Chronic myeloproliferative neoplasms

Sex and degree of severity influence the prognostic impact of anemia in primary myeloﬁbrosis: analysis based on 1109 consecutive patients

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Anemia is one of the major risk factors for survival in primary myeloﬁbrosis (PMF). Current prognostic models in PMF, including the International Prognostic Scoring System (IPSS) [1], dynamic IPSS (DIPSS) [2] and DIPSS-plus [3], all list hemoglobin level (Hb) of <10 g/dl as one of their risk variables. However, the 10 g/dl Hb threshold used in these prognostic models overlooks the signiﬁcant difference in Hb levels between men and women, and, also assumes similar prognostic weight between moderate and severe anemia; in regards to the latter, for example, red cell transfusion need confers an additional point of prognostic adversity in DIPSS-plus, which is independent of and in addition to what is accounted for a hemoglobin level of <10 g/dl [3]. The “dose-dependent” prognostic effect of anemia, and the effect of sex on the selection of prognostically relevant Hb thresholds, has previously been recognized in myelodysplastic syndromes (MDS) [4]. In the current study, we examined the severity-stratiﬁed and sex-specific prognostic contribution of anemia in PMF, in the context of other relevant clinical, cytogentic and molecular risk factors.

The current study was approved by the Mayo Clinic institutional review board and consent from research studies was obtained in all subjects. Consecutive cases of PMF were recruited from institutional databases of the Mayo Clinic, Rochester, MN, USA, where the 2016 World Health Organization (WHO) criteria for PMF were retrospectively applied to conﬁrm the diagnosis [5]. Special precaution was taken to avoid inclusion of patients with post-polycythemia vera myeloﬁbrosis; only one woman had a hemoglobin level of >16 g/dl (16.1 g/dl) and only one man had a hemoglobin level of >16.5 (16.7 g/dl); in both instances, review of clinical and bone marrow morphological features were fully consistent with PMF and there was no antecedent history of PV in either. Treatment was according to conventional standards utilized at the time of diagnosis. Clinical and laboratory data, including cytogentic information, were collected at the time of initial referral to our institution. Cytagenetic analysis and reporting was done according to the International System for Human Cytogenetic Nomenclature (ISCN) criteria [6]. The recently revised system was used for cytagenetic risk stratiﬁcation (Tefferi et al. personal communication, 17 December 2017; see Table 1 footnote). Survival was calculated from time of initial referral, commensurate with time of baseline data acquisition, including cytagenetic and mutation analysis. Standard statistical methods were used to determine signiﬁcance of differences among groups in the distribution of continuous or nominal variables. Overall survival data were prepared by the Kaplan–Meier method and compared by the log-rank test. Cox proportional hazard regression model was applied for multivariable analysis. P value < 0.05 was considered signiﬁcant. The Stat View (SAS Institute, Cary, NC, USA) statistical package was used for all calculations. Patients were stratiﬁed according to the severity of anemia: mild—Hb ≥ 10 g/dl but below sex-adjusted lower limit of normal; moderate—Hb between 8 g/dl and <10 g/dl; and severe—Hb < 8 g/dl or transfusion-dependent. Reference ranges for Hb, at our institution, were 13.5–17.5 g/dl for men and 12.0–15.5 g/dl for women.
A total of 1109 consecutive patients with PMF were considered (median age 65 years; 63% men) (Supplementary table 1). DIPSS risk distribution was 11% high, 43% intermediate—2, 34% intermediate-1 and 12% low. Among informative cases, karyotype was “very high risk” in 7%, unfavorable in 19% and favorable in 74%; driver mutation distribution was 66% JAK2, 16% CALR type 1/like, 3% CALR type 2/like, 5% MPL and 10% triple-negative; ASXL1, SRSF2 and U2AF1 mutation frequencies were 38, 14, and 16%, respectively (Supplementary table 1). Anemia, defined as a hemoglobin level of below the lower limit of normal, adjusted for sex, was present in 950 (86%)
patients and ranged in severity from mild (Hb ≥ 10 g/dl and less than sex-adjusted lower limit of normal) in 35%, to moderate (Hb ≥ 8 and <10 g/dl) in 14%, to severe (Hb < 8 g/dl or transfusion-dependent) in 37%; the prevalence of anemia in general, mild anemia and severe anemia for men vs. women were 90 vs. 78%, 38 vs. 29% and 38 vs. 35%, respectively (p < 0.001). As outlined in supplementary table 1, and not unexpectedly, patients with moderate or severe anemia displayed older age distribution, lower leukocyte and platelet counts, higher circulating blast percentage, higher prevalence of constitutional symptoms, and higher clinical and cytogenetic risk categories. Also, consistent with previous observations, U2AF1 mutations clustered with severe anemia and lower frequency of type 1/like CALR mutations [7]. On the other hand, as was evident in supplementary table 1, there was no difference in the prevalence of marked leukocytosis or SF3B1 or high molecular risk mutations, including ASXL1 and SRSF2.

