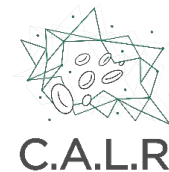


**Immunotherapy  
in Hematological  
Malignancies 2023**

May 18-19-20, 2023

# Artificial Intelligence in Hematology

*MATTEO DELLA PORTA, MD  
Humanitas Research Hospital, Milan Italy*



# Artificial Intelligence (AI) for precision medicine

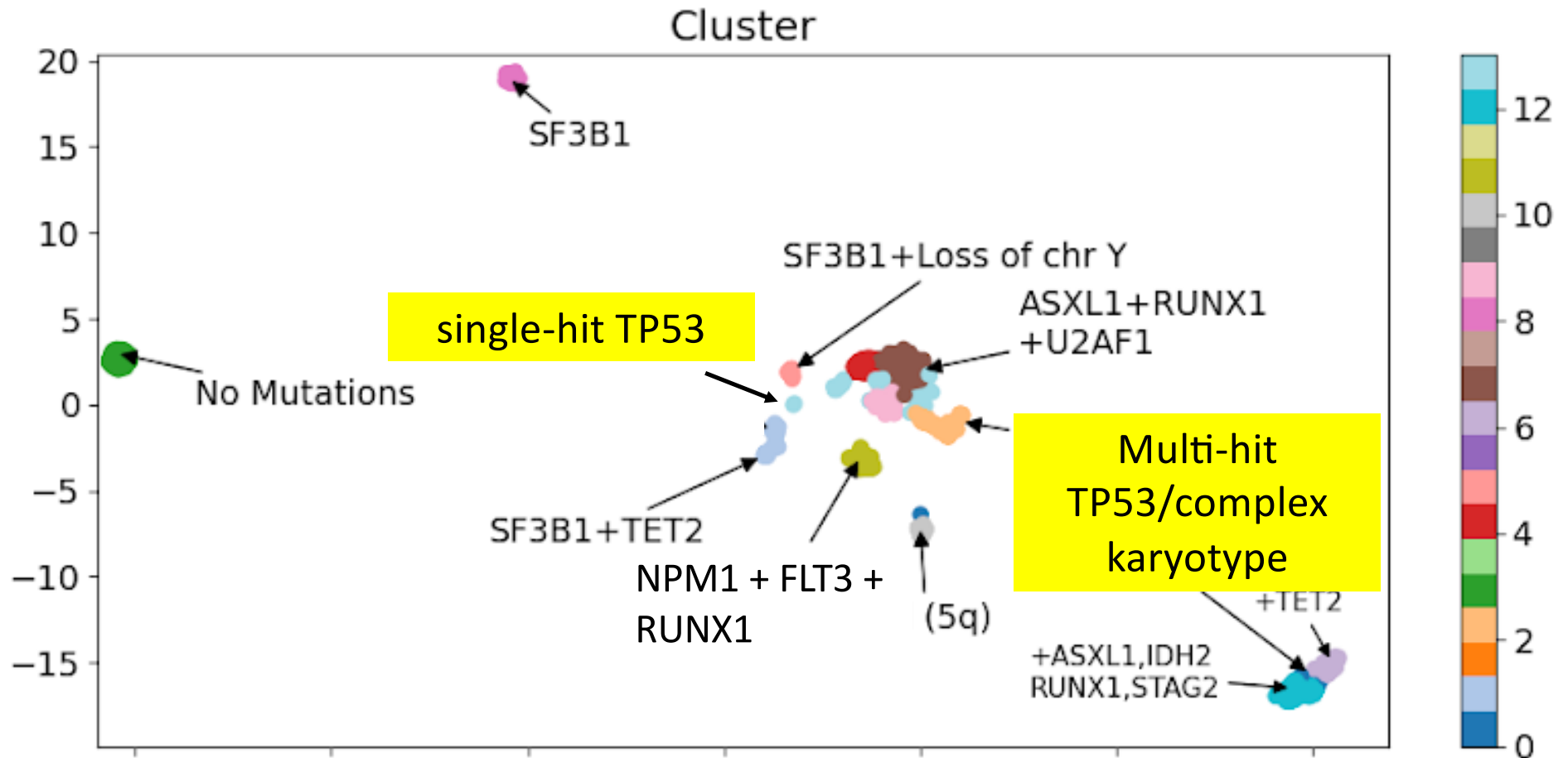
## 1- Machine Learning



## 2- Generative AI



# Molecular classification of MDS by machine learning methods (AI)



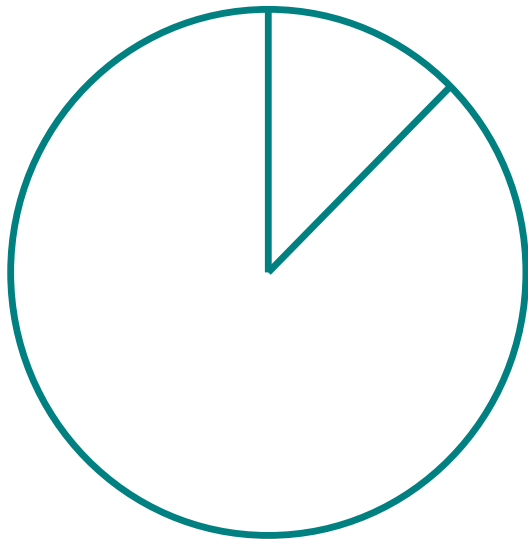
Bersanelli et al. *J Clin Oncol* 2021; D'Amico et al, *J Clin Oncol CCI* 2023, in press

## A new category of high-risk MDS is defined according to p53 dysfunction

SHAP (Shapley\_Additive\_Explanations) was used to explain the classification model by computing the contribution of each feature

