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Livio Pagano, Jon Salmanton-García, Francesco Marchesi, Ola Blennow, Maria Gomes da Silva, Andreas Glenthøj, Jaap van Doesum, Yavuz Bilgin, Alberto Lopez-Garcia, Federico Itri, et al.

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American Society of Hematology 2021 L Street NW, Suite 900, Washington, DC 20036 Phone: 202-776-0544 | Fax 202-776-0545 editorial@hematology.org

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Livio Pagano (Hematology, Fondazione Policlinico Universitario Agostino Gemelli - IRCCS, Rome, Italy, Italy) Jon Salmanton-García (University Hospital of Cologne, Germany) Francesco Marchesi (IRCCS Regina Elena National Cancer Institute, Italy) Ola Blennow (Department of Infectious Diseases, Karolinska University Hospital, Sweden) Maria Gomes da Silva (Portuguese Institute of Oncology, Portugal) Andreas Glenthøj (Copenhagen University Hospital - Rigshospitalet, Denmark) Jaap A. van Doesum (University Medical Center Groningen, Groningen, the Netherlands, Netherlands) Yavuz Bilgin (Admiraal de Ruijter Hospital, Netherlands) Alberto Lopez-Garcia (Fundacion Jimenez Diaz University Hospital, Health Research Institute IIS-FJD, Spain) Federico Itri (San Luigi Gonzaga Hospital - Orbassano, Orbassano, Italy, Italy) Raquel Nunes Rodrigues (Departamento de Hematologia, Instituto Português de Oncologia, Lisboa, Portugal, Portugal) Barbora Weinbergerová (Masaryk University and University Hospital Brno, Czech Republic) Francesca Farina (San Raffaele Scientific Institute, Italy) Giulia Dragonetti (Fondazione Policlinico Universitario Agostino Gemelli - IRCCS, Italy) Caroline BERG VENEMYR (Department of Hematology, Copenhagen University Hospital - Rigshospitalet, Copenhagen, Denmark, Denmark) Jens VAN PRAET (AZ Sint-Jan Brugge-Oostende AV, Brugge, Belgium, Belgium) Ozren Jaksic (Dubrava University Hospital, Croatia) Toni VALKOVIĆ ("University Hospital Centre Rijeka, Rijeka, Croatia Croatian Cooperative Group for Hematological Diseases (CROHEM) Faculty of Medicine and Faculty of Health Studies University of Rijeka, Rijeka, Cro, Croatia, Republic of) Iker Falces-Romero (La Paz University Hospital, Spain) Sonia MARTÍN-PÉREZ (Hospital Nuestra Señora de Sonsoles, Ávila, Spain, Spain) Moraima Jiménez (Department of Hematology, Vall d'Hebron University Hospital, Experimental Hematology, Vall d'Hebron Institute of Oncology (VHIO), Vall d'Hebron Barcelona Hospital Campus, Barcelona, Spain, Spain) Julio DÁVILA-VALLS (Hospital Nuestra Señora de Sonsoles, Ávila, Spain, Spain) Martin SCHÖNLEIN (Department of Oncology, Hematology and Bone Marrow Transplantation with Section of Pneumology, University Medical Center Hamburg-Eppendorf, Hamburg, Germany, Germany) Emanuele Ammatuna (University Medical Center Groningen, Netherlands) Stef Meers (AZ KLINA, Brasschaat, Belgium, Belgium) Mario Delia (Azienda Universitaria Ospedaliera Consorziale - Policlinico Bari, Italy) Zlate STOJANOSKI (University Clinic of Hematology, Skopje, Republic of North Macedonia, North Macedonia, Republic of) Anna NORDLANDER (Department of Infectious Diseases, Karolinska University Hospital,, Sweden) Tobias Lahmer (Technical University of Munich, School of Medicine, University hospital rechts der Isar, Germany) László Imre Pinczés (University of Debrecen, Hungary) Caterina BUQUICCHIO (Ematologia con Trapianto, Ospedale Dimiccoli Barletta, Barletta, Italy, Italy) Klára PIUKOVICS (Department of Internal Medicine, South Division Faculty of Medicine University of Szeged, Szeged, Hungary, Hungary) Irati ORMAZABAL-VÉLEZ (Complejo Hospitalario de Navarra, Iruña-Pamplona, Spain, Spain) Nicola Fracchiolla (UOC Ematologia, Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico, Italy) Michail SAMARKOS (35. Laikon Hospital, Greece) Gustavo-Adolfo MÉNDEZ (Hospital Escuela de Agudos Dr. Ramon Madariaga, Argentina) José-Ángel Hernández-Rivas (Hospital Universitario Infanta Leonor, Spain) Ildefonso Espigado (Hospital Univesitario Virgen Macarena & Virgen del Rocío, Seville,, Spain) Martin CERNAN (University Hospital Olomouc, Olomouc, Czech Republic, Czech Republic) Verena Petzer (Innsbruck Medical University, Austria) Sylvain Lamure (Montpellier University Hospital, France) Roberta Di Blasi (Hopital Saint Louis, France) Joyce Marques de Almeida (Institute of Oncology Research, Switzerland) Michelina Dargenio ('Vito Fazzi' Hospital, Italy) Monika Biernat (Wroclaw Medical University, Poland) Mariarita Sciumè (Hematology Unit, Fondazione IRCCS Cà Granda Ospedale Maggiore Policlinico, Italy) Cristina de Ramón (Hematology Department, University Hospital of Salamanca, Institute of Biomedical Research of Salamanca (IBSAL), Spain) Nick DE JONGE (Amsterdam UMC, location AMC, Amsterdam, Netherlands, Netherlands) Josip BATINIĆ (University Hospital Centre Zagreb, Zagreb, Croatia, Croatia, Republic of) Avinash Aujayeb (Northumbria Healthcare NHS Foundation Trust, United Kingdom) Monia Marchetti (Az. OSP. SS ANTONIO e BIAGIO e CESARE ARRIGO, Italy) Guillemette Fouquet (52. Cochin Hospital, France) Noemi Fernández Escalada (Marques de Valdecilla University Hospital, Spain) Giovanni Zambrotta (Azienda Ospedaliera San Gerardo, Italy) Maria Vittoria SACCHI (Azienda Ospedaliera Nazionale SS. Antonio e Biagio e Cesare Arrigo, Alessandria, Italy, Italy) Anna Guidetti (Fondazione IRCCS Istituto Nazionale Tumori, Italy) Fatih DEMIRKAN (Dokuz Eylul University, Division of Hematology, Izmir, Turkey, Turkey) Lucia Prezioso (University of Parma, Italy) Zdenek Racil (Institute of Hematology and Blood Transfusion Prague, Czech Republic) Marcio Nucci (Universidade Federal do Rio de Janeiro, Brazil) Miloš MLADENOVIĆ (COVID hospital "Batajnica",

Serbia) Raphaël LIÉVIN (Hopital Saint Louis, Paris, France, France) Michaela HANÁKOVÁ ("Institute of Hematology and Blood Transfusion, Prague, Czech Republic Department of Physiology, Faculty of Medicine, Masaryk University, Brno, Czech Republic", Czech Republic) Stefanie GRÄFE (University Medical Center Hamburg-Eppendorf, Germany) Uluhan Sili (School of Medicine, Marmara University, Istanbul, Turkey, Turkey) Marina Machado (Clinical Microbiology and Infectious Diseases Department, Hospital General Universitario Gregorio Marañón. Instituto de Investigación Sanitaria Gregorio Marañón, Spain) Chiara Cattaneo (Hematology, ASST-Spedali Civili, Brescia, Italy) Tatjana ADŽIĆ-VUKIČEVIĆ (COVID hospital "Batajnica", Belgrade, Serbia, Serbia) Luisa Verga (Azienda Ospedaliera San Gerardo, Italy) Jorge Labrador (Hospital Universitario de Burgos, Burgos,) Laman RAHIMLI ("University of Cologne, Faculty of Medicine and University Hospital Cologne, Department I of Internal Medicine, Excellence Center for Medical Mycology (ECMM), Cologne, Germany University of Cologne, Faculty of Medicine and University Hospital Cologne, Cologne Excellence Cluster on Cellular Stress Responses in Aging-Associated Diseases (CECAD), Cologne, Germany", Germany) Matteo BONANNI ("Hematology Unit, Fondazione Policlinico Universitario Agostino Gemelli - IRCCS, Rome, Italy Hematology Unit, Università Cattolica del Sacro Cuore, Rome, Italy", Italy) Francesco Passamonti (University of Insubria, Italy) Antonio Pagliuca (King's College Hospital, United Kingdom) Paolo Corradini (Fondazione IRCCS Istituto Nazionale dei Tumori, University of Milano, Italy) Martin Hoenigl (University of California San Diego, United States) Philipp Koehler (University Hospital Cologne, Germany) Alessandro Busca (A.O.U. Città della Salute e della Scienza, Italy) Oliver Cornely (Uniklinik Köln, Germany)

Abstract:

Limited data have been published on the epidemiology and outcomes of breakthrough COVID-19 in patients with hematological malignancy (HM) after anti-SARS-CoV-2 vaccination.

Adult HM who received at least one dose of anti-SARS-CoV-2 vaccine and diagnosed with breakthrough COVID-19 between January 2021 and March 2022 and registered in EPICOVIDEHA were included in this analysis.

