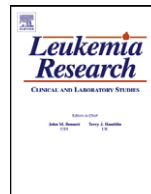




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Invited review

The lower risk MDS patient at risk of rapid progression

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ABSTRACT

Most patients with myelodysplastic syndrome (MDS) are classified at diagnosis as having a low/INT-I or INT-II/high risk disease, based on the classical International Prognostic Scoring System (IPSS) criteria. The low/INT-I risk patients are usually managed mildly with supportive care, including red blood cell (RBC) transfusions, erythroid stimulating agents (ESAs), other cytokines (G-CSF, platelet stimulating agents), as well as thalidomide and lenalidomide. Some patients receive immunosuppressive therapy, and iron chelation is indicated in iron overloaded patients. Aggressive approach (hypomethylating agents, chemotherapy and stem cell transplantation) is usually not applied in such patients.

Occasionally, we observe a “low risk” patient with rapid progression of disease and poor outcome. Can we identify demographic, clinical, laboratory, cellular-biological and/or molecular parameters that can predict “poor prognostic features” (PPF) in “low risk” MDS patients?

Clinical and laboratory parameters have been reported to be associated with poor prognosis, in addition to the known “classical” IPSS criteria. These include older age, male gender, poor performance status, comorbidities, degree of anemia, low absolute neutrophil count (ANC) and platelet counts, RBC transfusion requirements, high serum ferritin, high LDH, bone marrow (BM) fibrosis, increased number of BM CD34+ cells and multi-lineage dysplasia. Certain immunophenotypes (low CD11b, high HLA-Dr, CD34, CD13 and CD45), clonal granulocytes, multiple chromosomal abnormalities, chromosomal instability, short telomeres and high telomerase activity were also reported as PPF. Studies of apoptosis identified Bcl-2 expression and high caspase 3 as PPF, while the reports on survivin expression have been confusing.

Recent exciting data suggest that methylation of p15 INK4b and of CTNNA1 (in 5q–), high level of methylation of other genes, absence of the TET2 mutation, down regulation of the lymphoid enhancer binding factor 1 (LEF1), mutation of the polycomb-associated gene ASXL1 and a specific 6-gene signature in gene expression profiling – are all associated with poor prognosis in MDS.

Do we have data suggesting a different treatment for “low risk” MDS patients displaying PPF? Two teams, the combined Nordic-Italian and the GFM groups have reported an improved survival with ESAs. The GFM has achieved prolonged survival with iron chelation. Recently, encouraging data with survival advantage in azacitidine-treated patients have been published, including a few INT-I patients. Finally, data suggest that low/INT-I MDS patients who undergo stem cell transplantation (SCT) do better than INT-II/high risk patients).

In summary, some patients, classified as “low risk MDS” carry PPF. An appropriate therapeutic approach is indicated. Future updated classifications and prospective trials may lead to a better outcome.

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Patients are usually diagnosed as having myelodysplastic syndrome (MDS), based on well known recognized criteria [1–7]. This is commonly followed by prognostic staging, in order to plan the therapeutic approach. The commonly used prognostic system is the International Prognostic Scoring System (IPSS) [8]. The IPSS is based on three classical criteria, i.e. blast percentage in the bone marrow (BM), originally proposed by the FAB classification [9], cytogenetics (three types: favorable, poor, and intermediate), and the number of affected cytopenias. Using these criteria, each patient is given a score, according to which he is categorized as belonging to one of the four IPSS groups, i.e. low risk (LR), intermediate-I (INT-I), intermediate-II (INT-II) and high risk (HR) MDS.

Usually, the patients classified as LR and INT-I IPSS, are referred to as “lower risk MDS” (LrMDS) and offered relatively mild and conservative treatments. These include supportive care, such as red blood cell (RBC) transfusions [1,2,4–7], erythroid stimulating agents (ESAs) [10–12] and granulocyte – macrophage colony stimulating factors (GM-CSF) [11–13]. Recently, thrombopoietic agents [14], thalidomide [15] and lenalidomide have been introduced [16–18]. Some patients, especially those with hypocellular BM receive immunosuppressive therapy [19] and iron chelation is indicated in patients with iron overload [20,21]. More aggressive therapies such as hypomethylating agents [22–24], chemotherapy and stem cell transplantation (SCT) are usually not administered to patients with LrMDS [25,26].