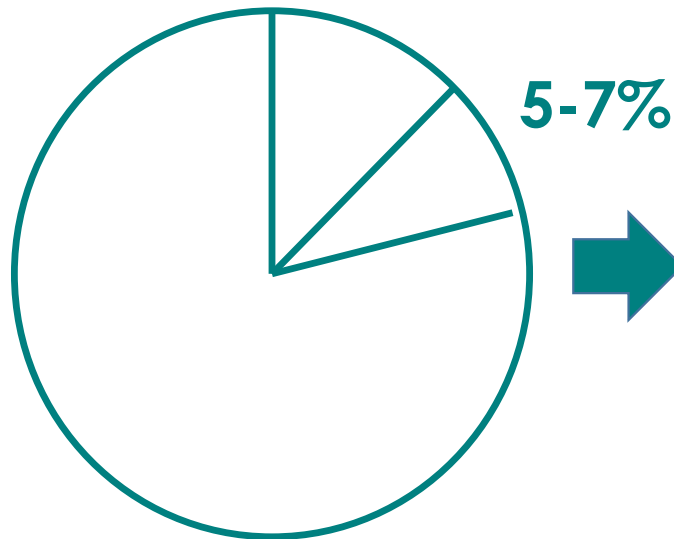
MDS with TP53-mutations

10-12%

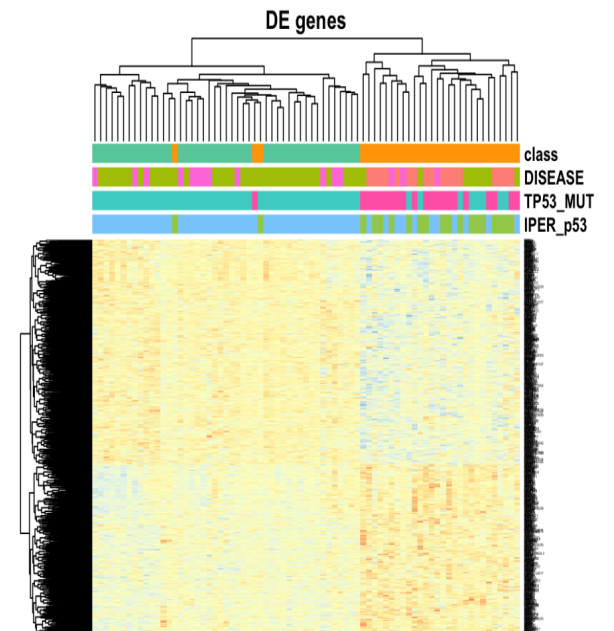


MDS with p53 dysfunction without TP53 mutations

5-7%



RNA sequencing



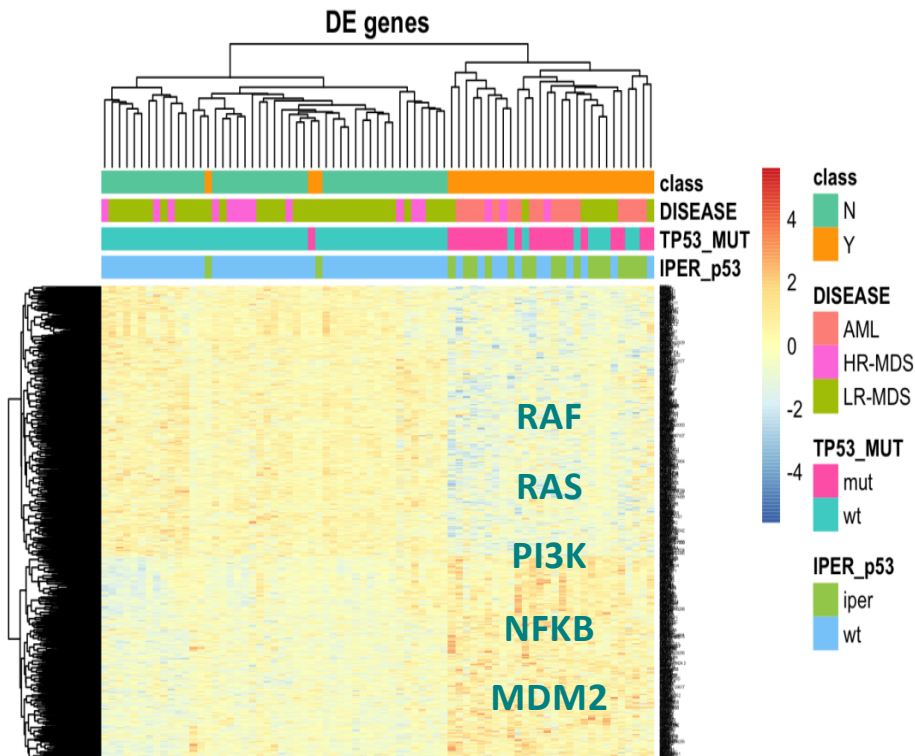
*Bersanelli et al. J Clin Oncol 2021*

*Riva et al. Blood 2022;140:4001-4; Zampini et al, manuscript in preparation*

HUMANITAS

## A new category of high-risk MDS is defined according to p53 dysfunction

RNA sequencing of CD34+ progenitor cells from  
236 MDS patients



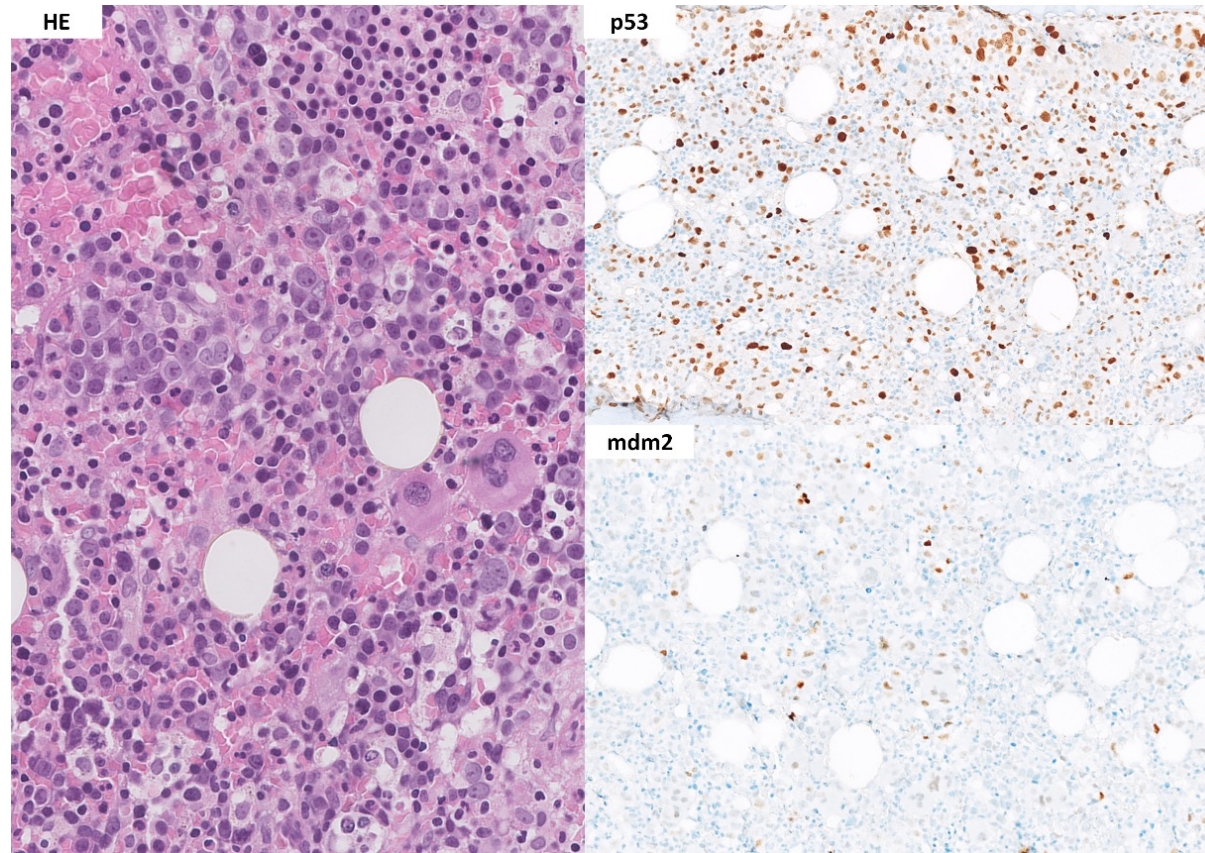
Evidence of impaired T cell and NK maturation and  
function in MDS with p53 dysfunction

