EFI
GENEVA
2024

37TH EUROPEAN IMMUNOGENETICS & HISTOCOMPATIBILITY CONFERENCE

MAY 20–23, 2024
GENEVA
SWITZERLAND

Unveiling Diversity, Nurturing Transplant Bonds, Orchestrating immunity

FINAL PROGRAM

CONTENT

WELCOME ADDRESS

Dear scientists, Dear EFI members,

It is with great pleasure and excitement that we welcome each one of you to the 37th Conference of the European Federation for Immunogenetics (EFI) in Geneva.

As we come together for this significant gathering, it is a testament to our shared commitment to advancing the field of Immunogenetics. Over the four days of the conference, we will have the privilege of engaging with cutting-edge research, insightful discussions, and networking opportunities. The scientific program is designed to cover a spectrum of topics, from the latest technological advancements to the ethical considerations that underscore our work in Immunogenetics. Our distinguished lineup of speakers, experts, and panelists will share their expertise, providing invaluable insights that will inspire and challenge our thinking.

Don't miss the final state-of-the-art lecture to navigate through our universe and beyond by learning more on exoplanets, a big field of expertise at the University of Geneva.

A conference of this size is not possible to arrange without the support from a number of faithful sponsors, who have prepared very interesting symposia. We sincerely thank all of them for their financial and scientific contributions.

We also hope that you will take time to visit and enjoy the international and peaceful atmosphere of Geneva, the smallest of the biggest capitals, which hosts international organisations, such as the European headquarters of the UN, the Red Cross, the WHO and a dozen of non-governmental organisations.

Hereby, we would like to express our gratitude to the EFI Executive Committee, the EFI Scientific Committee, the EFI Education Committee, to the members of the Organizing Committee as well as our PCO. They have all provided us with invaluable help and support in preparing the scientific program, organizing teaching sessions, evaluating abstracts and preparing the Conference.

Thank you for being an integral part of EFI 2024 in Geneva.

Here's to an inspiring and transformative conference!

From the local organizing committee
Sylvie Ferrari-Lacraz and Jean Villard

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WE WOULD LIKE TO THANK THE FOLLOWING PARTNERS FOR THEIR SUPPORT

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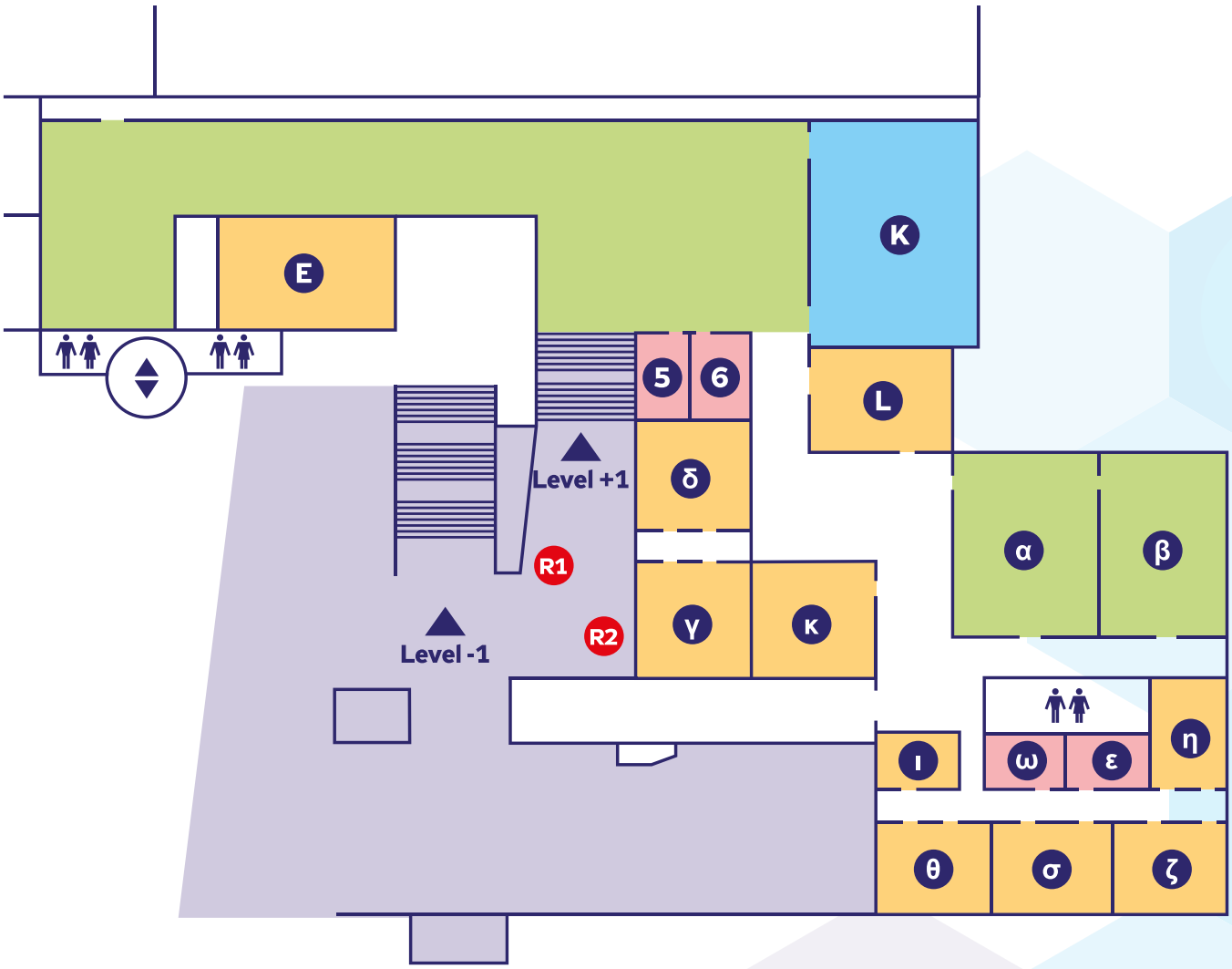
CONFERENCE INFORMATION

CONFERENCE VENUE
Palexpo Geneva
www.palexpo.ch

First floor



K
Conference hall

α β
Poster Session – Alpha, Beta

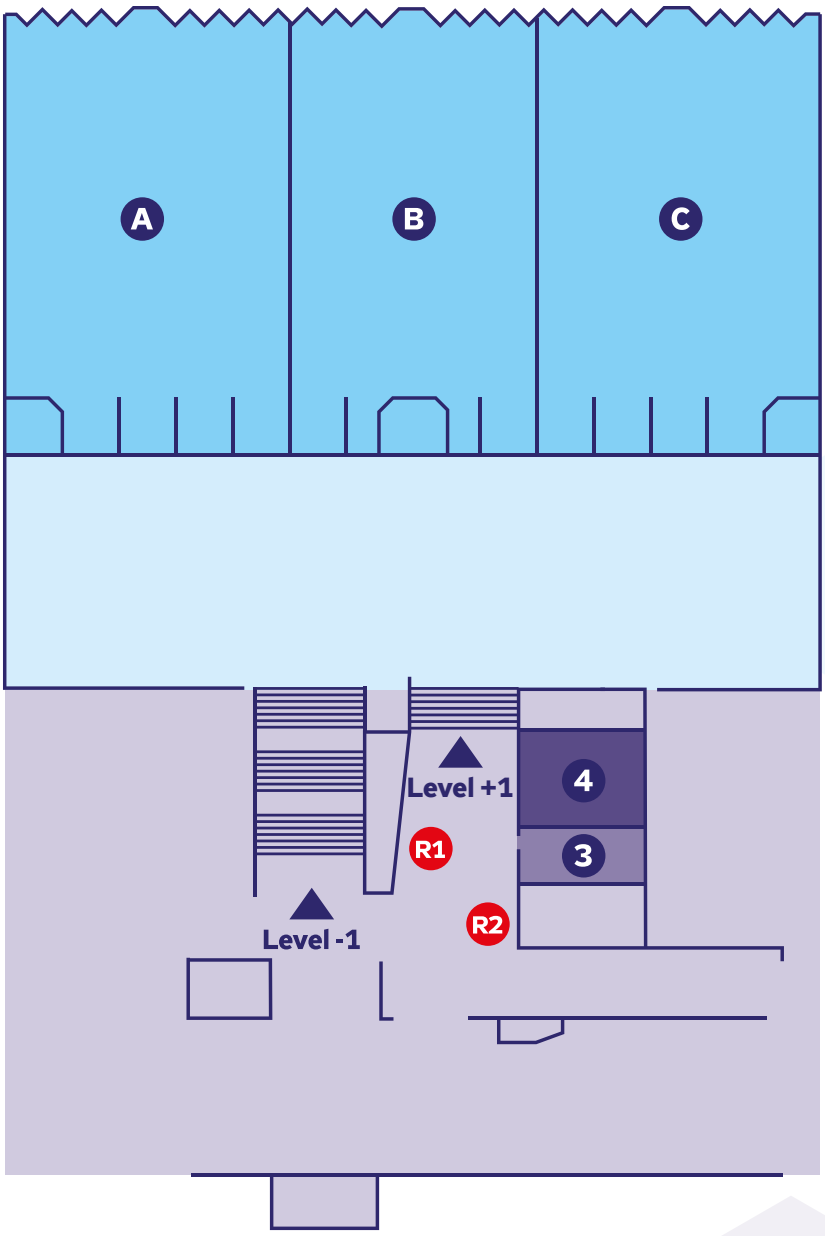
ε ω 5 6
Company private lounges – Epsilon, Omega, 5, 6

γ δ ζ η θ ι κ σ E L
Conference meeting rooms – Gamma, Delta, Zeta, Eta, Theta, Iota, Kappa, Sigma, E, L

R1
Self check-in kiosks

R2
On-site registration, bag delivery

Ground floor



A B C
Conference halls

3
Speakers´ Ready Room

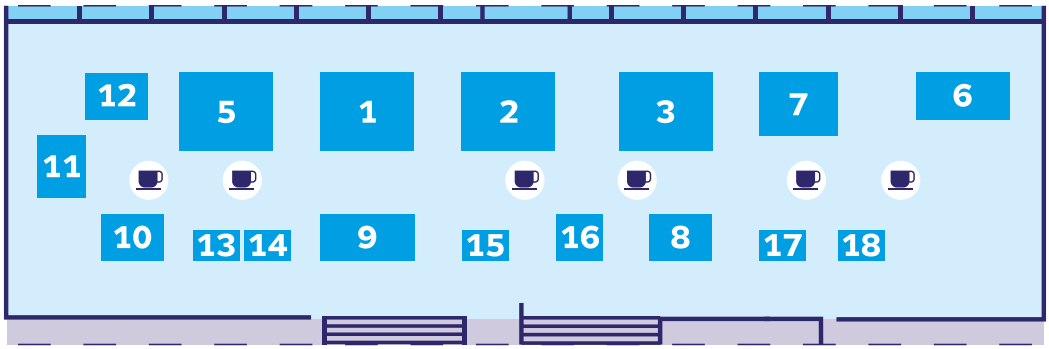
4
Room 4

Exhibition area

R1
Self check-in kiosks

R2
On-site registration, bag delivery

EXHIBITION PLAN



List of Exhibitors

1 GenDx
2 CareDx, Inc.
3 Thermo Fisher Scientific
5 Werfen
6 DiagnoSeq
7 Hansa Biopharma
8 Devyser
9 Omixon
10 DKMS Life Science Lab gGmbH
11 BAG Diagnostics GmbH
12 HistoGenetics
13 Pirche AG
14 Protrans medizinische diagnostische Produkte GmbH
15 EFI Office
16 inno-train Diagnostik GmbH
17 JETA Molecular
18 STEMCELL Technologies

FREE WI-FI

SSID: EFI2024
Password: Geneva2024

EFI 2024 MOBILE APPLICATION

With the mobile application, you will have access to the detailed program and all the important information related to the conference.

Download the mobile app in the App Store / Google Play.

EFI SOCIAL MEDIA

Follow EFI on Social Media

https://www.facebook.com/EFI2024
https://twitter.com/i/flow/login?redirect_after_login=%2FConferenceEfi
https://www.linkedin.com/company/efi-conference-2024/

As for EFI conference 2024 use #EFI2024.

REGISTRATION

SELF-CHECK-IN

On-site registration and issue of badges will take place at self-check-in stations. You will receive a QR code before the conference via email. You can print the QR code or use your mobile device to obtain your badge.

On-site Registration Fee

Member	760 EUR
Non-member	860 EUR
Technician	490 EUR
Student	490 EUR
Retired	490 EUR
One-day fee (May 21/22)	530 EUR
One-day fee (May 23)	430 EUR
Distributor	270 EUR
Accompanying Person	160 EUR

Registration fee includes 8,1 % VAT.

The Registration Fee Includes:

- Access to the conference and all its sessions (not included for Accompanying Persons)
- Welcome Cocktail
- Opening ceremony
- Coffee and lunch breaks
- Wine and cheese poster session

The distributor fee allows for access to the exhibition only. Access to the sessions need a full conference ticket.

The One-day Registration Fee Includes:

- Access to the conference and all sessions taking place on the selected day
- Coffee and lunch breaks

The Accompanying person's Registration Fee includes:

- Access to the building and exhibition
- Welcome Cocktail
- Wine and cheese poster session
- Accompanying persons do not have access to scientific sessions

REGISTRATION DESK

Opening Hours

Monday, May 20, 2024	08:00–20:00
Tuesday, May 21, 2024	07:30–18:30
Wednesday, May 22, 2024	08:00–18:30
Thursday, May 23, 2024	08:00–14:00

NAME BADGE

All delegates will receive a name badge upon registration. Everyone is kindly requested to wear his/her badge when attending the conference.

DELEGATE
SPEAKER
ONE-DAY FEE (MAY 21,22)
ONE-DAY FEE (MAY 23)
ORGANIZER
PARTNER / EXHIBITOR
ACCOMPANYING PERSON

CERTIFICATE OF ATTENDANCE

All delegates will receive a certificate of attendance after the conference by email.

INSTRUCTIONS FOR SPEAKERS (ORAL)

Please prepare your presentation in 16:9 format (screen is in format ratio 16:9).

We recommend to save your PowerPoint presentation using PPT(X) format.