A total of 1548 cases were included, mainly with lymphoid malignancies (1181 cases, 76%). After viral genome sequencing in 753 cases (49%), Omicron variant was prevalent (517, 68.7%). Most of the patients received at least two vaccine doses before COVID-19 (1419, 91%), mostly mRNA-based (1377, 89%). Overall, 906 patients (59%) received specific treatment for COVID-19. After 30-days follow-up from COVID-19 diagnosis, 143 patients (9%) died. The mortality rate in patients with Omicron variant was of 7.9%, comparable to that reported for the other variants. The 30-day mortality rate was significantly lower than in the pre-vaccine era (31%). In the univariable analysis, older age (p<0.001), active HM (p<0.001), severe and critical COVID-19 (p=0.007 and p<0.001, respectively) were associated with mortality. Conversely, patients receiving monoclonal antibodies, even for severe or critical COVID-19, had a lower mortality rate (p<0.001). In the multivariable model, older age, active disease, critical COVID-19 and at least 2-3 comorbidities were correlated with a higher mortality, whereas the administration of monoclonal antibodies, alone (p<0.001) or combined with antivirals (p=0.009), was observed protective.

While mortality is significantly lower than in the pre-vaccination era, breakthrough COVID-19 in HM is still associated with considerable mortality. Death rate was lower in patients who received monoclonal antibodies, alone or in combination with antivirals.

EPICOVIDEHA (www.clinicaltrials.gov; National Clinical Trials identifier NCT04733729) is an international open web-based registry for patients with HMs infected with SARS-CoV-2.

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Livio PAGANO,1*# Jon SALMANTON-GARCÍA,2* Francesco MARCHESI,3* Ola BLENNOW, 4 Maria GOMES DA SILVA, 5 Andreas GLENTHØJ, 6 Jaap VAN DOESUM, 7 Yavuz M. BILGIN,⁸ Alberto LÓPEZ-GARCÍA,⁹ Federico ITRI,¹⁰ Raquel NUNES RODRIGUES, 11 Barbora WEINBERGEROVÁ, 12 Francesca FARINA, 13 Giulia 7 DRAGONETTI, 14 Caroline BERG VENEMYR, 15 Jens VAN PRAET, 16 Ozren JAKSIC, 17 Toni VALCOVIĆ, ¹⁸ Iker FALCES-ROMERO, ¹⁹ Sonia MARTÍN-PÉREZ, ²⁰ Moraima JIMÉNEZ, ²¹ Julio DÁVILA-VALLS,²² Martin SCHÖNLEIN,²³ Emanuele AMMATUNA,²⁴ Stef MEERS,²⁵ 10 Mario **DELIA**, ²⁶ Zlate **STOJANOSKI**, ²⁷ Anna **NORDLANDER**, ²⁸ Tobias **LAHMER**, ²⁹ László 11 Imre PINCZÉS, 30 Caterina BUQUICCHIO, 31 Klára PIUKOVICS, 32 Irati ORMAZABAL-12 VÉLEZ, 33 Nicola FRACCHIOLLA, 34 Michail SAMARKOS, 35 Gustavo-Adolfo MÉNDEZ, 36 13 José-Ángel HERNÁNDEZ-RIVAS, 37 Ildefonso ESPIGADO, 38 Martin CERNAN, 39 Verena 14 PETZER, 40 Sylvain LAMURE, 41 Roberta DI BLASI, 42 Joyce MARQUES DE ALMEDIA, 43 15 Michelina DARGENIO, 44 Monika M. BIERNAT, 45 Mariarita SCIUMÈ, 46 Cristina DE 16 RAMÓN, 47 Nick DE JONGE, 48 Josip BATINIĆ, 49 Avinash AUJAYEB, 50 Monia 17 MARCHETTI,⁵¹ Guillemette FOUQUET,⁵² Noemí FERNÁNDEZ,⁵³ Giovanni ZAMBROTTA,⁵⁴ 18 Maria Vittoria SACCHI,⁵⁵ Anna GUIDETTI,⁵⁶ Fatih DEMIRKAN,⁵⁷ Lucia PREZIOSO,⁵⁸ 19 Zdeněk RÁČIL, 59 Marcio NUCCI, 60 Miloš MLADENOVIĆ, 61 Raphaël LIÉVIN, 62 Michaela 20 HANÁKOVÁ, 63 Stefanie GRÄFE, 64 Uluhan SILI, 65 Marina MACHADO, 66 Chiara 21 CATTANEO, 67 Tatjana ADŽIĆ-VUKIČEVIĆ, 68 Luisa VERGA, 69 Jorge LABRADOR, 70 Laman 22 RAHIMLI, 71 Matteo BONANNI, 72 Francesco PASSAMONTI, 73 Antonio PAGLIUCA, 74 Paolo 23 CORRADINI, 75 Martin HOENIGL, 76 Philipp KOEHLER, 77 Alessandro BUSCA, 78 Oliver A. 24 CORNELY⁷⁹ 25

^{*} L.P., J.S.G., and F.M. are joint first authors.

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26

28 Affiliations

- 29 1. Hematology Unit, Fondazione Policlinico Universitario Agostino Gemelli IRCCS,
- Rome, Italy
- Hematology Unit, Università Cattolica del Sacro Cuore, Rome, Italy
- 32 2. University of Cologne, Faculty of Medicine and University Hospital Cologne,
- Department I of Internal Medicine, Excellence Center for Medical Mycology (ECMM),
- 34 Cologne, Germany
- University of Cologne, Faculty of Medicine and University Hospital Cologne, Cologne
- 36 Excellence Cluster on Cellular Stress Responses in Aging-Associated Diseases
- 37 (CECAD), Cologne, Germany
- 38 3. Hematology and Stem Cell Transplant Unit, IRCCS Regina Elena National Cancer
- 39 Institute, Rome, Italy
- 40 4. Department of Infectious Diseases, Karolinska University Hospital, Stockholm,
- 41 Sweden
- 42 5. Portuguese Institute of Oncology, Lisbon, Portugal
- 43 6. Department of Hematology, Copenhagen University Hospital Rigshospitalet,
- 44 Copenhagen, Denmark
- 45 7. University Medical Center Groningen, Groningen, Netherlands
- 46 8. ADRZ, Goes, Netherlands
- 47 9. Fundación Jimenez Diaz University Hospital, Health Research Institute IIS-FJD,
- 48 Madrid, Spain
- 49 10. San Luigi Gonzaga Hospital Orbassano, Orbassano, Italy
- 50 11. Portuguese Institute of Oncology, Lisbon, Portugal
- 51 12. University Hospital Brno Department of Internal Medicine, Hematology and Oncology,
- 52 Brno, Czech Republic
- 53 13. IRCCS Ospedale San Raffaele, Milan, Italy
- 54 14. Hematology Unit, Fondazione Policlinico Universitario Agostino Gemelli IRCCS,
- 55 Rome, Italy
- 56 15. Department of Hematology, Copenhagen University Hospital Rigshospitalet,
- 57 Copenhagen, Denmark
- 58 16. Department of Nephrology and Infectious diseases, AZ Sint-Jan Brugge-Oostende AV,
- 59 Brugge, Belgium
- 60 17. University Hospital Dubrava, Zagreb, Croatia
- 61 18. CHC Rijeka, Rijeka, Croatia
- 62 19. La Paz University Hospital, Madrid, Spain
- 63 20. Hospital Nuestra Señora de Sonsoles, Ávila, Spain

- 64 21. Department of Hematology, Vall d'Hebron Hospital Universitari, Experimental
- Hematology, Vall d'Hebron Institute of Oncology (VHIO), Vall d'Hebron Barcelona
- 66 Hospital Campus, Barcelona, Spain
- 67 Departament de Medicina, Universitat Autònoma de Barcelona, Bellaterra, Spain
- 68 22. Hospital Nuestra Señora de Sonsoles, Ávila, Spain
- 69 23. Department of Oncology, Hematology and Bone Marrow Transplantation with Section
- 70 of Pneumology, University Medical Center Hamburg-Eppendorf, Hamburg, Germany
- 71 24. University Medical Center Groningen, Groningen, Netherlands
- 72 25. AZ KLINA, Brasschaat, Belgium
- 73 26. Azienda Ospedaliero-Universitaria Consorziale Policlinico di Bari, Bari, Italy
- 74 27. University Clinic of Hematology, Skopje, North Macedonia
- 75 28. Department of Infectious Diseases, Karolinska University Hospital, Stockholm,
- 76 Sweden
- 77 29. Medizinische Klinik II, Klinikum rechts der Isar, TU München, Munich, Germany
- 78 30. Division of Hematology, Department of Internal Medicine, University of Debrecen,
- 79 Debrecen, Hungary
- 80 31. Ematologia con Trapianto, Ospedale Dimiccoli Barletta, Barletta, Italy
- 81 32. Department of Internal Medicine, South Division Faculty of Medicine University of
- 82 Szeged, Szeged, Hungary
- 83 33. Complejo Hospitalario de Navarra, Iruña-Pamplona, Spain
- 84 34. Hematology Unit, Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico,
- 85 Milan, Italy
- 86 35. Laikon Hospital, Athens, Greece
- 87 36. Hospital Escuela de Agudos Dr. Ramón Madariaga, Posadas, Argentina
- 88 37. Hospital Universitario Infanta Leonor, Madrid, Spain
- 89 38. Department of Hematology, University Hospital Virgen Macarena University Hospital
- 90 Virgen del Rocío, Instituto de Biomedicina de Sevilla (IBIS / CSIC), Universidad de
- 91 Sevilla (Departamento de Medicina), Seville, Spain
- 92 39. University Hospital Olomouc, Olomouc, Czech Republic
- 93 40. Department of Hematology and Oncology, Medical University of Innsbruck, Innsbruck,
- 94 Austria
- 95 41. CHU Montpellier, Montpellier, France
- 96 42. Hopital Saint Louis, Paris, France
- 97 43. Istituto Oncologico della Svizzera Italiana, Bellinzona, Switzerland
- 98 44. Ospedale Vito Fazzi, Lecce, Italy
- 99 45. Department of Haematology, Blood Neoplasms, and Bone Marrow Transplantation,
- 100 Wroclaw Medical University, Wroclaw, Poland

