## SUPPLEMENTARY METHODS

## Model building strategy in the modelisation of effects of time since diagnosis and age on the excess mortality rate and on statistical cure.

We explain here the strategy used to model excess mortality and the effect of covariates (time since diagnosis, age) on : i) excess mortality using the flexible model proposed by Nelson *et al.*<sup>13</sup> and ii) statistical cure using the flexible cure model proposed by Andersson *et al.*<sup>14</sup>. These two models are fitted to the logarithm of the cumulative excess mortality using restricted cubic spline function and based on continuous modelling of the baseline hazard. Compared to the model proposed by Nelson *et al.*, the flexible cure model proposed by Andersson *et al.*: i) added a condition to force the cumulative excess mortality rate to be constant after the last knot of time since diagnosis; and ii) add a supplemental knot to model the Non-Linear effect of the baseline hazard at the 99<sup>th</sup> percentiles of the observed death times. Estimations from the model with the cure assumption and without cure assumption were graphically compared to assess adequacy of net survival curves with the non-parametric estimator proposed by Pohar-Perme and to assess adequacy of excess mortality rate with those obtained using a model based on a step function to model baseline hazard (**Supplementary Figure 1-11**).

One model was performed independently for each AML-subtype in both sex. Time since diagnosis and age are considered as continuous function. The model assuming a time-dependent and non-linear effect of age was privileged when the number of cases was sufficient. When it was not the case, the model was based on a non-linear and a proportional-hazard effect of age.

In situation where conditions fixed was accepted (number of cases at diagnosis higher than 500 or numbers of deaths at 10-years after AML diagnosis higher than 130), functions used in the modelisation were defined here:

	Function	Kind of function —	Location of the knot			
Effect			Flexible excess mortality model	Flexible cure model		
Baseline hazard	f	Restrictive Cubic spline	According to AML subtype and Sex	According to AML subtype and Sex		
Linearity of age	g	Quadratic spline	At mean age	At mean age		
Time-Dependent effect of age	h	Restrictive Cubic spline	At the 1st, 25 <sup>th</sup> , 50 <sup>th</sup> , 75 <sup>th</sup> . 95 <sup>th</sup> percentiles and 12 years after the diagnosis	Supplemental knot at 99th percentile		

If the number of cases at diagnosis less than 500 and numbers of death at 10-years after AML diagnosis was less than 130, we performed new strategy reducing the number of model parameters as regards to the number of events at 10-years. So, we eliminate a parameter until to respecting the criteria of "one parameter per 10 deaths" in order as follow: 1) drop the 95<sup>th</sup> percentiles to model baseline hazard (i.e. 11

parameters in the model); 2) drop the 75<sup>th</sup> percentiles to model baseline hazard (i.e 9 parameters in the model); 3) to model age with a non-linear and proportional effect (i.e. 6 parameters in the model)

AML-subtypes with less than 70 deaths until 10 years were not retained in survival analysis.

The model retained for each AML-subtype and sex (and flexible functions used) are presented in the following table with on the left column, final model retained using the flexible excess mortality model and on the right column, model retained using the flexible cure model.

