

**Table S1.** Analysis of covariates associated with the disease free survival. P of the univariate analysis is the p value of the Log rank test. HR is the value of the hazard ratio. 95% CI is the 95% confident interval of the hazard ratio. Data of 130 patients were complete and were included in the Cox proportional-hazards regression.

DFS	Univariate analysis			Cox regression		
	p	HR	95% CI	p	HR	95% CI
<b>Age &gt; 50y</b>	0.86	1.04	0.66-1.64	>0.1		
<b>NPM+ FLT3-ITD-</b>	0.005	0.49	0.30-0.81	0.001	0.36	0.34-0.84
<b>normal karyotype</b>	0.42	0.82	0.51-1.33	>0.1		
<b>allograft</b>	0.17	0.73	0.46-1.15	0.006	0.53	0.19-0.65
<b>IDA treatment</b>	0.99	1.00	0.60-1.67	>0.1		
<b>WBC count &gt; 30G/L</b>	0.68	1.10	0.70-1.75	>0.1		
<b>DNMT3a mutation</b>	0.22	0.72	0.43-1.22	>0.1		

**Table S2.** Analysis of covariates associated with the overall survival. P of the univariate analysis is the p value of the Log rank test. HR is the value of the hazard ratio. 95% CI is the 95% confident interval of the hazard ratio. Data of 156 patients were complete and were included in the Cox proportional-hazards regression.

OS	Univariate analysis			Cox regression		
	p	HR	95% CI	p	HR	95% CI
<b>Age &gt; 50y</b>	0.81	1.05	0.70-1.59	>0.1		
<b>NPM+ FLT3-ITD- normal karyotype</b>	0.003	0.49	0.31-0.79	0.003	0.37	0.19-0.71
<b>allograft</b>	0.04	0.64	0.42-0.98	0.049	0.61	0.37-0.99
<b>IDA treatment</b>	0.46	1.19	0.75-1.87	>0.1		
<b>WBC count &gt; 30G/L</b>	0.19	1.32	0.87-1.99	>0.1		
<b>DNMT3a mutation</b>	0.19	0.73	0.46-1.17	>0.1		