- 101 46. Hematology Unit, Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico,
- 102 Milan, Italy
- 103 47. Hematology Department, Hospital Universitario de Salamanca, Salamanca, Spain
- 104 IBSAL, Centro de Investigación del Cáncer-IBMCC (USAL-CSIC), Salamanca, Spain
- 105 48. Amsterdam UMC, location VUmc, Amsterdam, Netherlands
- 106 49. University Hospital Centre Zagreb, Zagreb, Croatia
- 107 Croatian Cooperative Group for Hematological Diseases (CROHEM), Croatia
- 108 Faculty of Medicine University of Zagreb, Zagreb, Croatia
- 109 50. Northumbria Healthcare, Newcastle, United Kingdom
- 110 51. Azienda Ospedaliera Nazionale SS. Antonio e Biagio e Cesare Arrigo, Alessandria,
- 111 Italy
- 112 52. Cochin Hospital, APHP, Paris, France
- 113 53. Hospital Universitario Marqués de Valdecilla, Santander, Spain
- 114 54. Azienda Ospedaliera San Gerardo Monza, Monza, Italy
- 115 Università Milano-Bicocca, Milan, Italy
- 116 55. Azienda Ospedaliera Nazionale SS. Antonio e Biagio e Cesare Arrigo, Alessandria,
- 117 Italy
- 118 56. Fondazione IRCCS Istituto Nazionale dei Tumori, Milan, Italy
- 119 57. Dokuz Eylul University, Division of Hematology, Izmir, Turkey
- 120 58. Hospital University of Parma Hematology and Bone Marrow Unit, Parma, Italy
- 121 59. Institute of Hematology and Blood Transfusion, Prague, Czech Republic
- Department of Physiology, Faculty of Medicine, Masaryk University, Brno, Czech
- 123 Republic
- 124 60. Department of Internal Medicine, Federal University of Rio de Janeiro, Rio de Janeiro,
- 125 Brazil
- 126 61. COVID hospital "Batajnica", Belgrade, Serbia
- 127 62. Hopital Saint Louis, Paris, France
- 128 63. Institute of Hematology and Blood Transfusion, Prague, Czech Republic
- 129 Department of Physiology, Faculty of Medicine, Masaryk University, Brno, Czech
- 130 Republic
- 131 64. Universitätsklinikum Hamburg Eppendorf, Hamburg, Germany
- 132 University of Cologne, Faculty of Medicine and University Hospital Cologne,
- Department I of Internal Medicine, Excellence Center for Medical Mycology (ECMM),
- 134 Cologne, Germany
- University of Cologne, Faculty of Medicine and University Hospital Cologne, Cologne
- 136 Excellence Cluster on Cellular Stress Responses in Aging-Associated Diseases
- 137 (CECAD), Cologne, Germany

- 138 65. Marmara University, Istanbul, Turkey
- 139 66. Clinical Microbiology and Infectious Diseases Department, Hospital General
- 140 Universitario Gregorio Marañón, Madrid, Spain
- 141 67. Hematology Unit, ASST-Spedali Civili, Brescia, Italy
- 142 68. COVID hospital "Batajnica", Belgrade, Serbia
- 143 69. Azienda Ospedaliera San Gerardo Monza, Monza, Italy
- 144 Università Milano-Bicocca, Milan, Italy
- 145 70. Department of Hematology, Hospital Universitario de Burgos, Burgos, Spain
- 146 71. University of Cologne, Faculty of Medicine and University Hospital Cologne,
- 147 Department I of Internal Medicine, Excellence Center for Medical Mycology (ECMM),
- 148 Cologne, Germany
- University of Cologne, Faculty of Medicine and University Hospital Cologne, Cologne
- 150 Excellence Cluster on Cellular Stress Responses in Aging-Associated Diseases
- 151 (CECAD), Cologne, Germany
- 152 72. Hematology Unit, Fondazione Policlinico Universitario Agostino Gemelli IRCCS,
- 153 Rome, Italy
- 154 Hematology Unit, Università Cattolica del Sacro Cuore, Rome, Italy
- 155 73. Department of Medicine and Surgery, University of Insubria and ASST Sette Laghi,
- 156 Ospedale di Circolo of Varese, Varese, Italy
- 157 74. Department of Hematological Medicine, King's College Hospital NHS Foundation
- 158 Trust, Kings College London & Anthony Nolan, United Kingdom
- 159 75. University of Milan and Fondazione IRCCS Istituto Nazionale dei Tumori, Milan, Italy
- 160 76. Division of Infectious Diseases and Global Public Health, Department of Medicine,
- University of California San Diego, San Diego, CA, United States
- 162 Clinical and Translational Fungal-Working Group, University of California San Diego,
- La Jolla, CA, United States
- 164 Division of Infectious Diseases, Department of Internal Medicine, Medical University of
- 165 Graz, Graz, Austria
- 166 77. University of Cologne, Faculty of Medicine and University Hospital Cologne,
- 167 Department I of Internal Medicine, Excellence Center for Medical Mycology (ECMM),
- 168 Cologne, Germany
- University of Cologne, Faculty of Medicine and University Hospital Cologne, Center for
- 170 Molecular Medicine Cologne (CMMC), Cologne, Germany
- 171 University of Cologne, Faculty of Medicine and University Hospital Cologne, Cologne
- 172 Excellence Cluster on Cellular Stress Responses in Aging-Associated Diseases
- 173 (CECAD), Cologne, Germany
- 174 78. Stem Cell Transplant Center, AOU Citta' della Salute e della Scienza, Turin, Italy

1/5	19.	Offiversity of Cologne, Faculty of Medicine and Offiversity Hospital Cologne,
176		Department I of Internal Medicine, Excellence Center for Medical Mycology (ECMM),
177		Cologne, Germany
178		University of Cologne, Faculty of Medicine and University Hospital Cologne, Chair
179		Translational Research, Cologne Excellence Cluster on Cellular Stress Responses in
180		Aging-Associated Diseases (CECAD), Cologne, Germany
181		University of Cologne, Faculty of Medicine and University Hospital Cologne, Clinical
182		Trials Centre Cologne (ZKS Köln), Cologne, Germany
183		University of Cologne, Faculty of Medicine and University Hospital Cologne, Center for
184		Molecular Medicine Cologne (CMMC), Cologne, Germany
185		German Centre for Infection Research (DZIF), Partner Site Bonn-Cologne, Cologne,
186		Germany

ORCID numbers

Livio PAGANO	
Livio.Pagano@unicatt.it	0000-0001-8287-928X
Jon SALMANTON-GARCÍA	0000 0000 0700 0007
jon.salmanton-garcia@uk-koeln.de	0000-0002-6766-8297
Francesco MARCHESI	0000-0001-6353-2272
francesco.marchesi@ifo.it	0000 0001 0000 2212
Ola BLENNOW ola.blennow@regionstockholm.se	0000-0002-7167-7882
Maria GOMES DA SILVA	
mgsilva@ipolisboa.min-saude.pt	0000-0002-6993-2450
Andreas GLENTHØJ	0000-0003-2082-0738
andreas.glenthoej@regionh.dk Jaap VAN DOESUM	
j.a.van.doesum@umcg.nl	0000-0003-0214-3219
Yavuz M. BILGIN	
y.bilgin@adrz.nl	0000-0003-4854-5424
Alberto LÓPEZ-GARCÍA	0000 0000 5254 5204
alberto.lgarcia@quironsalud.es	0000-0002-5354-5261
Federico ITRI	0000-0002-3532-5281
federico.itri@unito.it	0000-0002-0002-0201
Barbora WEINBERGEROVÁ	0000-0001-6460-2471
Weinbergerova.Barbora@fnbrno.cz Francesca FARINA	
farina.francesca@hsr.it	0000-0002-5124-6970
Giulia DRAGONETTI	0000 0003 1775 6333
dragonettigiulia@gmail.com	0000-0003-1775-6333
Caroline BERG VENEMYR	0000-0003-2082-0738
caroline.berg.venemyr.01@regionh.dk Jens VAN PRAET	
Jens.VanPraet@azsintjan.be	0000-0002-7125-7001
Ozren JAKSIC	
ojaksic@kbd.hr	0000-0003-4026-285X
Iker FALCES-ROMERO	0000-0001-5888-7706
falces88@gmail.com	0000 0001 0000 7700
Moraima JIMÉNEZ maria.jimenez@yhebron.net	0000-0003-1444-8562
Martin SCHÖNLEIN	
m.schoenlein@uke.de	0000-0002-1010-0975
Zlate STOJANOSKI	0000 0004 7500 0050
stojanoskiza@t.mk	0000-0001-7502-8356
László Imre PINCZÉS	0000-0003-0453-1709
pinczeslaszloimre@gmail.com	0000-0000-0400-1709
Caterina BUQUICCHIO	0000-0002-3683-5953
<u>caterinabuquicchio@libero.it</u> Klára PIUKOVICS	
piukovics.klara@gmail.com	0000-0003-4480-3131
Irati ORMAZABAL-VÉLEZ	0000-0003-1141-5546
irati.ormazabal.velez@gmail.com	333 333 1141 3340
Nicola FRACCHIOLLA nicola.fracchiolla@policlinico.mi.it	0000-0002-8982-8079
Michail SAMARKOS	
msamarkos@med.uoa.gr	0000-0001-9630-9712