- Immune checkpoint overexpression (PD-L1) at the stem cell level
- Reduced numbers of cytotoxic T cells
- Expansion of myeloid-derived suppressor cells (MDSCs)
- Expansion of regulatory T cells (Tregs).
- Impaired NK maturation and function

*Sallman DA Blood (2020) 136 (24): 2812–2823*

*Riva et al. Blood 2022;140:4001–4; Zampini et al, manuscript in preparation*

## A new category of high-risk MDS is defined according to p53 dysfunction



# 2021 WHO guidance on ethics and governance of AI for health

We have to address three important topics, deemed as essential for a **right deployment of AI in hematology**:

- **Transparency of models.** We have to provide a good understanding of the models (interpretability and explainability)
- **Reliability of models.** The main vulnerabilities of AI models are related to lack of generalizability. Therefore, extensive, independent validation of generated AI-models is required.
- **Protection of data and data sharing.** Innovative technologies such as federated learning procedures for data collection and analysis (without moving sensitive medical data from their original locations) are required to facilitate clinical implementability of AI solutions

1. *The World Health Organization. 2021 WHO guidance on ethics and governance of artificial intelligence for health. <https://www.who.int/publications/i/item/9789240029200>*

# Artificial Intelligence (AI) for precision medicine

## 1- Machine Learning



## 2- Generative AI





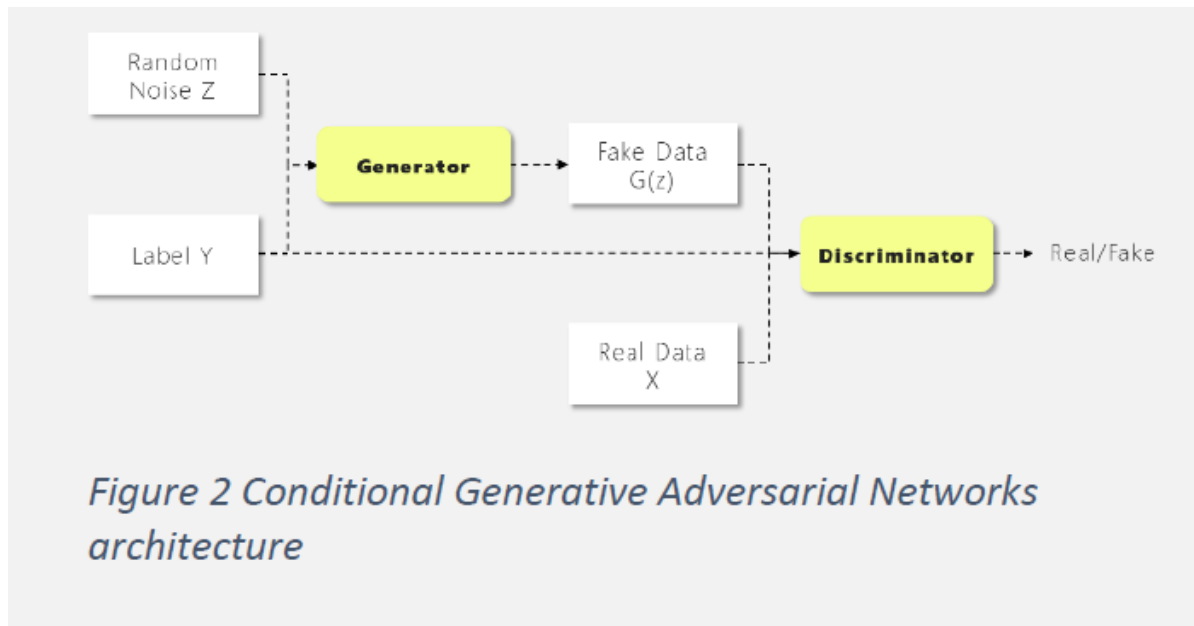
# Generation of Synthetic Data to accelerate Research & Development in MDS

- In MDS the first evidence of recurrent somatic mutations in splicing-related genes was published in 2011 and only 10 years later large patient populations ( $n > 2,000$ ) with comprehensive clinical and molecular information were available to test clinical significance and implementability of genomic screening
- The development of innovative data-driven digital health products and services, in fact, is currently slowed due to limited access to / availability of data. Additional challenges in health data include harmonization problems and data privacy (GDPR)
- **Synthetic data** are artificial data generated by an algorithm trained to learn all the essential characteristics of a real dataset. The new data are neither a copy nor a representation of the real data. Since they are not real data, they are not regulated by particular limitations so they can be easily accessed and shared.



# Generation of synthetic data to accelerate Research & Development in MDS

- Synthetic data can be generated by using neural networks (Generative Adversarial Networks, GAN).

