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1 **Letter to the editor (1500 words)**

2

3 **Clinical relevance of failed and missing cytogenetic analysis in acute myeloid leukaemia (AML)**

4

5 Chromosomal and genetic abnormalities are important prognostic factors in AML and most clinically  
6 relevant aberrations are detectable by cytogenetic analysis.(1, 2) One limitation of cytogenetics is  
7 failure due to a lack of analysable metaphases. Despite this shortcoming, chromosomal analysis  
8 remains the gold standard test for identifying abnormalities used to risk-stratify treatment because  
9 many abnormalities (e.g. those involving large chromosomal regions and a complex karyotype) can  
10 only be described in cytogenetic terms. Thus, failure to obtain a cytogenetic result impacts on risk  
11 stratification. In support of this suggestion, two recent reports concluded that failed and/or missing  
12 cytogenetic results were associated with an adverse prognosis.(3, 4) Our view is that assignment of  
13 risk on the basis of the absence of information is counterintuitive and potentially problematic.  
14 Therefore, we investigated the distribution and prognostic impact of failed and missing cytogenetic  
15 results in successive MRC AML trials.

16

17 Cytogenetic analysis of pre-treatment bone marrow or peripheral blood samples was performed  
18 locally, reviewed and collated by the Leukaemia Research Cytogenetics Group. Results were  
19 available from 10,685 patients (1-82 years old) recruited to successive trials (AML12, AML14, AML15,  
20 AML16) between 1995-2012.(1, 5-9) At diagnosis, patients recruited to AML14 and AML16 were  
21 classified, on the basis of presenting features, as suitable or unsuitable for intensive therapy. All  
22 studies were approved by the relevant ethics committees and informed consent was obtained in  
23 accordance with the Declaration of Helsinki.

24

25 Karyotypes were described according to ISCN.(10) If the regional cytogenetic laboratory received a  
26 sample within the diagnostic window (30 days prior to or 7 days after diagnosis) cytogenetic testing  
27 was deemed to have been attempted (“Sample”); otherwise cytogenetic analysis was classified as  
28 missing (“No sample”). Analysis was defined as “Successful” if a clonal chromosomal abnormality  
29 was detected or  $\geq 20$  normal metaphases were fully analysed; otherwise it was classified as  
30 “Failed”.(11)

31

32 Survival was calculated from trial entry to death or last follow-up. Patients were censored at  
33 31/10/2010 (AML12, AML14) or 01/01/2012 (AML15, AML16) when follow-up was complete for 95%  
34 of the patients. Survival rates were calculated and compared using the Kaplan-Meier method, log-  
35 rank test, and Cox regression model. Comparisons between groups were performed using logistic  
36 regression,  $\chi^2$  test or Wilcoxon rank-sum test. Multivariate logistic regression was used to determine  
37 predictors for missing and failed cytogenetics. As recruitment and eligibility changed by trial and  
38 diagnosis period, all odds (OR) and hazard (HR) ratios were adjusted for intensive (INT) versus non-  
39 intensive (NI) treatment and year of diagnosis. All P-values were two-tailed. Statistical analyses were  
40 performed using SAS v9.3 (SAS Institute Inc., Cary, NC, USA).

41

42 Cytogenetic analysis was attempted in 94% INT patients but in only 83% NI patients ( $p < 0.0001$ ).  
43 Among INT patients, cytogenetic analysis was attempted more frequently among patients for whom  
44 the result would affect treatment; that is, younger patients treated on AML12 or AML15 (Table 1).  
45 This correlated with the observation that cytogenetic uptake was lower, and did not vary by age or  
46 trial, among NI patients. Similarly, cytogenetic uptake was higher among patients with de novo or  
47 therapy-related AML than for those with an antecedent hematologic disease. These patients were  
48 likely to have had cytogenetic analysis carried out at the time of initial diagnosis and subsequent  
49 testing may have been deemed unnecessary. Surprisingly, the uptake of cytogenetic analysis