Most LrMDS patients experience a slowly progressive disease with a long course [8,27]. However, occasionally, we encounter a patient, who is classified as having LrMDS, yet progresses rapidly, i.e. displays decreasing counts, complications, possible leukemic transformation, and a short survival.

Who are these “LrMDS” patients with rapid disease progression? Can we identify them at diagnosis upon classification or earlier? And if so, what are the additional poor prognostic features (PPF), in addition to the known classical IPSS criteria that are used to identify these patients? If and when we identify these “LrMDS” patients, who are probably not low risk, should we attempt an alternative therapeutic approach to achieve better outcome? Do we have data to support such an alternative treatment? This review will address these questions.

1. Poor prognostic features (PPF) in lower risk MDS – diagnostic tools

Studies, summarized in Table 1, have identified a list of parameters which are not used in the IPSS classification, but may have prognostic relevance. Starting with the simple clinical and demographic markers, which can be applied in every practice, one can review the original IMRAW/IPSS data. Kao et al. [28] re-examined the data on 816 MDS patients, which served for the original IPSS classification, and concluded that hemoglobin (Hb), but not neutrophile or platelet counts, was a reliable predictor for overall survival but not for time to leukemia conversion. Kantarjian et al. [29] analyzed the data on 1915 MDS patients, including 507 patients with primary MDS, treated at the MD Anderson Cancer Center. They found that older age, poor performance status, anemia, low platelet count and prior transfusion need – were all predictors of poor outcome. They also proposed a new risk model, based on these prognostic parameters. A recent Chinese prospective analysis on 435 patients, reported that age >60 year, ANC <1000/mm³ and Hb below 9 g/dl were PPF [30]. The Dusseldorf MDS registry confirmed older age, especially > 50yr, as PPF [31]. The German-Austrian MDS study group has recently summarized data on 897 patients with primary MDS and found in their retrospective analysis that older age (>66 year) and male gender were associated with poor prognosis [32]. The

Table 1
Poor prognostic features (PPF) in lower risk MDS (LrMDS).

Class of markers	PPF	Refs.
Clinical/demographic	Older age	[29–32]
	Gender (male)	[32]
	Poor performance status	[29]
	Co-morbidities	[33]
	Transfusion needs	[29,34–37]
	Iron overload	[35–37]
Lab values	High serum ferritin	[37]
	Hb ↓	[28–30,37]
	PLT ↓	[29]
	ANC ↓	[30]
Bone marrow (BM)	High LDH	[27]
	BM fibrosis	[39]
	CD34+ clusters	[39–41]
	Multi-lineage dysplasia	[35,36,39–41]
Immunophenotyping	Normal/high cellularity	[42]
	↑ HLA-Dr	[43]
	Low CD11b	[43]
	↑ CD34	[44]
	↑ CD13	[44]
	↑ CD45	[44]
Clonality	Flow score	[45]
	Clonal granulocytes	[41]
Cytogenetics	Additional chromosomal abnormalities	[8,46]
	Chromosomal instability	[47]
Telomeres	Short telomeres	[48–51]
	High telomerase activity	[49,52–54]
Apoptosis	↑ BCL2	[56]
	↑ Caspase 3	[57]
	Survivin (???)	[58–60]
	Cell senescence (PIG INK4)	[61]
Genetic/epigenetic/molecular	P15 INK4b methylation	[62,63]
	CTNNA1	[64]
	High methylation	[65]
	Unmutated TET2	[66]
	LEF1 down regulation	[67]
	ASXL1 mutation	[68]
6-gene poor risk signature	[69]	

Austrian group has emphasized that co-morbidity, as used by the hematopoietic-stem cell transplantation-specific co-morbidity index (HCT-CI) and Charlson co-morbidity index (CCI), were additional PPF [33].