Gustavo-Adolfo MÉNDEZ	0000-0003-0514-7004
mendez.doc@gmail.com	3000 0000 0011 1001
Ildefonso ESPIGADO Ildefonso.espigado.sspa@juntadeandalucia.es	0000-0002-4043-6613
Verena PETZER	0000-0002-9205-1440
verena.petzer@i-med.ac.at	
Monika M. BIERNAT	0000-0003-3161-3398
monika.biernat@am.wroc.pl	0000 0000 0101 0000
Mariarita SCIUMÈ mariarita.sciume@policlinico.mi.it	0000-0001-7958-4966
Cristina DE RAMÓN	0000-0002-8167-6410
cristinaderamonsanchez@gmail.com	
Nick DE JONGE ni.dejonge@amsterdamumc.nl	0000-0002-9901-0887
Josip BATINIĆ	
batinic.josip@gmail.com	0000-0001-5595-9911
Monia MARCHETTI	
moniamarchettitamellini@gmail.com	0000-0001-7615-0572
Giovanni ZAMBROTTA	
giovannizambrotta92@gmail.com	0000-0002-8612-2994
Maria Vittoria SACCHI	
mariavittoria.sacchi@ospedale.al.it	0000-0001-8133-3357
Fatih DEMIRKAN	0000-0002-1172-8668
fatih.demirkan@deu.edu.tr	
Lucia PREZIOSO	0000-0003-1660-4960
lprezioso@ao.pr.it	0000 0000 1000 1000
Zdeněk RÁČIL	
Zdenek.Racil@uhkt.cz	0000-0003-3511-4596
zdenek.racil@gmail.com	
Marcio NUCCI	0000-0003-4867-0014
mnucci@hucff.ufrj.br	0000-0003-4007-0014
Uluhan SILI	
uluhan@hotmail.com	0000-0002-9939-9298
uluhan.sili@marmara.edu.tr	
Marina MACHADO	
marinamachadov@gmail.com	0000-0002-8370-2248
Chiara CATTANEO	
chiara.cattaneo@asst-spedalicivili.it	0000-0003-0031-3237
Luisa VERGA	
luisa.verga@libero.it	0000-0002-0868-3358
Francesco PASSAMONTI	
francesco.passamonti@uninsubria.it	0000-0001-8068-5289
francesco.passamonti@asst-settelaghi.it	0000-0001-0000-0209
Antonio PAGLIUCA	
	0000-0003-2519-0333
antonio.pagliuca@kcl.ac.uk	
Paolo CORRADINI	0000-0002-9186-1353
paolo.corradini@unimi.it	-
Martin HOENIGL	0000-0002-1653-2824
hoeniglmartin@gmail.com	
Philipp KOEHLER	0000-0002-7386-7495
philipp.koehler@uk-koeln.de	
Alessandro BUSCA	0000-0001-5361-5613
abusca@cittadellasalute.to.it	2000 0001 0001 0010
Oliver A. CORNELY	0000-0001-9599-3137
oliver.cornely@uk-koeln.de	0000-0001-0000-0101

- 188 Collaborators (to be listed in PubMed)
- 189 Laura SERRANO, José-María RIBERA-SANTA SUSANA, Joseph MELETIADIS, Panagiotis
- 190 TSIRIGOTIS, Nicola COPPOLA, Malgorzata MIKULSKA, Nurettin ERBEN, Caroline BESSON,
- 191 Maria MERELLI, Tomás-José GONZÁLEZ-LÓPEZ, Jorge LOUREIRO-AMIGO, Carolina
- 192 GARCÍA-VIDAL, Elizabeth DE KORT, Annarosa CUCCARO, Sofia ZOMPI, Florian REIZINE,
- 193 Olimpia FINIZIO, Rémy DULÉRY, Maria CALBACHO, Ghaith ABU-ZEINAH, Sandra MALAK,
- 194 Przemyslaw **ZDZIARSKI**, Gina **VARRICHIO**, Athanasios **TRAGIANNIDIS**, Gaëtan **PLANTEFEVE**,
- 195 Rafael **DUARTE**, François **DANION**, Maria Chiara **TISI**, Ioanna **SAKELLARI**, Meinholf
- 196 KARTHAUS, Ana GROH, Monica FUNG, Ziad EMARAH, Omar-Francisco CORONEL-AYALA,
- 197 Louis Yi Ann CHAI, Mathias BREHON, Valentina BONUOMO, Dominik WOLF, Jana WITTIG,
- 198 Maria **VEHRESCHILD**, Mario Virgilio **PAPA**, Julia **NEUHANN**, María-Josefa **JIMÉNEZ-LORENZO**,
- 199 Jan GROTHE, Eleni GAVRIILAKI, Ramón GARCÍA-SANZ, Nicole GARCÍA-POUTÓN, Shaimaa
- 200 Saber EL-ASHWAH, Matthias EGGERER, Raul CORDOBA, Gökçe Melis ÇOLAK, Elena
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- 212 # Corresponding Author
- 213 Prof. Livio Pagano, MD
- Fondazione Policlinico Universitario A. Gemelli IRCCS Università Cattolica del Sacro Cuore
- 215 Largo Francesco Vito 1, 00168 Rome (Italy)
- 216 E-mail: livio.pagano@unicatt.it

217 Key points

- Mortality rate in hematologic malignancy patients with breakthrough COVID-19 is about 9%,
 lower than in the pre-vaccination era
- Patients who received monoclonal antibodies, alone or combined with antivirals, show a better clinical outcome

Abstract

Limited data have been published on the epidemiology and outcomes of breakthrough COVID-19 in patients with hematological malignancy (HM) after anti-SARS-CoV-2 vaccination.

Adult HM who received at least one dose of anti-SARS-CoV-2 vaccine and diagnosed with breakthrough COVID-19 between January 2021 and March 2022 and registered in EPICOVIDEHA were included in this analysis.

A total of 1548 cases were included, mainly with lymphoid malignancies (1181 cases, 76%). After viral genome sequencing in 753 cases (49%), Omicron variant was prevalent (517, 68.7%). Most of the patients received at least two vaccine doses before COVID-19 (1419, 91%), mostly mRNA-based (1377, 89%). Overall, 906 patients (59%) received specific treatment for COVID-19. After 30-days follow-up from COVID-19 diagnosis, 143 patients (9%) died. The mortality rate in patients with Omicron variant was of 7.9%, comparable to that reported for the other variants. The 30-day mortality rate was significantly lower than in the pre-vaccine era (31%). In the univariable analysis, older age (p<0.001), active HM (p<0.001), severe and critical COVID-19 (p=0.007 and p<0.001, respectively) were associated with mortality. Conversely, patients receiving monoclonal antibodies, even for severe or critical COVID-19, had a lower mortality rate (p<0.001). In the multivariable model, older age, active disease, critical COVID-19 and at least 2-3 comorbidities were correlated with a higher mortality, whereas the administration of monoclonal antibodies, alone (p<0.001) or combined with antivirals (p=0.009), was observed protective.

While mortality is significantly lower than in the pre-vaccination era, breakthrough COVID-19 in HM is still associated with considerable mortality. Death rate was lower in patients who received monoclonal antibodies, alone or in combination with antivirals.

Introduction

Coronavirus disease 19 (COVID-19) is a life-threatening infection in patients with hematologic malignancies (HM), associated with severe clinical presentation and high risk of death. In April 2020, the European Hematology Association – Scientific Working Group Infectious in Hematology (EHA-SWG) opened the EPICOVIDEHA registry to collect all adult patients with HM that developed COVID-19. It aimed to describe the epidemiology, risk factors, and reported a mortality rate of 31.2% among 3801 patients. In December 2020, nearly one year after the first described COVID-19 case, vaccines against the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) were approved and became available first for high-risk patients, including HM. The recently published recommendations from the European Conference of Infections in Leukemia (ECIL-9) identify the critical role of mRNA-based vaccines in the fight against COVID-19 and recommend their use in HM, although they may have more limited efficacy amongst severely immunocompromised patients.

We collected data on adult HMs who developed breakthrough COVID-19 to assess the vaccine efficacy and the potential role of new emergent treatments against SARS-CoV-2. Our preliminary data, regarding the first 113 patients included, showed a significant decrease in the overall mortality rate in the post-vaccination era (12.4%), which was, however, still remarkably higher compared to the rate observed in the overall population.¹⁰ To date, few reports have been published about severity and outcomes of breakthrough COVID-19 in patients with cancer in general¹¹⁻¹² and HMs specifically,¹³ all showing high rates of severe clinical presentation, hospitalization and death among these patients. This suggests that HMs require close monitoring and increased medical attention when COVID-19 is diagnosed, regardless of previous anti-SARS-CoV-2 vaccine.

In this study, we analyzed the epidemiology and outcome of breakthrough COVID-19 in a large cohort of HMs and evaluated anti-SARS-CoV-2 treatment received by the patients.