	Flexible excess mortality model (Nelson's model)		Flexible cure model (Andersson's model)		
MEN	Final model retained	Baseline hazard	Final model retained	Baseline hazard Supplémental knots	
Acute promyelocytic leukemia (APL)	f(Time) + g(Age) + h(Time) x Age	at 1st, 25th, 50th, 75th, percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	
AML with recurrent cytogenetic abnormalities (AML-RCA)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 25th, 50th percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	
AML with myelodysplasia-related changes (AML-MRC)	f(Time) + g(Age)	at 1st, 25 <sup>th</sup> , 50 <sup>th</sup> , 75 <sup>th</sup> , 95 <sup>th</sup> percentiles and 12 years	f(Time) + g(Age)	at 99 <sup>th</sup> percentile	
Therapy-related AML; NOS (t-AML)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 50th percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 25 <sup>th</sup> , 75 <sup>th</sup> , 95 <sup>th</sup> , 99 <sup>th</sup> percentile	
Pure erythroid leukemia (AML-M6)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 50th, 95th percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 25 <sup>th</sup> , 75 <sup>th</sup> , 99 <sup>th</sup> percentile	
Acute Myelomonocytic leukemia (AML-M4)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 25th, 50th, 75th, 95th percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	
Acute myeloid leukemia with minimal differentiation (AML-M0)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 25th, 50th, 75th, 95th percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	
Acute Myeloid leukemia without maturation (AML-M1)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 25th, 50th, 75th, 95th percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	
Acute Myeloid leukemia with maturation (AML-M2)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 25 <sup>th</sup> , 50 <sup>th</sup> , 75 <sup>th</sup> , 95 <sup>th</sup> percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	
Acute monoblastic and monocytic leuaemia (AML-M5)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 25th, 50th, 75th. 95th percentiles and 12 year	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	
Acute Myeloid leukemia, not otherwise specified (AML-NOS)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 25 <sup>th</sup> , 50 <sup>th</sup> , 75 <sup>th</sup> , 95 <sup>th</sup> percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	
WOMEN					
Acute promyelocytic leukemia (APL)	f(Time) + g(Age)	at 1st, 25th, 50th, percentiles and 12 years	f(Time) + g(Age)	at 99 <sup>th</sup> percentile	
AML with recurrent cytogenetic abnormalities (AML-RCA)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 25 <sup>th</sup> , 50 <sup>th</sup> percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	
AML with myelodysplasia-related changes (AML-MRC)	f(Time) + g(Age)	at 1st, 25 <sup>th</sup> , 50 <sup>th</sup> , 75 <sup>th</sup> , 95 <sup>th</sup> percentiles and 12 years	f(Time) + g(Age)	at 99 <sup>th</sup> percentile	
Therapy-related AML; NOS (t-AML)	f(Time) + g(Age)	at 1st, 25 <sup>th</sup> , 50 <sup>th</sup> , 75 <sup>th</sup> , 95 <sup>th</sup> percentiles and 12 years	f(Time) + g(Age)	at 99 <sup>th</sup> percentile	
Pure erythroid leukemia (AML-M6)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 50 <sup>th</sup> , 95 <sup>th</sup> percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	
Acute Myelomonocytic leukemia (AML-M4)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 25 <sup>th</sup> , 50 <sup>th</sup> , 75 <sup>th</sup> , 95 <sup>th</sup> percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	
Acute myeloid leukemia with minimal differentiation (AML_M0)	f(Time) + g(Age)	at 1st, 25 <sup>th</sup> , 50 <sup>th</sup> , 75 <sup>th</sup> , 95 <sup>th</sup> percentiles and 12 years	f(Time) + g(Age)	at 99 <sup>th</sup> percentile	
Acute Myeloid leukemia without maturation (AML-M1)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 25 <sup>th</sup> , 50 <sup>th</sup> , 75 <sup>th</sup> , 95 <sup>th</sup> percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	
Acute Myeloid leukemia with maturation (AML_M2)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 50 <sup>th</sup> percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 25th, 75th, 95th, 99th percentile	
Acute monoblastic and monocytic leukemia (AML-M5)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 50 <sup>th</sup> percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	
Acute Myeloid leukemia, not otherwise specified (AML-NOS)	$f(Time) + g(Age) + h(Time) \times Age$	at 1st, 25 <sup>th</sup> , 50 <sup>th</sup> , 75 <sup>th</sup> , 95 <sup>th</sup> percentiles and 12 years	f(Time) + g(Age) + h(Time) x Age	at 99 <sup>th</sup> percentile	

In the following, LL stands for "Log-Linear", NL for "Non-Linear", TD for "Time-Dependent", PH for "Proportional-Hazard". Even if the model on which estimates are based mainly correspond to the full model adjusted on the age at diagnosis with NL and TD effect, we explored all the same the shape of effects of age using a model-building strategy on inference about non-linear (i.e. risk not constant between tow modality of a covariable) and time-dependent effect (i.e. risk not constant over time since diagnosis).

