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A phase II, multicentre trial of decitabine in higher-risk chronic myelomonocytic leukemia

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(Article begins on next page)

1 **Decitabine in higher-risk chronic myelomonocytic leukemia: results from a**
2 **phase 2 Italian study**

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16

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22 MDS/MPN

23 Somatic mutations

24 **Abstract**

25 Chronic myelomonocytic leukemia (CMML) is a complex clonal hematological disorder classified
26 among myelodysplastic/myeloproliferative neoplasms (MDS/MPNs). Prognosis is poor and there is a
27 lack of effective treatments. The hypomethylating agent decitabine has shown activity against MDS
28 and elderly acute myeloid leukemia, but there is little data focusing specifically on its efficacy in
29 CMML. In this prospective, phase 2 Italian study, CMML patients received i.v. decitabine 20
30 mg/m²/day on Days 1 to 5 of a 28-day treatment cycle. Response was evaluated after 4 and 6 cycles;
31 patients responding at the end of 6 cycles could continue treatment with decitabine. Forty-three
32 patents were enrolled; >50% were high-risk according to four CMML-specific scoring systems. In the
33 intent-to-treat population (n = 42), the overall response rate after 6 cycles was 47.6%, with 7
34 complete responses (16.6%), 8 marrow responses (19%), 1 partial response (2.4%) and 4
35 hematological improvements (9.5%). After a median follow-up of 51.5 months (range: 44.4-57.2),
36 median overall survival was 17 months, with responders having a significantly longer survival than
37 non-responders (P=0.02). Grade 3/4 anemia, neutropenia and thrombocytopenia occurred in 28.6%,
38 50% and 38% of patients, respectively. Decitabine appears to be an effective and well tolerated
39 treatment for patients with high-risk CMML.

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45 INTRODUCTION

46 Chronic myelomonocytic leukemia (CMML) is a complex clonal hematological disorder that is
47 classified by the World Health Organization (WHO) among myelodysplastic/myeloproliferative
48 neoplasms (MDS/MPNs).¹ The 2016 revision to the WHO classification of tumours of the
49 hematopoietic and lymphoid tissues describes three categories of CMML based on blast count:^{2,3}
50 CMML-0 (<2% peripheral blasts and <5% bone marrow blasts), CMML-1 (2–4% peripheral blasts
51 and/or 5–9% bone marrow blasts) and CMML-2 (5–19% peripheral blasts, 10–19% bone marrow
52 blasts and/or presence of Auer rods). Before this revision, patients with <2% peripheral blasts and
53 <5% bone marrow blasts were included in the CMML-1 category.⁴

54 Diagnosis can be difficult, requiring a combination of morphologic, histopathologic and cytogenetic
55 approaches.⁵ The WHO diagnostic criteria for CMML are:^{2,3} persistent monocytosis $\geq 1 \times 10^9/L$; no
56 Philadelphia chromosome or *BCR-ABL1* fusion gene; exclusion of primary myelofibrosis,
57 polycythemia vera and essential thrombocythemia; no *PDGFRA*, *PDGFRB* or *FGFR1*
58 rearrangements or *PCM1-JAK2* fusions if eosinophilia present; <20% blasts in peripheral blood and
59 bone marrow; dysplasia in one or more myeloid lineages. If myelodysplasia is absent or minimal, a
60 diagnosis of CMML can still be made if a cytogenetic abnormality is present in the hematopoietic
61 stem cell, or if monocytosis has persisted for more than three months with all other possible causes
62 excluded.

63 Significant heterogeneity makes prognosis in CMML difficult to estimate, but in general it is poor.
64 Commonly used for MDS, the original and revised International Prognostic Scoring System (IPSS)^{6,7}
65 are not suitable for CMML because they exclude patients with proliferative disease. Newer
66 prognostic models (such as the CMML-specific prognostic scoring system (CPSS),^{8,9} Groupe
67 Francophone de Myelodysplasies (GFM) model,¹⁰ and the Mayo Molecular Model¹¹) take
68 cytogenetics and somatic mutations into account. Very recently, an integrated prognostic scoring

69 system has been proposed that takes clinical parameters, cytogenetics and somatic mutations into
70 account.¹²

71 The only potentially curative treatment option for CMML is hematopoietic stem cell transplant
72 (HSCT), but this is not suitable for many patients because of their age and comorbidities. There are
73 currently no prospective data on the benefits and risks of HSCT in CMML. Management usually
74 focuses on supportive care and cytoreductive therapy, depending on whether the disease is
75 dysplastic or myeloproliferative.¹³ Hydroxyurea is currently a mainstay therapy for proliferative
76 disease.¹⁴

77 The hypomethylating agents (HMAs) azacitidine and decitabine have been shown to be active in
78 MDS patients in randomized phase 3 trials.¹⁵⁻¹⁷ However, the numbers of CMML patients in these
79 trials were limited, and their results were not reported separately. In two retrospective analyses of
80 decitabine, overall response rates (ORRs) ranged from 26 – 68%, and 2-year survival from 25 –
81 48%.^{18,19} In a prospective phase 2 study in which 39 CMML patients received 20 mg/m² decitabine
82 per day on days 1 – 5 of 28-day cycles, the ORR was 38% and 2-year survival was 48%.²⁰

83 The European Medicines Agency has approved azacitidine for the treatment of non-proliferative
84 CMML (white blood cell (WBC) count <12,000), but HMAs are not currently a licensed option for
85 treating proliferative forms. In Italy, several national societies recommend that patients with
86 myelodysplastic-type CMML and ≥10% bone marrow blasts should be managed with supportive
87 therapy in combination with HMAs.¹³ Alongside the lack of specific treatment options, CMML-
88 specific response criteria were not used in any clinical trials, having only been recently developed by
89 Savona *et al.*²¹

90 Here, we report the results of a prospective phase 2 study that assessed the efficacy and safety of
91 decitabine in Italian CMML patients.