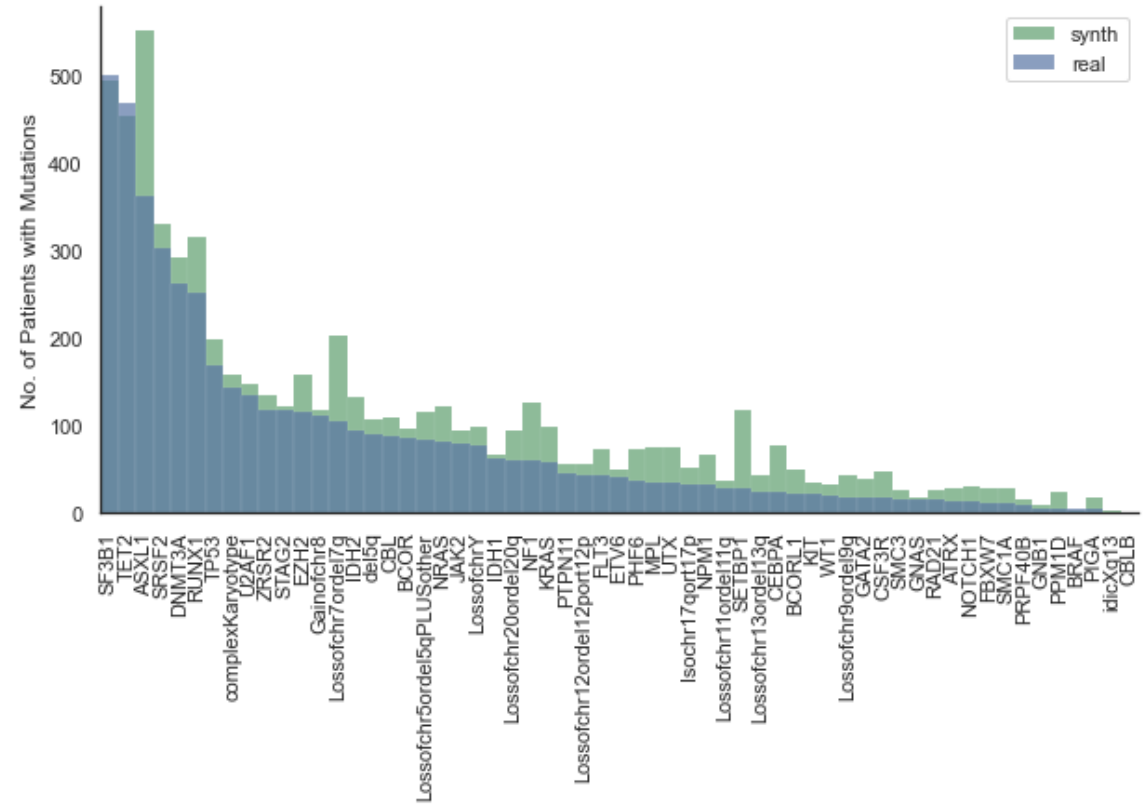
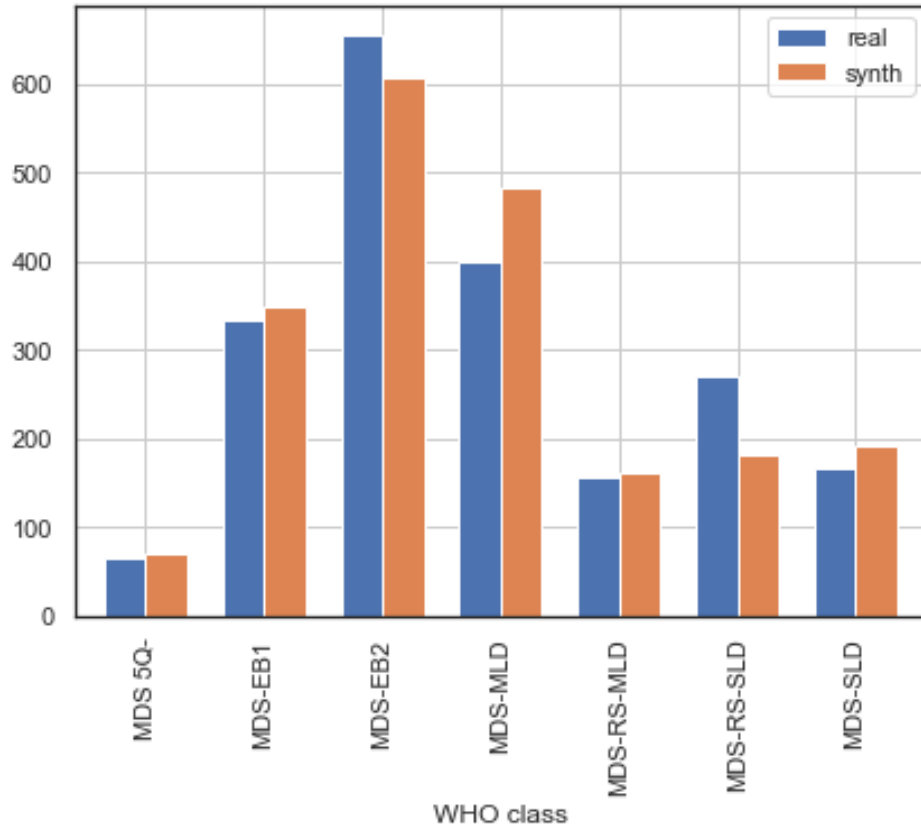


## Possible applications

- Data sharing (GDPR)
- Classes balance and resolution of missing information
- Data augmentation for learning/validation purpose
- Generation of new evidence



# Synthetic vs. Real Data: comparison of clinical and molecular features

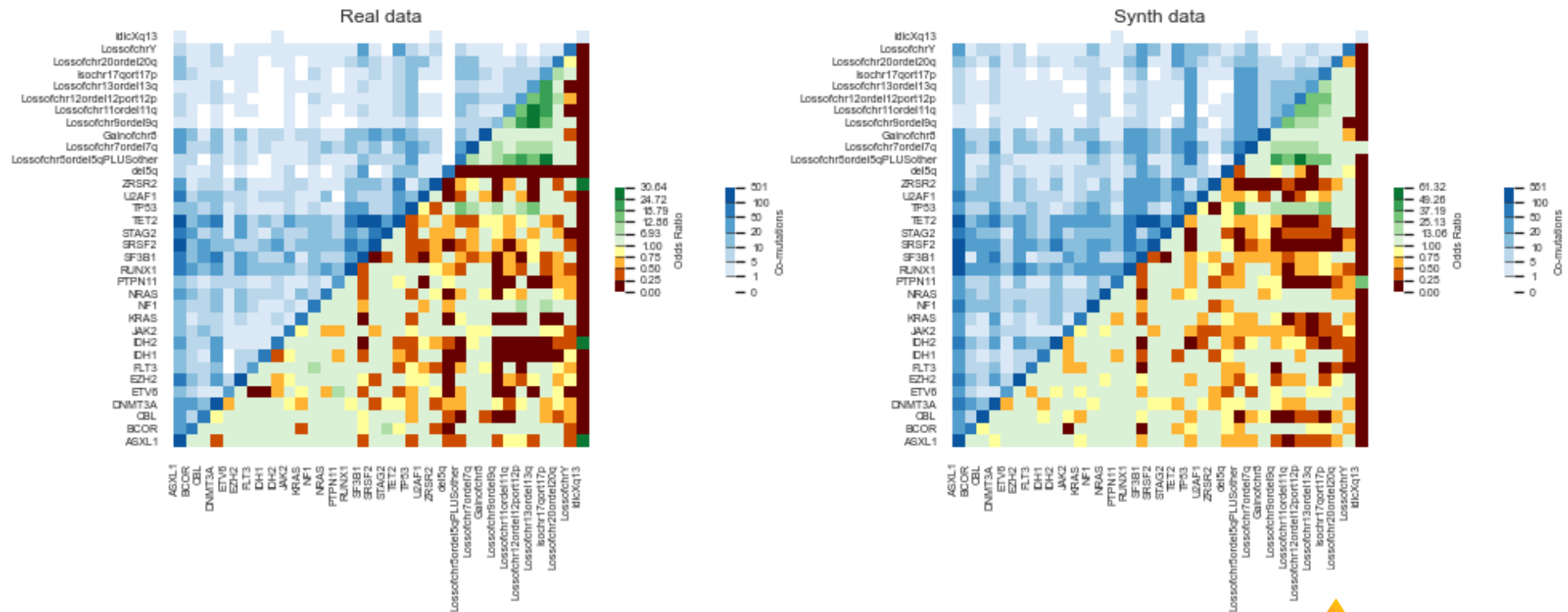


D'Amico S et al, J Clin Oncol CCI 2023, in press



# Synthetic vs. Real Data: pairwise association among genes and cytogenetic abnormalities

Pairwise associations among genes and cytogenetic abnormalities



D'Amico S et al, J Clin Oncol CCI 2023, in press



# Synthetic vs. Real Data: survival

COX models (overall survival)