In each AML subtype and sex, we used a likelihood ratio test to study:

1) the non-linear effect of age assuming firstly a TD effect of age and secondly a PH effect of age,

2) the time-dependent effect of age assuming firstly a NL effect of age and secondly a LL effect of age.

The effect was considered statistically significant with a cut off equal to 0.05.

Results of likelihood ratio test by AML subtype and sex, are presented in the following table with on the left column, effect of age retained using the flexible excess mortality model and on the right column, effect of age retained using the flexible cure model. Functions used in the modelisation are the same that those presented previously.

	Flexible excess mortality model (Nelson's model)				Flexible cure model (Andersson's model)			
P-value	Linear effect		Time dependent effect		Linear effect		Time dependent effect	
	NL	NL	TD	TD	NL	NL	TD	TD
	assuming TD	assuming PH	assuming NL	assuming LL	assuming TD	assuming PH	assuming NL	assuming LL
MEN								
Acute promyelocytic leukemia with t(15;17)(q22;q12) (APL)	0.017	0.024	0.014	0.019	0.013	0.021	0.002	0.004
AML with recurrent cytogenetic abnormalities (AML-RCA)	0.106	0.015	0.0676	0.011	0.018	0.011	0.208	0.122
AML with myelodysplasia-related changes (AML-MRC)	not tested	0.243	not tested	not tested	not tested	0.232	not tested	not tested
Therapy-related AML; NOS (t-AML)	0.500	0.355	0.042	0.031	0.386	0.305	0.155	0.123
Pure erythroid leukemia (AML-M6)	0.256	0.415	0.148	0.222	0.308	0.327	0.426	0.451
Acute Myelomonocytic leukemia (AML-M4)	0.009	0.001	0.339	0.070	0.008	0.002	0.185	0.037
Acute myeloid leukemia with minimal differentiation (AML-M0)	0.794	0.275	0.003	0.001	0.728	0.259	0.001	< 0.001
Acute Myeloid leukemia without maturation (AML-M1)	0.001	0.001	0.088	0.079	0.002	0.001	0.094	0.031
Acute Myeloid leukemia with maturation (AML-M2)	0.019	0.025	0.001	0.001	0.031	0.017	0.001	0.001
Acute monoblastic and monocytic leukemia (AML-M5)	0.048	0.003	0.028	0.002	0.013	0.003	0.127	0.027
Acute Myeloid leukemia, not otherwise specified (AML-NOS)	0.002	< 0,001	0.092	0.025	0.002	< 0.001	0.068	0.008
WOMEN								
Acute promyelocytic leukemia with t(15;17)(q22;q12) (APL)	not tested	0.104	not tested	not tested	not tested	0.066	not tested	not tested
AML with recurrent cytogenetic abnormalities (AML-RCA)	0.044	0.025	0.998	0.556	0.024	0.024	0.696	0.734
AML with myelodysplasia-related changes (AML-MRC)	not tested	0.944	not tested	not tested	not tested	0.946	not tested	not tested
Therapy-related AML; NOS (t-AML)	not tested	0.147	not tested	not tested	not tested	0.149	not tested	not tested
Pure erythroid leukemia (AML-M6)	0.926	0.992	0.706	0.739	0.985	0.995	0.777	0.785
Acute Myelomonocytic leukemia (AML-M4)	0.008	0.002	0.099	0.027	0.011	0.002	0.044	0.009
Acute myeloid leukemia with minimal differentiation (AML-M0)	not tested	0.388	not tested	not tested	not tested	0.378	not tested	not tested
Acute Myeloid leukemia without maturation (AML-M1)	< 0,001	0.003	0.034	< 0,001	< 0,001	0.002	0.028	0.001
Acute Myeloid leukemia with maturation (AML-M2)	0.008	0.016	0.055	0.103	0.028	0.007	0.151	0.039
Acute monoblastic and monocytic leukemia (AML-M5)	0.001	< 0,001	0.828	0.524	< 0,001	< 0,001	0.309	0.654
Acute Myeloid leukemia, not otherwise specified (AML-NOS)	0.317	0.092	0.096	0.032	0.272	0.086	0.037	0.012