Methods

Study design, patients, and procedures

From January 1st, 2021, until March 10th, 2022, participating institutions documented episodes of COVID-19 in their HMs that received anti-SARS-CoV-2 vaccination. Our analysis comprised data from the EPICOVIDEHA registry. EPICOVIDEHA (www.clinicaltrials.gov; National Clinical Trials identifier NCT04733729) is an international open web-based registry for patients with HMs infected with SARS-CoV-2.14 EPICOVIDEHA was approved by the local ethics committee of the Fondazione Policlinico Universitario Agostino Gemelli - IRCCS, Università Cattolica del Sacro Cuore of Rome, Italy (Study ID: 3226). When applicable, the respective local ethics committee of each participating institution have approved the project. EPICOVIDEHA methods have been described elsewhere.4,14 The electronic case report form (eCRF) is accessible online at www.clinicalsurveys.net (EFS Summer 2021, TIVIAN, Cologne, Germany). Each documented patient was reviewed and validated by infectious diseases and hematology experts from the coordination team. Inclusion criteria were: a) active HMs within the last five years before COVID-19 diagnosis, b) patients ≥18 years old, c) laboratory-based diagnosis of SARS-CoV-2 infection, and d) last vaccine dose 15 or more days before PCR confirmed SARS-CoV-2 infection. Data on baseline conditions pre-COVID-19 (i.e., age, sex, status of HM at COVID-19 diagnosis, factors predisposing for COVID-19), HM clinical management (i.e., last HM treatment strategy, vaccine type, spike protein concentration at diagnosis of COVID-19, COVID-19 diagnosis and management (i.e., reason for diagnostic test, symptoms at onset, stay during infection, treatments received for infection) and outcome (i.e., mortality, attributable mortality [assessed by the medical team in charge of the patient], last day of follow-up) were collected. Status of HM at COVID-19 onset and last follow up was defined as active (onset and refractory/resistant), stable disease or controlled (complete and partial response) based on the reports from the respective participating institution.

Study objectives

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The primary objective of this study was to assess the epidemiology and the outcome of HMs affected by breakthrough COVID-19. Secondary objectives were: 1) to estimate the relative frequency of disease severity, graded according to international standards in our patient population; 15-16 2) to evaluate the relative frequency of ICU admission among our patients; 3) to

evaluate the overall case-fatality rate; 4) to explore the impact of cancer treatment phase (induction, consolidation, maintenance, palliative, re-induction); 5) to explore the impact of vaccine doses administered to patient outcomes; 6) to explore the impact of COVID-19 treatment on patient outcomes. Moreover, data collected were compared with those reported in our previously published study performed in the pre-vaccine era by using the same registry.⁴

Sample size and statistical analysis

No *a priori* sample size calculation was performed for this analysis. Categorical variables are presented with frequencies and percentages, and continuous variables with median, interquartile range (IQR) and absolute range. Univariable Cox regression model was performed with variables suspected to play a role in the mortality of HM patients with COVID-19. Variables with a p-value ≤ 0.1 were considered for multivariable analysis. A multivariable Cox regression model was calculated with the Wald backward method. Mortality was analyzed by using Kaplan–Meier survival plots. Log-rank test was used to compare the survival probability of the patients included in the different models. A p-value ≤ 0.05 was considered statistically significant. No a priori sample size calculation was done for this exploratory study. SPSSv25.0 was employed for statistical analyses (SPSS, IBM Corp., Chicago, IL, United States). Patients with missing data in essential fields (i.e. HM, chemotherapeutic program, vaccination status, COVID-19 management or survival status) were considered as not valid and, then, excluded from the final analysis. Among the valid cases, if a value in a specific variable was missing or unknown, it is indicated as such in the descriptive analysis. Patients with missing data in a certain variable were excluded from regression analyses in case that variable was included into such analyses.

Data sharing statement

Requests for data sharing may be submitted to Livio Pagano (livio.pagano@unicatt.it)

Results

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Study population

A total of 94 centers in 26 countries, mainly from Europe, participated and registered 1583 cases. A list of enrolled cases from each participating country is available in the supplemental material (Fig. S1 and Fig. S2 panel A). Out of these 1583 cases, 35 were excluded since COVID-19 was diagnosed within 14 days from the first vaccine dose. Clinical characteristics of 1548 evaluable cases are reported in Table 1. Lymphoid malignancies were the largest subgroup, accounting 1181 cases (76.3%); the most frequently reported diagnosis was non-Hodgkin lymphoma (NHL, 549 cases). Among myeloid malignancies, the most frequent diagnosis was acute myeloid leukemia (AML, 140 cases). We found a significantly different distribution lymphoid/myeloid malignancies with that reported in pre-vaccination era (pre-vaccination lymphoid malignancies cases: 67.3% vs post-vaccination: 76.3%, p<0.001). At the time of COVID-19 diagnosis, most patients had a controlled malignancy (n=821, 53%), 322 (20.8%) a stable disease and the remaining 365 (23.6%) an active disease with 185 cases registered at HM onset. The most frequently reported last HM treatment was immuno-chemotherapy or immunotherapy alone (n=708, 42%), followed by targeted therapies (n=311, 20.1%) and conventional chemotherapy (n=234, 15.1%); 92 patients (5.9%) had received HSCT within six months before COVID-19 (allogeneic: 76; autologous: 16) and 8 had chimeric antigen receptor T cells (CAR-T) therapy. Most patients presented at least one comorbidity (60.7%) and 180 (11.6%) had a history of smoking; a complete list of comorbidities and associated clinical outcomes is available in the supplemental material (Table S1).

COVID-19 severity, variants and anti-SARS-CoV-2 spike proteins

COVID-19 was mild, severe, or critical in 39%, 32.9% and 9.8% of cases, respectively. Two-hundred eighty-three patients (18.3%) were asymptomatic and in most of them the diagnosis was made in screening programs (Table 1). We found a significantly lower rate of severe or critical cases compared to that we reported in pre-vaccination era (pre-vaccination: 2425/3801, 63.8% vs post-vaccination: 661/1545, 42.7%; p<0.001). Overall, 823 (53.2%) patients required

hospitalization and amongst them 152 (18.1%) required admission to intensive care (ICU). The hospitalization and ICU admission rate was significantly lower than reported in the pre-vaccination era (53.2% vs 73%; p<0.001 and 9.8% vs 18.1%; p<0.001, respectively). The asymptomatic cases percentage was of 18.3% (283/1548), similar to that reported in our previous publication with data from the pre-vaccine era (17.8%, 675/3801). Viral genomes were studied in 753 cases (48.6%), with the different *Omicron* variant as the most frequent viral strain (517/753, 68.7%). Most patients received two or three anti-SARS-CoV-2 vaccine doses (91%), mostly with mRNA-based technology (89%); only few patients (8.6%) received a vector-based vaccine and a minority of them an inactivated vaccine (Table 1, Fig. S2 panel B, C and D). Anti-SARS-CoV-2 spike protein IgG levels were analyzed in 244 (15.8%) fully vaccinated patients, 2-4 weeks after the last vaccine dose; among these patients, 109 (44.7%) presented an antibody response (optimal: 75, 30.7%; weak: 34, 13.9%), whereas the remaining 135 (55.3%) were non-responders. Most patients who did not have a serological response to vaccines were affected by lymphoid malignancies, as expected (126/135, 93.3%; Fig. 1).

COVID-19 treatments and risk factors for mortality

Overall, 906 patients (58.5%) received a specific treatment for COVID-19, whereas 642 (41.5%) were not treated, or received symptomatic therapies (non-steroidal anti-inflammatories, painkillers, antipyretics). Among patients who received a specific treatment for COVID-19, 311 (34.3%) were treated with monoclonal antibodies only, 246 (27.1%) with corticosteroids only, 218 (24.1%) with antivirals only, 108 (11.9%) with antiviral plus monoclonal antibodies and the remaining 23 with convalescent plasma. Details on COVID-19 treatments and outcomes are displayed in the supplemental material (Table S2). Overall day-30 mortality (i.e., from COVID-19 diagnosis) was 9.2% (143/1548 patients died); if we consider symptomatic patients only, the day-30 mortality rate was of 10.3% (130/1265 symptomatic patients died). The primary cause of death was COVID-19 in 97 patients (67.8%), a combination of both, COVID-19 and progressive HM in 39 cases (27.2%) and HM alone or combined with other reasons in the remaining 7 patients (4.8%). The mortality rate was significantly lower than that reported in pre-vaccine era (pre-vaccine 31.2%)

vs post-vaccine 9.2%; p<0.001). Looking at two of the largest patient cohorts (i.e. chronic lymphocytic leukemia, CLL and NHL) we evaluated the potential role of chemotherapeutic treatment type on mortality rate. In CLL patients, we did not observe any significant difference in terms of 30-days mortality rate among patients who had received immune-chemotherapy (13.4%), immunotherapy alone (12.5%) or new targeted therapies (16.1%). On the contrary, in NHL we did observe a slightly higher mortality rate for patients recently treated with CAR-T (20%), compared to those treated with immune-chemotherapy (8%), immunotherapy alone (14.3%) or targeted therapies (9.5%). The outcome of patients according to clinical characteristics, vaccine received and specific treatments against SARS-CoV-2 is detailed in Table 2. As shown in Fig. 2, we did not find any significant difference in terms of 30-day mortality rate among the different HM (p=0.693). in contrast to that observed in the pre-vaccination era in which we reported a higher number of fatalities in acute myeloid leukemia/myelodysplastic syndrome patients. In univariable analysis, the factors associated with a worse mortality rate were older age (p<0.001), active HM disease (p<0.001), and presence of 2-3 comorbidities (p<0.001) severe and critical COVID-19 (p=0.007) and p<0.001, respectively) (Table 3). Referring to the age, patients younger than 60 years showed a more favorable outcome (30-days mortality rate: 2.6%), compared with patients aged 60-69 years (7%), 70-79 years (14.8%) and 80 years or more (19.6%) (p<0.001). Conversely, we observed a better clinical outcome for patients who received monoclonal antibodies (with or without antivirals; Fig. 3). Analyzing the severity of COVID-19 presentation, a better clinical outcome was observed in patients treated with monoclonal antibodies alone for asymptomatic, mild, or severe disease and with monoclonal antibodies combined with antivirals in critical cases (Fig. 4). We did not find differences in terms of outcome according to the number of vaccine doses received; however, a slightly better clinical outcome was evident among patients who received three to four doses versus one to two doses (p=0.040, Table 3). We did not observe differences in survival when sorting patients according to viral strain detected (p=0.664; Fig. 5), or post-vaccine anti-spike IgG levels (Table 2).