Probability of OS stratified by IPSS-R

**Real data:**

Global Concordance: 0.779; Std.err:0.013

Partial Concordance of risk components:

	Clinical	CNA	Demographics	Genetics
<b>concordant</b>	0.711	0.569	0.630	0.782
<b>std(c-d)</b>	0.013	0.011	0.013	0.013

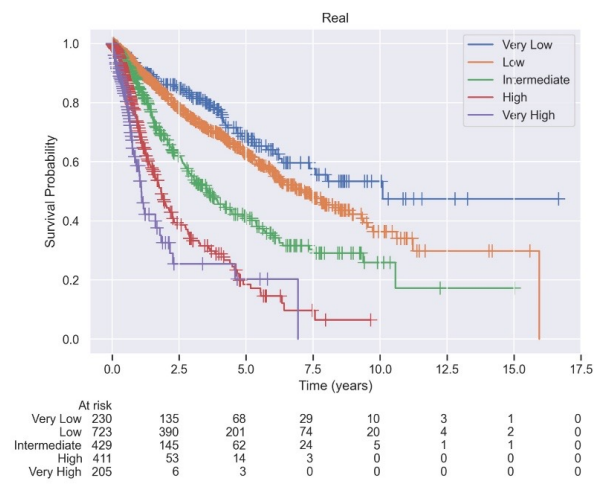
**Synthetic data:**

Global Concordance: 0.822; Std.err:0.013

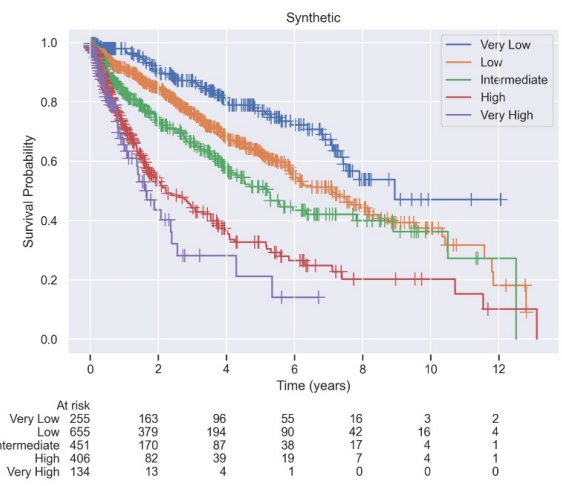
Partial Concordance of risk components:

	Clinical	CNA	Demographics	Genetics
<b>concordant</b>	0.732	0.536	0.646	0.746
<b>std(c-d)</b>	0.013	0.011	0.013	0.013

**REAL DATA**

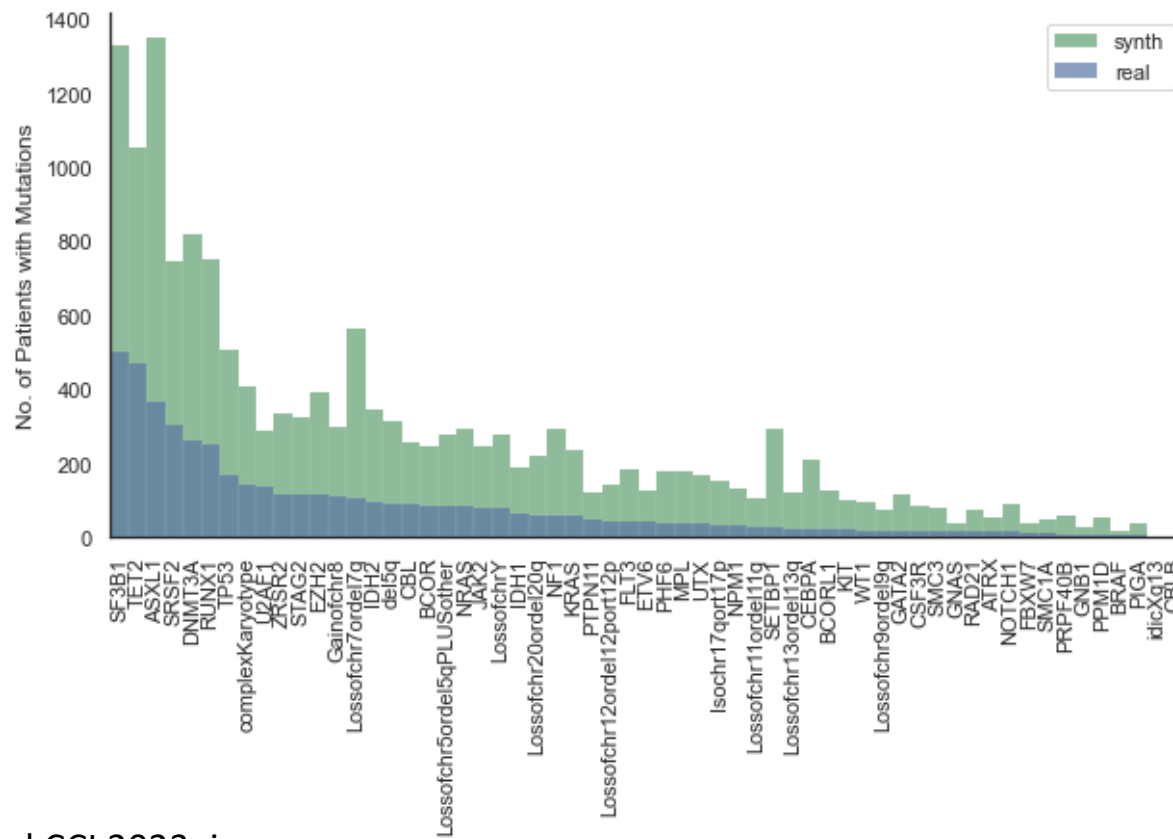


**SYNTHETIC DATA**



# Synthetic vs. Real Data: data augmentation

Data augmentation: form 2043 to 5000 patients



D'Amico S et al, J Clin Oncol CCI 2023, in press



# Performance of Synthetic Data

## DEMOGRAPHIC, CLINICAL AND SURVIVAL DATA



**92.1 %**

### SYNTHETIC CLINICAL FITNESS

Evaluated with distribution plot, Principal Component Analysis and correlation matrices.

## GENOMIC DATA



**90.2 %**

### SYNTHETIC GENOMIC FITNESS

Evaluated with mutation frequencies and pairwise association.

