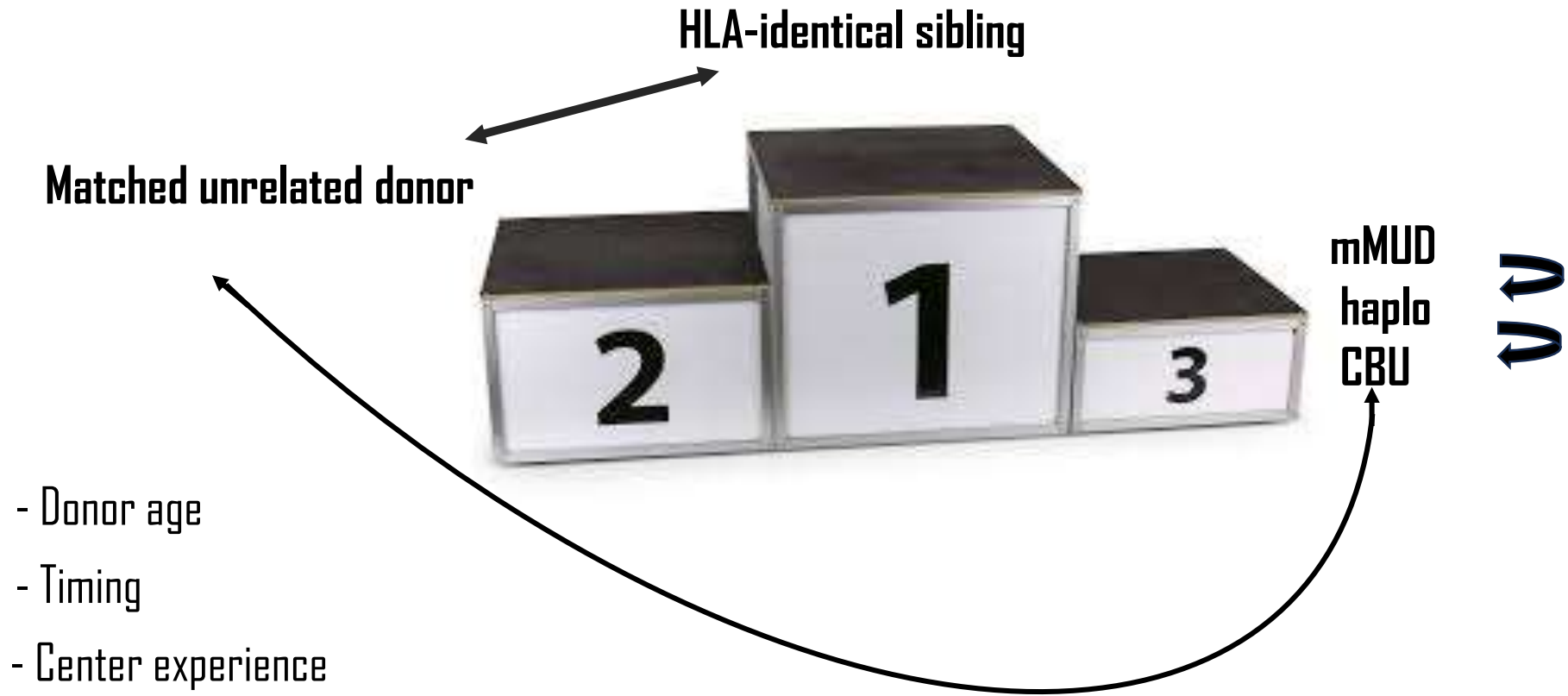


L'uso dell'intelligenza artificiale nella valutazione della compatibilità HLA ai fini della selezione del miglior donatore di cellule staminali: creazione di un modello prototipo e sviluppi futuri

Roberto Crocchiolo

XXX Congresso Nazionale AIBT - 11 ottobre 2024



Stem cell donor selection: factors

HLA

Patient-donor matching
Donor-specific anti-HLA Abs

non-HLA

Donor age
Gender
ABO
CMV
Parity
Registry



Aim of the study

- To develop a calculator of 2-y OS according to the type of stem cell donor, to support the transplant physician during the selection for a pre-determined patient