## Supplementary Figure 1-11:

## Adequacy of flexible excess model and flexible cure model in each Acute Myeloid Leukemia subtype by sex: A) in Men and B) in Women using:

- 1) Estimation of probability of net survival compared with those obtained using the non-parametric estimator of Pohar-Perme to estimate:
  - Probability of net survival (%) over time since diagnosis for all age (Top left graph),
  - One-year net survival and CI95% over age at diagnosis in years (Top middle graph),
  - Five-year net survival and CI95% over age at diagnosis in years (Top right graph).

Note that for Pohar-Perme estimator, net survival was estimated separately per age-group in 10 year intervals;

- 2) Estimation of excess mortality rate compared with those obtained using a model with a step function to model baseline hazard for estimation of excess mortality rate over time from diagnosis in three age-groups:
  - for patient of [15-44] years old (Bottom left graph),
  - for patient of [45-64] years old (Bottom middle graph),
  - for patient equal or more than 65 years old (Bottom right graph).

The flexible excess and cure models were adjusted, respectively, on the median age of each age-group and excess mortality was model using step function for baseline hazard separately in each age-group without other covariates.

Supplementary Figure 1 - Adequacy of flexible excess model and flexible cure model in acute promyelocytic leukemia (APL): A) Men and B) Women. Net survival (%) over time since diagnosis for all age (Top left graph). Net survival (and CI95%) over age at diagnosis in years at 1-year (Top middle graph) and 5-year (Top right graph). Excess mortality rate in deaths per person-year over time since diagnosis in years in [15-44] age-group (Bottom left graph), in [45-64] age-group (Bottom middle graph) and in 65 years old and more (Bottom right graph).



Supplementary Figure 2 - Adequacy of flexible excess model and flexible cure model in acute myeloid leukemia with recurrent cytogenetic abnormalities (AML-RCA): A) Men and B) Women. Net survival (%) over time since diagnosis for all age (Top left graph). Net survival (and CI95%) over age at diagnosis in years at 1-year (Top middle graph) and 5-year (Top right graph). Excess mortality rate in deaths per person-year over time since diagnosis in years in [15-44] age-group (Bottom left graph), in [45-64] age-group (Bottom middle graph) and in 65 years old and more (Bottom right graph).



Supplementary Figure 3 - Adequacy of flexible excess model and flexible cure model in acute myeloid leukemia with myelodysplasia-related changes (AML-MRC): A) Men and B) Women. Net survival (%) over time since diagnosis for all age (Top left graph). Net survival (and CI95%) over age at diagnosis in years at 1-year (Top middle graph) and 5-year (Top right graph). Excess mortality rate in deaths per person-year over time since diagnosis in years in [15-44] age-group (Bottom left graph), in [45-64] age-group (Bottom middle graph) and in 65 years old and more (Bottom right graph).



Supplementary Figure 4 - Adequacy of flexible excess model and flexible cure model in therapy-related acute myeloid leukemia, NOS (t-AML): A) Men and B) Women. Net survival (%) over time since diagnosis for all age (Top left graph). Net survival (and CI95%) over age at diagnosis in years at 1-year (Top middle graph) and 5-year (Top right graph). Excess mortality rate in deaths per person-year over time since diagnosis in years in [15-44] age-group (Bottom left graph), in [45-64] age-group (Bottom middle graph) and in 65 years old and more (Bottom right graph).