50 decreased marginally across successive trials and by period of diagnosis. However, it should be  
51 noted that recruitment rates and patterns changed significantly over this period with a greater  
52 number of smaller regional hospitals participating in later trials. Across the whole cohort,  
53 multivariate analysis revealed that secondary disease (OR 2.08 (95% confidence interval 1.73-2.49),  
54  $p < 0.0001$ ), white blood cell count (WBC) (0.82 per 10-fold increase (0.74-0.92),  $p = 0.0002$ ), and age  
55 (1.10 per decade (1.04-1.16)  $p = 0.0009$ ) were the most significant predictors of cytogenetic testing.  
56 Similar results were obtained when INT and NI patients were examined separately; although age was  
57 not significant in the latter group (Table 1). Among INT patients, a lack of cytogenetic testing was  
58 associated with an inferior OS: 27% v 38%, HR = 1.41 (1.26-1.58),  $p < 0.0001$  (Figure 1A). However,  
59 this effect was restricted to younger adults (OS 35% v 45%, 1.41 (1.20-1.64),  $p < 0.0001$ ), and not  
60 observed among children (65% v 66%, 0.95 (0.43-2.09),  $p = 0.9$ ) or older adults (11% v 14%, 1.07  
61 (0.92-1.24),  $p = 0.4$ ) (Figure 1B) ( $p$  value for heterogeneity = 0.01). Similar results were obtained when  
62 the analysis was adjusted for age, WBC, secondary disease and performance status.

63

64 The frequency of cytogenetic testing among NI patients was similar to the Swedish study (4) which  
65 excluded NI patients (83% v 80%). Interestingly, cytogenetics was not used to guide therapy in  
66 Sweden during the study timeframe, which may explain the low uptake of cytogenetic testing;  
67 similar to the rate among NI patients in this study. Lazarevic et al concluded that patients without  
68 cytogenetic testing had an inferior outcome; similar to that for high risk cytogenetic patients.  
69 However, the survival of patients with and without cytogenetic testing was similar (28% v 22%). In  
70 contrast, we found that the association between inferior outcome and the uptake of cytogenetics  
71 was only significant among younger adults. Moreover, among younger adults the survival of patients  
72 without cytogenetic testing (35%) was closer to those with intermediate rather than high risk  
73 cytogenetics (33% and 12%, respectively).(1) In this study, a lack of cytogenetic testing was  
74 associated with other high risk features (age and secondary disease) which are established  
75 prognostic factors. Although cytogenetic testing was associated with an inferior outcome in

76 multivariable analyses (HR 1.13 (1.01-1.23)  $p=0.04$ ) the size of the effect was diminished indicating  
77 that other factors like secondary disease are also important. The Swedish study did not report the  
78 frequency of secondary disease and it is likely to be higher in a population-based study than a clinical  
79 trial. Collectively these findings indicate that there are numerous factors governing the uptake of  
80 cytogenetic testing at the time of diagnosis; many of which are also likely to impact on survival. Also,  
81 there are likely to be additional factors that cannot be examined in centralised retrospective studies.

82

83 A successful cytogenetic result was obtained in 90% cases and there was no difference according to  
84 treatment intensity (Table 1). Among NI patients, there were no significant predictors of cytogenetic  
85 failure whereas age and increasing WBC correlated with higher cytogenetic failure rates among INT  
86 patients. The variation in failure rate by trial was linked to age as AML14 and AML16, trials for older  
87 adults, had the highest failure rates. Multivariate logistic regression analysis revealed that the key  
88 predictor of cytogenetic failure was age (OR 1.14 per decade (1.09-1.19),  $p<0.0001$ ) and, to a lesser  
89 extent, WBC (OR 1.11 per 10-fold increase (1.01-1.22),  $p=0.04$ ). The link between age and  
90 cytogenetic failure could be explained by the increasing frequency of normal karyotype with age(12)  
91 and the fact that the threshold used to distinguish normal and failed cytogenetic result has shifted  
92 over time.(11) The link between cytogenetic failure and high WBC may be due to overcrowding of  
93 accumulated blasts within the bone marrow leading to inhibition of cell division, an observation  
94 which has often been made within routine preparation of leukaemic samples (unpublished  
95 observation). There was no association between cytogenetic failure and survival either overall  
96 (Figure 1C) or within different age groups for INT or NI patients. The OS rates for children, young  
97 adults and older adults treated intensively with successful and failed cytogenetics was: 38% v 37%,  
98 1.04 (0.95-1.13),  $p=0.4$ ; 45% v 48%,  
99 0.94 (0.84-1.05),  $p=0.3$ ; 15% v 13%, 0.99 (0.87-1.12),  $p=0.9$ , respectively. In contrast, the SWOG and  
100 Swedish studies (3, 4) concluded that cytogenetic failure was associated with an inferior outcome.