HLA-identical sibling



2-y OS = X

Matched unrelated donor



2-y OS = Y

**mMUD
haplo
CBU**



2-y OS = Z

Stefania Cacace, PoliMI

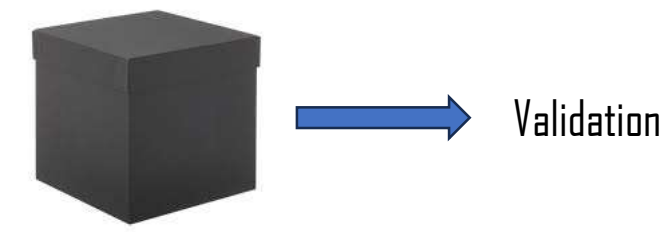
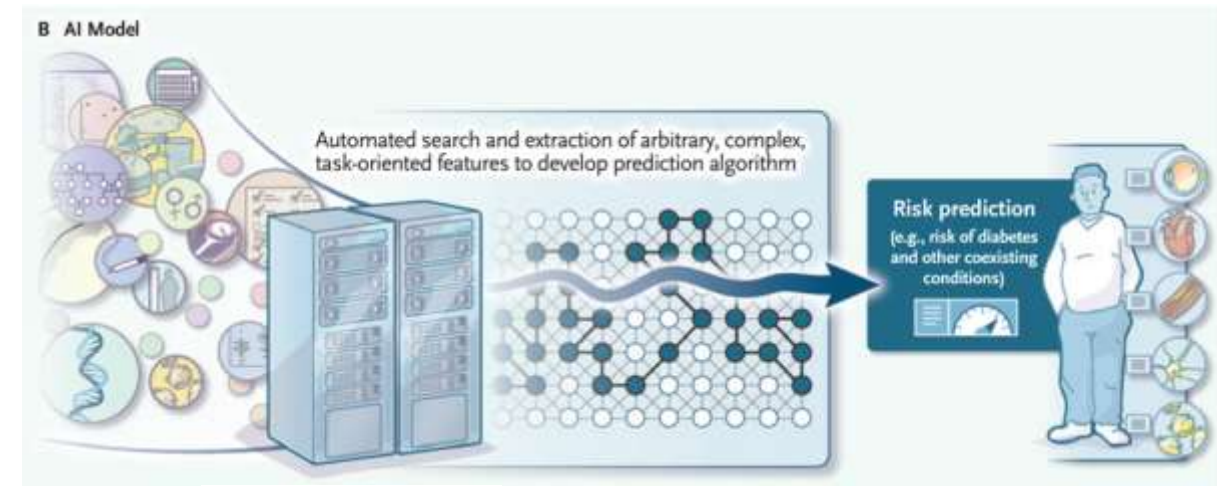


Regression analysis

AI models

$$Y_p = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots + \beta_k x_k + \sigma \Phi^{-1}(p)$$

1. Y_p : Represents either the failure time or the logarithm of the failure time, depending on the distribution of the data.
2. $\beta_0, \beta_1, \dots, \beta_k$: Regression coefficients.
3. x_1, x_2, \dots, x_k : predictor variables.
4. σ : A scale parameter (for most distributions) or shape parameter (for Weibull distribution).
5. $\Phi^{-1}(p)$: The p -th quantile of the standardized life distribution.



Real-world data obtained from clinical electronic charts
of two transplant centers (ICH, AO Vittorio Emanuele), interval Jul 2010 - Jan 2022

Patient characteristics	Number of patients
Number of patients	737
Gender	
Male	431
Female	306
Median age at transplantation	48
HLA match	
Identical sibling	218
Matched unrelated	198
Haploidentical	321
Diagnosis*	
Group 1	306
Group 2	114
Group 3	317
Median Karnofsky score	87.7
Karnofsky score >90% at HSCT	446
Positive CMV serostatus	684
Negative CMV serostatu	53
Total comorbidities	
No comorbidity	423
1 comorbidity	256
>2 comorbidities	58

Donor characteristics	Number of donors
Median age at donation	40
Gender	
Male	500
Female	331
missing	6
Positive CMV serostatus	556
Negative CMV serostatus	181