92

93 **METHODS**

94 Study design and patients

95 This was an open-label, phase 2 study carried out at 15 centres across Italy between April 2010 and
96 October 2011. Patients aged ≥ 18 years with a diagnosis of CMML according to WHO criteria,⁴ an
97 Eastern Cooperative Oncology Group (ECOG) performance status ≥ 2 , and a life expectancy ≥ 6
98 months were eligible to enter the study. Patients with a WBC count $\geq 12\ 000/\text{mm}^3$ were required to
99 have IPSS intermediate-2 risk. Those with a WBC count $< 12\ 000/\text{mm}^3$ had to have at least two of the
100 following: blast cells $> 5\%$ in bone marrow; a cytogenetic abnormality other than $t(5;12)(q33;p13)$;
101 anaemia (i.e. Hb < 10 g/dl); thrombocytopenia (i.e. platelets $< 100\ 000/\text{mm}^3$); splenomegaly (> 5 cm
102 below the costal margin); extramedullary localization. Patients with a myeloproliferative or
103 myelodysplastic syndrome other than CMML, and those who had acute blastic transformation of
104 CMML with bone marrow blast cells $> 20\%$ were excluded. Other exclusion criteria included eligibility
105 for allogenic stem cell transplant with an identified donor; CMML with $t(5;12)$ or *PDGFBR*
106 rearrangement; intensive chemotherapy in the last 3 months; previous treatment with a HMA.
107 Patients received i.v. decitabine (Dacogen®; Janssen Pharmaceutica NV, Beerse, Belgium) 20
108 mg/m²/day on Days 1 to 5 of a 28-day treatment cycle. Discontinuation was allowed at the patient's
109 request, or if they experienced progression with blastic transformation, grade 3/4 toxicity according
110 to National Cancer Institute criteria (except cytopenia), or other changes in their condition that the
111 investigator felt warranted removal of the patient from the study.

112 After a minimum of 4 treatment cycles, patients were assessed for response to treatment.
113 Responders were defined as patients who achieved hematological improvement (HI) or better
114 according to International Working Group 2006 criteria;²² these patients continued treatment for a
115 further 2 cycles. Minor responders and patients with stable disease were allowed to continue in the
116 study at the investigator's discretion. Patients with progressive disease were discontinued from the

117 study. Patients who completed all 6 treatment cycles were eligible to receive maintenance
118 treatment with decitabine. After completion of, or discontinuation from, the study, patients were
119 followed-up every 4 months.

120 The study was carried out in accordance with the Declaration of Helsinki. All patients provided
121 written informed consent and all participating trial sites gained approval from the relevant local
122 ethics committee. This study is registered on ClinicalTrials.gov (NCT01251627).

123

124 Objectives and outcome measures

125 The primary aim of the study was to assess the efficacy of decitabine in the treatment of CMML. The
126 primary outcome measure was the ORR, defined as the proportion of patients achieving a complete
127 response (CR), marrow CR (mCR), partial response (PR) or HI. Secondary outcome measures
128 included: overall survival (OS); event-free survival (EFS); duration of response; the number of blood
129 and platelet transfusions; the number of days of hospitalization; safety.

130

131 Somatic mutations

132 Bone marrow samples were collected before treatment and DNA was extracted from unsorted
133 mononuclear cells. The methods used for mutational sequencing have been published previously.²³
134 Briefly, target regions (exons plus splice junctions) were captured and 500 ng of DNA from each
135 sample was quantified and sequenced using paired-end sequencing. Sequences were aligned to the
136 human genome and the Genome Analysis Toolkit²⁴ was used to perform further local indel alignment
137 and base-quality score recalibration, and to generate single nucleotide variation and indel calls.
138 Variants with functional consequence on genes were annotated and their presence identified in
139 dbSNP 137, the 1000 Genomes Project, ESP6500 (the National Heart, Lung and Blood Institute GO
140 Exome Sequencing Project), and COSMIC 67.

141

142 Statistical analyses

143 Conventional treatments for CMML (hydroxyurea or etoposide) give ORRs of no more than 15%. This
144 study was designed to detect a clinically relevant 20% increase in ORR with decitabine (i.e. from 15%
145 to 35%) with 85% power and a significance level of 0.05. The planned sample size for this single-
146 stage Fleming-A'Hern phase 2 design was 39 patients. Achievement of OR by ≥ 11 patients after 6
147 cycles was to be considered sufficient to justify further investigation. To take into account losses to
148 follow-up for time-to-event endpoints, the sample size was increased by 10%, and 43 patients were
149 enrolled.

150 Primary efficacy and safety data were analyzed for the intent-to-treat (ITT) population, i.e. all
151 patients who received at least one dose of decitabine. Discrete variables were summarized by
152 frequency and percentage. Continuous variables were summarized by mean and standard deviation
153 (SD) or median and interquartile range (IQR).