Although many felt for years that MDS patients who require regular blood transfusions represent a “poor prognostic” disease, Cazzola and Malcovati [34] demonstrated that transfusion dependent MDS patients do worse than MDS patients who are transfusion free. The same Pavia team later on, comparing their data on more than 400 MDS patients with the Dusseldorf registry, reported that transfusion dependence, iron overload, and multi-lineage (as opposed to uni-lineage) dysplasia predicted poor outcome [35,36]. Based on transfusion requirements they proposed an updated version of the IPSS system – WHO classification – based prognostic scoring system (WPSS). A recent retrospective analysis of 137 patients from the Czech Republic, confirmed that transfusion dependence, Hb <8 g/dl, and high serum ferritin level (>2000 mg/dl) were associated with poor prognosis [37]. Germing et al. [27], reported that high serum LDH can also serve as PPF.

Regarding more complex parameters, the Pavia team retrospectively reviewed the BM samples of 301 patients and concluded that BM fibrosis, the presence of CD34+ cell clusters (a reminder of the old “Abnormal Localization of Immature Progenitors, ALIP, as suggested by Tricot et al. [38]), and multi-lineage (as opposed to uni-lineage) dysplasia were associated with poor prognosis [39].

This was confirmed by others [40,41]. Yue et al. [42] reported that hypocellular BM correlated with a favorable outcome, compared with normal or hypercellular BM.

Immunophenotyping, both for diagnosis and for predicting prognosis of MDS patients, has attracted several groups. We found that high expression of HLA-Dr, and low CD11b expression predicted early leukemic transformation [43]. More recently, high expression of CD34, CD13, and CD45 was reported as PPF [44]. Van de Loosdrecht et al. [45] studied the immunophenotyping of CD34+ BM cells from 50 MDS patients. They found high degree of aberration of myelomonocytic antigens to which they gave a flow score, correlated with prognosis. Clonal granulocytes were also reported to predict poor outcome [41].

Cytogenetics has always been a field for investigation in these diseases, for diagnosis, staging and for predicting prognosis. In the original IPSS classification multiple chromosomal abnormalities were considered as PPF [8]. This has recently been confirmed by the MD Anderson experience with 2743 patients [46]. Chromosomal instability, as could be expected, may also predict rapid disease progression [47].

Several studies have focused on telomeres. In summary, short telomeres, resulting in genetic instability [48–51], coupled with high telomerase activity (associated with a high proliferation rate) [49,52–54], both correlate with poor prognosis.

Apoptosis was found to be increased in early MDS, and decreased in later phases of the disease [55]. Several groups have studied apoptosis-associated markers as predictors of disease course. High expression of the anti-apoptotic protein BCL-2 [56], and of caspase 3 [57], predicted poor prognosis. The current data on the apoptosis inhibitor survivin are inconclusive [58–60]. Cell senescence, determined by P16 INK4a expression, was also found to be PPF [61].

Recent epigenetic studies have generated exciting molecular data. Methylation of p15 INK4b was found to predict poor outcome [62,63]. In MDS patients with the 5q– abnormality, methylation of the promoter of CTNNA1 correlated with poor prognosis [64].

Shen et al. [65], have recently screened 24 MDS patients for promoter CpG methylation of 24 genes and identified aberrant hypermethylation at 10 genes. They then performed quantitative methylation analysis by bisulfite pyrosequencing of the identified genes in 317 patient samples and assessed relations between methylation and clinical outcome. While methylation frequencies of individualized genes ranged from 7 to 70%, by applying an individual methylation z score based on all genes for each patient, they found that higher methylation correlated with a shorter median survival (12.3 vs 17.5 m) and a shorter progression free survival (6.4 vs 14.9 m).

The TET2 mutation has recently gained attention. Kosmider et al. [66], have found that not only this genetic abnormality is common in MDS (23%), but also 5-year overall survival and 3-year leukemia-free survival were significantly shorter in patients carrying non-mutated TET2 compared with patients carrying the mutated gene: 18 vs 77, and 64 vs 89%, respectively.

Pellagatti et al. [67], studied the granulopoiesis regulator lymphoid enhancer binding factor 1, LEF1, and found that its down regulation was associated with poor prognosis. This group has also reported that ASXL1 mutation could serve as a molecular marker for disease progression [68].