## ALL DATA



**70.6 %**

### PRIVACY PRESERVABILITY

Evaluated considering the possibility of tracing real data from synthetic ones.

# Generation of Synthetic Data to accelerate translational research in Hematology

The NEW ENGLAND JOURNAL of MEDICINE

ORIGINAL ARTICLE

## Somatic *SF3B1* Mutation in Myelodysplasia with Ring Sideroblasts

E. Papaemmanuil, M. Cazzola, J. Boulton, L. Malcovati, P. Vyas, D. Bowen, A. Pellagatti, J.S. Wainscoat, E. Hellstrom-Lindberg, C. Gambacorti-Passerini, A.L. Godfrey, I. Rapado, A. Cvejic, R. Rance, C. McGee, P. Ellis, L.J. Mudie, P.J. Stephens, S. McLaren, C.E. Massie, P.S. Tarpey, I. Varela, S. Nik-Zainal, H.R. Davies, A. Shlien, D. Jones, K. Raine, J. Hinton, A.P. Butler, J.W. Teague, E.J. Baxter, J. Score, A. Galli, M.G. Della Porta, E. Travaglino, M. Groves, S. Tauro, N.C. Munshi, K.C. Anderson, A. El-Naggar, A. Fischer, V. Mustonen, A.J. Warren, N.C.P. Cross, A.R. Green, P.A. Futreal, M.R. Stratton, and P.J. Campbell for the Chronic Myeloid Disorders Working Group of the International Cancer Genome Consortium

**6 patients**  
**2011**

NEJM  
Evidence

Published June 12, 2022  
NEJM Evid 2022; 1 (7)  
DOI: [10.1056/EVIDoa2200008](https://doi.org/10.1056/EVIDoa2200008)

ORIGINAL ARTICLE

## Molecular International Prognostic Scoring System for Myelodysplastic Syndromes

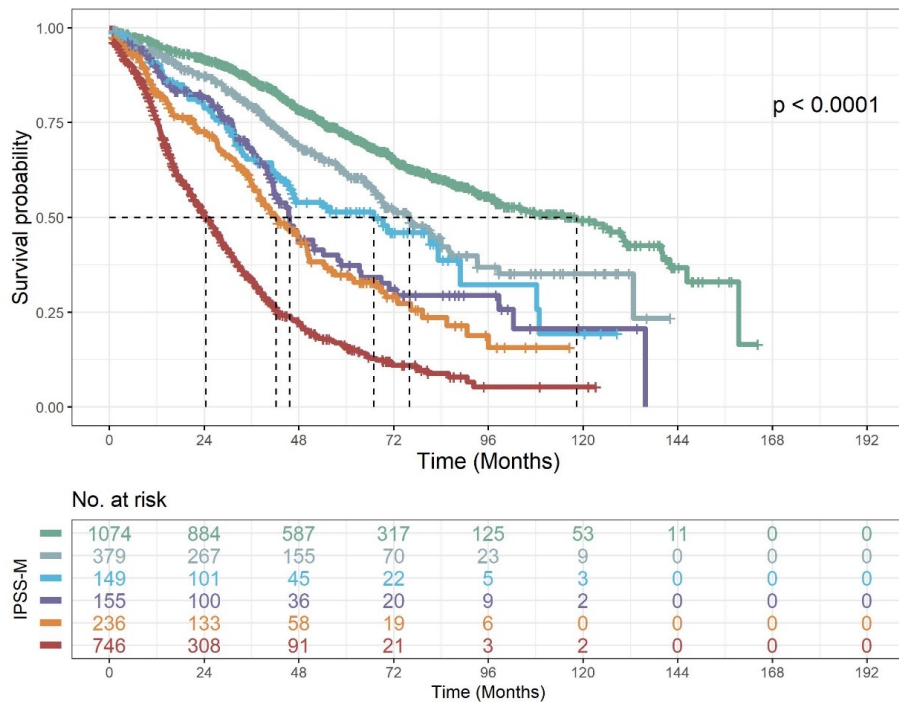
Elsa Bernard, Ph.D.,<sup>1</sup> Heinz Tuechler, Peter L. Greenberg, M.D.,<sup>2</sup> Robert P. Hasserjian, M.D.,<sup>3</sup> Juan E. Arango Ossa, M.S.,<sup>1</sup> Yasuhiro Nannya, M.D., Ph.D.,<sup>4,5</sup> Sean M. Devlin, Ph.D.,<sup>1</sup> Maria Creignou, M.D.,<sup>6</sup> Philippe Pinel, M.S.,<sup>1</sup> Lily Monnier, M.S.,<sup>1</sup> Gunes Gundem, Ph.D.,<sup>1</sup> Juan S. Medina-Martinez, M.S.,<sup>1</sup> Dylan Domenico, B.S.,<sup>1</sup> Martin Jädersten, M.D., Ph.D.,<sup>6</sup> Ulrich Germing, M.D.,<sup>7</sup> Guillermo Sanz, M.D., Ph.D.,<sup>8,9,10</sup> Arjan A. van de Loosdrecht, M.D., Ph.D.,<sup>11</sup> Olivier Kosmider, M.D., Ph.D.,<sup>12</sup> Matilde Y. Follo, Ph.D.,<sup>13</sup> Felicitas Thol, M.D.,<sup>14</sup> Lurdes Zamora, Ph.D.,<sup>15</sup> Ronald F. Pinheiro, Ph.D.,<sup>16</sup> Andrea Pellagatti, Ph.D.,<sup>17</sup> Harold K. Elias, M.D.,<sup>18</sup> Detlef Haase, M.D., Ph.D.,<sup>19</sup> Christina Ganster, Ph.D.,<sup>19</sup> Lionel Ades, M.D., Ph.D.,<sup>20</sup> Magnus Tobinsson, M.D., Ph.D.,<sup>6</sup> Laura Palomo, Ph.D.,<sup>21</sup> Matteo Giovanni Della Porta, M.D.,<sup>22</sup> Akifumi Takaori-Kondo, M.D., Ph.D.,<sup>23</sup> Takayuki Ishikawa, M.D., Ph.D.,<sup>24</sup> Shigeru Chiba, M.D., Ph.D.,<sup>25</sup> Senji Kasahara, M.D., Ph.D.,<sup>26</sup> Yasushi Miyazaki, M.D., Ph.D.,<sup>27</sup> Agnes Viale, Ph.D.,<sup>28</sup> Kety Huberman, B.S.,<sup>28</sup> Pierre Fenaux, M.D., Ph.D.,<sup>29</sup> Monika Belickova, Ph.D.,<sup>29</sup> Michael R. Savona, M.D.,<sup>30</sup> Virginia M. Klinek, M.D.,<sup>18</sup> Fabio P. S. Santos, M.D., Ph.D.,<sup>31</sup> Jacqueline Boulton, Ph.D.,<sup>17</sup> Ioannis Kotsianidis, M.D., Ph.D.,<sup>32</sup> Valeria Santini, M.D.,<sup>33</sup> Francesc Solé, Ph.D.,<sup>21</sup> Uwe Platzbecker, M.D.,<sup>34</sup> Michael Heuser, M.D.,<sup>14</sup> Peter Valent, M.D.,<sup>35,36</sup> Kazuma Ohyashiki, M.D., Ph.D.,<sup>37</sup> Carlo Finelli, M.D.,<sup>38</sup> Maria Teresa Voso, M.D.,<sup>39</sup> Lee-Yung Shih, M.S.,<sup>40</sup> Michaela Fontenay, M.D., Ph.D.,<sup>12</sup> Joop H. Jansen, Ph.D.,<sup>41</sup> José Cervera, M.D., Ph.D.,<sup>42</sup> Norbert Gattermann, M.D.,<sup>7</sup> Benjamin L. Ebert, M.D., Ph.D.,<sup>43</sup> Rafael Bejar, M.D., Ph.D.,<sup>44</sup> Luca Malcovati, M.D.,<sup>45</sup> Mario Cazzola, M.D.,<sup>45</sup> Seishi Ogawa, M.D., Ph.D.,<sup>4,46,47</sup> Eva Hellström-Lindberg, M.D., Ph.D.,<sup>6</sup> and Elli Papaemmanuil, Ph.D.<sup>1</sup>