101 However, they compared patients whose samples failed cytogenetic testing to those stratified by  
102 cytogenetic risk. There is no biological reason why patients with failed cytogenetics should differ  
103 from those with successful cytogenetics; in fact there is evidence to the contrary.(13) Hence a  
104 successful versus failed comparison is the most informative analysis. The survival of patients with  
105 successful and failed cytogenetics in the SWOG and Swedish studies were not very different (21% v  
106 16% and 28% v 25% respectively). Cytogenetic failure was higher in our study (~10%) than the SWOG  
107 and Swedish studies (6% and 3%) because we used a definition based on the likelihood of detecting  
108 a clonal chromosomal abnormality.(11) Applying this stricter definition would move cases from the  
109 intermediate risk to the failed category; hence would not have altered the conclusions from the  
110 other studies. The factors governing cytogenetic failure are not fully understood but sample  
111 transport and processing are likely to be more important than underlying biological factors.(13, 14)  
112 Hence there is no rationale as to why cytogenetic failure should be linked to outcome.

113

114 Given the importance of genetics in guiding therapy in AML, the reasons for not sending a sample for  
115 analysis do warrant further investigation; but this must be done prospectively and more detailed  
116 information about the diagnostic environment needs to be collected. The results of this large study  
117 coupled with a re-examination of the previous studies do not support the conclusion that missing  
118 nor failed cytogenetics are reliable or, indeed, appropriate prognostic markers.

119

120

#### 121 **Conflict-of-interest disclosure**

122 The authors declare no competing financial interests.

123

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130

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132 A.V.M analysed and interpreted data. D.G. and R.K.H. provided clinical and follow-up data; A.K.B.  
133 was the Chief Investigator on all trials. C.J.H. and A.V.M. provided financial and administrative  
134 support. A.V.M wrote the manuscript with input and approval from all other authors.

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147

148 REFERENCES

149

- 150 1. Grimwade D, Hills RK, Moorman AV, Walker H, Chatters S, Goldstone AH, *et al.* Refinement  
151 of cytogenetic classification in acute myeloid leukemia: determination of prognostic  
152 significance of rare recurring chromosomal abnormalities among 5876 younger adult  
153 patients treated in the United Kingdom Medical Research Council trials. *Blood* 2010 Jul 22;  
154 **116**(3): 354-365.
- 155
- 156 2. Papaemmanuil E, Gerstung M, Bullinger L, Gaidzik VI, Paschka P, Roberts ND, *et al.* Genomic  
157 Classification and Prognosis in Acute Myeloid Leukemia. *N Engl J Med* 2016 Jun 9; **374**(23):  
158 2209-2221.
- 159
- 160 3. Medeiros BC, Othus M, Estey EH, Fang M, Appelbaum FR. Unsuccessful diagnostic  
161 cytogenetic analysis is a poor prognostic feature in acute myeloid leukaemia. *British journal*  
162 *of haematology* 2014 Jan; **164**(2): 245-250.
- 163
- 164 4. Lazarevic V, Horstedt AS, Johansson B, Antunovic P, Billstrom R, Derolf A, *et al.* Failure  
165 matters: unsuccessful cytogenetics and unperformed cytogenetics are associated with a  
166 poor prognosis in a population-based series of acute myeloid leukaemia. *European journal of*  
167 *haematology* 2015 May; **94**(5): 419-423.
- 168
- 169 5. Burnett AK, Hills RK, Hunter AE, Milligan D, Kell WJ, Wheatley K, *et al.* The addition of  
170 gemtuzumab ozogamicin to low-dose Ara-C improves remission rate but does not  
171 significantly prolong survival in older patients with acute myeloid leukaemia: results from  
172 the LRF AML14 and NCRI AML16 pick-a-winner comparison. *Leukemia* 2013 Jan; **27**(1): 75-  
173 81.
- 174
- 175 6. Burnett AK, Milligan D, Goldstone A, Prentice A, McMullin MF, Dennis M, *et al.* The impact of  
176 dose escalation and resistance modulation in older patients with acute myeloid leukaemia  
177 and high risk myelodysplastic syndrome: the results of the LRF AML14 trial. *British journal of*  
178 *haematology* 2009 May; **145**(3): 318-332.
- 179
- 180 7. Burnett AK, Russell NH, Hills RK, Hunter AE, Kjeldsen L, Yin J, *et al.* Optimization of  
181 chemotherapy for younger patients with acute myeloid leukemia: results of the medical  
182 research council AML15 trial. *Journal of clinical oncology : official journal of the American*  
183 *Society of Clinical Oncology* 2013 Sep 20; **31**(27): 3360-3368.
- 184
- 185 8. Gibson BE, Webb DK, Howman AJ, De Graaf SS, Harrison CJ, Wheatley K, *et al.* Results of a  
186 randomized trial in children with Acute Myeloid Leukaemia: medical research council AML12  
187 trial. *British journal of haematology* 2011 Nov; **155**(3): 366-376.
- 188
- 189 9. Grimwade D, Walker H, Oliver F, Wheatley K, Harrison C, Harrison G, *et al.* The importance of  
190 diagnostic cytogenetics on outcome in AML: analysis of 1,612 patients entered into the MRC  
191 AML 10 trial. The Medical Research Council Adult and Children's Leukaemia Working Parties.  
192 *Blood* 1998 Oct 1; **92**(7): 2322-2333.