Each presenter will be given a time slot of 10 minutes in total. Each oral presentation should not exceed 8 minutes. It will be followed by 2 minutes discussion.

All speakers are requested to strictly keep their allocated time slots. Session chairs will enforce the schedule.

SPEAKERS' READY ROOM

All session halls are equipped with standard PowerPoint presentation facilities.

All presentations will be networked to the appropriate room "Speakers' Ready Room" (Office 6, located on the first floor of the venue).

All speakers are asked to submit their presentations to the Speakers' Ready Room at least 1 hour prior to the session you present in.

Early morning presentation:

Please submit your presentation the day before.

We kindly ask you to hand in your presentation to the technical staff in the Speakers' Ready Room on-site via an external drive. While doing so, we encourage speakers to verify their presentation. This will ensure no formatting errors.

IT support will be available in all the conference rooms during live sessions. All speakers are kindly requested to use provided PC onsite.

Please be present in the session room 15 minutes before the start of your session and follow the instructions from the Chairs and/or technician.

During your presentation, a remote control will be available for controlling your presentation. Please observe the time allotted to you as you will not be permitted to speak for a longer time.

At the end of the conference, all presentations will be deleted from the presentation system and computers on-site.

Opening hours of the Speakers' Ready Room

Monday, May 20, 2024	15:30–20:00
Tuesday, May 21, 2024	07:30–18:00
Wednesday, May 22, 2024	08:00–18:30
Thursday, May 23, 2024	07:30–12:30

INSTRUCTIONS FOR POSTER PRESENTERS

The poster section is located in the foyer of the 1st floor and in the Alpha + Beta rooms.

All posters will be presented in a "paper" format (printed posters) and will be organized according to the paper ID numbers.

- Posters must be prepared in PORTRAIT style (**please DO NOT prepare a landscape poster**). The recommended dimensions (for a good legibility) are **84.1 × 118.9 cm wide**. (format A0). Do NOT exceed the overall dimensions of the poster board (180 cm high by 95 cm wide).
- Posters will be displayed throughout the whole conference.
- Posters should be attached to the boards (smooth surface) with stickers, which will be supplied by the organisers.
- Fixing material (adhesive gum, standard tape) will be available in the Posters Area
- The poster boards will be numbered by the organizers.

The number of your poster can be found in the List of posters. The presentation number assigned to your poster should not be placed on your poster.

POSTER CREATION

The header of the Conference can be downloaded here (JPG format, PDF format), and the instructions on layout, font style, and margins are in the attached preview. The preview is informative only. The text, illustrations, etc. should be big enough to be read from a distance of 1.5 meters, and the message of the poster should be clear and understandable.

Please print your poster and bring it with you to the Conference. **No** poster printing service is provided by the organizers. Please **DO NOT** prepare a landscape poster.

A SPECIAL SESSION

Poster Session viewing is scheduled on Tuesday, May 21 from 18:00–20:00

Be at your poster during this poster session to discuss your work with scientists visiting your poster. A Selection of posters will be assessed for best posters awards. The best posters awards will be announced during the closing ceremony on Thursday, May 23, from 12:00.

Poster mounting time:

Monday, May 20, 2024	16:00–19:30
Tuesday, May 21, 2024	07:30–10:00

Poster removal time:

Wednesday, May 22, 2024	17:00–19:00

We ask the presenters to remove their posters during the removal time, otherwise they will be removed and discarded by the organizer.

ORGANISERS AND COMMITTEES

LOCAL ORGANIZING COMMITTEE

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Valérie Olivier, *Switzerland*
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SPEAKERS

Plenary 1: Solid Organ Transplantation (How to Transplant Highly Sensitized Patients)
Lori West, *Alberta Transplant Institute, Canada*
Mats Bengtsson, *Uppsala University Hospital, Sweden*
Pernille Bundgaard Koefoed-Nielsen, *Aarhus University Hospital Denmark*

Plenary 2: Hematopoietic stem cell transplantation (New therapies for acute leukemia: HSCT, CAR-T, Bispec-Ab)
Jakob Passweg, *USB, Basel*
Catherine Thieblemont, *APHP, Saint Louis*
Federico Simonetta, *HUG, Geneva*

Plenary 3: HLA Immunogenetics and Population Genetics a combined SIP-EFI session
Alicia Sanchez-Mazas, *UNIGE*
Diogo Meyer, *Sao Paulo, Brazil*
Paul Norman, *CU Anschutz, Denver, Colorado*

Plenary 4: Giving life-bone marrow / solid organ donors
Johannes Schetelig, *Dresden UniversityKlinik and DKMS*
Constanca Figueiredo, *Medizine Ochshule Hannover*
Efstratios (Stratos) Chatzixiros, *World Health Organization (Switzerland)*

Plenary 5: Antigen presentation and neoantigens
Giacomo Oliveira, *Harvard, Boston*
Andrea Sotto, *Human Technopole, Milan*
Michal Bassani-Steinberg, *CHUV*

SFHI session
Thomas Fehr, *Kantonspital Chur*
Fadi Issa, *Oxford, UK*
Tifanie Blein, *Institut Imagine, Paris*

Network immune cell session
Mikaël Pittet, *UNIGE*
Camilla Jandus, *UNIGE*
David Gfeller, *UNIL*

DP special session
Katharina Fleischhauer, *Essen, Germany*
Julien Racle, *UNIL*
Effie Petersdorf, *Fredhutchinson, Seattle*
Stéphane Buhler, *HUG, Geneva, Switzerland*

Teaching KIR
Paul Norman, *CU Anschutz, Denver, Colorado*
Alice Koenig, *CHU Lyon, France*
Stéphane Buhler, *HUG, Geneva, Switzerland*

Teaching transfusion
Deborah Sage, *NHS, London, UK*
Anthony Poles, *NHS, Bristol, UK*
Agnes Basire, *EFS, Marseille*

Teaching HLA and disease
Gwendaline Guidicelli, *CHU Bordeau, France*
Frantisek Mrazek, *Olomuc University Hospital, Czech Republic*
Deborah Pritchard, *Wales, UK*

Young professionals session
Sandra Tafulo, *Portuguese Institute for Blood and Transplantation, Porto, Portugal*
Nicolas Vince, *Nantes University, France*
Lovis Christian, *HUG, Switzerland*

Closing ceremony
Monika Lendl, *UNIGE*

PROGRAM AT GLANCE

TIME	MONDAY, MAY 20, 2024
	Hall B
07:30	Registration
17:30	17:30–19:15 Opening Ceremony

TIME	TUESDAY, MAY 21, 2024			
	Hall A	Hall B	Hall C	Hall K
07:30	Registration			
08:30		08:30–10:00 **Plenary session I** Solid Organ Transplantation (How to Transplant Highly Sensitized Patients)		
10:00	Coffee Break			
10:30	10:30–12:00 **Abstract session 1** Immunotherapy, gene therapy & NK cells	10:30–12:00 Special joint SFHI session – Tolerance	10:30–12:00 **Abstract session 2** Bioinformatics, data analysis in immunogenetics	10:30–12:00 **Teaching session 1** HLA and transfusion
12:00	Lunch			
12:10	12:10–12:50 **Industry symposium** Werfen			
13:10		13:10–13:50 **Industry symposium** CareDx		
14:00	14:00–15:30 **Abstract session 3** New technologies in Immunogenetics	14:00–15:30 Special joint session with the Geneva University – The network of immune cells in health and disease	14:00–15:30 **Abstract session 4** Immunogenetics in organ transplantation 1	14:00–15:30 **Teaching session 2** HLA and diseas
15:30	Coffee Break			
16:00		16:00–17:30 **Plenary session II** Hematopoietic stem cell transplantation (New therapies for acute leukemia: HSCT, CAR-T, Bispec-Ab)		
17:30	17:30–18:10 **Industry symposium** Omixon	17:30–18:00 Serological nomenclature: update	17:30–18:10 **Industry symposium** Hansa Biopharma	
18:10	18:10–20:00 Wine and Cheese Poster viewing Session – Poster area on the 1st Floor & Alpha + Beta rooms			

TIME	WEDNESDAY, MAY 22, 2024			
	Hall A	Hall B	Hall C	Hall K
08:00	Registration			
08:30		**08:30–10:00** **Plenary session III** HLA Immunogenetics and Population Genetics a combined SIP-EFI session		
10:00	Coffee Break			
10:30	**10:30–12:00** **Abstract session 5** Immunogenetics in organ transplantation 2	**10:30–12:00** Special session – HLA DP models and beyond	**10:30–12:00** **Abstract session 6** MHC evolution, population genetics	
12:00	Lunch			
12:10	**12:10–12:50** **Industry symposium** Thermo Fisher Scientific			
13:10	**13:10–13:50** **Industry symposium** GenDx			
14:00	**14:00–15:30** **Abstract session 7** Hematopoietic stemcell transplantation	**14:00–15:30** Young EFI professional special session – AI and big data in immunogenetics	**14:00–15:30** **Abstract session 8** Autoimmunity, Infection, Reproduction & Cancer	**14:00–15:30** **Teaching session 3** KIR in solid organ and hematopoietic stem cell transplantation
15:30	Coffee Break			
16:00		**16:00–17:30** **Plenary session IV** Giving life- bone marrow / solid organ donor		
17:30		**17:30–18:00** EFI Medals Ceremony		
18:00		**18:00–19:00** EFI General Assembly		

TIME	THURSDAY, MAY 23, 2024
	Hall B
07:30	Registration
08:30	**08:30–10:00** Best abstract session
10:00	Coffee break
10:30	**10:30–12:00** **Plenary session V** Antigen presentation and neoantigens
12:00	**12:00–13:00** **Closing Ceremony**

DETAILED PROGRAM

ASSOCIATED MEETINGS

SUNDAY, MAY 19, 2024

Room α + β (Alpha + Beta)

09:00–17:00 **Inspectors workshop**
Chairpersons *Blanka Vidan-Jeras, Christien Voorter, Sabine Scherer*

Rooms ι (Iota), ω (Omega), Office 5, Office 6

09:00–17:00 **ESHI diploma examination**

MONDAY, MAY 20, 2024

Rooms θ (Theta), ω (Omega), ι (Iota), σ (Sigma), ζ (Zeta), ε (Epsilon), η (Eta)

08:30–16:30	**Executive committee meeting**	**θ (Theta)**
	External proficiency testing committee meeting	**ω (Omega)**
	IT & bioinformatics committee meeting	**ι (Iota)**
	Accreditation committee meeting	**σ (Sigma)**
	Standards committee meeting	**ζ (Zeta)**
	Education committee meeting	**ε (Epsilon)**
	Scientific committee meeting	**η (Eta)**
Chairpersons	*Chairs of each committee*	

SCIENTIFIC PROGRAM

MONDAY, MAY 20, 2024

Room γ (Gamma)

12:00–13:30 **Population Genetics Working Group (Open Meeting)**

Chairpersons *A. Sanchez-Mazas, José Nunes*

Hall B

17:30–19:15 **Opening Ceremony**

17:30–18:00 **Welcome addresses**
EFI President, Dean of the Faculty, Health Minister

18:00–18:15 **HLA award**
Steven Marsh

18:15–18:45 **Julia Bodmer award**
Luca Vago

18:45–19:15 **Ceppellini lecture**
Ann-Margaret Little

Exhibition Hall

19:30–21:30 **Welcome reception**

TUESDAY, MAY 21, 2024

Hall B

08:30–10:00 **Plenary session I**
Solid Organ Transplantation (How to transplant highly sensitized patients)
Chairpersons *Valérie Dubois, Sebastiaan Heidt*

08:30–09:00 **New approach to ABO – Incompatible transplantation**
Lori West
09:00–09:30 **Is Imlifidase the new magic bullet for highly sensitized recipients?**
Mats Bengtsson
09:30–10:00 **10 years of the Scandiatransplant acceptable mismatch program**
Pernille Bundgaard Koefoed-Nielsen

PARALLEL SESSIONS

Hall B

10:30–12:00 **Special joint SFHI session**
Tolerance
Chairpersons *Paul Rouzaire, Alice Aarnink*

10:30–11:00 **Mixed chimerism for induction of renal allograft tolerance**
Thomas Fehr
11:00–11:30 **Development and assessment of T cell therapies for transplantation**
Fadi Issa
11:30–12:00 **Genetic engineering to empower Treg therapy**
Tifanie Blein

Room K

10:30–12:00 **Teaching session 1**
HLA and transfusion
Chairpersons *Deborah Sage, Agnès Basire*

10:30–11:00 **Human platelet antigens - Definition and laboratory investigation**
Anthony Poles
11:00–11:30 **Fetal and neonatal alloimmune thrombocytopenia diagnosis and management: a commissioner point of view**
Agnes Basire
11:30–12:00 **Platelet refractoriness: laboratory diagnosis and patient management**
Deborah Sage

Hall A

10:30–12:00 **Abstract session 1: Immunotherapy, gene therapy & NK cells**
Chairpersons *Britta Eiz-Vesper, Roberto Crocchiolo*

Hall C

10:30–12:00 **Abstract session 2: Bioinformatics, data analysis in immunogenetics**
Chairpersons *Martin Maiers, James Robinson*

Room θ (Theta)

12:00–13:00 **EC + ECC meeting**

Hall A

12:10–12:50 **Industry symposium: Werfen**

Hall B

13:10–13:50 **Industry symposium: CareDx**

Room σ (Sigma)

13:30–14:30 **CME/CPD Launch meeting**

PARALLEL SESSIONS

Hall B

14:00–15:30 **Special joint session with the Geneva University**
The network of immune cells in health and disease
Chairpersons *Nicole Mifsud, Frans Claas*

14:00–14:30 **The neutrophil and macrophage networks**
Mikaël Pittet
14:30–15:00 **The ILC network**
Camilla Jandus
15:00–15:30 **The T/TCR network**
David Gfeller

Room K

14:00–15:30 **Teaching session 2**
HLA and disease
Chairpersons *Gwendaline Guidicelli, Deborah Pritchard*

14:00–14:30 **Best practices for HLA and Genetic Testing in H&I laboratories: SFHI Guidelines**
Gwendaline Guidicelli
14:30–15:00 **Current position of disease associations with the HLA system in clinical diagnostics**
Frantisek Mrazek
15:00–15:30 **HLA genetics and Coeliac disease**
Deborah Pritchard