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In the multivariable model older age, active disease, eritical COVID-19 and 2-3 comorbidities were the factors significantly correlated with a higher mortality, whereas receiving anti-SARS-CoV-

2 treatment with monoclonal antibodies alone or combined with antivirals was independently associated with a lower mortality (HR: 0.155, 95%CI: 0.077-0.313; p<0.001 - HR: 0.407, 95%CI: 0.206-0.803; p=0.010, respectively) (Table 3). Survival and severity according to vaccine doses administration and post-vaccine anti-spike IgG levels are shown in Fig. 6 and Fig. 1, respectively.

Discussion

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In the pre-vaccination era, several studies reported a high COVID-19 mortality in HM.¹⁻⁴ From December 2020, anti-SARS-CoV-2 vaccines have been administered in cancer patients, including those with HM.⁷⁻⁸ Most published studies in HMs confirmed the efficacy and safety of vaccines, particularly those using mRNA, however, most showing less efficacy in patients with lymphoid malignancies treated with immunosuppressive drugs.¹⁷⁻²²

The current study was performed in a large cohort of vaccinated HMs to evaluate epidemiology, risk factors for adverse clinical outcome and treatments of breakthrough COVID-19. We found a predominance of lymphoid malignancies, higher than observed in our previous survey during the pre-vaccine era; this difference might be explained by the lower efficacy of vaccines in this patient population, as further suggested by the high rate of serological non-responders among patients with lymphoid malignancies when evaluating anti-spike IgG levels These data are consistent with those in a recent report describing COVID-19 breakthrough infections in a large HM patient cohort, mostly consisting of patients with lymphoid malignancies. 13 Advanced age, presence of comorbidities and active HM were confirmed in the present study as factors that negatively influenced clinical outcome and survival; these were the same risk factors that had previously been reported in the pre-vaccination era. 1-4 Interestingly, in our study, the underlying malignancy did not have a significant impact on survival, which was different from our previous experience in non-vaccinated patients, where AML and myelodysplastic syndrome were associated with higher mortality risk. A potential explanation for this difference might be the better efficacy of anti-SARS-CoV-2 vaccines in myeloid malignancies. 23-25 than in lymphoid malignancies; 17-22 however, we may hypothesize new specific anti-SARS CoV-2 drugs and better COVID-19 management to be particularly important for patients with AML at risk of increased mortality if urgent chemotherapy is delayed. Similarly, as reported by other studies¹³, we did not find any significant difference in terms of mortality among different treatments received for HM. As expected, severe and critical COVID-19 had a worse clinical outcome than mild ones, showing a strong correlation with an increased mortality rate both in univariable and multivariable analysis. Given the vaccine protection, the occurrence of respiratory symptoms, hospitalization rate and

severe-critical clinical presentations were significantly lower than in the pre-vaccination era, even though still strongly higher compared to the overall population.²⁶⁻²⁹ However, it is worth underlining that about 20% of patients were asymptomatic and SARS-CoV-2 infection was detected in screening programs. Interestingly, this percentage is analogous to that reported in our published study referring to the pre-vaccination era.4 Unfortunately, it is not possible to estimate the true incidence of breakthrough infections nor the true number of asymptomatic patients with our data as only patients with COVID19 were included in the registry: we are of course aware this is a potential selection bias, hypothetically hampering the reliability of our results. To the best of our knowledge, only few studies evaluated the incidence and cumulative COVID-19 risk among vaccinated cancer patients, thus showing an increased risk in HM patients compared with the overall population. 30-32 In particular, Lee and coworkers recently published a nice population-based test-negative casecontrol study in the United Kingdom, evaluating COVID-19 breakthrough infections among a huge number of vaccinated cancer patients and healthy controls. The authors showed that the vaccine effectiveness at 3-6 months after the second dose was lower in the cancer cohort than in the control population and among cancer patients was lower in HM patients, especially those affected by leukemia and lymphoma. Very recently, an Italian study evaluated the immunogenicity and clinical efficacy of anti-SARS-CoV-2 vaccine in HM patients on 365 patients. The authors showed an overall incidence of breakthrough infections of 2.98 per 10000 person-days, significantly lower in post-vaccine seropositive patients, whereas a clear correlation between T-cellular immunity response and risk of post-vaccine infection has not been found.³³

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In our study, we reported an overall 30-day-mortality rate of 9.2%, mainly driven by COVID-19 infection as a direct or contributing factor which is significantly lower than in the pre-vaccination era. 1-4 Moreover, the 30-days mortality rate in symptomatic patients only was of 10.3%. The success of vaccination strategies is likely a major factor in the reported improvement, but not the only factor; a better COVID-19 management and the less severity of newer variants may have played a significant role as well. Previous reports suggest that COVID19 management (e.g., steroids, etc.) have also impacted outcomes. Newer variants may be less severe. Data reported in our study are coincident to other recently published reports that showed a significant mortality rate

of COVID-19 breakthrough infections amongst cancer patients¹¹⁻¹² or more specifically among those affected by HM.¹³

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In our study, we collected data about viral genotyping in about half of patients; among those, the most prevalent variant was *Omicron*, accounting for more than 2/3 of patients. These data are not surprising if we consider the large number of registered patients between late 2021 and early 2022, months in which the *Omicron* variant was rapidly spreading throughout Europe.³⁴ Interestingly, we did not find any significant difference in terms of severity of clinical presentation and mortality rate between *Omicron* and other variants, matching to other small recently published reports on HMs,³⁵⁻³⁶ but different to reports in immunocompetent patients in which *Omicron* presents with better outcome than other variants.^{34,37}

The vast majority of patients enrolled in our study received two or three vaccine doses; comparing clinical presentation and outcomes, we did not find consistent data supporting a better clinical outcome for patients who had received a higher number of vaccine doses, even though a slight difference in deaths proportion was observed comparing those who received 1-2 vs 3-4 doses. However, in multivariable analysis, the number of doses did not significantly impact on the overall 30-day-mortality. Several studies highlighted the role of a third vaccine dose as capable of restoring the immune response in serologically less responsive HM patients. 38-39 However, there are insufficient data to consider patients with low anti-spike antibody titers at high risk of worse outcomes. Indeed, in our study we did not find any differences in terms of outcomes stratifying patients according to serological response after 2-4 weeks from the last vaccine dose. By using World Health Organization international standards (BAU/mL), we did not find a significantly better survival for patients with optimal response, compared to those with weak or no response, although these data were only available in a small percentage of patients (16%). This lack of direct correlation between serological response and survival might be at least in part explained by the putative role of anti-SARS-CoV-2 induced cellular immunity, as suggested by several studies, 23-24,40 since the presence of memory T-cells might control the infection and prevent severe COVID-19 even if high titers of long-lasting neutralizing antibodies are not elicited. 41 However, since a recently published study did not find a clear correlation between post-vaccine T-cell immunity and vaccine clinical efficacy,33 further studies are warrant to better understand this aspect. Another possible explanation is related to the role of the specific anti-SARS-CoV-2 treatments (i.e. monoclonal antibodies, antivirals) that could have partially balanced the lack of protection of serological non responders. Indeed, from our survey, monoclonal antibodies with or without antivirals showed a high clinical activity irrespective of COVID-19 severity, showing the best efficacy when administered as single agents in asymptomatic mild and severe patients, and when administered in combination with antivirals in critical ones. The role of monoclonal antibodies in mitigating the negative impact of weak vaccine responses is supported by a recent randomized trial evaluating their role in immunocompetent people without serological response.⁴² Moreover, our multivariable model confirmed the positive impact on 30-day mortality risk for patients who had received monoclonal antibodies alone or combined with antivirals. We are aware that the present study has limitations due to the retrospective observational design and the possible selection bias due to the large number of participating institutions. Moreover, viral genotyping and serological data were not available for all enrolled patients and we did not know whether COVID-19 was first diagnosed in hospital or in the community, a potential key information for discriminating patient risk and infection natural history. Further prospective studies better evaluating the role of vaccine response in HM are needed.

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In conclusion, our survey has shown that vaccination and novel COVID-19 treatments have brought significant improvements in terms of mortality in HMs. To further improve the prognosis of these patients, the role of additional booster vaccine doses, and the role of prophylactic monoclonal antibodies in patients with an ineffective response to vaccination should be investigated.

Acknowledgments

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Author contribution

LP served as the principal investigator. JSG and FM served as project manager and research assistant, respectively. LP, JSG, and FM contributed to study design, study supervision, and data interpretation and wrote the paper. AB, PC, MH, PK, AP, FP, AOC and LP conceived the registry idea. LP, JSG and FM did the statistical plan, analysis and interpreted the data. All the authors recruited participants and collected and interpreted data. All authors contributed to manuscript writing and review of the manuscript. All authors agreed to be accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved.

