lung fibrosis, renal fibrosis, scleroderma, etc.^{3,10,79}. There are no adequate data to clarify the mechanism through which MCs exert their fibrotic action. We did not have enough patients to study the possible relation of BMMCs in MDS with BM fibrosis.

CONCLUSION

An increase in BMMCs in MDS was found and they were higher in the advanced stages of the disease. To clarify whether they contribute to disease progression further studies are required with MC cultures and clonality investigation in various stages of the disease.

SIGNIFICANCE STATEMENT

The microenvironment plays an important role in MDS evolution. Studying the factors of the microenvironment that contribute to malignant clone growth and evolution and the use of proper agents targeting the microenvironment might help in delaying disease progression.

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