At the time of this writing, 823 (74%) deaths, 82 (7.4%) leukemic transformations, and 53 (4.8%) stem cell transplants were recorded and median follow-up time for living patients was 6.1 years. In univariate analysis, all grades of anemia (i.e., mild, moderate, and severe) predicted shortened survival with HR (95% CI) of 3.4 (2.7–4.3) for severe, 2.1 (1.6–2.8) for moderate, and 1.4 (1.1–1.7) for

Fig. 1  a Survival data on 1109 patients with primary myelofibrosis stratified by the degree of anemia. b Survival data on 694 men with primary myelofibrosis stratified by the degree of anemia. c Survival data on 415 women with primary myelofibrosis stratified by the degree of anemia
mild (Table 1; Fig. 1a). Prognostic relevance was sustained for all grades of anemia during multivariable analysis, with respective HR (95% CI) of 2.5 (1.7–3.8), 1.7 (1.1–2.8) and 1.7 (1.1–2.5) for severe, moderate, and mild degrees of anemia; other variables of significance in the multivariable prognostic model included the revised cytogenetic risk stratification \((p < 0.001)\), driver mutational status \((p < 0.001)\), ASXL1 \((p < 0.001)\) and SRSF2 \((p = 0.001)\) mutations, and all clinical variables used in DIPSS or DIPSS-plus (Table 1). However, sex-specific analysis revealed that the prognostic contribution of mild anemia (i.e. Hb ≥ 10 g/dl but below sex-adjusted lower limit of normal) was evident only in men (HR: 1.7, 95% CI 1.2–2.4) and not in women (HR: 1.0, 95% CI: 0.7–1.4), whereas the impact of severe or moderate anemia were apparent in both sexes (Fig. 1b, c); severe vs no anemia HRs (95% CI) were 3.9 (2.8–5.6) for males and 2.9 (2.1–4.0) for females, while the corresponding figures for moderate vs no anemia were 2.6 (1.8–3.9) for males and 1.7 (1.2–2.6) for females.

In univariate analysis, severe anemia (HR: 2.8, 95% CI: 1.3–6.0) and moderate anemia (HR: 2.7, 95% CI: 1.2–6.2) were both associated with inferior leukemia-free survival; other variables of significance for leukemia-free survival, in univariate analysis, included platelets <100 \times 10^9/l, circulating blasts ≥1%, triple-negative driver mutational status, ASXL1 mutations, SRSF2 mutations, revised cytogenetic risk category, DIPSS and DIPSS-plus \((p < 0.05\) in all instances). However, in multivariable analysis, prognostic relevance of degree of anemia was no longer apparent \((p = 0.24)\) whereas cytogenetic risk (HR: 3.6, 95% CI: 1.2–10.7 for very high risk karyotype and HR: 2.4, 95% CI: 1.2–5.0 for unfavorable karyotype), SRSF2 mutations (HR: 5.4, 95% CI: 3.0–10.2), ASXL1 mutations (HR: 2.0, 95% CI 1.1–3.5), platelets <100 \times 10^9/l (HR: 2.4, 95% CI: 1.3–4.4) and circulating blasts ≥1% (HR: 2.0, 95% CI: 1.1–3.5) remained significant.

In the current series of patients with PMF, anemia defined as a Hb level of below sex-adjusted lower limit of normal, was present in 86% of the patients. In this regard, it is important to note that there was no intential selection bias and that we did not distinguish between pre and overtly fibrotic PMF [5]. It is conceivable that the incidence of anemia might have been lower in patients with prefibrotic PMF and in those evaluated at the time of initial diagnosis. Therefore, one should be careful in the interpretation of the results from the current study as they might not be as applicable in the setting of IPSS or prefibrotic PMF.

The current study suggests that the hemoglobin threshold level of 10 g/dl, conventionally used for prognostication in PMF, might be deficient as it fails to account for severity of anemia and effect of sex on hemoglobin levels. More specifically, we found that mild anemia, defined as a hemoglobin level of ≥10 g/dl, but below the sex-adjusted lower limit of normal, independently predicted shortened survival in men, but not in women (Fig. 1b, c). Also of note, severe anemia, defined as a Hb level of <8 g/dl or transfusion-dependence, was associated with a more than 1.5-fold increase in risk of death, compared to that observed for moderate anemia, in both sexes. In this regard, it is important to note that the differential prognostic impact of anemia severity was already recognized in the context of DIPSS-plus, where adverse point allocation for anemia included two points for patients requiring red cell transfusions vs. only one point for those not requiring transfusions but displayed a hemoglobin level of <10 g/dl [3]. Furthermore, in a post hoc analysis of receiver operating characteristic (ROC) curve, the Hb threshold for severe anemia was determined at 8 g/dl for women and 9 g/dl for men. Therefore, the impact of sex on the prognostic relevance of anemia in PMF might not be restricted to only patients with mild anemia. We conclude that future prognostic models in PMF should employ sex-specific Hb thresholds, as is currently practiced in MDS [8], and also consider assigning prognostic scores for anemia that are proportional to its severity.

**Author contributions:** All authors have reviewed and approved the manuscript. CAH reviewed the pathology data. RPK reviewed the cytogenetic data. NG, AP, and AT contributed patients and assisted in the data extraction. MN and MM assisted in the data extraction, and analysis and preparation of tables and figures. TLL was in charge of molecular studies and analysis. AT developed the study concept and design, contributed patients, assisted in the data extraction, performed statistical analysis, and wrote the paper.

**Compliance with ethical standards**

**Conflict of interest** The authors declare that they have no conflict of interests.

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**Lymphoma**

Genomic profiling reveals spatial intra-tumor heterogeneity in follicular lymphoma

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Follicular lymphoma (FL) is an incurable B-cell malignancy characterized by advanced stage disease and a heterogeneous clinical course, with high-risk groups including those that transform to an aggressive lymphoma, or progress early (within 2 years) following treatment. Recent sequencing studies have established the diverse genomic landscape and the temporal clonal dynamics of FL [1–7]; however, our understanding of the degree of spatial or intra-tumor heterogeneity (ITH) that exists within an individual patient is limited. In contrast, multi-site profiling in solid organ malignancies has demonstrated profound ITH impacting mechanisms of drug resistance and compromising