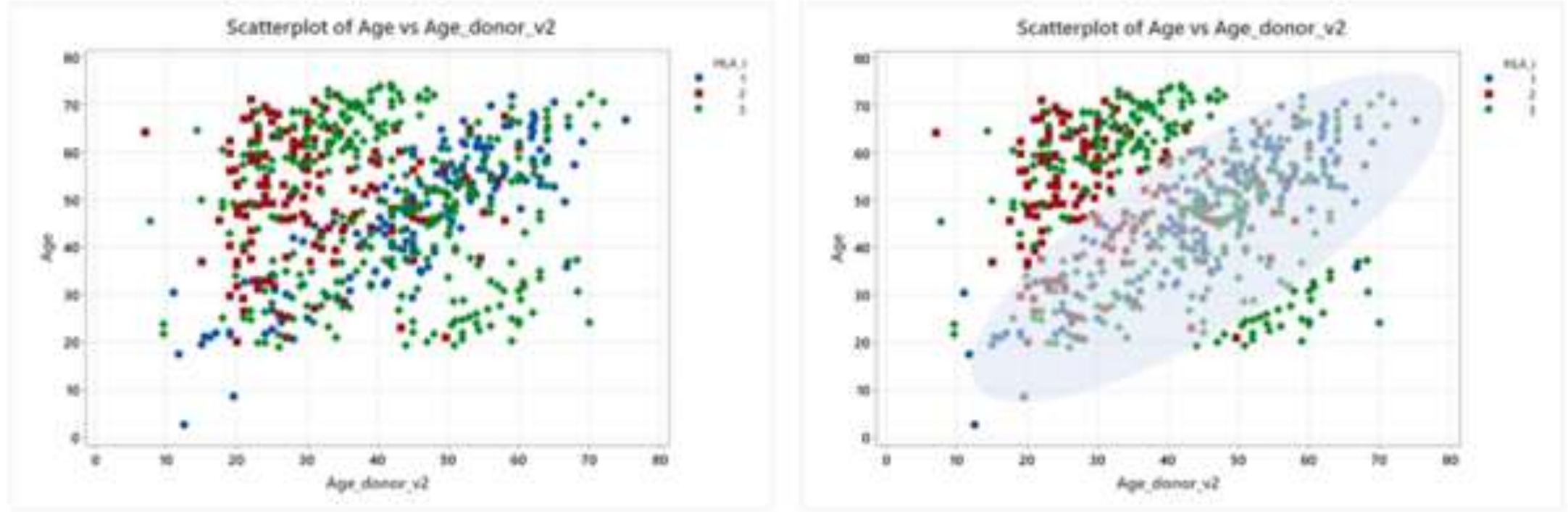




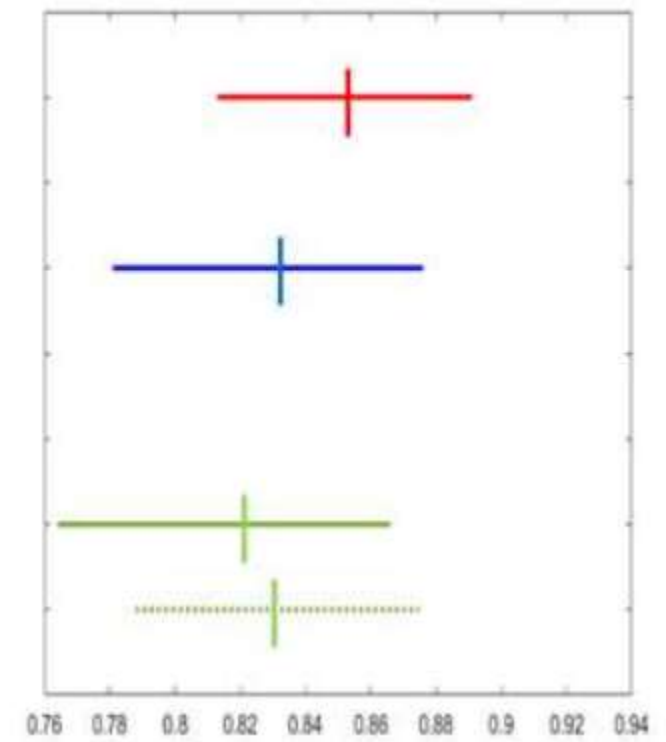


Figure 1. Scatterplots of patient and donors ages by HLA match types. The X axis represents the age of the donor, while the Y axis the age of the patient. Each point represents a donor-patient pair and is color-coded: HLA-identical sibling (blue), unrelated donor (red), haploidentical donor (green).

	Age	Diagnosis	HLA	Sorror	Age_donor	2y-OS	CI_l	CI_u
	45	AL in CR1	HLA-id sibling	0	45	0.856	0.813	0.891
	45	AL in CR1	MUD 10/10	0	30	0.834	0.781	0.876
	45	AL in CR1	haplo	0	20	0.821	0.764	0.866
	45	AL in CR1	haplo	0	45	0.836	0.788	0.875





n = 22,241 HSCTs



Random survival forest

<https://mednexus.org/doi/full/10.1097/CM9.0000000000001539>

Extreme gradient boosting [XGB]

<https://www.nature.com/articles/s41598-021-86327-7>

Deep learning

<https://ojs.aaai.org/index.php/AAAI/article/view/11842>

Bayesian hierarchical Cox Proportional model

<https://www.nature.com/articles/s41598-021-03645-6>

Bayesian Additive Regression Trees (BART)

https://www.researchgate.net/publication/363058671_Bayesian_Additive_Regression_Trees_for_Predicting_Colon_Cancer_Methodological_Study_Validity_Study

Conclusions

- Feasibility of developing a tool (calculator) that may help transplant physicians in the decision-making process → prototype
- AI models = regression
- Novelty → all donor types
- AI is expected to outperform with large dataset (ongoing)

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Alice Fumagalli ⁴, Maria Pia Azzaro ³, Sebastiana Toscano ³ and Quirico Semeraro ²**

- ¹ Servizio di Immunoematologia e Medicina Trasfusionale, ASST Grande Ospedale Metropolitano Niguarda, Milano, Italy
- ² Dipartimento di Meccanica, Politecnico di Milano, Milano, Italy
- ³ Ematologia, Azienda Ospedaliera Vittorio Emanuele, Catania, Italy
- ⁴ Ematologia, Istituto Clinico Humanitas, Rozzano, Italy

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