Hall A

14:00–15:30 **Abstract session 3: New technologies in Immunogenetics**
Chairpersons *Eric Spierings, Jakob Nilsson*

Hall C

14:00–15:30 **Abstract session 4: Immunogenetics in organ transplantation 1**
Chairpersons *Caroline Wehmeier, Claudia Lehmann*

Hall B

16:00–17:30 **Plenary session II**
Hematopoietic stem cell transplantation
(New therapies for acute leukemia: HSCT, CAR-T, Bispec-Ab)

Chairpersons *Kay Poulton, Pietro Crivello*

16:00–16:30 **HSCT**
Jakob Passweg

16:30–17:00 **B-specific antibody therapy**
Catherine Thieblemont

17:00–17:30 **CAR-T cells**
Federico Simonetta

Hall B

17:30–18:00 **Serological nomenclature: update**

Chairperson *Marcelo Fernández-Viña*

Hall C

17:30–18:10 **Industry symposium: Hansa Biopharma**

Hall A

17:30–18:10 **Industry symposium: Omixon**

Poster Hall

18:10–20:00 **Poster viewing session**

WEDNESDAY, MAY 22, 2024

Hall B

08:30–10:00 **Plenary session III**
HLA immunogenetics and population genetics a combined SIP-EFI session
Chairpersons *Neema Mayor, Alicia Sanchez-Mazas*

08:30–09:00 **Using genomic data to understand the nature and timescale of selection on HLA genes**
Diogo Meyer
09:00–09:30 **Archaic introgression enhanced recognition of HLA-A by first nations Oceanian Natural Killer cells**
Paul Norman
09:30–10:00 **Unveiling HLA population diversity: evolutionary meanings and clinical implications**
Alicia Sanchez-Mazas

PARALLEL SESSIONS

Hall B

10:30–12:00 **Special session**
HLA DP models and beyond
Chairpersons *Jill Hollenbach, Pierre-Antoine Gourraud*

10:30–11:00 **The importance of Immunopeptidomics data to analyze the immune response**
Julien Racle
11:00–11:30 **The immunopeptidome in HCT – lessons learned from HLA-DP**
Katharina Fleischhauer
11:30–12:00 **MHC class II haplotypes**
Effie Petersdorf

Hall A

10:30–12:00 **Abstract session 5: Immunogenetics in organ transplantation 2**
Chairpersons *Valérie Olivier, Jean-Luc Taupin*

Hall C

10:30–12:00 **Abstract session 6: MHC evolution, population genetics**
Chairpersons *Natasja de Groot, Steven Mack*

Room 4

12:00–13:15 **Presidents meeting**

Hall A

12:10–12:50 **Industry symposium: Thermo Fisher Scientific**
13:10–13:50 **Industry symposium: GenDx**

PARALLEL SESSIONS

Hall B

14:00–15:30 **Young EFI professional special session**
AI and big data in immunogenetics
Chairpersons *Timo Olieslagers, Arianne Brandsma*

14:00–14:30 **AI and Big data: friend or foe?**
Christian Lovis
14:30–15:00 **Endeavour in HLA research, an exciting winding path**
Nicolas Vince
15:00–15:30 **Scientific career path; inspiration, collaboration and commitment**
Sandra Tafulo

Room K

14:00–15:30 **Teaching session 3**
KIR in solid organ and hematopoietic stem cell transplantation
Chairpersons *Paul Norman, Stéphane Buhler*

14:00–14:30 **The KIR system**
Paul Norman
14:30–15:00 **KIRs in solid organ transplantation**
Alice Koenig
15:00–15:30 **KIRs in hematopoietic stem cell transplantation**
Stéphane Buhler

Hall A

14:00–15:30 **Abstract session 7: Hematopoietic stem-cell transplantation (HSCT)**
Chairpersons *Nina Svetlitzky, Katarzyna Bogunia-Kubik*

Hall C

14:00–15:30 **Abstract session 8: Autoimmunity, Infection, Reproduction & Cancer**
Chairpersons *Danillo Augusto, Fatma Oğuz*

Hall B

16:00–17:30 **Plenary session IV**
Giving life- bone marrow / solid organ donors
Chairpersons *Joannis Mytilineos, TBA*

16:00–16:30 **The DKMS studies**
Johannes Schetelig
16:30–17:00 **One organ for every recipient**
Constanca Figueiredo
17:00–17:30 **The view and strategy of WHO**
Efstratios (Stratos) Chatzixiros

Hall B

17:30–18:00 **EFI medals ceremony**
Chairperson *Ann-Margaret Little*

Hall B

18:00–19:00 **EFI General Assembly**

THURSDAY, MAY 23, 2024

Hall B

08:30–10:00 **Best abstract session**
Chairpersons *Marco Andreani, Luca Vago*

10:30–12:00 **Plenary session V**
Antigen presentation and neoantigens
Chairpersons *Lotte Wieten, Esteban Arrieta-Bolaños*

10:30–11:00 **Deciphering the properties of antitumor T cells in solid cancers**
Giacomo Oliveira

11:00–11:30 **Lymphatic vessels as immunomodulators in cancer**
Stéphanie Hugues

11:30–12:00 **Antigen discovery for the development of personalized cancer immunotherapy**
Michal Bassani-Sternberg

12:00–12:30 **Closing Ceremony**
Chairpersons *Marco Andreani, Sylvie Ferrari-Lacraz*

12:00–12:15 **19th IHIW presentation**

12:15–12:30 **Jon Van Rood award & best abstracts award**
Committee chair: Steven Marsh
Best poster awards
Committee chair: Stefan Schaub

12:30–13:00 **Lecture**
The world of exoplanets
Monika Lendl

Remarks
Welcome to Prague and final conclusions

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AUTOIMMUNITY, INFECTION, REPRODUCTION & CANCER

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Susan Ott[1], Ngoc Giang Le[1], Nanine de Groot[1], Marit van der Wiel[1], Jesse Mittertreiner[1], Natasja de Groot[1], Jesse Bruijnesteijn[1] and Ronald E. Bontrop[1]
[1]Biomedical Primate Research Centre, Netherlands

P2 | HLA allele frequencies and susceptibility to Bullous Pemphigoid in a group of 116 Italian patients

Marco Andreani[1], Feliciana Mariotti[2], Anna Pira[2], Giuseppe Testa[1], Priscilla Caputi[2], Mariarosa Battarra[1], Tiziana Galluccio[3], Franco Locatelli[4] and Giovanni Di Zenzo[2]
[1]Laboratorio di Immunogenetica dei Trapianti, Ospedale Pediatrico Bambino Gesù, Roma, Italy; ; [2]Laboratorio di Biologia Molecolare e Cellulare, Istituto Dermopatico dell'Immacolata (IDI)-IRCCS, Roma, Italy; [3]Laboratorio di Immunogenetica dei Trapianti, Ospedale Pediatrico Bambino Gesù, Roma, Italy; [4]Dipartimento Onco-Ematologia e Terapia Cellulare e Genica, Ospedale Pediatrico Bambino Gesù, Roma, Italy

P3 | Genetic Variants associated with Narcolepsy

Steven Jervis[1], Kay Poulton[1], Antony Payton[2] and Arpana Verma[2]
[1]Manchester Transplantation Laboratory, United Kingdom; [2]University of Manchester, United Kingdom

P4 | NKG2D receptor and MICA and MICB ligand polymorphism in CALR mutation – driven Myeloproliferative Neoplasms

Milena Ivanova[1], Angelina Mladenova[1], Gergana Tsvetkova[2], Evgueniy Hadjiev[2] and Velizar Shivarov[3]
[1]Department of Clinical Immunology, Alexandrovska University Hospital, Medical University Sofia, Sofia, Bulgaria; [2]Department of Clinical Hematology, Alexandrovska University Hospital, Medical University Sofia, Sofia, Bulgaria; [3]Department of Experimental Research, Medical University Pleven, Pleven, Bulgaria

P5 | A non-invasive method for fetal trophoblast HLA typing in early pregnancy

Liseanne J. van T. Hof[1], Hanneke M. Kapsenberg[1], Jos J.M. Drabbels[1], Michael Eikmans[1] and Marie-Louise L.P. van der Hoorn[2]
[1]Department of Immunology, Leiden University Medical Center, Leiden, Netherlands; [2]Department of Obstetrics and Gynaecology, Leiden University Medical Center, Leiden, Netherlands

P6 | Differential Modulation of Mutant CALR and JAK2 V617F-Driven Oncogenesis by HLA Genotype in Myeloproliferative Neoplasms

Velizar Shivarov[1], Gergana Tsvetkova[2], Ilina Micheva[3], Evgueniy Hadjiev[2], Jasmina Petrova[4], Anela Ivanova[4], Galja Madjarova[4] and Milena Ivanova[5]
[1]Department of Experimental Research, Medical University Pleven, Pleven, Bulgaria; [2]Department of Clinical Hematology, Alexandrovska University Hospital, Medical University Sofia, Sofia, Bulgaria; [3]Department of Clinical Hematology, Saint Marina University Hospital, Medical University Varna, Varna, Bulgaria; [4]Department of Physical Chemistry, Faculty of Chemistry and Pharmacy, Sofia University "St. Kl. Ohridski",Bulgaria, Bulgaria; [5]Department of Clinical Immunology, Alexandrovska University Hospital, Medical University Sofia, Sofia, Bulgaria

P7 | Effects of HLA-G molecules in Primary Biliary Cholangitis: The Sardinian Experience

Caterina Mereu[1], Michela Lorrai[1], Stefano Mocci[1], Roberto Littera[2], Michela Miglianti[3], Celeste Sanna[1], Michela Murgia[1], Meropi Plousiou[1], Chiara Cocco[1], Alessia Mascia[4], Marina Serra[4], Sara Lai[2], Erika Giuressi[2], Maria Conti[5], Cinzia Balestrieri[5], Giancarlo Serra[5], Francesco Pes[5], Teresa Zolfino[6], Andrea Perra[3], Luchino Chessa[3] and Sabrina Rita Giglio[2]
[1]Medical Genetics Unit, Department of Medical Sciences and Public Health, University of Cagliari, Cagliari, Italy; [2]Medical Genetics Unit, R. Binaghi Hospital, Local Public Health and Social Care Unit of Cagliari, Cagliari, Italy; [3]Department of Medical Sciences and Public Health, University of Cagliari, Cagliari, Italy; [4]Section of Pathology, Department of Biochemical Sciences, University of Cagliari, Cagliari, Italy; [5]Liver Unit, University Hospital, Monserrato, Cagliari, Italy; [6]Gastroenterology Unit, ARNAS Brotzu, Cagliari, Italy

P8 | Role of HLA-G in Sardinian Idiopatic Pulmonary Fibrosis patients

Michela Lorrai[1], Caterina Mereu[1], Stefano Mocci[1], Roberto Littera[2], Silvia Deidda[3], Celeste Sanna[1], Michela Murgia[1], Meropi Plousiou[1], Gaia Maria Tosone[1], Alessia Mascia[4], Marina Serra[4], Sara Lai[2], Erika Giuressi[2], Andrea Perra[5] and Sabrina Rita Giglio[2]
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P9 | Islet Antibodies Among North Indian Type 1 Diabetes Patients: Prevalence and Persistence Pattern

Uma Kanga[1], Shreya Sharma[2], Alpesh Goyal[2] and Nikhil Tandon[2]
[1]Department of Transplant Immunology and Immunogenetics, All India Institute of Medical Sciences, India; [2]Department of Endocrinology and Metabolism, All India Institute of Medical Sciences, India

*Among this T1D cohort, 74.9% were HLA-DRB1*03 positive while only 30.6% were DRB1*04 positive. Significant associations of DRB1*03 alleles with presence of GAD antibodies and DRB1*04 alleles with ZnT8 antibody positivity were observed. Antibody persistence did not reflect any HLA association. A unique antibody profile was observed in this T1D cohort.*

P10 | Nanopore sequencing reveals high resolution HLA alleles associated to autoimmune neutropenia in early childhood

Kirstine Kløve-Mogensen[1], Thure Mors Haunstrup[1], Tania Nicole Masmas[2], Andreas Glenthøj[3], Petter Höglund[4], Henrik Hasle[5], Kaspar Rene Nielsen[1] and Rudi Steffensen[1]
[1]Department of Clinical Immunology, Aalborg University Hospital, Aalborg, Denmark; [2]Department of Pediatrics and Adolescent Medicine, Copenhagen University Hospital, Rigshospitalet, Copenhagen, Denmark; [3]Department of Hematology, Copenhagen University Hospital – Rigshospitalet, Copenhagen, Denmark; [4]Department of Medicine Huddinge, Karolinska Institute, Stockholm, Sweden; [5]Department of Pediatrics and Adolescent Medicine, Aarhus University Hospital, Aarhus, Denmark

P11 | Detection and identification of platelet antibodies using a Luminex bead assay

Nataša Katalinić[1], Tajana Crnić Marčetić[2], Sandra Šever Šušnjar[2], Marijana Duhović[2] and Sanja Balen[1]
[1]Tissue Typing Laboratory, CHC Rijeka; Department of Clinical Laboratory Diagnostics, Medical Faculty Rijeka, Croatia; [2]Tissue Typing Laboratory, Clinical Hospital Center Rijeka, Croatia

P12 | Distinguishing Anti-3-hydroxy-3-methylglutaryl-CoA reductase Immune-Mediated Necrotizing Myopathy from Non-Immune-Mediated Statin Myotoxicity: The Role of HLA-DRB1*11:01 Association

Diana Prieto-Peña[1], Nerea Mota-Pérez[2], J. Gonzalo Ocejo[2], Cristina Corrales-Selaya[1], Carmen García-Ibarbia[3], Verónica Pulito-Cueto[4], Raquel López-Mejías[4], Ricardo Blanco[1] and José Luis Hernández[3]
[1]Rheumatology Department, Hospital Universitario Marqués de Valdecilla, IDIVAL, Santander, Spain; [2]Immunology Department, Hospital Universitario Marqués de Valdecilla, IDIVAL, Santander, Spain; [3]Internal Medicine Department, Hospital Universitario Marqués de Valdecilla, IDIVAL, Santander, Spain; [4]IDIVAL Health Research Institute of Cantabria, Santander, Spain

*Myotoxicity and controls showed no significant differences in HLA-DRB1*11 allele distribution. Our findings suggest a strong association between HLA-DRB1*11, particularly HLA-DRB1*11:01, and anti-HMGCR IMNM, not observed in non-immune-mediated statin myotoxicity. Identifying HLA-DRB1*11:01 may help identify those at high risk of anti-HMGCR IMNM.*

P13 | The impact of HLA-DRB1 alleles in a Hellenic Pediatric Onset Multiple Sclerosis cohort: Implications on clinical and neuroimaging profile.