154 OS was defined as the time from enrolment to death from any cause or last follow-up evaluation.
155 EFS was defined as the time from enrolment to progression, transformation to acute myeloid
156 leukemia (AML), or death from any cause. Duration of response was defined as the time from clinical
157 response to progression, transformation to AML or death from any cause. These time-to-event
158 endpoints were analyzed using the Kaplan-Meier method.

159 Adverse events (AEs) were reported by type and grade according to the Common Terminology
160 Criteria for Adverse Events (version 3.0).

161

162 **RESULTS**

163 Patients

164 Between April 2010 and October 2011, 43 patients were enrolled at 15 sites across Italy. The ITT
165 population included 42 patients; their baseline characteristics are shown in Table 1. Most patients
166 were male (71.4%) and two-thirds had proliferative CMML. Between 76% and 93% of patients were
167 high- or intermediate-risk (depending on the prognostic scoring system; Table 1), and approximately
168 one-third had an *ASXL1* mutation.

169 Figure 1 shows the flow of patients through the study. The median number of treatment cycles was
170 6 (range: 1–34). Twenty-six patients (62%) received all 6 cycles; the most common reasons for
171 discontinuation were treatment failure (n = 9) and death (n = 5).

172 Somatic mutations

173 The results of the analysis of the most frequent somatic mutation found in CMML were not possible
174 in all cases in this study, and have already been published previously in a study in which a subset of
175 the CMML patients treated with decitabine was analysed for methylation pattern, gene expression
176 profile and presence of somatic mutations²³. The incidence and type of mutation at diagnosis is
177 presented in Table 1. The most frequent mutations as expected were those of *SRSF2* (45.2%), *TET2*
178 (38.1%) and *ASXL1* (35.7%).

179 There was no correlation between the presence of a single mutation and pattern of response to
180 decitabine.

181

182 Response rates

183 In the ITT population, the ORR was 47.6% (Table 2). Patients with CMML-1 had a higher ORR than
184 those with CMML-2 (53.8% vs. 37.5%, respectively). The ORR was also higher in patients with
185 dysplastic CMML (64.3%) than in those with proliferative CMML (39.3%).

186 For the ITT population, the binominal proportion (95% CI) of ORR after 6 cycles was 0.40 (0.25-0.56).
187 The lower limit of the CI was superior to the null hypothesis (i.e. that ORR = 0.15) and the *P*-value
188 was <0.0001, indicating that decitabine is significantly more effective than standard therapy. **[Please**
189 **advise if this paragraph is to remain. If so, do the data need updating – this was taken from the**
190 **original results document from December 2015].**

191

192 Survival and progression

193 The median duration of follow-up was 51.5 months (range: 44.4–57.2). Median OS was 17 months
194 (Figure 2A). The 1-year, 2-year and 3-year OS rates were 66.7%, 33.3% and 28.6%, respectively.
195 Patients who responded to treatment had a significantly longer OS than those who did not (*P* = 0.02;
196 Figure 2B). Median EFS was 8 months (Figure 3). The most common event was death, which
197 occurred in 36 patients (85.7%). Thirty-two patients (76.2%) progressed and 24 (57.1%) had
198 transformation to AML. The 1-year, 2-year and 3-year EFS rates were 35.7%, 21.4% and 19.1%,
199 respectively. The median duration of response was 10 months (Figure 4). At 1-year, 52.6% of
200 responders were still responding to treatment; the corresponding figures at 2 and 3 years were
201 42.1% and 26.3%, respectively.

202

203 Transfusions and hospitalizations

204 At baseline, 18 patients (42.9%) required transfusions. During the treatment period, transfusions
205 were carried out during 117 cycles; 39 patients needed at least one transfusion. During follow-up, 21
206 patients needed at least one transfusion.

207 During the treatment period, 9 patients had a total of 24 scheduled hospital admissions. The median
208 length of hospitalization was 6.5 (range 1.0-31.0) days. Five patients each had one unscheduled
209 hospital admission; the median length of their hospital stay was 7.0 (range 1.0-21.0) days. During
210 follow-up, 10 patients had a total of 21 hospital admissions.

211

212 Safety

213 The most common AEs were haematological: thrombocytopenia, anemia and neutropenia (Table 3).
214 In 50% of cases, anemia was grade 3 or 4. More than three-quarters of thrombocytopenia cases, and
215 over 80% of neutropenia cases were grade 3 or 4. The most common non-hematological AEs were
216 gastrointestinal; all of these were grade 1 or 2. Two patients had grade 5 AEs: one cardiac event and
217 one bleeding event. Thirty-six patients died: 5 during the six-month study period and 31 during
218 follow-up. In 29 of these patients, MDS was the cause of death.

219

220 **DISCUSSION**

221 CMML is a disease that is difficult to diagnose and has a poor prognosis. There are currently no
222 effective treatments for patients who are unsuitable for HSCT. Dysplastic and proliferative forms of
223 CMML are likely to require different treatment approaches. Current recommendations are to treat
224 dysplastic CMML with supportive care plus azacitidine, and proliferative CMML with cytoreductive
225 therapy to control proliferation and reduce organomegaly.¹³

226 In our study, decitabine induced a response in approximately half of patients, with responders
227 having a survival advantage over non-responders. Although patients with CMML-2 and those with
228 proliferative disease had lower response rates than those with CMML-1 or dysplastic disease, the
229 results in these subgroups were encouraging. Decitabine was well tolerated in our elderly cohort;
230 the incidence and type of AEs were as expected.