Supplementary Figure 5 - Adequacy of flexible excess model and flexible cure model in Pure erythroid leukemia (AML-M6) in men. Net survival (%) over time since diagnosis for all age (Top left graph). Net survival (and CI95%) over age at diagnosis in years at 1-year (Top middle graph) and 5-year (Top right graph). Excess mortality rate in deaths per person-year over time since diagnosis in years in [15-44] age-group (Bottom left graph), in [45-64] age-group (Bottom middle graph) and in 65 years old and more (Bottom right graph).



Supplementary Figure 6 - Adequacy of flexible excess model and flexible cure model in Acute Myelomonocytic Leukemia (AML-M4): A) Men and B) Women. Net survival (%) over time since diagnosis for all age (Top left graph). Net survival (and CI95%) over age at diagnosis in years at 1-year (Top middle graph) and 5-year (Top right graph). Excess mortality rate in deaths per person-year over time since diagnosis in years in [15-44] age-group (Bottom left graph), in [45-64] age-group (Bottom middle graph) and in 65 years old and more (Bottom right graph).



Supplementary Figure 7 - Adequacy of flexible excess model and flexible cure model in acute myeloid leukemia with minimal differentiation (AML-MO): A) Men and B) Women. Net survival (%) over time since diagnosis for all age (Top left graph). Net survival (and CI95%) over age at diagnosis in years at 1-year (Top middle graph) and 5-year (Top right graph). Excess mortality rate in deaths per person-year over time since diagnosis in years in [15-44] age-group (Bottom left graph), in [45-64] age-group (Bottom middle graph) and in 65 years old and more (Bottom right graph).



Supplementary Figure 8 - Adequacy of flexible excess model and flexible cure model in acute myeloid leukemia without maturation (AML-M1): A) Men and B) Women. Net survival (%) over time since diagnosis for all age (Top left graph). Net survival (and CI95%) over age at diagnosis in years at 1-year (Top middle graph) and 5-year (Top right graph). Excess mortality rate in deaths per person-year over time since diagnosis in years in [15-44] age-group (Bottom left graph), in [45-64] age-group (Bottom middle graph) and in 65 years old and more (Bottom right graph).



Supplementary Figure 9 - Adequacy of flexible excess model and flexible cure model in acute myeloid leukemia with maturation (AML-M2): A) Men and B) Women. Net survival (%) over time since diagnosis for all age (Top left graph). Net survival (and CI95%) over age at diagnosis in years at 1-year (Top middle graph) and 5-year (Top right graph). Excess mortality rate in deaths per person-year over time since diagnosis in years in [15-44] age-group (Bottom left graph), in [45-64] age-group (Bottom middle graph) and in 65 years old and more (Bottom right graph).



Supplementary Figure 10 - Adequacy of flexible excess model and flexible cure model in acute monoblastic and monocytic leukemia (AML-M5): A) Men and B) Women. Net survival (%) over time since diagnosis for all age (Top left graph). Net survival (and CI95%) over age at diagnosis in years at 1-year (Top middle graph) and 5-year (Top right graph). Excess mortality rate in deaths per person-year over time since diagnosis in years in [15-44] age-group (Bottom left graph), in [45-64] age-group (Bottom middle graph) and in 65 years old and more (Bottom right graph).



Supplementary Figure 11 - Adequacy of flexible excess model and flexible cure model in Acute Myeloid Leukemia, not otherwise specified (AML-NOS): A) Men and B) Women. Net survival (%) over time since diagnosis for all age (Top left graph). Net survival (and CI95%) over age at diagnosis in years at 1-year (Top middle graph) and 5-year (Top right graph). Excess mortality rate in deaths per person-year over time since diagnosis in years in [15-44] age-group (Bottom left graph), in [45-64] age-group (Bottom middle graph) and in 65 years old and more (Bottom right graph).