Charalampos Skarlis[1], Nikolaos Markoglou[2], Maria Gontika[3], Petros Prapas[1], Maria Kotsari[1], Artemios Artemiadis[4], Maria-Roser Pons[5], Leonidas Stefanis[2], Marinos Dalakas[6], George Chrousos[7] and Maria Anagnostouli[2]
[1]Immunogenetics Laboratory, First Department of Neurology, School of Medicine, NKUA, Aeginition University Hospital, Greece; [2]First Department of Neurology, School of Medicine, NKUA, Aeginition University Hospital, Greece; [3]Penteli Children's Hospital, Greece; [4]Neurology Department, Cyprus University, Greece; [5]First Department of Pediatrics, School of Medicine, NKUA, Agia Sofia University Hospital, Greece; [6]Neuroimmunology Laboratory, Department of Pathophysiology School of Medicine, NKUA, Greece; [7]Clinical and Translational Research Unit in Endocrinology, NKUA, Greece

P14 | Availability of CD36 negative platelets and the importance of CD36 Donor Screening

Sarah Petermann[1], Brigitte Flesch[2], Beate Kirchharz[3], Martina Wessiepe[4], Svenja Woestmann[1] and Alexander Carbol[1]
[1]German Red Cross Blood Service Rhineland-Palatinate and Saarland, Bad Kreuznach, Germany; [2]German Red Cross Blood Service West, Hagen, Germany; [3]German Red Cross Blood Service West, Ratingen, Germany; [4]University Hospital RWTH Aachen, Transfusion Medicine, Aachen, Germany

P15 | Investigation of the interaction of HLA-G with natural killer inhibitory receptors in colorectal cancers

Ezgi Dincer[1], Fatma Kaya Dagistanli[2], Kivanc Derya Peker[3], Damlanur Sakiz[4], Demet Kivanc Izgi[1], Hayriye Senturk Ciftci[1] and Fatma Savran Oguz[1]
[1]Istanbul University, Istanbul Faculty of Medicine, Institute of Health Sciences, Department of Medical Biology, Turkey; [2]Istanbul University-Cerrahpasa, Cerrahpasa Faculty of Medicine, Department of Medical Biology, Turkey; [3]Hisar Intercontinental Hospital, Department of General Surgery, Turkey; [4]University of Health Sciences Bakırkoy Dr. Sadi Konuk Training and Research Hospital, Pathology Clinic, Turkey;

P16 | Potential actionable somatic variants in Chronic Lymphocytic Leukemia

Gurvinder Kaur[1], Ayushi Jain[1], Kamaljeet Singh[1], Lingaraja Jena[1], Ajay Gogia[2], Atul Sharma[2] and Ritu Gupta[1]
[1]Lab Oncology, Dr BRAIRCH, All India Institute of Medical Sciences, New Delhi, India; [2]Medical Oncology, Dr BRAIRCH, All India Institute of Medical Sciences, New Delhi, India

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Daria Shmelkova[1], Daria Stashkevich[1], Tatiana Suslova[1], Inessa Devald[1] and Alexandra L. Burmistrova[1]
[1]Chelyabinsk State University, Russia;

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Cigdem Kekik[1], Sonay Temurhan[1], Yeliz Ogret[1], Demet Kivanc Izgi[1], Behnoush Nasr Zanjani[2], Fatma Betul Oktelik[3], Fatma Savran Oguz[1], Murat Kose[4] and Gunnur Deniz[3]
[1]Department of Medical Biology, HLA Laboratory, Istanbul Faculty of Medicine, Istanbul University, Istanbul, Turkey; [2]Institute of Graduate Studies in Health Sciences, Istanbul University, Istanbul, Turkey; [3]Department of Immunology, Aziz Sancar Institute of Experimental Medicine, Istanbul University, Istanbul, Turkey; [4]Department of Internal Medicine, Division of General Internal Medicine, Istanbul Faculty of Medicine, Istanbul, Turkey

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Ritu Aggarwal[1], Tanvi Bhatia[1], Arshiya Mehra[1] and Vanita Suri[2]
[1]Department of Immunopathology, PGIMER, Chandigarh, India; [2]Department of Obstetrics and Gynecology, PGIMER, Chandigarh, India

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Martin Petrek[1], Katerina Sikorova[2], Lenka Kocourkova[3], Martin Dobias[2] and Jana Petrkova[1]
[1]University Hospital Olomouc and Faculty of Medicine and Dentistry Palacky University Olomouc, Czechia; [2]University Hospital Olomouc, Czechia; [3]Faculty of Medicine and Dentistry Palacky University Olomouc, Czechia

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Irina Pavlova[1], Elena Kuzmich[1], Elena Shilova[1], Tatyana Glazanova[1] and Ludmila Bubnova[1]
[1]Russian Research Institute of Hematology and Transfusiology, Russia

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Daria Stashkevich[1], Tatiana Suslova[1] and Alexandra L. Burmistrova[1]
[1]Chelyabinsk State University, Russia;

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Eulalia Catamo[1], Valentina Bazzo[2], Valentina Moro[2], Antonietta Robino[1], Luana Aldegheri[1], Andrea Conti[1], Gianluca Tornese[1] and Elena Bevilacqua[2]
[1]Institute for Maternal and Child Health—IRCCS "Burlo Garofolo", Trieste , Italy; [2]Tissue Typing Laboratory, Transfusion Medicine Department, University Hospital (ASUGI), Trieste, Italy

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Vasiliki Kitsiou[1], Elisavet Kontou[1], Petros Mantzios[1], Glykeria Tsouka[2], Stella Pomoni[1], Theofilos Athanassiades[1], Katerina Tarassi[1], Vasileios Vougas[3], Maria Darema[2] and Alexandra Tsirogianni[1]
[1]Immunology-Histocompatibility Department, Evangelismos General Hospital, Athens, Greece; [2]Nephrology Department, Evangelismos General Hospital, Athens, Greece; [3]Department of General Surgery and Transplant Unit, Evangelismos General Hospital, Athens, Greece

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Svetlana Balandina[1], Svetlana Belyaeva[1] and Daria Stashkevich[1]
[1]Chelyabinsk State University, Russia;

BIOINFORMATICS, DATA ANALYSIS IN IMMUNOGENETICS

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Julien Paris[1], Sonia Bourguiba-Hachemi[1], Romain Casey[2], Nicolas Vince[1] and Pierre-Antoine Gourraud[1]
[1]Nantes Université, CHU Nantes1, INSERM, CR2TI, UMR 1064, F-44000, Nantes, France; [2]OFSEP, Research Center in Neurosciences of Lyon, INSERM 1028 and CNRS UMR 5292, F-69003 Lyon, France

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Nickolai Alexandrov[1], Ting Wang[1], Brian Nadon[1], Lindley Blair[1], Yuki Saito[1] and David Sayer[2]
[1]Thermo Fisher Scientific Inc., United States; [2]Thermo Fisher Scientific Inc., Australia

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Liza Huijse[1], Solomon M. Adams[2], Joshua N. Burton[2], Julianne K. David[2], Russell S. Julian[2], Galit Meshulam-Simon[2], Harry Mickalide[2], Bersabeh D. Tafesse[2], Verónica Calonga-Solís[3], Ivan Rodrigo Wolf[3], Ashby J. Morrison[4], Danillo G. Augusto[3] and Solomon Endlich[2]

[1]Base5 Genomics, Inc., Netherlands; [2]Base5 Genomics, Inc., United States; [3]The University of North Carolina at Charlotte, United States; [4]Stanford University, United States

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Ngoc Giang Le[1], Jesse Mittertreiner[1], Susan Ott[1], Nanine de Groot[1], Marit van der Wiel[1], Natasja de Groot[1], Jesse Bruijnesteijn[1] and Ronald E. Bontrop[1]

[1]Biomedical Primate Research Centre, Netherlands;

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Ryan Nickens[1], Livia Tran[2], Leamon Crooms[3] and Steven Mack[2]

[1]Lafayette College, Easton, PA, United States; [2]University of California, San Francisco, United States; [3]University of Arizona, United States

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Henry Loeffler-Wirth[1], Claudia Lehmann[2], Nils Lachmann[3] and Ilias Doxiadis[2]

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Henry Loeffler-Wirth[1], Nils Lachmann[2], Claudia Lehmann[3] and Ilias Doxiadis[3]

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Cinthia Alves[1] and Eduardo Donadi[1]

[1]Ribeirão Preto Medical School, University of São Paulo, Brazil;

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Jonna Clancy[1], Silja Tammi[2], Jukka Partanen[2] and Satu Koskela[1]

[1]Finnish Red Cross Blood Service Biobank, Finland; [2]Finnish Red Cross Blood Service, R&D, Finland

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Enrique Alanis[1], Diana Alcantara[2], Alicia Sanchez-Mazas[1] and José Manuel Nunes[1]

[1]Department of Genetics and Evolution, University of Geneva, Switzerland; [2]Brighton and Sussex Medical School, United Kingdom

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Michaela Agapiou[1], Richard H.L. Natarajan[2], Claudia Rutt[3], Sumati Misra[3], James Robinson[1], Nezih Cereb[3] and Steven G.E. Marsh[1]

[1]Anthony Nolan Research Institute, London, United Kingdom; UCL Cancer Institute, London, United Kingdom; [2]Anthony Nolan Research Institute, London, United Kingdom; [3]DATRI Blood Stem Cell Donors Registry, Chennai, India, United Kingdom

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Susanne Seitz[1], Karen Ende[1], Katharina Daniel[2], Vinzenz Lange[2] and Alexander H. Schmidt[1]

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P40 | PRIMUS: A distributed web application architecture to benchmark sets of analytics based on reuse of multi-source clinical research data in multiple sclerosis under the umbrella of precision medicine

Igor Faddeenkov[1], Stanislas Demuth[1], Joëlle Martin-Gauthier[1], Julien Paris[1], Gilles Edan[2], David Laplaud[1] and Pierre-Antoine Gourraud[1]

[1]Nantes Université, CHU Nantes, INSERM, CR2TI, UMR 1064, ITUN, Nantes, France; [2]Department of Neurology, University Hospital of Rennes, France

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Panagiotis Mallis[1], Alexandra Siorenta[1], Erasmia Stamathioudaki[1], Vasiliki Vrani[1] and Georgios Paterakis[1]

[1]Immunology Department & National Tissue Typing Center, General Hospital of Athens "G. Gennimatas", Athens, Greece

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Franz Fenninger[1], Bronwyn Baillie[1], Karen Sherwood[1], Jenny Tran[1], James Lan[1] and Paul Keown[1]

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Enrique Alanis[1], Alicia Sanchez-Mazas[1] and José Manuel Nunes[1]

[1]Department of Genetics and Evolution, University of Geneva, Switzerland

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Cinthia Alves[1], Rafaela Miranda Barbosa[1] and Eduardo Donadi[1]

[1]Ribeirão Preto Medical School, University of São Paulo, Brazil;

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Benedict Matern[1] and Matthias Niemann[2]

[1]PIRCHE AG, Netherlands; [2]PIRCHE AG, Germany

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Deniz Akdemir[1] and Yung-Tsi Bolon[1]

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Shahram Aliyari[1], Zahra Salehi[2], Benedikt Brors[3], Kaveh Kavousi[4] and Mohammad Hossein Norouzi-Beirami[5]

[1]University of Tehran, Kish International Campus, Iran; [2]Tehran University of Medical Sciences, Iran; [3]German Cancer Research Center DKFZ Heidelberg, Germany; [4]Institute of Biochemistry and Biophysics, Iran; [5]Islamic Azad University, Iran

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Laura Bungener[1], Niels Kouprie[1], Magdalena Huberts-Kregel[1], Annechien Lambeck[1], Bart-Jan Kroesen[1] and Bouke Hepkema[1]

[1]Transplantation Immunology, Dept. of Laboratory Medicine, University Medical Center Groningen, University of Groningen, Netherlands

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Matěj Röder[1], Vít Svojše[2], Tomáš Kebrle[2], Kateřina Jáklová[1], Jiří Gurka[3], Petr Raška[3] and Antonij Slavčev[1]
[1]Department of Immunogenetics, Institute for Clinical and Experimental Medicine, Prague, Czechia; [2]Bindworks S.R.O., Czechia; [3]Information Technology Division, Institute for Clinical and Experimental Medicine, Prague, Czechia

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Luis Cristovao Porto[1], Angela Maria Santos[1], Danielle Secco[1], Raiana Barbosa[2], Romulo Oliveira[1], Antonio Carlos Campos-de-Carvalho[2], Adriana Carvalho[2], Daniellii Oliveira[3] and Tais Brunswick[2]
[1]Rio de Janeiro State University, Brazil; [2]Federal University of Rio de Janeiro, Brazil; [3]National Cancer Institute, Brazil

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Manuel Quirno Costa[1], Maria Belén Balekjian[1], Santiago Meinardo Cambra[2] and Gustavo Piccinelli[3]
[1]Laboratorio Nacional de Inmunogenética – INCUCAI, Argentina; [2]Freelance Programmer, Argentina; [3]Centro Provincial de Histocompatibilidad – CUCAIBA, Argentina

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Asimina Fylaktou[1], Georgios Lioulios[2], Georgios Petasis[1], Artemis Maria Iosifidou[3], Myrto Aikaterini Iosifidou[3], Georgia Chatzika[1], Evaggelia Sidira[1], Fani Makrovasili[1], Maria Stangou[4] and Ioannis Theodorou[5]
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P53 | Advancing HLA typing on nanopore sequencing platforms: A novel phasing-based bioinformatics software for high resolution and accuracy

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P54 | Laboratory reporting of DQB1*03:05 (DQ8) and DQB1*03:19 (DQ7) in UK NEQAS for H&I's external proficiency testing scheme for Coeliac Disease

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P55 | Performance of participants of UK NEQAS for H&I pharmacogenetics schemes over the last 5 years

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P56 | Enhancing organ transplantation outcomes through standards-compliant software solutions

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P57 | Avocado as a supplement in Afro-Colombian patients with obesity, diabetes and who express the HLA-DQB1*02:01

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HEMATOPOIETIC STEM-CELL TRANSPLANTATION (HSCT)

P58 | Effect of Killer-cell Immunoglobulin-like Receptors and their cognate HLA ligands on tumor disease-free survival after allogeneic hematopoietic stem cell transplantation

Jacek Nowak[1], Agnieszka Witkowska[1], Agnieszka Malinowska[1], Klaudia Nestorowicz-Kałużna[2], Elżbieta Graczyk-Pol[1], Anna Flaga[1], Urszula Szlendak[1], Anna Wnorowska[1], Aleksandra Wawiórko[1], Wiktoria Szwajkowska[1] and Agnieszka Gawron[1]
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P59 | HLA-loss relapse detection using NGS data from standard HLA genotyping in haploidentical stem cell transplantation

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P60 | HLA Loss detection after Haplo-identical Hematopoietic Stem Cell Transplantation using a Next Generation Sequencing based specific approach.