231 The ORR in our study is slightly higher than that of 38% reported in a previous study in 39 CMML
232 patients conducted by the GFM.²⁰ Whereas we found CMML-1 patients to be more likely to respond
233 to treatment, the GFM CMML study showed the opposite, with 50% of CMML-2 patients responding,
234 compared with 17.6% of CMML-1 patients. Median OS was similar in our study and the GFM CMML
235 study, but 2-year OS was lower in our study (33.3% vs. 48%).

236 Previous studies in mixed cohorts of patients with MDS and related malignancies have linked
237 increased response to HMAs to mutations in *TET2*^{25,26} (particularly when *ASXL1* is not mutated²⁶) and
238 *DNTM3A*.²⁵ However, in a cohort of 40 patients from the present study, we found that no somatic
239 mutation, including *ASXL1*, was predictive of response to decitabine in CMML.²³ Likewise, the GFM
240 CMML study also found no association between response to decitabine and mutational status.²⁰ This
241 may indicate a difference in the impact of mutational status between patients with CMML and those
242 with other myeloid malignancies. In addition, the studies showing an association between somatic
243 mutations and response included patients who received azacitidine, as well as patients who received
244 decitabine.^{25,26}

245 Although somatic mutations did not differentiate responders from non-responders in our cohort, we
246 found a pattern of 167 differentially methylated regions of DNA that was predictive of response.²³
247 Using this, we developed an epigenetic classifier that can accurately predict response to decitabine
248 at the time of diagnosis. It can take several cycles of treatment before it becomes apparent whether
249 the patient will respond or not; this classifier would allow potential non-responders to be identified
250 early and put onto an alternative treatment, rather than having to endure months of fruitless
251 treatment with decitabine.

252 Many of the patients in our cohort were high risk according to the prognostic scoring systems used,
253 so we could not determine whether high- and low-risk patients have a differential sensitivity to
254 decitabine. The median OS was 17 months, which compares favourably with best supportive care
255 and hydroxyurea.^{9,10} Such et al used a cohort of patients receiving best supportive care when
256 developing the CPSS prognostic scoring system. Patients who fell into the high-risk category had a
257 median OS of 5-9 months.⁹ Patients classed as high-risk according to the GFM prognostic scoring
258 system had a median OS of 14.4 months.¹⁰ Most patients in this latter study were receiving best
259 supportive care, but hydroxyurea and HMAs were also used. High-risk patients according to the
260 Mayo Molecular Model had a median OS of 16 months; the authors do not report what treatment(s)
261 the patients were receiving.¹¹

262 Decitabine appears to be an effective treatment for patients with high-risk CMML, including those
263 with proliferative disease. Further research is needed to determine whether there is a difference in
264 response between low- and high-risk patients. Owing to the rarity of CMML, large, specific trials can
265 be difficult to conduct. However, we are currently conducting (within the guidance of the European
266 MDS Studies Coordination Office) an international, randomized, phase 3 trial comparing decitabine
267 (\pm hydroxyurea) with hydroxyurea in patients with advanced proliferative CMML (ClinicalTrials.gov
268 identifier: NCT 02214407). The results of this trial will provide further important insights into the

269 efficacy of decitabine as a treatment for CMML, particularly in patients with proliferative disease, for
270 whom treatment with HMAs is currently not a licensed option.

271

272

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278

279 **REFERENCES**

- 280 1. Jaffe ES, Harris NL, Stein H, Vardiman JW (eds). World Health Organization Classification of
281 Tumours. Pathology and Genetics of Tumours of Haematopoietic and Lymphoid Tissues. IARC:
282 Lyon, France, 2001.
- 283 2. Swerdlow SH et al (eds). World Health Organization Classification of Tumours of
284 Haematopoietic and Lymphoid Tissues. IARC: Lyon, France, 2016. In press.
- 285 3. Arber DA, Orazi A, Hasserjian R, Thiele J, Borowitz MJ, Le Beau MM et al. The 2016 revision to
286 the World Health Organization classification of myeloid neoplasms and acute leukemia. *Blood*
287 2016; **127**: 2391-405.
- 288 4. Swerdlow SH, Campo E, Harris NL, Jaffe ES, Pileri SA, Stein H et al (eds). World Health
289 Organization Classification of Tumours of Haematopoietic and Lymphoid Tissues. IARC: Lyon,
290 France, 2008.
- 291 5. Parikh S, Tefferi A. Chronic myelomonocytic leukemia: 2013 update on diagnosis, risk
292 stratification, and management. *Am J Hematol* 2013; **88**: 968-74.
- 293 6. Greenberg P, Cox C, LeBeau MM, Fenau P, Morel P, Sanz G et al. International scoring system
294 for evaluating prognosis in myelodysplastic syndromes. *Blood* 1997; **89**: 2079-88.
- 295 7. Greenberg PL, Tuechler H, Schanz J, Sanz G, Garcia-Manero G, Solé F et al. Revised international
296 prognostic scoring system for myelodysplastic syndromes. *Blood* 2012; **120**: 2454-65.
- 297 8. Such E, Cervera J, Costa D, Solé F, Vallespí T, Luño E et al. Cytogenetic risk stratification in
298 chronic myelomonocytic leukemia. *Haematologica* 2011; **96**: 375-83.
- 299 9. Such E, Germing U, Malcovati L, Cervera J, Kuendgen A, Della Porta MG et al. Development and
300 validation of a prognostic scoring system for patients with chronic myelomonocytic leukemia,
301 *Blood* 2013; **121**: 3005-15.