Disclosure of conflicts of interest

526 All the authors have no disclosures to declare for this submitted paper.

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652 **Tables**

Table 1. Clinical characteristics of 1548 vaccinated HM patients who developed COVID-19

		0/
Cov	<u> </u>	%
Sex Female/male	661/887	42.7/57.3
	001/007	42.7/37.3
Age	66 (EE 7E)	[40 06]
Median (y.o.) (IQR) [range]	66 (55 - 75)	
<50/>50 y.o.	301/1247	19.5/80.5
Comorbidities	000/040	20.2/00.7
None/ 1-2-3 comorbidities	608/940	39.3/60.7
Smoking history	180	11.6
Malignancy	4404	70.0
Lymphoid malignancies	1181	76.3
Acute lymphoid leukemia	64	4.1
Chronic lymphoid leukemia	211	13.6
Hodgkin lymphoma	65	4.2
Non-Hodgkin lymphoma	549	35.5
Low grade	289	18.7
High grade	260	16.8
Multiple myeloma	275	17.8
Amyloid light-chain amyloidosis	10	0.6
Hairy cell leukemia	7	0.5
Myeloid malignancies	356	23.0
Acute myeloid leukemia	140	9.0
Chronic myeloid leukemia	44	2.8
Essential thrombocythemia	18	1.2
Myelodysplastic syndromes	93	6.0
Low-intermediate risk	69	4.5
High risk	23	1.5
Myelofibrosis	39	2.5
Polycythemia vera	16	1.0
Systemic mastocytosis	6	0.4
Aplastic anemia	11	0.7
Malignancy status before COVID-19		
Controlled disease	821	53.0
Complete remission	524	33.9
Partial remission	297	19.2
Stable disease	322	20.8
Active disease	365	23.6
Onset	185	12.0
Refractory/Resistant	180	11.6
Unknown	40	2.6
Last malignancy treatment		
alloHSCT	76	4.9
autoHSCT	16	1
CAR-T	8	0.5
Chemotherapy	-	
Conventional chemotherapy	234	15.1
Demethylating agents	80	5.2
Immunotherapy	146	5.7
Immuno-chemotherapy	562	36.3
Targeted therapy	311	20.1
Supportive measures	36	2.3
ouppointe incasules	30	۷.5

Table 1. Clinical characteristics of 1548 vaccinated HM patients who developed COVID-19

	n	%
No treatment	136	8.8
Vaccination		
One dose	129	8.3
Two doses (or J&J)	770	49.7
Three doses	639	41.3
Four doses	10	0.6
Type of vaccine		
mRNA	1377	89.0
BioNTech/Pfizer	1121	72.4
Moderna COVE	256	16.5
Vector-based	133	8.6
AstraZeneca Oxford	99	6.4
Sputnik	13	0.8
J&J – Janssen	21	1.4
Inactivated	38	2.5
CoronaVac Sinovac	21	1.4
Sinopharm	17	1.1
Spike protein dosage after vaccination (*)		
No response	135	8.7
Weak response	34	2.2
Optimal response	75	4.8
Not tested	1304	84.2
COVID-19 infection		
Wild type	40	2.6
Alpha (α)	34	2.2
Beta (β)	1	0.1
Delta (δ)	161	10.4
Omicron (o)	517	33.4
Not tested	795	51.4
Severity		
Asymptomatic	283	18.3
Mild infection	604	39.0
Severe infection	509	32.9
Critical infection	152	9.8
Symptomatology at onset		
Asymptomatic	306	19.8
Pulmonary	528	34.1
Pulmonary + extrapulmonary	400	25.8
Extrapulmonary	314	20.3
Stay during COVID-19		
Hospital	823	53.2
ICU	152	9.8
Home	800	51.7

653 **HM:** hematologic malignancy; **IQR:** interquartile range; **HSCT:** hematopoietic stem cell

transplantation; CART: chimeric antigen receptor T-cells; ICU: intensive care unit.

657 coronavirusdisease-covid-19)

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^{655 (*)} Referring to World Health Organization international standards, BAU/mL

⁽https://www.who.int/news-room/feature-stories/detail/standardization-of-vaccines-for-

Table 2. Outcome of vaccinated HM patients who developed COVID-19

	Δ	Alive		Dead	
	n	%	n		%
Outcome at 30 days					
Alive	1405	90.8			
Dead			143	9.2	
Reason for death					
COVID-19			97	67.8	
COVID-19 + HM			39	27.2	
HMs +/- other reasons			7	4.8	
Sex	50.4	00.4	70	40.0	
Female	591	89.4	70	10.6	ns
Male	814	91.8	73	8.2	
Age	46	100.0	0	0.0	
18-25 years old	46 250	100.0	0	0.0	
26-50 years old	250 585	98.0 94.2	5 36	2.0 5.8	
51-69 years old	524	94.2 83.7	102	16.3	<0.001
Over 70 years old Comorbidities	524	03.1	102	10.3	
No comorbidities	581	95.6	27	4.4	
1 comorbidity	471	91.5	44	8.5	
2 comorbidities	223	84.8	40	15.2	
3 or more comorbidities	130	80.2	32	19.8	<0.001
Smoker or ex-smokers	158	87.8	22	12.2	
Malignancies	100	07.0		12.2	
Lymphoid malignancies	1070	92.8	111	7.2	
Acute lymphoid leukemia	62	96.9	2	3.1	
Chronic lymphoid leukemia	186	88.2	25	11.8	
Hodgkin lymphoma	63	96.9	2	3.1	
Non-Hodgkin lymphoma	497	90.5	52	9.5	
Low grade	261	90.3	28	9.7	
High grade	236	90.8	24	9.2	
Multiple myeloma	246	89.5	29	10.5	
Amyloid light-chain amyloidosis	10	100.0	0	0.0	
Hairy cell leukemia	6	85.7	1	14.3	
Myeloid malignancies	324	91.0	32	9.0	
Acute myeloid leukemia	127	90.7	13	9.3	
Chronic myeloid leukemia	43	97.7	1	2.3	
Essential thrombocythemia	18	100.0	0	0.0	
Myelodysplastic syndromes	81	87.1	12	12.9	
Low-intermediate risk	63	91.3	6	8.7	
High risk	18	78.3	5	21.7	ns
Myelofibrosis	34	87.2	5	12.8	
Polycythemia vera	15	93.8	1	6.3	
Systemic mastocytosis	6	100.0	0	0.0	
Aplastic anemia	11	100.0	0	0.0	
Malignancy status Controlled disease	768	93.5	53	6.5	
Complete remission	768 505	93.5 96.4	53 19	3.6	
Partial remission	263	96.4 88.6	34	3.6 11.4	
Stable disease	203 294	91.3	28	8.7	
Active disease	307	96.3	58	3.7	
Onset	165	90.3 89.2	20	10.8	<0.001
Refractory/Resistant	142	78.9	38	21.1	~U.UU1
กษาเลบเบา y/ กษรเรเลาแ	142	10.9	30	۷۱.۱	

Table 2. Outcome of vaccinated HM patients who developed COVID-19

	Alive		De	p-value	
	n %		n	%	
Unknown	36	90.0	4	10.0	
Last malignancy treatment before COVID-19					
alloHSCT	72	94.8	4	5.2	
autoHSCT	16	100.0	0	0.0	
CAR-T	6	75.0	2	25.0	
Conventional chemotherapy	215	90.6	91.9	8.1	
Demethylating agents	73	90.5	7	9.5	
Immuno-chemotherapy	512	91.2	50	8.8	
Immunotherapy	78	87.6	11	12.3	
Targeted therapy	279	89.8	32	10.2	no
Supportive measures	28	77.8	8	22.2	ns
No treatment	126	92.6	10	7.4	
SARS-CoV-2 vaccination before COVID-19 (*)					
One dose	115	89.1	14	10.9	
Two doses	689	89.5	81	10.5	
Three doses	591	91.9	48	8.1	no
Four doses	10	100.0	0	0.0	ns
Type of SARS-CoV-2 vaccine					
mRNA	1250	90.8	127	9.2	
BioNTech/Pfizer	1011	90.2	110	9.8	
Moderna COVE	239	93.4	17	6.6	
Vector-based	123	92.5	10	7.5	
AstraZeneca Oxford	91	91.9	8	8.1	
Sputnik	13	100.0	0	0.0	
J&J - Janssen	19	90.5	2	9.5	ns
Inactivated	32	84.3	6	15.7	
CoronaVac Sinovac	18	85.7	3	14.3	
Sinopharm	14	82.4	3	17.6	
Spike protein dosage after vaccination (**)					
No response	118	87.4	17	12.6	
Weak response	31	91.2	3	8.8	ns
Optimal response	71	94.7	4	5.3	110
Not tested	1185	90.9	119	9.1	
COVID-19 variant					
Wild type	36	90.0	4	10.0	
Alpha	30	88.2	4	11.8	
Beta	1	100.0	0	0.0	
Delta	141	87.6	20	12.4	
Omicron	476	92.1	41	7.9	ns
Not tested	721	90.7	74	9.3	
COVID treatment	040	00.0	0.4	^ -	
No specific treatment reported	618	96.3	24	3.7	
Antivirals + monoclonal antibodies	98	90.7	10	9.3	
Antivirals	186	85.3	32	14.7	
Corticosteroids	185	75.2	61	24.8	40 00d
Monoclonal antibodies	302	97.1	9	2.9	<0.001
Plasma	16	69.6	7	30.4	
COVID-19 infection	070	05.5	40	4.5	
Asymptomatic	270	95.5	13	4.5	0.000
Mild infection	581 456	96.1	23	3.9	0.002
Severe infection	456	89.6	53	10.4	

Table 2. Outcome of vaccinated HM patients who developed COVID-19

	Δ	live	De	p-value	
	n	%	n		%
Critical infection	98	64.5	54	35.5	
COVID-19 symptoms					
Pulmonary	473	89.6	55	10.4	
Pulmonary + extrapulmonary	349	87.3	51	12.8	0.002
Extrapulmonary	297	94.6	17	5.4	
Asymptomatic	286	93.5	20	6.5	

- 658 **HM:** hematologic malignancy; **IQR:** interquartile range; **HSCT:** hematopoietic stem cell
- transplantation; CART: chimeric antigen receptor T-cells; ICU: intensive care unit; ns: not
- 660 statistically significant.
- 661 (*) 1-2 doses vs 3-4 doses p-value: 0.040
- 662 (**) Referring to World Health Organization international standards, BAU/mL
- (https://www.who.int/news-room/feature-stories/detail/standardization-of-vaccines-for-
- 664 coronavirusdisease-covid-19).