193  
194 10. Shaffer LG, McGowan-Jordan J, Schmid M. An International System for Human Cytogenetic  
195 Nomenclature (ISCN). Basel: S. Karger; 2013.

196  
197 11. Swansbury GJ. The proportion of clonal divisions varies in different hematologic  
198 malignancies. *Cancer Genetics and Cytogenetics* 1998; **104**: 139-145.

199  
200 12. Moorman AV, Roman E, Kane EV, Dovey GJ, Cartwright RA, Morgan GJ. Karyotype and age in  
201 acute myeloid leukaemia: Are they linked? *Cancer Genetics and Cytogenetics* 2001; **126**(2):  
202 155-161.

203  
204 13. Cox MC, Panetta P, Venditti A, del Poeta G, Maurillo L, Tamburini A, *et al.* Fluorescence in  
205 situ hybridization and conventional cytogenetics for the diagnosis of 11q23+/mll+  
206 translocation in leukaemia. *British Journal of Haematology* 2003; **121**(6): 953-955.

207  
208 14. Hawkins JM, Secker-Walker LM. Evaluation of cytogenetic samples and pertinent technical  
209 variables in adult acute lymphocytic leukemia. *Cancer Genetics and Cytogenetics* 1991; **52**:  
210 79-84.

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212 Table and Figure Legends

213 Table 1: Demographics and clinical features for 10,685 patients treated on consecutive UK MRC  
214 acute myeloid leukaemia trials.

215 Figure 1: Overall survival of MRC AML intensively treated patients according the presence or absence  
216 of cytogenetic analysis (A) and for older adults (B) and by the success of cytogenetic analysis (C).  
217 Survival rates are at 5 years for intensively treated patients.