- 302 10. Itzkyson R, Kosmider O, Renneville A, Gelsi-Boyer V, Meggendorfer M, Morabito M et al.
303 Prognostic score including gene mutations in chronic myelomonocytic leukemia. *J Clin Oncol*
304 2013; **31**: 2428-36.
- 305 11. Patnaik MM, Itzkyson R, Lasho TL, Kosmider O, Finke CM, Hanson CA et al. ASXL1 and SETBP1
306 mutations and their prognostic contribution in chronic myelomonocytic leukemia: a two-center
307 study of 466 patients. *Leukemia* 2014; **28**: 2206-12.
- 308 12. Elena C, Gallì A, Such E, Meggendorfer M, Germing U, Rizzo E et al. Integrating clinical features
309 and genetic lesions in the risk assessment of patients with chronic myelomonocytic leukemia.
310 *Blood* 2016; **128**: 1408-17.
- 311 13. Onida F, Barosi G, Leone G, Malcovati L, Morra E, Santini V et al. Management
312 recommendations for chronic myelomonocytic leukemia: consensus statement from the SIE,
313 SIES, GITMO groups. *Haematologica* 2013; **98**: 1344-52.
- 314 14. Patnaik MM, Tefferi A. Chronic myelomonocytic leukemia: focus on clinical practice. *Mayo Clin*
315 *Proc* 2016; **91**: 259-72.
- 316 15. Silverman LR, Demakos EP, Peterson BL, Kornblith AB, Holland JC, Odchimar-Reissig R et al.
317 Randomized controlled trial of azacitidine in patients with the myelodysplastic syndrome: a
318 study of the cancer and leukemia group B. *J Clin Oncol* 2002; **20**: 2429-40.
- 319 16. Fenaux P, Mufti G, Hellstrom-Lindberg E, Santini V, Finelli C, Giagounidis A et al. Efficacy of
320 azacitidine compared with that of conventional care regimens in the treatment of higher-risk
321 myelodysplastic syndromes: a randomised, open-label, phase III study. *Lancet Oncol* 2009; **10**:
322 223-32.
- 323 17. Kantarjian H, Issa J-PJ, Rosenfeld CS, Bennett JM, Albitar M, DiPersio J et al. Decitabine improves
324 patient outcomes in myelodysplastic syndromes. Results of a phase III randomized study.
325 *Cancer* 2006; **106**: 1794-80.
- 326 18. Aribi A, Borthakur G, Ravandi F, Shan J, Davisson J, Cortes J et al. Activity of decitabine, a
327 hypomethylating agent, in chronic myelomonocytic leukemia. *Cancer* 2007; **109**: 713-7.

- 328 19. Wijermans PW, Rüter B, Baer MR, Slack JL, Saba HI, Lübbert M. Efficacy of decitabine in the
329 treatment of patients with chronic myelomonocytic leukemia (CMML). *Leuk Res* 2008; **32**: 587-
330 91.
- 331 20. Braun T, Itzkyson R, Renneville A, de Renzis B, Dreyfus F, Laribi K et al. Molecular predictors of
332 response to decitabine in advanced chronic myelomonocytic leukemia: a phase 2 trial. *Blood*
333 2011; **118**: 3824-31.
- 334 21. Savona MR, Malcovati L, Komrokji R, Tiu RV, Mughal TI, Orazi A et al. An international
335 consortium proposal of uniform response criteria for myelodysplastic/myeloproliferative
336 neoplasms (MDS/MPN) in adults. *Blood* 2015; **125**: 1857-65.
- 337 22. Cheson BD, Greenberg PL, Bennett JM, Lowenberg B, Wijermans PW, Nimer SD et al. Clinical
338 application and proposal for modification of the International Working Group (IWG) response
339 criteria in myelodysplasia. *Blood* 2006; **108**: 419-25.
- 340 23. Meldi K, Qin T, Buchi F, Droin N, Sotzen J, Micol J-P et al. Specific molecular signatures predict
341 decitabine response in chronic myelomonocytic leukemia. *J Clin Invest* 2015; **125**: 1857-72.
- 342 24. DePristo MA, Banks E, Poplin R, Garimella KV, Maguire JR et al. A framework for variation
343 discovery and genotyping using next-generation DNA sequencing data. *Nat Genet* 2011; **43**:
344 491-8.
- 345 25. Traina F, Visconte V, Elson P, Tabarrokhi A, Jankowska AM, Hasrouni E et al. Impact of molecular
346 mutations on treatment response to DNMT inhibitors in myelodysplasia and related neoplasms.
347 *Leukemia* 2014; **28**: 78-87.
- 348 26. Bejar R, Lord A, Stevenson K, Bar-Natan M, Pérez-Lagada A, Zaneveld J et al. *TET2* mutations
349 predict response to hypomethylating agents in myelodysplastic syndrome patients. *Blood* 2014;
350 **124**: 2705-12.
- 351
- 352

353 **FIGURE LEGENDS**

354 **Figure 1.** Patient disposition. Flow of patients from enrolment to time of analysis.

355 **Figure 2.** Overall survival. Kaplan-Meier curves showing overall survival in (A) the ITT population and
356 (B) responders vs. non-responders. Vertical lines denote censored patients.

357 **Figure 3.** Event-free survival. Kaplan-Meier curves showing event-free survival in the ITT population.
358 Vertical lines denote censored patients.