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P61 | DSA determination and eplet mismatches analysis in pediatric patients affected by hemoglobinopathies treated with haploidentical hematopoietic stem cell transplantation

Paola Giustiniani[1], Federica Galaverna[2], Pietro Merli[2], Antonio Giuseppe Bianculli[1], Marco Becilli[2], Roberto Carta[2], Emilia Boccieri[2], Maria Troiano[1], Rita Maria Pinto[2], Mariarosa Battarra[1], Marco Andreani[1] and Franco Locatelli[2]
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P62 | Eplets mismatched analysis in a group of pediatric patients affected by hematological malignant diseases treated with haploidentical HSCT

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P63 | HLA-DQ heterodimers and graft failure after haploidentical stem cell transplantation in patients with acute leukemia

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P64 | Validation and implementation of HLA loss relapse detection by NGS-based HLA typing

Elena Gómez Massa[1], Sílvia Mantecón-Ferrer[1], Laura Mongay Berdet[1], Josefa Caro Martínez[1], M. Luz Uria-Oficialdegui[2], Marta Peña Domingo[3], Maria Huguet Mas[4], Cecilia Gonzalez[5], Francesc Rudilla-Salvador[1] and María José Herrero[1]
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P65 | Impact of KIR genotype on clinical outcome of Hematopoietic Stem Cell transplants: a single center experience

Francesco Ingrassia[1], Alice Pecoraro[1], Maria Blando[1], Alessia Angela Corica[1], Floriana Di Paola[1], Giuseppe Davì[1], Rosalba Bavetta[1], Floriana Bruno[1], Serena Mistretta[1], Stefania Tringali[2], Roberta Fedele[1], Luca Castagna[2] and Valentina Cappuzzo[1]
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P66 | The impact of donor-specific antibodies presence on the outcome post-allogeneic hematopoietic stem cell transplantation: a survey from a single center

Sabrina Giammarco[1], Patrizia Chiusolo[2], Filippo Frioni[1], Elisabetta Metafuni[1], Maria Assunta Limongiello[3], Eugenio Galli[3], Federica Sorá[2], Andrea Bacigalupo[2], Elvira Poggi[4], Mariano Antonio Feccia[5], Annarita Manfreda[5] and Simona Sica[2]
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P67 | Impact of HLA mismatches at the molecular level on the clinical evolution of haploidentical hematopoietic stem cell transplantation with post-transplant cyclophosphamide

Francisco Javier Gil-Etayo[1], Jairo Eduardo Niño-Ramírez[2], Marta Fonseca[1], Pilar Terradillos-Sánchez[2], Isabel Jiménez Hernaz[3], Ariadna Vicente Parra[3], Ana Balanzategui[3], Daniel Arroyo-Sánchez[1], Beatriz García Martín[1], Yasmín Roldán[4], Francisco Boix[5], Miguel Alcoceba[6], Estefania Pérez[1], Mónica Cabrero[1], Almudena Navarro-Bailón[1], Fermín Sánchez-Guijo[1], Ramón García Sanz[3], Lucía López-Corral[1] and Amalia Tejeda[3]
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P68 | Inferior survival is observed among adult HSCT recipients having pre-transplant anti-HLA antibodies even though they are not directed against the donor

Antonio Milano[1], Giuliana Lando[2], Giulia Di Maggio[2], Giorgia Cornacchini[2], Giovanni Grillo[2], Silvano Rossini[2] and Roberto Crocchiolo[2]

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P69 | Management of ABO-incompatible HSCT: 2021-2023 data analysis

Chiara Sindici[1], Donatella Londero[1], Silvia Borgobello[1], Elisa Cecchini[1], Celeste Cervellin[1], Ilaria Sandron[1] and Giovanni Barillari[1]

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P70 | Maternal uniparental disomy of chromosome 6 (upd(6) mat) in a patient affected by acute myelogenous leukemia awaiting HSC transplant

Franco Papola[1], Alessandra Zoli[2], Attilio Olivieri[3], Raffaella Azzarone[1], Olaida Valdez[1], Maria Grazia Tupone[1], Stefano Agolini[2], Ilaria Scortechini[3], Elia Trinchini[1], Stefano Scipione[1], Simona Scacchi[1], Giovanni Rombolà[4] and Carla Cervelli[1]

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P71 | Post-Transplant Chimerism Monitoring – Real-Time qPCR or STR?

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P72 | Donor telomere length and telomerase reverse transcriptase gene polymorphism may affect the outcome of allogeneic hematopoietic stem cell transplantation in children – a preliminary study

Marta Dratwa[1], Barbara Wysoczanska[1], Marek Ussowicz[2], Blanka Rybka[2], Renata Ryczan-Krawczyk[2], Krzysztof Kalwak[2] and Katarzyna Bogunia-Kubik[3]

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P73 | Genetic predisposition to hematologic malignancies in patients undergoing allogeneic hematopoietic stem cell transplantation: case reports

Alice Brogi[1], Sara Iozzi[1], Beatrice Boschi[1], Luisa Candita[1], Giulia Carignani[1], Giulio Capecchi[2], Francesco Mannelli[2], Paola Guglielmelli[2], Chiara Nozzoli[3], Simona Palchetti[1], Mauro Leucio Mattei[1], Martina Betti[1], Elisabetta Pelo[1] and Irene Donnini[3]

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P74 | Standardization of evaluation of cytokine gene polymorphisms by next generation sequencing

Selma Zenia D'Silva[1], Manisha Tambe[1], Shreya Kupekar[1] and Meenakshi Singh[1]

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P75 | Improved haploidentical donor matching including low expression HLA loci in the immunogenetic analysis using Next-Generation Sequencing

Wiktoria Wypych[1], Dominika Moskalik-Kierat[1], Martyna Brzoza[1], Anna Wasilewska[1], Mateusz Mrozowski[1], Agnieszka Jałbrzykowska[1], Ewa Karakulska-Prystupiuk[2], Piotr Kacprzyk[2], Agnieszka Tomaszewska[2], Grzegorz Basak[2] and Marcelina Grabowska[1]

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P76 | Detection and monitoring of donor-specific antibodies in a haploidentical stem cell transplant recipient: a case report

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P77 | Analysis of the significance of donor chimerism at +14 days after haploidentical hematopoietic stem cell transplantation

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P78 | Changes of the immunogenic profile of an acute myeloid leukemia relapsed patient with partial loss of heterozygosity in HLA genes after haploidentical transplantation of hematopoietic stem cells

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P79 | Partial loss of heterozygosity in HLA genes in patient with relapsed acute lymphoblastic leukemia

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P80 | Mismatched unrelated donors for hematopoietic stem cell transplantation in children with inborn errors of immunity: selection criteria and the Newcastle experience

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IMMUNOGENETICS IN ORGAN TRANSPLANTATION

P81 | On the road to epitope matches and mismatches – with epiTOol

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Following kidney transplantation although up to 40 epitope mismatches were present. In contrast, recipients with only nine EMM had produced antibodies towards the donor. Defining epitopes is helpful for understanding the humoral response but it is not a matter of numbers but a matter of quality and could be a useful tool for donor selection in the future.

P82 | Clinical relevance of isolated preformed HLA-DP donor specific antibodies on the outcome of kidney transplantation

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P83 | Interference of cold agglutinins and/or cryoglobulins in antibody diagnostics for kidney transplantation: a case study

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P84 | High levels of BCMA transcript expression prior to transplantation, increased plasmablast, lymphocyte B cell class-switched levels and viral loading are associated with early CMV reactivation in renal recipients

Marina Fernandez[1], Rafael Alfaro[1], Maria Jose Alegria-Marcos[1], Jose Antonio Galian[1], Helios Martinez-Banaclocha[1], Carmen Botella[1], Alfredo Minguela[1], Maria Rosa Moya-Quiles[1], Santiago Llorente[1], Isabel Legaz[2] and Manuel Muro[1]
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P85 | Imlifidase for Kidney Transplantation of Highly Sensitized Patients with a Positive Crossmatch: The French Consensus Guidelines

Gwendaline Guidicelli[1], Paolo Malvezzi[2], Lucile Amrouche[3], Dany Anglicheau[3], Gilles Blancho[4], Sophie Caillard[5], Marine Freist[6], Nassim Kamar[7], Carmen Lefaucheur[8], Christophe Mariat[9], Alice Koenig[10], Johan Noble[2], Olivier Thaunat[10], Antoine Thierry[11], Jean-Luc Taupin[8], Dominique Bertrand[12] and Lionel Couzi[1]
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P86 | Effect of peri-transplantation circumstances on the amelioration of cellular immunity following kidney transplantation

Lambros Vagiotas[1], Georgios Lioulios[1], Efstratios Kasimatis[1], Aliki Xochelli[1], Anna Boukla[1], Georgia Chatzika[1], Despoina Asouchidou[1], Margarita Samali[1], Nikolaos Antoniadis[1], Georgios Tsoulfas[1], Maria Stangou[1] and Asimina Fylaktou[1]

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P87 | Detection of donor-derived cell-free DNA in sequential kidney transplanted patients

Linnéa Pettersson[1], Lukas Frischknecht[2], Sofia Westerling[1], Hamid Ramezanali[1], Lukas Weidmann[3], Kai Castrezana Lopez[3], Thomas Schachtner[3] and Jakob Nilsson[3]

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P88 | Role of HLA matching and donor specific antibody development in long-term survival, acute rejection and cardiac allograft vasculopathy

Dario Costa[1], Antonietta Picascia[1], Vincenzo Grimaldi[1], Cristiano Amarelli[2], Andrea Petraio[2], Anna Levi[1], Mario Di Donato[1], Anna Virginia Adriana Pirozzi[1], Carmela Fiorito[1], Giusi Moccia[1], Aurora Gallo[1], Claudio Marra[2], Marisa De Feo[2], Francesco Cacciatore[3], Ciro Maiello[2] and Claudio Napoli[1]

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P89 | Correlation between anti-angiotensin II receptor 1 and anti-endothelin type A receptor 1 in kidney transplanted pediatric patients

Antonio Giuseppe Bianculli[1], Paola Giustiniani[1], Annalisa Guagnano[1], Andrea Di Luzio[1], Francesca Besi[1], Raffaella Labbadia[2], Luca Antonucci[2], Andrea Cappoli[2], Isabella Guzzo[2] and Marco Andreani[1]

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P90 | Acute allograft dysfunction and glomerular microangiopathy in kidney transplant recipients in absence of donor specific anti-HLA antibodies: a case series

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P91 | Biomarkers of Innate Immunity and Immunological Susceptibility to Viral Infection in Patients with Alcoholic Cirrhosis

Isabel Legaz[1], Elena Navarro-Noguera[1], Aurelia Collados-Ros[1], Rosana Gonzalez-Lopez[2], Jose Miguel Bolarin[1] and Manuel Muro[2]

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P92 | Non-HLA antibodies in highly sensitized recipients on the kidney waiting list

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P93 | Deceased kidney donor virtual crossmatch introduction in north Italian transplant program (NITp) area

Elena Longhi[1], Loretta Crespiatico[1], Viviana Sioli[1], Vittoria Caporale[1], Alessia Comino[1], Francesca Drago[1], Alejandro Espadas de Arias[1], Miriam Ramondetta[1], Augusto Tagliamacco[1], Marco Guarene[1], Denise Bertola[1], Caterina Brambilla[1], Nicoletta Cagni[1], Sara Capogreco[1], Ludovica Chidichimo[1], Michela Grassi[1], Annalisa Innocente[1], Mario Macchiagodena[1], Barbara Speringo[1], Nemania Suvajac[1], Mara Tivelli[1] and Tullia Maria De Feo[1]

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P94 | A personalized delisting strategy enables successful kidney transplantation in highly sensitized patients with preformed donor-specific anti-HLA antibodies.