Finally, the Stanford team has performed detailed molecular analysis with gene expression profile in CD34+ BM cells from MDS patients [69]. Studying 40,000 CDNAs chip arrays, they detected 1175 genes that were differentially expressed. Moreover, they identified six genes (RPL23, RPS4x, RPS25, RPS19, KLK3 and TPP2), four ribosomal and two enzymes, all over expressed in patients who later on progressed and transformed to acute leukemia, thus providing a “6-gene-poor-risk-signature”.

2. Poor prognostic features (PPF) in lower risk MDS – therapeutic options

Once, a patient with LrMDS but with a predicted poor survival is identified, should he be treated differently? Do we have data to suggest an alternative approach?

Although phase III controlled trials have not been published, we do have little evidence suggesting that an active approach in some LrMDS patients may attenuate the course of disease.

Erythroid stimulating agents (ESAs) have been a pivotal anti-anemic treatment for LrMDS patients. We [70–73], as well as others, have suggested that ESAs confer immunomodulatory anti-neoplastic effect(s) as well. Several trials have suggested that ESAs-treated cancer [74] and MDS [75] patients benefit from prolonged survival compared with patients who do not receive ESAs. Recently, the Nordic group has compared the outcome of their 121 MDS patients treated with recombinant erythropoietin (EPO) and G-CSF, with the outcome of 237 untreated MDS patients from the Pavia Cohort [13]. Increased overall survival was observed in EPO + G-CSF treated LrMDS patients ($p=0.033$), but not in patients with higher risk MDS. The GFM data are similar [76].

Iron overload, aggravated by repeated blood transfusions, has been associated with organ damage, suggesting poor prognosis [20,21]. But, has iron chelation therapy (ICT) resulted in a better prognosis? No prospective comparative data are available; however, two reports summarizing retrospective cohort data have suggested a survival benefit with ICT. The GFM analysis reported a positive survival impact of ICT in regularly transfused patients with MDS: the median survival of the whole MDS group was 63 months, with 115 month survival for the ICT-treated patients, compared with only 51 months ($p<0.0001$) for MDS patients with no ICT [77]. A Canadian cohort series observed similar results [78].

As mentioned, hypomethylating agents such as decitabine and 5-azacitidine are usually offered to higher risk MDS patients [22,23]. The recently published AZA-001 trial reported survival advantage in 179 azacitidine-treated MDS patients, compared with the control group, treated with conventional therapy (chemotherapy, low dose cytarabine, or supportive treatment). The overall survival was 24.5 months for the azacitidine group but only 15.0 months for the conventional group ($p=0.0001$) [24]. While most patients in the AZA-001 trial were higher risk MDS patients, analyzing the data reveals that 18 patients (5%) were in fact, MDS patients classified as INT-I. Although no subset analysis has been available on that subgroup, it is tempting to hypothesize that they also might benefit from such treatment.

Finally, can stem cell transplantation (SCT) improve survival in LrMDS with PPF? Again, no prospective trial comparing SCT with less aggressive approach in this patient subpopulation has been published. But, the recently published Italian – GITMO experience with SCT outcome (1990–2006) in MDS patients, reports on 5-year overall survival of 80%, 5-year probability of relapse of only 9% and 5 year transplant-related mortality of only 14% for refractory anemia (RA) patients, and 57, 22 and 39% respectively, for refractory cytopenia (RC or RCMD) patients. These results are better than those obtained with other patients (RAEB-1, RAEB-2, and AML post-MDS) [79]. This does not mean that SCT is recommended to all LrMDS patients with or without PPF. But it suggests, that if a more aggressive than the usually applied approach is considered – it might be relatively non-toxic, and sometimes successful [25,26,79].

In summary, some MDS patients, although classified as “lower risk MDS” (LrMDS), and probably offered a mild therapeutic approach – are not really “lower risk”. Such patients, despite being grouped as LrMDS experience a rapid complicated progressive course with or without leukemic transformation and a dismal prog-

nosis. A list of poor prognostic features (PPF) described in this review, including clinical and laboratory values, others such as BM parameters, and more recently established molecular markers – can identify those LrMDS patients, and predict rapid course and poor outcome. Although, no prospective convincing data are available, emerging information suggests that a more aggressive approach might improve the outcome. Obviously, future prospective comparative trials will have to test this hypothesis, and will probably lead both to a revised updated classification and to a more individualized treatment.

Conflict of interest

All authors have no conflict of interest to declare.

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