359 **Figure 4.** Duration of response. Kaplan-Meier curves showing duration of response. Vertical lines
360 denote censored patients.

361

362 **Table 1.** Baseline characteristics (ITT population)

Patients, n	42
Median age, years (range)	71.5 (42-84)
Male/female, n (%)	30/12 (71.4/28.6)
CMML-1/CMML-2 ^a , n (%)	26/16 (61.9/38.1)
dCMML/pCMML, n (%)	14/28 (33.3/66.7)
Hb g/dl, median (IQR)	9.8 (9.1-11.0)
AMC x 10 ⁹ /l, median (IQR)	3.39 (2.23-7.25)
WBC x 10 ⁹ /l, median (IQR)	18.6 (13.9-28.1)
PLT x 10 ⁹ /l, median (IQR)	54.5 (34.0-75.0)
Bone marrow blasts %, median (IQR)	6.0 (3-12)
<i>Cytogenetics, n (%)</i>	
Altered	12 (28.6)
Normal	28 (66.7)
Not evaluable	2 (4.7)
Splenomegaly, n (%)	22 (52.4)
Hepatomegaly, n (%)	19 (45.2)
Lymphadenomegaly, n (%)	6 (14.3)
Marrow fibrosis, n (%)	13 (30.9)
<i>ASXL1, n (%)</i>	
Mutated	15 (35.7)
Not evaluable	4 (9.5)
<i>SRSF2, n (%)</i>	
Mutated	19 (45.2)
Not evaluable	5 (11.9)
<i>TET2, n (%)</i>	
Mutated	16 (38.1)
Not evaluable	5 (11.9)
<i>P53, n (%)</i>	
Mutated	3 (7.1)
Not evaluable	5 (11.9)

MMM prognostic risk categories, n (%)

High risk	13	(30.9)
Int-2	14	(33.3)
Int-1	10	(23.8)
Not evaluable	5	(12.0)

CPSS prognostic risk categories, n (%)

High risk	3	(7.1)
Int-2	20	(47.6)
Int-1	15	(35.7)
Low	2	(4.8)
Not evaluable	2	(4.8)

Mayo prognostic risk categories, n (%)

High risk	26	(61.9)
Int	13	(30.6)
Low	3	(7.1)

GFM prognostic risk categories, n (%)

High	14	(33.3)
Int	18	(42.9)
Low	10	(23.8)

363 Abbreviations: AMC, absolute monocyte count; CMML, chronic monomyelocytic leukemia; dCMML, dysplastic
364 CMML; CPSS, CMML-specific prognostic scoring system; GFM, Groupe Francophone de Myelodysplasies; Hb,
365 hemoglobin; IQR, interquartile range; MMM, Mayo Molecular Model; PLT, platelet; pCMML, proliferative
366 CMML; WBC, white blood cells. Percentages may not total 100 owing to rounding. ^aDefined according to the
367 2008 edition of the WHO classification of tumours of the hematopoietic and lymphoid tissues.

368 **Table 2.** Overall clinical response (end of cycle 6 or at early withdrawal)

	Number (%) of patients				
	ITT (n = 42)	CMML-1 ^a (n = 26)	CMML-2 ^a (n = 16)	dCMML (n = 14)	pCMML (n = 28)
ORR	20 (47.6)	15 (57.6)	5 (31.25)	9 (64.3)	11 (39.3)
CR	7 (16.6)	5 (19.2)	2 (12.5)	3 (21.4)	4 (14.3)
mCR	8 (19.0)	6 (23.1)	2 (12.5)	4 (28.6)	4 (14.3)
PR	1 (2.4)	0 (0.0)	1 (6.2)	0 (0.0)	1 (3.5)
HI	4 (9.5)	4 (15.3)	0 (0.0)	2 (14.2)	2 (7.2)
SD	9 (21.4)	4 (15.3)	5 (31.3)	0 (0.0)	9 (32.1)
PD	13 (31.0)	7 (26.9)	6 (37.5)	5 (35.7)	8 (28.6)

369 Abbreviations: CR, complete response; dCMML, dysplastic CMML; HI, haematological improvement; ORR, overall response rate;

370 mCR, marrow CR; pCMML, proliferative CMML; PD, progressive disease; PR, partial remission; SD, stable disease. Percentages may not total 100 owing to rounding.