218

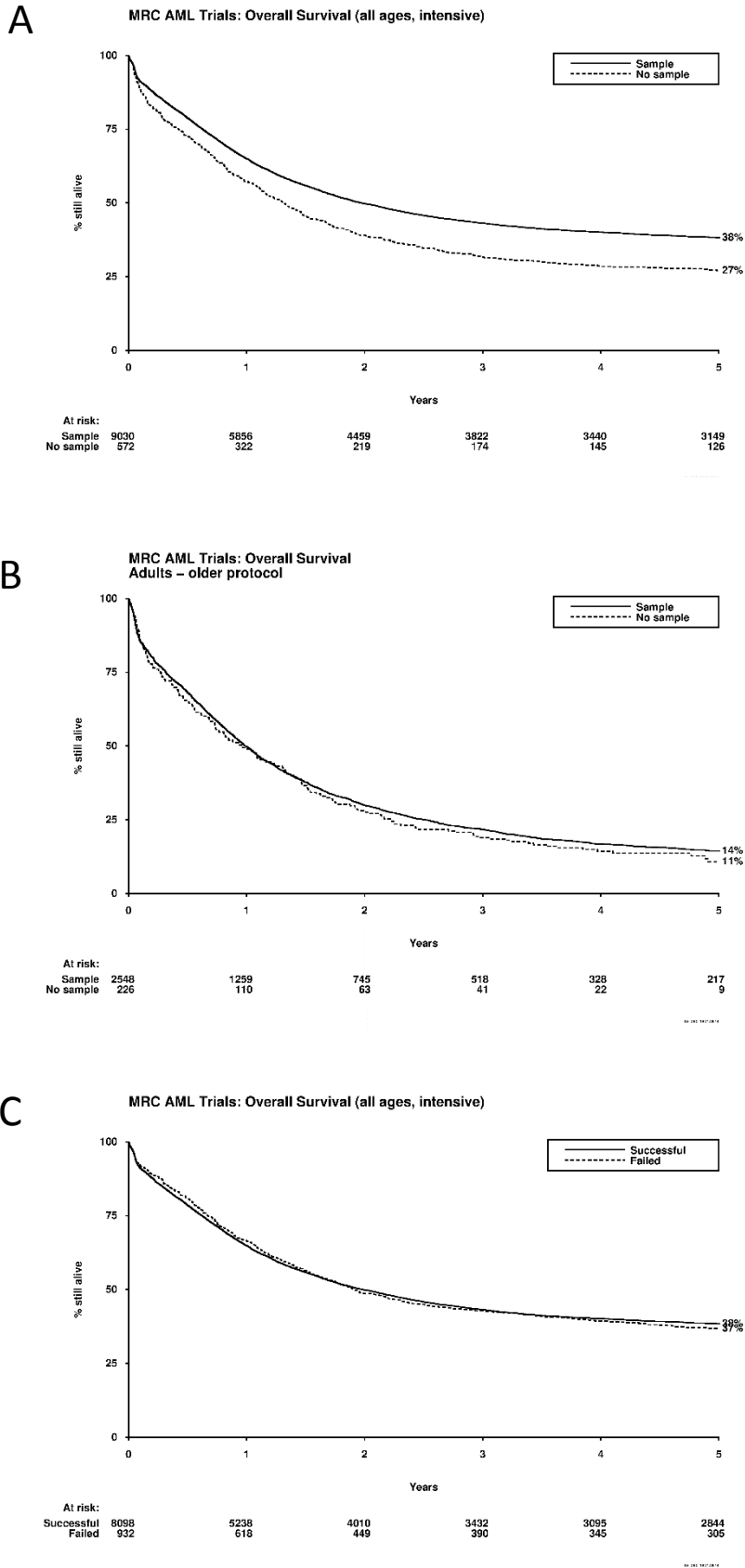


Figure 1: Overall survival of MRC AML intensively treated patients according the presence or absence of cytogenetic analysis (A) and for older adults (B) and by the success of cytogenetic analysis (C). Survival rates are at 5 years for intensively treated patients.

**Table 1: Demographics and clinical features for 10,685 patients treated on consecutive UK MRC acute myeloid leukaemia †**