Sandra García Jiménez[1], Estela Paz-Artal[1], Hernando Trujillo[1], Natalia Polanco[1], Maria J Castro[1], Manuel J Del Rey[1], Angel Alfocea[1], Enrique Morales[1], Esther Gonzalez[1], Amado Andres[1] and Esther Mancebo[1]

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P95 | Imlifidase desensitization in highly-HLA sensitized patients with positive Cross-Match: first experience in Parma

Silvia Giuliodori[1], Madga Benecchi[1], Pamela Berni[1], Chiara Foroni[1], Claudia Labate[1], Roberta Merli[1], Valentina Sgobba[1], Stefania Bardini[1], Francesca Lobascio[1], Elisa Magni[1], Joseph Manduca[1], Jessica Parrotta[1], Irene Pezzani[1], Elena Russello[1], Raffaele Troiano[1], Umberto Maggiore[2] and Paola Zanelli[1]

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P96 | Case report: simultaneous combined liver-kidney transplant. Focus on immunological assessment

Paola Zanelli[1], Silvia Giuliodori[1], Giorgia Comai[2], Matteo Ravaioli[3], Irene Pezzani[1], Giovanni Rombolà[1] and Gaetano La Manna[2]

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P97 | Desensitization-Resistant Eplet-Specific HLA Antibodies

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P98 | Differences of the subpopulations of T-lymphocytes between long-term and recent kidney transplant recipients

Evangelos Memmos[1], Georgios Lioulios[2], Efstratios Kasimatis[3], Aliki Xochelli[4], Lambros Vagiotas[5], Vasiliki Nikolaidou[4], Nikolaos Antoniadis[5], Georgios Tsoulfas[5], Maria Stangou[3] and Asimina Fylaktou[4]

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P99 | The reduction of T and B regulatory lymphocytes in long-term kidney transplant recipients

Evangelos Memmos[1], Efstratios Kasimatis[2], Georgios Lioulios[3], Aliki Xochelli[4], Lambros Vagiotas[5], Vasiliki Nikolaidou[4], Nikolaos Antoniadis[5], Georgios Tsoulfas[5], Maria Stangou[2] and Asimina Fylaktou[4]

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P100 | Pre-transplant flow cytometric crossmatch in patients undergoing Rituximab treatment: employment of Pronase and anti-CD20

Mario Macchiagodena[1], Barbara Speringo[1], Miriam Ramondetta[1], Viviana Sioli[1], Augusto Tagliamacco[1], Alessia Comino[1], Caterina Brambilla[1], Nicoletta Cagni[1], Annalisa Innocente[1], Sara Capogreco[1], Loretta Crespiatico[1] and Elena Longhi[1]

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P101 | Deciphering HLA Antibody Reactivity Patterns: A Cluster-Based Analysis of SAB Assay Data

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P102 | Can Molecular HLA Mismatch Scores Predict Antibody-Mediated Rejection in Desensitized Kidney Transplant Recipients?

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P103 | Clinical utility of 1:16 serum dilution as a predictor of response to therapeutic plasma exchange for HLA antibody-mediated rejection treatment and overall survival in lung transplant recipients: A two center study

Mohamed Elrefaei[1], Tathagat Narula[1], Francisco Alvarez[1], Elizabeth Godbey[1], Gerard Criner[2], Francis Cordova[2], Norihisa Shigemura[3], Yoshiya Toyoda[3] and Olga Timofeeva[4]

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P104 | Characterization of sHLA-AlexaFluor647-conjugates for accurate determination of serum HLA antibody concentration and affinity

Matthias Reist[1], Rico Buchli[2], Sean R. A. Devenish[3], Dominik Meinel[4], Sebastiaan Heidt[5], Canthia S.M. Kramer[5], Suzanne Bezstarosti[5], Rosa G.M. Lammerts[6], Stefan Schaub[7] and Gideon Hönger[1]

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P105 | High Levels of Complement-Binding Donor-Specific Anti-HLA Antibodies: Are They Always Pathogenic?

Katrin Hacke[1], Firas Marayati[2], Sami Alasfar[2], Hasan A. Khamash[2], Lavanya Kodali[2], Pooja Budhiraja[2], Girish K. Mour[2], Bassam G. Abu Jawdeh[2] and Andrés Jaramillo[1]

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P106 | Effect of HLA-B -21 dimorphism on cellular response after COVID-19 vaccination in patients with renal disorders

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P107 | Could donor-derived cell-free DNA level be considered a potential biomarker of subclinical graft rejection or early organ damage in kidney transplant recipients?

Marina Fernandez-Gonzalez[1], Victor Jimenez-Coll[1], Carmen Botella[1], Rosana Gonzalez-Lopez[1], Maria Jose Alegria-Marcos[1], Carlos Sanchez-Rodriguez[1], Maria Rosa Moya-Quiles[1], Jose Antonio Galian[1], Santiago Llorente-Viñas[1], Isabel Saura-Luján[1], Francisco Morales-Caravaca[1], Gloria Martinez-Gomez[1] and Manuel Muro[1]

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P108 | A critical retrospective analysis of Flow Cytometry Crossmatch in the emergency context in order to optimize the available resources

Cláudia Silva[1], Paula Aires[1], Pedro Ramoa[1], Paula Santo[1] and Paula Xavier[1]

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P109 | Positive crossmatch in ABO-incompatible neonatal heart transplant due to transplacental transfer of maternal HLA antibodies

Cristiana Caorsi[1], Gabriele Maria Togliatto[1], Claudia Maria Rosso[1], Sara Tontoni[2], Rossella Chidichimo[2], Enrico Aidala[3], Carlo Pace Napoleone[3] and Silvia Deaglio[2]

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P110 | Comparison of results of crossmatch tests from peripheral blood cells with crossmatch tests from secondary lymphoid organs cells in kidney transplantation

Aggeliki Vittoraki[1], Georgios Lioulios[2], Alexandra Siorenta[1], Margarita Samali[3], Chrysanthi Tsirligkani[1], Aliki Xochelli[3], Aikaterini Anastasiou[3], Vasiliki Nikolaidou[3], Athina Aikaterini Nikolaou[1], Georgios Paterakis[1] and Asimina Fylaktou[3]

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P111 | Donor derived cell-free DNA: clinical utility in surveillance strategy for heart transplant

Elvira Poggi[1], Vincenzo Dinallo[2], Sabina Maria Bruno[2], Silvia Sinopoli[2], Giada Distefano[3], Fabio Sbaraglia[3], Giulio Cacioli[3], Federico Ranocchi[3], Rachele Adorisio[4], Antonio Amodeo[5] and Mariano Antonio Feccia[6]

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P112 | Diagnosis and treatment of humoral rejection in heart-transplant patients: single center experience

Elvira Poggi[1], Annarita Manfreda[2], Silvia Sinopoli[2], Ivan Gabrielli[2], Vincenzo Dinallo[2], Simona Caporali[2], Sabina Maria Bruno[2], Federico Ranocchi[3], Giada Distefano[3], Fabio Sbaraglia[3] and Mariano Antonio Feccia[3]

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P113 | Correlation between DSA and kidney transplant prognosis

Hyun-Ji Lee[1], Shingyeop Hwang[2], Hanbin Jeong[2], Soohyun Lee[2], Junhyun Gu[2] and Kyunghwa Shin[2]

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P114 | Luminex single antigen assay on 1:10 diluted serum predicts the drop in anti-HLA antibodies before desensitization

Dominique Bertrand[1], Charlotte Laurent[1], Fabienne Farce[2] and Rangolie Kaveri[2]

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P115 | A comprehensive comparative assessment of mean fluorescence intensity of Luminex single antigen bead tests between laboratories and commercial platforms; a report from the Italian histocompatibility network

Antonina Piazza[1], Giovanni Rombolà[2], Dario Ciappi[3], Maria Chiara de Stefano[4], Umberto Maggiore[5], Sara Iozzi[6], Andrea Ricci[4], Franco Papola[7] and Massimo Cardillo[4]

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P116 | Exploring false positive reactions in Anti-HLA antibody identification via Single Antigen testing

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P117 | Urine Biomarker CXCL10 Bead-Based Detection Assay for Monitoring Clinical Transplant Rejection

Julie Nguyen[1], Michael Trinh[1], Elizeh Hasan[1], Chien-Yu Chen[1], Rui Pei[1] and Dave Lowe[1]

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P118 | Evaluation of serotype representation in different Luminex single antigen kits regarding the serotype frequencies in 2,021 deceased donors

Renato de Marco[1], Alberto Cardoso Martins Lima[1], João Henrique Campos[1] and Maria Gerbase-DeLima[1]

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P119 | Strategy for delisting prohibited HLA antigens to increase the possibilities of kidney transplantation of hypersensitized patients

Juanfran Luchoro-Cerda[1], José Luis Caro[1], David Cucchiari[1], Juan Torres[1], Esther Mancebo[2], María José Pérez-Saez[3], Marta Crespo[3], Amado Andrés-Belmonte[2], Francesc Moreso[4], Oriol Bestard[4], Fritz Diekmann[1] and Eduard Palou[1]

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P120 | Immunogenetic Profiling in Living Donor Kidney Transplantation: Insights from DSAs and Pronase-Treated Flow Cytometry Crossmatch

Luís Ramalhete[1], Ana Teixeira[2], Cristiana Teixeira[2], Maria Isabel Silva[2], Paula Almeida[2], Alice Lima[2] and Ruben Araujo[3]

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P121 | Optimizing Kidney Re-transplantation Outcomes: Validation of a Highly Sensitive Assay for Monitoring of Donor-Derived Cell-Free DNA

Sofia Westerling[1], Natasa Sikanic[1], Venkat Talla[1] and Linnéa Pettersson[1]
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P122 | HLA-DQA1*03:02-DQB1*03:03 is the dominant immunogenic heterodimer for post-transplant HLA-DQ *de novo* DSA development in a cohort of Chinese kidney transplant patients

Xiangjun Liu[1], Zhaoru Huang[2] and Wenjun Shang[2]
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P123 | Cross-platform optimization of (d)dPCR assays intended for donor-derived cell-free DNA monitoring

Bram Luiken[1], Lara van der Klugt[1], Bo Rood[1], Bart Valkenburg[1], Freek Manders[1], Eleni Draina[1] and Sake van Wageningen[1]
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P124 | Playing Hide and Seek with HLA-DRB3/4/5 Data in Kidney Transplantation Reports in Argentina

Manuel Quirno Costa[1]
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P125 | Analysis of the influence of IgM on the results of CDC-XM in patients with chronic kidney disease

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P126 | Kinetics of donor-derived cell-free DNA in the early kidney post-transplantation phase

Mauro Leucio Mattei[1], Serena Rossinelli[1], Marilù Bartiromo[2], Martina Betti[1], Sara Iozzi[1], Eleonora Pizzo[2], Glenda Cancila[2], Alice Brogi[1], Elisabetta Pelo[1], Lino Cirami[2], Sergio Serni[3] and Simona Palchetti[1]
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P127 | HLA mismatch and circulating donor-specific antibodies predict long term graft loss after kidney transplantation: a retrospective study from Campania region, Italy

Mariagrazia Strozziero[1], Dario Costa[1], Giuditta Benincasa[1], Vincenzo Grimaldi[1], Paride De Rosa[2], Giovanni Valeriani[2], Michele Santangelo[3], Rosa Carrano[3], Sara Pacilio[4], Francesco Cacciatore[4] and Claudio Napoli[1]
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P128 | Performance of Devyser's NGS-based Assay for Monitoring donor-derived cell-free DNA in Kidney Allografted Patients

Annette Plesner[1], Soeren Schwartz Soerensen[1] and Helle Bruunsgaard[1]
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P129 | Effect of serum dilution in the patient with antibody-mediated rejection undergoing therapeutic plasma exchange

Olaida Valdez[1], Maria Grazia Tupone[1], Cervelli Carla[1], Alessandra Panarese[2], Anna Rughetti[3], Carla Battistoni[1], Daniela Pulcinelli[1], Vittoriano Torrelli[1], Ivana Abruscio[1] and Franco Papola[1]
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P130 | The interpretation of the anti-HLA-A3 appearance after treatment with anti-thymocyte globulin

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P131 | CDC vs Luminex in HLA class I antibody detection – 12 years' experience in Tissue Typing Laboratory, Rijeka

Tajana Crnić Marčetić[1], Nataša Katalinić[2], Aida Mujić Franić[1], Helena Kurtović[1], Ines Šimac Sušanj[1], Hrvoje Rimac[1] and Sanja Balen[2]
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P132 | Optimizing a protocol for diluting patient serum with EDTA as a pre-treatment step toward suppressing the prozone effect in the detection of anti-HLA antibodies

Daniel Iguasnia Portilla[1], Ricardo Cuesta[2], Rocío Navarro[3], Manuel Rodríguez[3], Luisa Mas[3], Rafaela Sánchez[3], Isabel Terrón[3], Rosa Granell[3], Estefanía Solaz[3], Dolores Planelles[3], Luis-Hidalgo Mar[3], Castro Emma[3], Luis Larrea[3], Cristina Arbona[3] and Francisco Boix[3]

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P133 | Comparison of Two Single Antigen Bead Assays for Detection of Anti-HLA Antibodies and assessment of their complement-binding capacity

Asimina Fylaktou[1], Georgios Lioulios[2], Georgios Petasis[1], Marianthi Papachristou[1], Margarita Samali[1], Artemis Maria Iosifidou[3], Myrto Aikaterini Iosifidou[3], Aikaterini Anastasiou[1], Maria Stangou[4] and Ioannis Theodorou[5]

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P134 | First experiences with detection of dd-cfDNA in heart transplant patients in North Macedonia

Meri Kirijas[1], Marija Gjerakarovska Radovikj[2], Kristina Stamatovska[1], Boban Dobrevski[1], Gorjan Milanovski[1], Teodora Brnjarchevska Blazevska[1], Tamara Savevska[1] and Aleksandar Petlichkovski[1]

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P135 | MicroRNAs as Potential Graft Rejection or Tolerance Biomarkers and Their Dilemma in Clinical Routines Behaving like Devilish, Angelic, or Frightening Elements

Marina Fernandez-Gonzalez[1], Maria Jose Alegria-Marcos[1], Santiago Llorente[1], Rosana Gonzalez-Lopez[1], Jose Antonio Galian[1], Helios Martinez-Banaclocha[1], Carmen Botella[1], Maria Rosa Moya-Quiles[1], Alfredo Minguela[1], Isabel Legaz[1] and Manuel Muro[1]

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P136 | Validation of the Flow cytometry crossmatch technique with the BD FACSLyric and calculation of new cut-off values

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P137 | Challenges in Immunological evaluation of Four Sensitized Patients

Mahendra Mishra[1] and Aarti Nagpal[1]

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P138 | Analysis of HLA-A, -B and -DRB1 genes in living donor kidney transplantation in Montenegro

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P139 | HLA-antibodies in solid organ transplantation: Eplet analysis of a paired kidney cohort

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IMMUNOTHERAPY, GENE THERAPY & NK CELLS

P140 | NK Cell-Based Adoptive Immunotherapy for Controlling HLA class II Antibody-Mediated Rejection in Transplantation

Carolt Arana[1], Ainhoa Garcia-Busquets[2], Sergi Betriu[3], Eduard Palou[3], Fritz Diekmann[1] and Jordi Rovira[2]