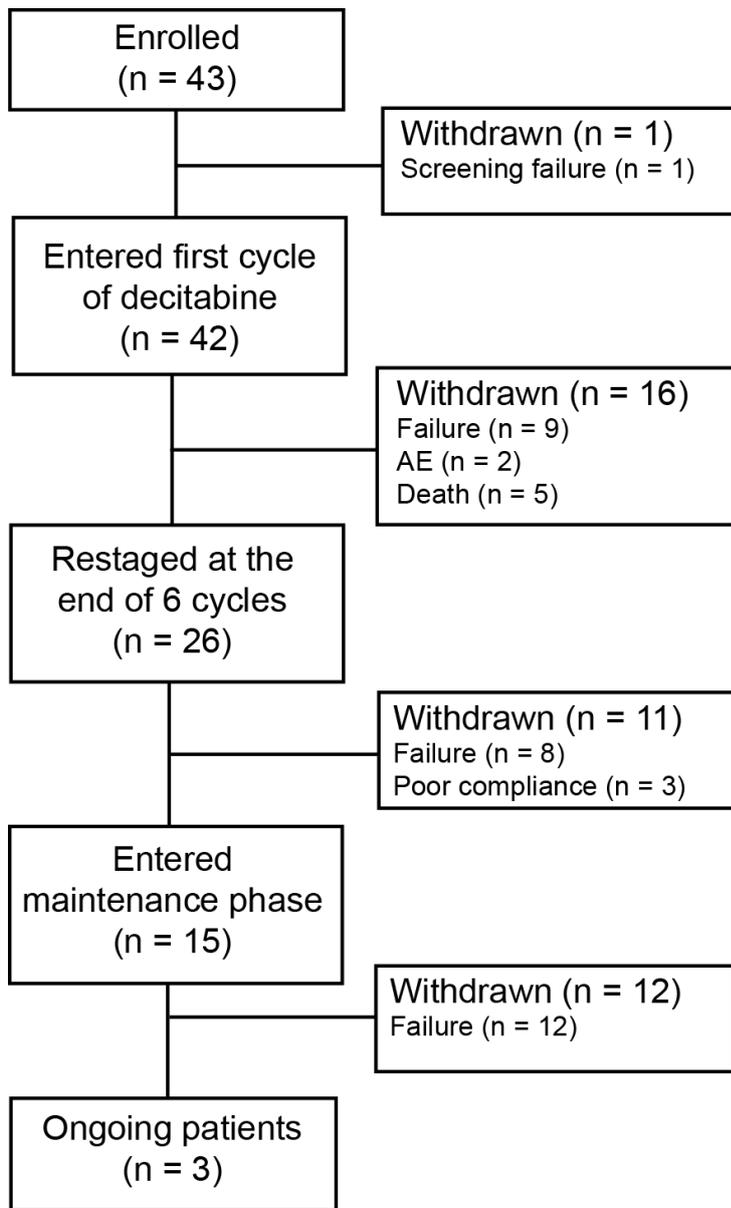
371 ^aDefined according to the 2008 edition of the WHO classification of tumours of the hematopoietic and lymphoid tissues.

372 **Table 3.** Adverse events (ITT population; n = 42)

	Number (%) of patients		
	<i>Any grade</i>	<i>Grade 3</i>	<i>Grade 4-5</i>
Anemia	24 (57.1)	11 (26.2)	1 (2.4)
Thrombocytopenia	27 (64.3)	4 (9.5)	17 (63.1)
Neutropenia	19 (45.2)	7 (18.7)	9 (21.4)
Cardiac	2 (4.8)	-	1 ^a (2.4)
Neurological	1 (2.4)	-	-
Gastrointestinal	10 (23.8)	-	-
Hepatic	2 (4.8)	1 (2.4)	-
Documented infection	6 (14.3)	2 (4.8)	1 (2.4)
Bleeding	9 (21.4)	-	1 ^a (2.4)
Febrile neutropenia	2 (4.8)	2 (4.8)	-
Other	8 (19.0)	1 (2.4)	-