		Intensively treated patients						Non-intensively treated patients					
		No sample	Sample	<i>p</i>	Successful	Failed	<i>p</i>	No sample	Sample	<i>p</i>	Successful	Failed	<i>p</i>
Total, n (%)		583 (6)	9085 (94)		8151 (90)	934 (10)		176 (17)	841 (83)		754 (90)	87 (10)	
Sex, n (%)	Female	260 (6)	4175 (94)	0.5	3754 (90)	421 (10)	0.6	61 (15)	347 (85)	0.1	304 (88)	43 (12)	0.1
	Male	323 (6)	4910 (94)		4397 (90)	513 (10)		115 (19)	494 (81)		450 (91)	44 (9)	
Age (years), n (%)*	<15	21 (3)	722 (97)	<.0001	677 (94)	45 (6)	<.0001	-	-	0.3	-	-	0.8
	15-29	46 (5)	899 (95)		832 (93)	67 (7)		-	-		-	-	
	30-39	39 (4)	1016 (96)		919 (90)	97 (10)		1^	0		1^	0	
	40-49	64 (4)	1416 (96)		1266 (89)	150 (11)		-	-		-	-	
	50-59	133 (6)	2061 (94)		1857 (90)	204 (10)		4 (40)	6 (60)		5 (83)	1 (17)	
	60-69	190 (8)	2190 (92)		1929 (88)	261 (12)		37 (22)	129 (78)		115 (89)	14 (11)	
	70-79	88 (10)	768 (90)		661 (86)	107 (14)		97 (15)	546 (85)		490 (90)	56 (10)	
	80+	2 (13)	13 (87)		10 (77)	3 (23)		38 (19)	159 (81)		143 (90)	16 (10)	
WBC x10 <sup>9</sup> /L *,n (%)	0-9.9	329 (7)	4600 (93)	0.0003	4146 (90)	453 (10)	0.0001	110 (19)	465 (81)	0.12	412 (89)	53 (11)	0.6
	10-49.9	166 (6)	2505 (94)		2276 (91)	229 (9)		45 (15)	252 (85)		231 (92)	21 (8)	
	50-99.9	38 (4)	971 (96)		858 (88)	113 (12)		15 (15)	86 (85)		78 (91)	8 (9)	
	100+	43 (5)	911 (95)		779 (86)	132 (14)		6 (14)	37 (86)		33 (86)	5 (14)	
Performance Status*, n (%)	0	366 (6)	5593 (94)	0.9	5014 (90)	579 (10)	0.2	65 (19)	275 (81)	0.3	246 (89)	30 (11)	>0.95
	1	168 (6)	2603 (94)		2356 (91)	247 (9)		87 (17)	424 (83)		383 (90)	41 (10)	
	2+	46 (6)	770 (94)		668 (87)	102 (13)		24 (15)	141 (85)		125 (89)	16 (11)	
Diagnosis, n (%)	De Novo	455 (5)	8170 (95)	<.0001	7340 (90)	830 (10)	0.3	111 (15)	627 (85)	0.002	556 (89)	71 (11)	0.11
	Secondary	128 (12)	915 (88)		811 (89)	104 (11)		65 (23)	214 (77)		198 (93)	16 (7)	
Type of secondary, n (%)	AHD	95 (14)	607 (86)	0.04; 0.01**	542 (89)	65 (11)	0.03; 0.16**	13 (20)	53 (80)	0.5; 0.4**	141 (94)	9 (5)	0.3;0.12**
	t-AML	5 (5)	98 (95)		92 (94)	6 (6)		50 (25)	150 (75)		9 (82)	2 (18)	
	Not stated	28 (12)	210 (88)		177 (84)	33 (16)		2 (15)	11 (85)		48 (91)	5 (9)	
Trial, n (%)	AML12	134 (4)	3270 (96)	<.0001	2982 (91)	288 (9)	<.0001	-	-	0.03	-	-	0.5
	AML14	78 (7)	1044 (93)		887 (85)	157 (15)		36 (13)	239 (87)		217 (91)	22 (9)	
	AML15	219 (6)	3259 (94)		2941 (90)	318 (10)		-	-		-	-	
	AML16	152 (9)	1512 (91)		1341 (89)	171 (11)		140 (19)	602 (81)		537 (89)	65 (11)	
Period of diagnosis*, n (%)	1995-99	89 (4)	2193 (96)	<.0001	1996 (91)	197 (9)	0.15	1 (3)	37 (97)	0.005	29 (78)	8 (22)	0.7
	2000-04	189 (6)	3100 (94)		2760 (89)	340 (11)		25 (15)	146 (85)		137 (94)	9 (6)	
	2005-09	238 (7)	3100 (93)		2776 (90)	324 (10)		99 (17)	468 (83)		417 (89)	51 (11)	
	2010-12	67 (9)	692 (91)		619 (89)	73 (11)		51 (21)	190 (79)		171 (90)	19 (10)	

\* test for trend; \*\* excluding not stated. Some children did not have WHO PS (not valid); ^ This 36 year old patient was deemed unsuitable for non-intensive treatment and was treated on AML16