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P141 | Genetic engineering of limbs during *ex vivo* machine perfusion to overcome the unsolved hurdle of rejection
Suganya Annadurai[1], Sevval Besli[1], Tamina Rother[1], Yuliia Yuzefovych[1], Nicco Krezdorn[1], Rainer Blasczyk[1] and Constanca Figueiredo[1]
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P142 | Elucidating the sequence features driving KIR diversification
Jesse Bruijnesteijn[1], Marit van der Wiel[1], Nanine de Groot[1], Ngoc Giang Le[1], Natasja de Groot[1] and Ronald E. Bontrop[1]
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P143 | Cytomegalovirus (CMV) Intra-uterine transmission: involvement of KIR receptors-HLA ligand interactions
Rosalia Cacciatore[1], Annamaria Pasi[1], Chiara Fornara[2], Ilaria Sbarsi[1], Paola Bergamaschi[1], Carmen Tania Prezioso[1], Carolina Radaelli[1], Chiara Bottazzi[1], Enrica Paudice[1], Milena Furione[1], Alessia Arossa[1], Piera D'Angelo[1], Fausto Baldanti[1], Arsenio Spinillo[3], Cesare Perotti[1] and Daniele Lilleri[1]
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P144 | Post-injection monitoring of CAR-T cells targeting CD19 in the treatment of hematologic malignancies: quantitative PCR versus flow cytometry
Judith Desoutter[1], Lucie Thirache[1], Cécilia Da Costa[1], Magalie Joris[2], Lavinia Merlusca[2], Lina Mustapha[3], Christele Ossart[4], Julien Lion[1] and Nicolas Guillaume[1]
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P145 | Short Term Follow up of Soluble Immune Checkpoints in Renal Transplantation
Cemil Pehlivanoğlu[1], Başak Aru[1], Ali Osman Gürol[2] and Gülderen Yanıkkaya Demirel[1]
[1]Faculty of Medicine, Immunology Department, Yeditepe University, Istanbul, Turkey; [2]Department of Immunology, Aziz Sancar Institute of Experimental Medicine, Istanbul University, Istanbul, Turkey

P146 | Analysis of HLA matching between deceased donors and cord blood units from a national bank network as a basis for potential platforms for chimerism-based immune tolerance after solid organ transplantation
Roberto Crocchiolo[1], Letizia Lombardini[2], Nicoletta Sacchi[3], Ilaria Lombardi[2], John Blake[4], David Allan[4], Mohamad Sobh[4], Francesca Puoti[2], Silvia Trapani[2], Anna Maria Gallina[3], Marco Sacchi[5], Simonetta Pupella[6], Paola Bergamaschi[7], Silvano Rossini[1] and Massimo Cardillo[2]
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P147 | Possible Role of KIR3DL2/HLA-A*11:01 Interaction in SARS-CoV-2 Infection Severity
Maria Grazia Tupone[1], Vincenza Cofini[2], Carla Cervelli[1], Raffaella Azzarone[1], Olaida Valdez[1], Maria Scimitarra[1], Barbara Spaziani[1], Stefano Necozione[1], Antonia Petrucci[3], Stefania Melena[4], Michela Falco[5] and Franco Papola[1]
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P148 | Potential co-partnership in predisposition to Covid-19: KIR-HLA class I pairs control NK cells activity and T cells discriminate between HLA-C C1 and C2 groups in Antiviral Immunity
Carmen Tania Prezioso[1], Annamaria Pasi[2], Rosalia Cacciatore[2], Ilaria Sbarsi[1], Carolina Radaelli[1], Paola Bergamaschi[1], Chiara Bottazzi[1], Enrica Paudice[1], Federica Zavaglio[1], Sara Bozzini[1], Federica Meloni[1], Lorenzo Cavagna[1], Fausto Baldanti[3], Daniele Lilleri[1] and Cesare Perotti[1]
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P149 | KIR haplotyping using allele-level KIR typing results from NGSengine
Max Diekman[1], Sam Stokman[1], Freek Manders[1], Sake van Wageningen[1], Bram Luiken[1] and Loes van de Pasch[1]
[1]GenDx, Netherlands

P150 | TLR1 gene polymorphism in SARS-CoV-2 bilateral pneumonia
Alexander Evdokimov[1], Ekaterina Peredelskaya[1], Daria Stashkevich[1], Svetlana Belyaeva[2], Mikhail N. Vavilov[2] and Tatiana Suslova[2]
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P151 | Allele-level characterization of KIR gene polymorphism in healthy elderly populations from Bulgaria, Romania and Turkey

Bushra Al Hadra[1], Tsvetelin Lukanov[1], Ileana Constantinescu[2], Fatma Oguz[3], Dimitri Apostol[2], Yeliz Ogret[3], Anastasiya Mihaylova[1] and Elissaveta Naumova[1]

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P152 | Distribution of KIR genes in a Romanian cohort

Andreea Mirela Caragea[1], Radu Ioan Ursu[1], Alexandra Elena Constantinescu[1], Ion Maruntelu[1], Adriana Talangescu[1], Mirela Maria Iacob[2], Maria Tizu[1], Alexandra Cojocaru[1], Radu Alexandru Truica[1], Larisa Denisa Visan[2], Mariana Steluta Surugiu[2], Corina Rotarescu[2] and Constantinescu Ileana[2]

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P153 | Killer cell immunoglobulin-like receptor genes and their HLA ligands in southern Tunisia

Sirine Louati[1], Aida Charfi[1], Imen Daoud[1], Lilia Gaddour[1], Faiza Hakim[1], Ines Kammoun[1], Bakhta Mallek[1], Fadia Oualha[1], Arwa Kamoun[2] and Nadia Mahfoudh[1]

[1]Immunology and Histocompatibility Department, University Hedi Chaker Hospital, Sfax, Tunisia; [2]Immunology Department, Renal pathology research laboratory LR19ES11, University Hedi Chaker Hospital, Sfax, Tunisia

P154 | KIR/HLA Pairs in Southern Tunisia

Sirine Louati[1], Aida Charfi[1], Imen Daoud[1], Lilia Gaddour[1], Faiza Hakim[1], Ines Kammoun[1], Bakhta Mallek[1], Fadia Oualha[1], Arwa Kamoun[2] and Nadia Mahfoudh[1]

[1]Histocompatibility and Immunology Laboratory, Hedi Chaker Hospital, Tunisia; [2]Immunology Department, Renal pathology research laboratory LR19ES11, University Hedi Chaker Hospital, Sfax, Tunisia

MHC EVOLUTION, POPULATION GENETICS

P155 | Conserved extended 8.1 ancestral haplotype – the polymorphism beyond it

Marija Maskalan[1], Danijela Svilicic[1], Katarina Stingl Jankovic[1], Marija Burek Kamenaric[1], Zorana Grubic[1] and Renata Zunec[1]

[1]University Hospital Centre Zagreb, Croatia

P156 | The strength of the humoral immune response to mRNA SARS-CoV-2 vaccination is influenced by HLA type of the vaccinee

Sendi Montanic[1], Sabina Kunilo Jamnik[1], Sonja Vuletić[1], Urška Rahne Potokar[1] and Blanka Vidan Jeras[1]

[1]Blood Transfusion Center of Slovenia, Slovenia

P157 | Allelic HLA-DPA1 ~ DPB1 haplotype analysis in a large Chinese population of 584 families

Xiangjun Liu[1], Yanling Zhang[2] and Junbo He[3]

[1]BFR Diagnostics, China; [2]Department of Hematology, Hebei Yanda Lu Daopei Hospital, Langfang, Hebei, China; [3]The First Affiliated Hospital of Zhengzhou University, Zhengzhou, China

P158 | Complement MHC Bf alleles show trans species evolution between man and chimpanzee

Antonio Arnaiz-Villena[1], Ignacio Juarez[1], Christian Vaquero-Yuste[1], Tomas Lledo[1], Jose Manuel Martin-Villa[1] and Fabio Suarez-Trujillo[1]

[1]Dept Immunology, University Complutense, Medical School, Madrid, Spain

P159 | Genotypic Frequencies of Mutations Associated with Alpha-1 Anti-trypsin Deficiency in Unrelated Bone Marrow Donors from the Murcia Region Donor Registry in the Southeast of Spain

Irene Cuenca[1], Carmen Botella[1], Maria Rosa Moya-Quiles[1], Jose Antonio Galian[1], Helios Martinez-Banaclocha[1], Marina Fernandez[1], Alfredo Minguela[1], Isabel Legaz[2] and Manuel Muro[1]

[1]Immunology Service. University Clinical Hospital "Virgen de la Arrixaca". Murcia. Spain, Spain; [2]Department of Legal and Forensic Medicine, University of Murcia (UMU), Murcia, Spain

P160 | High-throughput genotyping of HLA-E, HLA-F and HLA-G and analysis of allele frequency distributions in Croatia

Lucija Jukic[1], Marija Maskalan[1], Marija Burek Kamenaric[1], Katarina Stingl Jankovic[1], Zorana Grubic[1], Sanja Salamunovic[1] and Renata Zunec[1]

[1]University Hospital Centre Zagreb, Tissue Typing Centre Zagreb, Croatia

P161 | HLA association study with bipolar disorder in South Tunisian population

Nadia Khlif[1], Fatma Guermazi[2], Aida Charfi[1], Rihab Ouali[2], Arwa Kamoun[1], Ines Feki[2], Faiza Hakim[1], Lilia Gaddour[1], Ines Kammoun[1], Fedya Walha[1], Jawaher Masmoudi[2] and Nadia Mahfoudh[1]
[1]Histocompatibility Laboratory, Hedi Chaker Hospital, Sfax, Tunisia; [2]Psychiatry A Department, Hedi Chaker University Hospital, Sfax, Tunisia

P162 | HLA-B, -C, -DRB1, -DQA1, and -DQB1 alleles involved in genetic predisposition to psoriasis, and psoriasis associated with psoriatic arthritis in East Croatian patients

Saška Marczi[1], Mirjana Suver Stević[1], Anđelka Bugarin[1], Elizabeta Knezović[1] and Marina Samardžija[1]
[1]University Hospital Osijek, Croatia

P163 | Genetic and Anthropological Relationships between Canary Islands and Azores Islands: The Saharo-Canarian Circle

Antonio Arnaiz-Villena[1], Ignacio Juarez[1], Christian Vaquero-Yuste[1], Tomas Lledo[1] and Fabio Suarez-Trujillo[1]
[1]Department of Immunology, Medicine School, University Complutense of Madrid, Spain

P164 | The influence of HLA-DRB1 and HLA-DQB1 on post-transfusion alloimmunization with red blood cell antigens in Polish population

Agnieszka Rzeszotarska[1], Anna Piotrowska[1] and Jolanta Korsak[1]
[1]Military Institute of Medicine, Poland
*blood cells (multi-responders). Additionally, a more frequent combined occurrence of HLA-DRB1*07:01 and DRB1*15:01 alleles was found in the group of people producing multi-specific antibodies to red blood cells (multi-responders).*

P165 | HLA-G 3´UTR 14bp indel polymorphism in Southern Tunisian Bipolar Disorder patients

Aida Charfi[1], Nadia Khlif[1], Fatma Guermazi[2], Imen Daoud[1], Rihab Ouali[2], Ines Feki[2], Lilia Gaddour[1], Faiza Hakim[1], Fedya Walha[1], Ines Kammoun[1], Jawaher Masmoudi[2], Arwa Kamoun[1] and Nadia Mahfoudh[1]
[1]Histocompatibility and immunology laboratory, Hedi Chaker Hospital, Tunisia; [2]Psychiatry A Department, Hedi Chaker University Hospital, Sfax, Tunisia

P166 | Managing Possible New Alleles in Immunogenetics Laboratory of the Portuguese Institute of Blood and Transplantation, Porto

Maria Peixoto[1], Susana Oliveira[1], Filomena Mendes[1], Vasco Guerra[1], Rita Ferreira[1], Manuel Dias[1] and Fátima Freitas[1]
[1]Portuguese Institute of Blood and Transplantation of Porto, Portugal

P167 | Linkage disequilibrium between MICA-129Met/Val and HLA-C1/C2 in the Russian population of the Chelyabinsk region of the Russian South Urals

Mikhail N. Vavilov[1], Tatiana Suslova[1] and Alexandra L. Burmistrova[1]
[1]Chelyabinsk State University, Russia

P168 | HLA alleles and haplotypes in a Sudanese population and their relationship with Mediterraneans and East to West demic diffusion

Fabio Suarez-Trujillo[1], Sayda El-Safi[2], Christian Vaquero-Yuste[1], Tomas Lledo[1], Ignacio Juarez[1], Jose Manuel Martin-Villa[1] and Antonio Arnaiz-Villena[1]
[1]Dept Immunology, University Complutense, Medical School, Madrid, Spain; [2]Ibn Sina Specialized Hospital, Khartoum, Sudan

P169 | Autonomic Study and Influence of UV Radiation on Evolution

Carlos Parga-Lozano[1]
[1]Research and Innovation Center CIIS and Fundacion Universitaria del Area Andina, Colombia

P170 | Identification of HLA alleles involved in immune thrombotic thrombocytopenic purpura patients from Turkey

Cevat Ilteris Kikili[1], Demet Kivanc Izgi[2], Damla Ortaboz[3], Hayriye Senturk Ciftci[2], Mustafa Nuri Yenerel[4], Meliha Nalcaci[4], Muhlis Cem Ar[3], Fatma Savran Oguz[2] and Sevgi Kalayoglu Besisik[4]
[1]Istanbul University, Istanbul Faculty of Medicine, Department of Internal Medicine, Turkey; [2]Istanbul University Medicine of Faculty Department of Medical Biology, Turkey; [3]Istanbul University-Cerrahpaşa, Cerrahpaşa Faculty of Medicine, Department of Hematology, Turkey; [4]Istanbul University, Istanbul Faculty of Medicine, Department of Adult Hematology, Turkey

P171 | The frequency of HLA-B*78:01 allele in the southern Tunisian population

Aida Charfi[1], Arwa Kamoun[1], Imen Daoud[1], Faiza Hakim[1], Ines Kammoun[1], Bakhta Mallek[1], Lilia Gaddour[1] and Nadia Mahfoudh[1]
[1]Immunology and histocompatibility Department, University Hospital Hedi Chaker, Sfax, Tunisia

P172 | Distribution of HLA antigens among patients with acute myeloid leukemia in the Republic of Kazakhstan

Aida Turganbekova[1], Zhulduz Zhanzakova[1], Dana Baimukasheva[1], Zhazira Saduakas[1], Didara Khamitova[1] and Sanya Abdrakhmanova[1]
[1]Republican Scientific and Production Center for Transfusiology, Astana, Kazakhstan

P173 | HLA-A, -B, -C, -DRB1, and -DQB1 allele and haplotype frequencies: An analysis of umbilical cord blood units at the Sichuan Cord Blood Bank

Qiang Chen[1], Xingjie Li[2] and Jue Wang[3]
[1]Institute of Blood Transfusion, CAMS & PUMC and Sichuan Cord Blood Bank, China; [2]Sichuan Cord Blood Bank, China; [3]Institute of Blood Transfusion, CAMS & PUMC, China

P174 | Identification of three novel HLA class I alleles, A*24:556N, B*55:129, and C*01:02:87, found in Chinese cord blood units

Qiang Chen[1], Xingjie Li[2] and Jue Wang[3]
[1]Institute of Blood Transfusion, CAMS & PUMC and Sichuan Cord Blood Bank, China; [2]Sichuan Cord Blood Bank, China; [3]Institute of Blood Transfusion, CAMS & PUMC, China

P175 | Detection of the HLA A*02:275 allele, which was included in the HLA nomenclature in September 2022, in only one patient scheduled for hematopoietic stem cell transplantation: a single center experience.