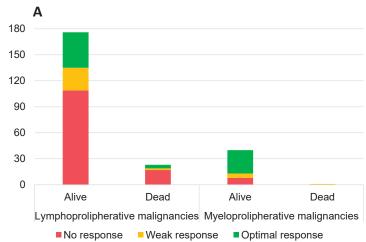
Table 3. Univariable and multivariable analysis of factors influencing mortality at 30 days

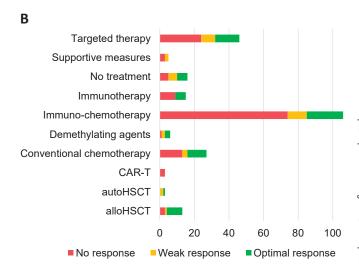
	Univariable				Multivariable				
	p HR 95 CI			р					
	value	1111	Lower	Upper	value	1111	Lower	Upper	
Sex									
Female	-	-	-	-					
Male	0.148	0.785	0.566	1.090					
Age	<0.001	1.059	1.044	1.075	<0.001	1.042	1.024	1.061	
Malignancy status at COVID-19 diagnosis									
Controlled disease	-	-	-	-	-	-	-	-	
Stable disease	0.183	1.364	0.863	2.157	0.767		0.647	1.806	
Active disease	<0.001	2.494	1.718	3.619	0.001	1.981	1.305	3.008	
Baseline malignancy									
Aplastic anemia	-	-	-	-					
Lymphoid malignancies		3032.714							
Myeloid malignancies	0.876	2974.523	0.000						
Comorbidities									
0-1 comorbidities	-	-	-	-	-	-	-	-	
≥ 2 comorbidities	< 0.001	2.802	2.019	3.889	0.027	1.503	1.050	2.229	
Type of last vaccination									
mRNA	-	-	-	-					
Vector-based	0.359	0.740	0.389	1.409					
Inactivated	0.122	1.907	0.841	4.326					
SARS-CoV-2									
Omicron	-	-	-	-					
Alpha	0.800	1.142	0.409	3.190					
Beta	0.960	0.000	0.000						
Delta	0.210	1.408	0.825	2.403					
Wild type	0.758	1.175	0.421	3.281					
Not tested	0.399	1.179	0.805	1.726					
Vaccine doses before COVID-19									
One dose	-	-	-	-					
Two doses	0.870	1.049	0.595	1.849					
Three or more doses	0.637	0.866	0.478	1.572					
Serological response before COVID-19									
No response	-	-	-	-					
Weak response	0.632	0.740	0.217	2.529					
Optimal response	0.124	0.425	0.143	1.264					
COVID-19 treatment									
Corticosteroids	-	-	-	-	-	-	-	-	
Antivirals + monoclonal antibodies	0.001	0.333	0.171	0.651	0.010	0.407	0.206	0.803	
Antivirals	0.010	0.570	0.372	0.874	0.099	0.680	0.431	1.075	
Monoclonal antibodies	<0.001	0.123	0.061	0.247	<0.001	0.155	0.077	0.313	
Plasma	0.852	1.077	0.493	2.355	0.243	1.605	0.726	3.549	

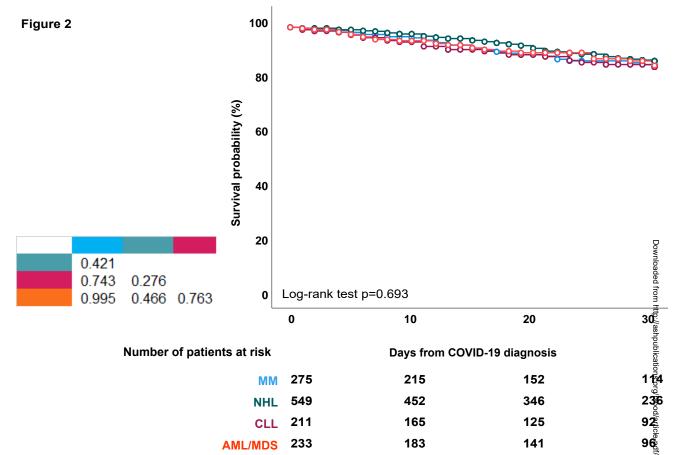
665 Figure legends

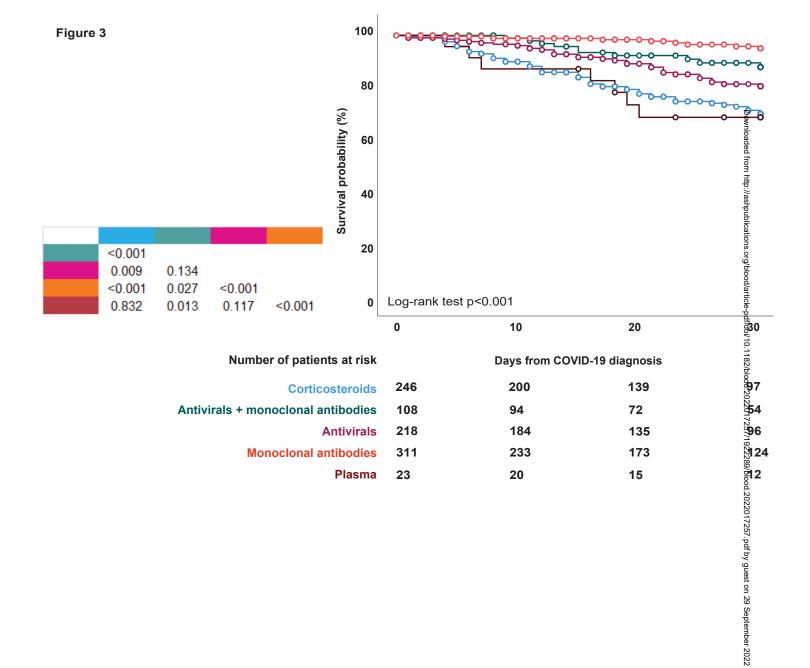
- 666 Figure 1. Patient distribution by serological response after last COVID-19 vaccination before
- 667 COVID-19. Panel A) By baseline malignancy; Panel B) By last treatment for hematological
- 668 malignancy immediately before COVID-19
- 669 *Figure 2.* Survival probability by most prevalent underlying condition.
- 670 Figure 3. Survival probability of patients by COVID-19 treatment.
- 671 Figure 4. Survival probability by COVID-19 treatment and COVID-19 severity. Panel A)
- 672 Asymptomatic patients; Panel B) Mild patients; Panel C) Severe patients; Panel D) Critical
- 673 patients.
- 674 *Figure 5.* Survival probability by SARS-CoV-2 variant.
- 675 Figure 6. Patient distribution by number of doses administered before COVID-19 and COVID-19
- 676 severity.

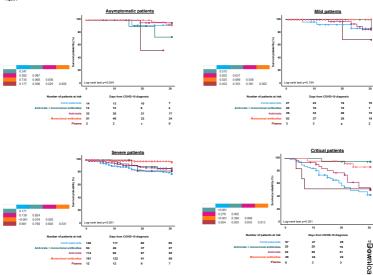
Figure 1











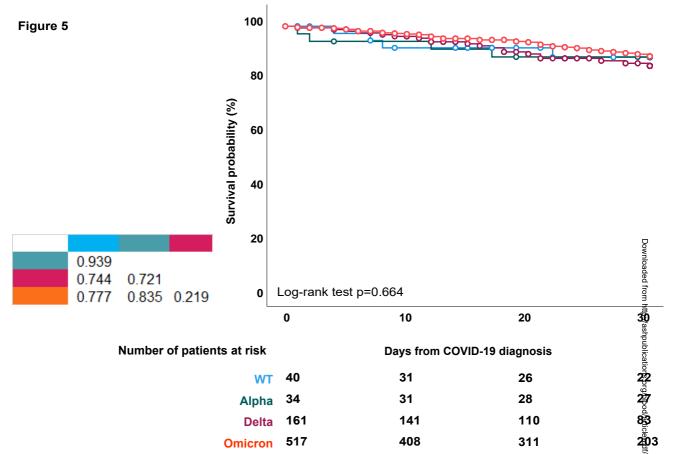


Figure 6