373 ^aGrade 5 event

374 **Figure 1.** Patient disposition

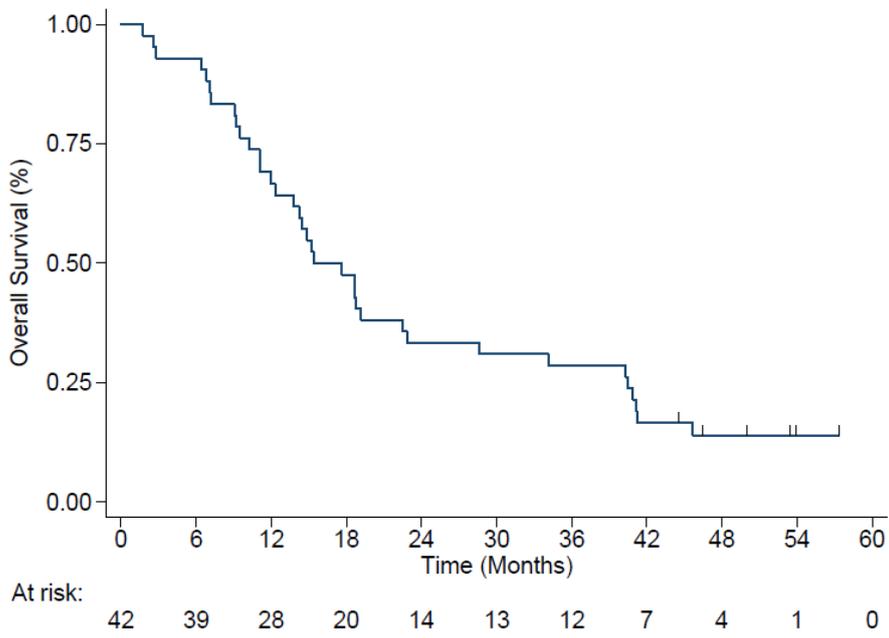


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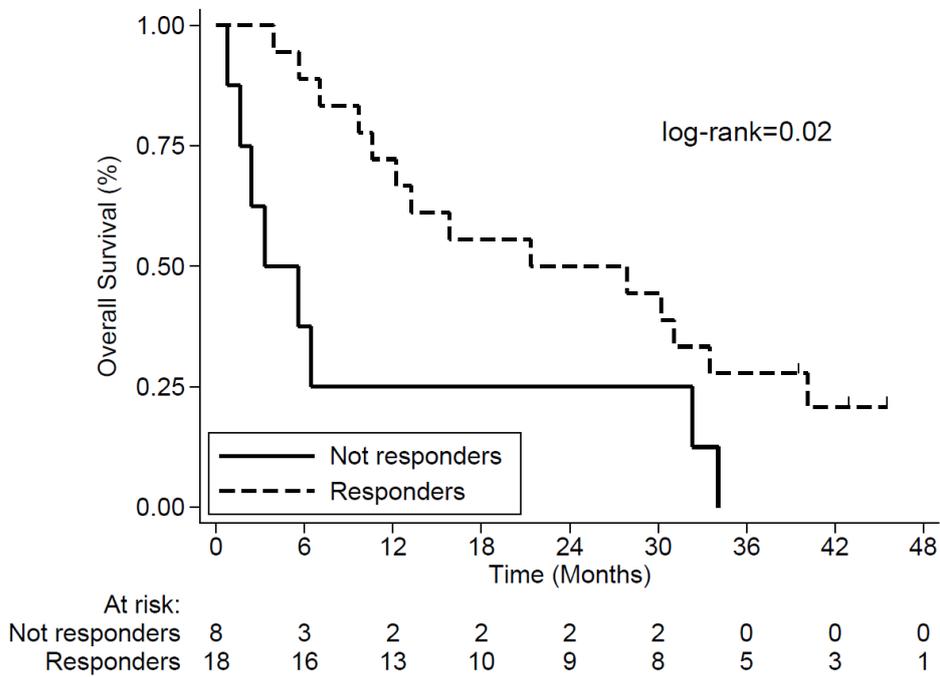
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377 **Figure 2.** Overall survival.

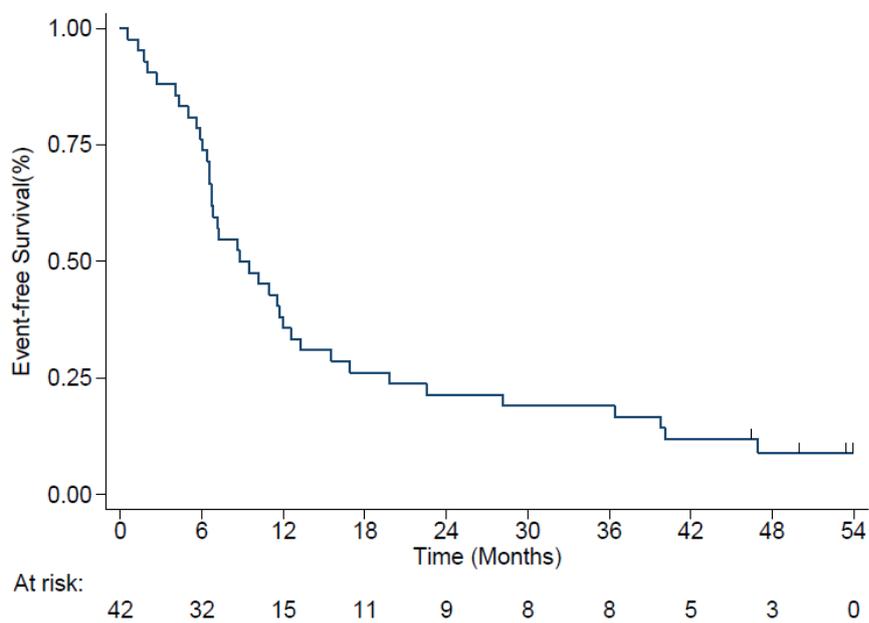
378 **A**



389 **B**



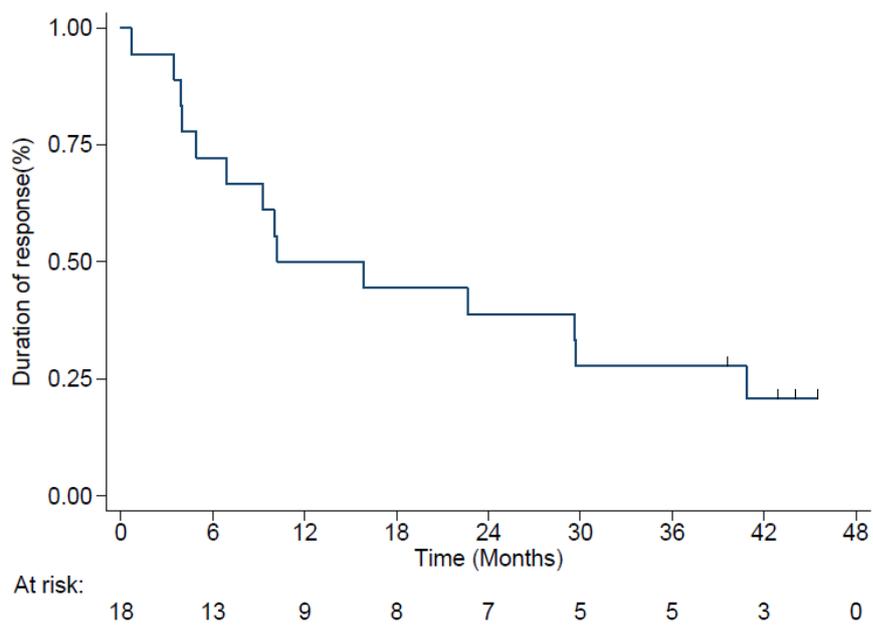
402 **Figure 3.** Event-free survival.



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404

405 **Figure 4.** Duration of response.



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