**2957 patients**  
**2022**

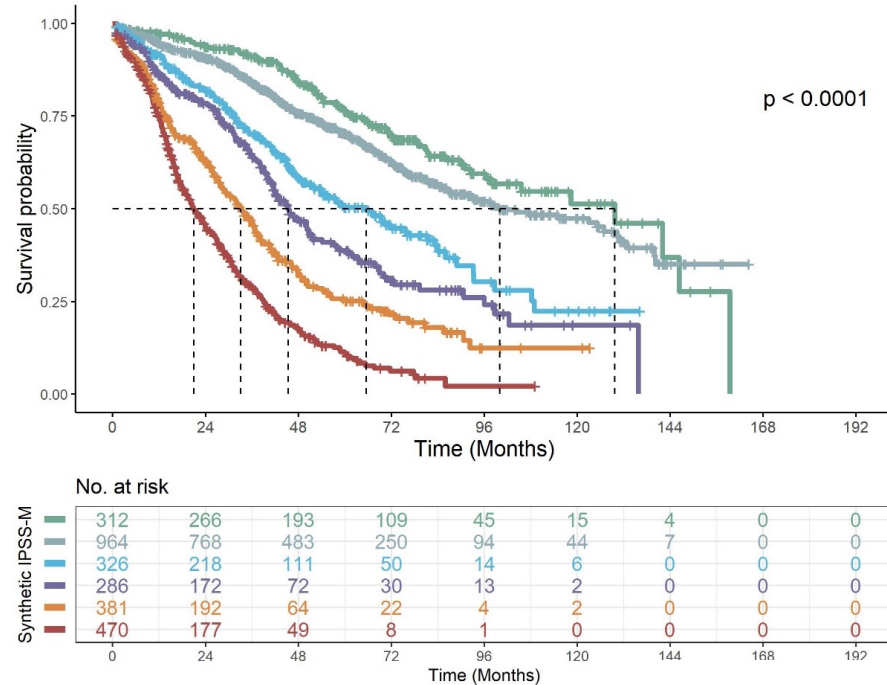


# Generation of Synthetic Data to accelerate translational research in Hematology

IPSS-M (real data, 2022)



Syntetic IPSS-M (synthetic data, 2013)



D'Amico S et al, J Clin Oncol CCI 2023, in press



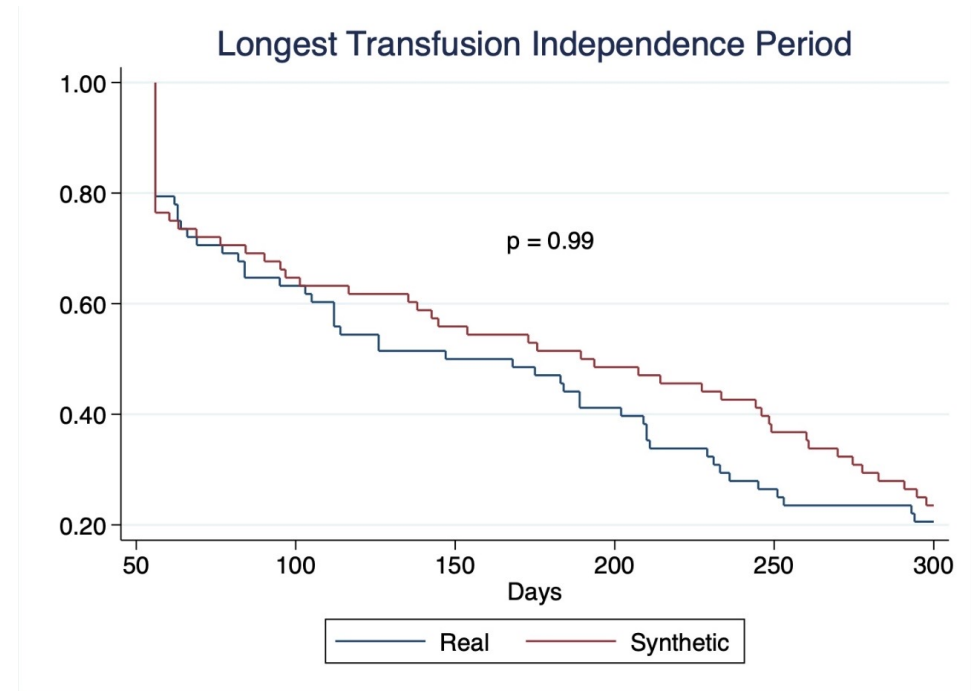
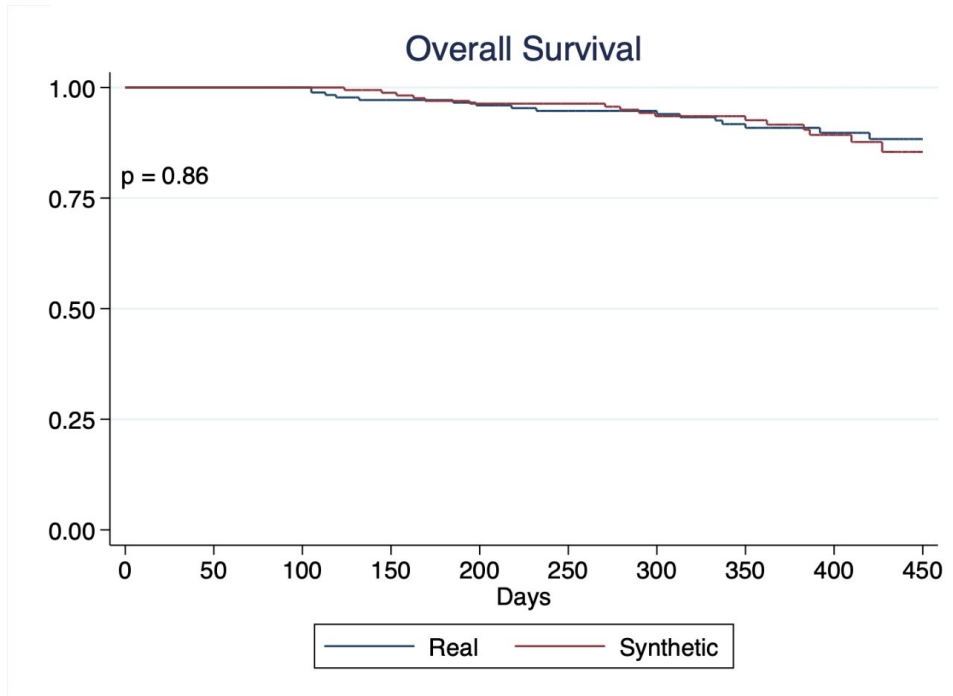
# Generation of Synthetic Data to accelerate clinical research in Hematology

Comparing endpoints of clinical trials using **real** and **synthetic** control arms. Real-world efficacy and safety of luspatercept in adult patients with transfusion-dependent anemia due to very low-, low and intermediate-risk myelodysplastic syndrome (MDS) with ring sideroblasts, who had an unsatisfactory response to or are ineligible for erythropoietin-based therapy: a multicenter study by Fondazione Italiana Sindromi Mielodisplastiche (FISIM)

## Primary endpoint

	Real data	Synthetic data	Pvalue
RBC-TI $\geq$ 8 weeks 1-24	56 (31,5)	56 (31,5)	1.0
Longest Transfusion Independence Period (weeks), median (range)	195 (56-490)	191 (56-490)	0.34
RBC-TI $\geq$ 8 weeks 1-48	68 (38,2)	61 (34,3)	0.50
RBC-TI $\geq$ 12 weeks 1-24	36 (20,2)	41 (23,0)	0.60
RBC-TI $\geq$ 12 weeks 1-48	51 (28,7)	46 (25,8)	0.63
Reduction $\geq$ 4 RBC	62 (34,8)	63 (35,4)	1.0
Reduction $\geq$ 50%	77 (43,3)	72 (40,4)	0.66
AML Evolution	4 (2,2)	6 (3,4)	0.75
Discontinued patients	74 (41,6)	82 (46,1)	0.64

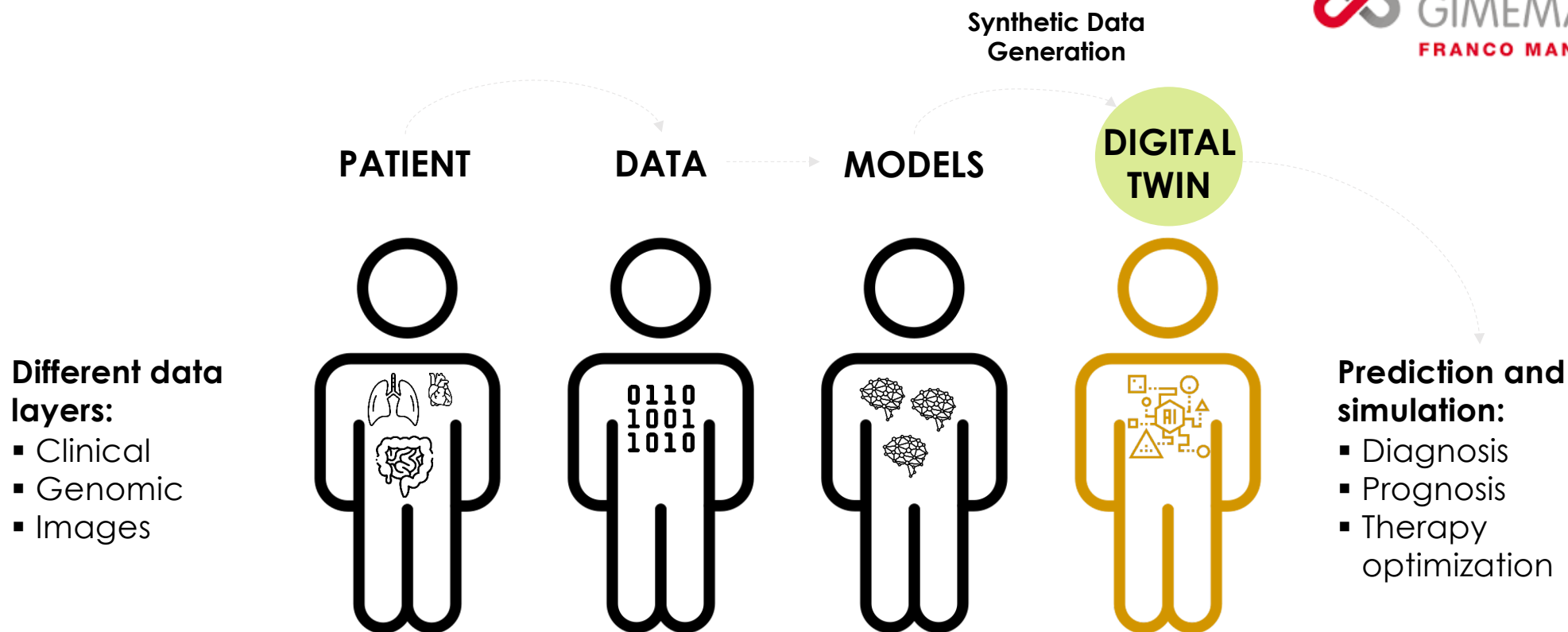
# Generation of Synthetic Data to accelerate clinical research in Hematology



Comparing endpoints of clinical trials using **real** and **synthetic** control arms. Real-world efficacy and safety of luspatercept in adult patients with transfusion-dependent anemia due to very low-, low and intermediate-risk myelodysplastic syndrome (MDS) with ring sideroblasts, who had an unsatisfactory response to or are ineligible for erythropoietin-based therapy: a multicenter study by Fondazione Italiana Sindromi Mielodisplastiche (FISIM)

D'Amico S et al, J Clin Oncol CCI 2023, in press

# From Synthetic data to Digital Twins





C.A.L.R.

## CENTER FOR ACCELERATING LEUKEMIA/LYMPHOMA RESEARCH

*Artificial Intelligence and real world data analysis to improve  
patient care and advance medical research in hematology*

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HUMANITAS  
UNIVERSITY

POLITECNICO  
MILANO 1885



### AI people

- Saverio D'Amico
- Elisabetta Sauta
- Gianluca Asti
- Victor Savevski
- Gastone Castellani

### Clinical team

- Luca Lanino
- Giulia Maggioni
- Erica Travaglino
- Alessia Campagna
- Marta Ubezio
- Antonio Russo
- Gabriele Todisco
- Armando Santoro

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