Sevim Gönen[1], Cemre Demir[2], Zühre Kaya[3] and Oğuz Söylemezoğlu[3]
[1]Gazi University Faculty of Medicine Tissue Typing (HLA) Laboratory, Turkey; [2]Diagnoseq Tissue Typing (HLA) Laboratory, Turkey; [3]Gazi Universit Faculty of Medicine Department of Pediatric Hematology, Turkey

NEW TECHNOLOGIES IN IMMUNOGENETICS

P176 | Revolutionizing high resolution HLA genotyping for transplantation assessment: validation, implementation and challenges of Oxford Nanopore Technologies' Q20+ sequencing

Dianne De Santis[1], Naser El-Lagta[1], Linh Truong[1], Fredrick Mobegi[1], Laila Gizzarelli[1], Irena Vukovic[1], Scott Bazley[1], Wong Zo Ee[1], Jess Rao[1], Nichola Hugo[1], Jacqueline Dey[1], Doddi Venkat[1] and Lloyd D'Orsogna[1]
[1]Department of Clinical Immunology, PathWest, Fiona Stanley Hospital, Australia

P177 | The development of full gene multiplex PCR based assay for the non-classical HLA-E, HLA-F, HLA-G, HLA-H, MICA, and MICB genes.

Dianne De Santis[1], Naser El-Lagta[1] and Lloyd D'Orsogna[1]
[1]Department of Clinical Immunology, PathWest, Fiona Stanley Hospital. University of Western Australia., Australia

P178 | Exploring ABO-histocompatibility: Luminex assay allows detection and characterization of endothelial-targeted ABO antibodies

Anne Halpin[1], Francis Leier[1], Bruce Motyka[1], Caishun Li[1], Simon Urschel[1], Jean Pearcey[1], Esme Dijke[1] and Lori West[1]
[1]University of Alberta, Alberta Precision Laboratories, Alberta Transplant Institute, Canada;

P179 | Comparative assessment of cytometry by time-of-flight and full spectral flow cytometry based on a 33-color antibody panel

Antonia Schäfer[1], Senan D'Almeida[2], Julien Dorier[3], Nicolas Guex[3], Jean Villard[1] and Miguel Garcia[3]
[1]Geneva University Hospital, Switzerland; [2]Viollier AG, Switzerland; [3]EPFL, Switzerland

P180 | Rapid HLA High-Resolution Genotyping Utilizing Nanopore DNA Sequencing Technology for Allocation of Organs from Deceased Donors

Diamanto Kouniaki[1], Katerina Tarassi[1], Vasiliki Kitsiou[1], Theofilos Athanassiades[1], Aikaterini Gkizori[1], Georgia Galaziou[1] and Alexandra Tsirogianni[1]
[1]Immunology and Histocompatibility Department, Evangelismos General Hospital, Greece

P181 | NGS-Pronto: High throughput, high resolution HLA typing using nanopore sequencing

Eline der Steege[1], Sjoerd Creutzburg[1], Bart Valkenburg[1], Anna Hill[1], Sake van Wageningen[1], Marcel van de Streek[1], Joris Albers[1], Loes van de Pasch[1] and Pascal van der Weele[1]
[1]GenDx, Netherlands

P182 | Validation of NGSTurbo typing with Nanopore sequencing for rapid and accurate HLA typing in deceased donor organ transplantation

Timo Olieslagers[1], Carmen de Voijs[1], Simone van der Linden[1], Mathijs Groeneweg[1], Christina Voorter[1] and Lotte Wieten[1]
[1]Maastricht University Medical Center, Netherlands

P183 | High-resolution HLA genotyping using nanopores – a multicenter study

Stéphane Buhler[1], Maja Nørgaard[2], Rudi Steffensen[3], Kirstine Kløve-Mogensen[4], Bjarne Kuno Møller[2], Rebecca Grossmann[5], Sylvie Ferrari-Lacraz[1] and Claudia Lehmann[5]
[1]Transplantation Immunology Unit and National Reference Laboratory for Histocompatibility, Dep. of Diagnostic, Geneva, Switzerland; [2]Department of Clinical Immunology, Aarhus University Hospital, Denmark; [3]Department of Clinical Immunology, Aalborg University Hospital, Aalborg, Denmark; [4]Dep. of Clinical Immunology, Aalborg University Hospital, Dep. of Clinical Medicine, Aalborg University, Denmark; [5]Laboratory for Transplantation Immunology, University Hospital Leipzig, Leipzig, Germany;

P184 | Contribution of circulating donor-derived DNA in the prediction of kidney transplant rejection

Thibault Pajot[1], Isabelle Top[1] and Vincent Elsermans[1]
[1]CHU de Lille, France

P185 | Overcoming Challenges in KIR Gene Typing: A Third Generation Nanopore Sequencing Approach

Laia Closa[1], Cristina Ambrós[1], Claudia Ulldemolins[2], Francisco Vidal[3] and María José Herrero[1]
[1]Histocompatibility and Immunogenetics Laboratory, Blood and Tissue Bank, Barcelona, Spain; [2]Department of Immunology, Hospital Clínic, Barcelona, Spain; [3]Congenital Coagulopathy Laboratory, Blood and Tissue Bank, Barcelona, Spain

P186 | HLA class I (HLA-A, -B, -C) Single Reaction, Full Gene, Unambiguous Genotyping by Technology Oxford Nanopore Sequencing

Vrushank Makwana[1], Clare Nevin[1], Ruifeng Zhang[1], Rohan Raval[1], Tom Browne[1] and Martin Howell[1]
[1]Service Development Laboratory, NHS Blood and Transplant, Colindale, United Kingdom

P187 | Assessing the Performance of NG-Mix, a Novel Next Generation Sequencing Option developed by the Reagent Production Unit of the French Blood Center

Fabien Francois[1], Béatrice Bardy[1], Philippe Moskovtchenko[1], Margot Lepage[1], Dan-Adrian Luscalov[1], Catherine Giannoli[1], Xavier Fournel[1], Celine Dard[1], Pierre Fournier[1] and Valérie Dubois[1]
[1]French Blood Center (Etablissement Français du Sang, EFS), France

P188 | Assessment of HLA Typing and Post-Transplant Monitoring Combined on a Single NGS Run

Amanda Willis[1], Curt Lind[1], Thomas Morris[1], Andrea Zimmerman[1], Thomas Hsiao[1] and Beata Kmiec[2]
[1]CareDx, United States; [2]CareDx, Sweden

P189 | Validation of AlloSeq cfDNA to detect Donor Derived Cell-Free DNA (dd-cfDNA) from renal transplant recipient (RTR) blood samples extracted using the Promega-Maxwell® RSC ccfDNA LV Plasma Kit

Selda Duman[1], Nicola Thal[2], Madalina Pinzaru[1], Sharon Vivers[3], Sandra Frater[1], Lisa Walsh[1], Franco Tavarozzi[1] and Raymond Fernando[2]
[1]Anthony Nolan Laboratories, London, United Kingdom; [2]Anthony Nolan Laboratories, London, UK and Solid Organ Group, Royal Free Hospital, London, United Kingdom; [3]Anthony Nolan Laboratories, London, UK and UCL Cancer Institute, UCL Campus, London UK, United Kingdom;

P190 | High-Resolution full gene HLA-DRB1 genotyping using Oxford Nanopore long read sequencing technology

Ruifeng Zhang[1], Rohan Raval[1], Clare Nevin[1], Vrushank Makawana[1], Tom Browne[1] and Martin Howell[1]
[1]NHS Blood and Transplant, United Kingdom

P191 | MICA & MICB – Are your samples up to the challenge?

Nicola Brosnan[1], Evelien Little[1], Selda Duman[1], Jade Kally[1], Sharon Vivers[2], Reetinder Grewal[1], Raymond Fernando[3], Lisa Walsh[1] and Franco Tavarozzi[1]
[1]Anthony Nolan Histocompatibility Laboratories, London, United Kingdom; [2]Anthony Nolan Histocompatibility Laboratories, London, UK; UCL Cancer Institute, UCL Campus, London UK, United Kingdom; [3]Anthony Nolan Histocompatibility Laboratories, London, UK; Solid Organ Group, Royal Free Hospital, London, United Kingdom

P192 | High Resolution HLA Typing Comparative Analysis of Blood Stem Cell and Solid Organ Donors and Recipients with Next Generation Sequencing and Real Time PCR

Chryso Pierides[1], Andri Papaloizou[1], Georgios Kallis[1], Maria Kyriakou[1], Efi Streppa[1] and Paul Costeas[1]
[1]The Karaiskakio Foundation, The Center for the Study of Haematological Malignancies, Cyprus;

P193 | Nanopore kit for the detection of new alleles
Pascal Pedini[1], Coralie Frassati[1], Agnès Basire[1], Maxime Raz[2], Jacques Chiaroni[1] and Christophe Picard[1]
[1]Immunogenetics Laboratory, Etablissement Français du Sang, Marseille, France; [2]Service de Transfusion Sanguine, Centre Hospitalier de Nouméa, Noumea, New Caledonia

P194 | Nanotype and nanopore sequencing – a rapid and reliable technology for routine HLA typing
Ana Moise[1], Ruxandra Caisan[1] and Irina Monica Dutescu[1]
[1]National HLA Laboratory, National Institute of Blood Transfusion Prof dr. C. T Nicolau, Bucharest, Romania

P195 | Deceased Donor HLA typing by Next Generation Sequencing – Ready for Prime time!
Cathi Murphey[1], Fernando Garcia[1], Ellie Saunders[1], Cody Murray[1] and Shannon Mesa[1]
[1]Southwest Immunodiagnostics, Inc., United States;

P196 | Comparing MFI and MCI in single antigen HLA antibody detection on alternative platforms (Luminex vs HISTO SPOT microarrays)
Murielle Verboom[1] and Kristin Launhardt[2]
[1]Institute of Transfusion Medicine and Transplant Engineering, Hannover Medical School, Hannover, Germany; [2]BAG Diagnostics GmbH, Germany

P197 | Utilization of PreSorb beads in removing nonspecific reactivities in HLA antibody test by Luminex solid phase single antigen beads assay: a single center experience
Zeying Du[1], Charles Williams[1], Rachel Bosai[1] and Mohamed Elrefaei[1]
[1]Mayo Clinic Florida, United States

P198 | Rapid and accurate monolocus HLA typing using Oxford Nanopore Technology: new routine for the "HLA and disease" activity at Bordeaux University Hospital
Mamy Ralazamahaleo[1], Océane Tenchon[2], Elodie Wojciechowski[1], Marine Cargou[1], Jonathan Visentin[2] and Gwendaline Guidicelli[1]
[1]Laboratoire d'Immunologie et Immunogénétique, Groupe Hospitalier Pellegrin, CHU de Bordeaux, France; [2]Laboratoire d'Immunologie et Immunogénétique, CHU de Bordeaux et CNRS, ImmunoConcEpT, Université de Bordeaux, France

P199 | Genotyping of six HLA-G SNPs by oligonucleotide ligation assay: optimization of the technique
Imen Daoud[1], Sirine Louati[2], Aida Charfi[2], Lilia Gaddour[2], Faiza Hakim[2], Ines Kammoun[2], Fadia Oualha[2], Bakhta Mallek[2], Arwa Kamoun[3] and Nadia Mahfoudh[2]
[1]Hedi Chaker Hospital of Sfax, Tunisia; [2]Immunology and Histocompatibility Department, University Hedi Chaker Hospital, Sfax, Tunisia; [3]Histocompatibility Department, Renal Pathology Research Laboratory LR19ES11, Hedi Chaker Hospital, Sfax, Tunisia;

P200 | Assessment of automated cell-free DNA extraction utilizing Promega Maxwell
Amanda Willis[1], Thomas Morris[1], Curt Lind[1] and Thomas Hsiao[1]
[1]CareDx, United States

P201 | Analytical and clinical validation of the One Lambda™ Devyser Accept cfDNA kit
Pascal Pedini[1], Nisem Cherouat[1], Alizée Sebastian[1], Benjamin Coiffard[2], Agnès Basire[1], Jacques Chiaroni[1], Martine Reynaud-Gaubert[2], Coralie Frassati[1] and Christophe Picard[1]
[1]Immunogenetics Laboratory, Etablissement Français du Sang, Marseille, France; [2]Lung Transplant Department, APHM, Marseille, France

P202 | Validation of NGS and intermediate resolution methods for HLA typing at the Laboratorio Nacional de Inmunogenética – INCUCAI of Argentina
Pablo Galarza[1], Maria Fernanda Yaunguzian[1], Lucas Monzon[1], Maria Belén Balekjian[1], Laura Aguerre[1], Cecilia Delfino[1], Richard Malan[1] and Carlos Soratti[1]
[1]INCUCAI, Argentina

P203 | NanoTYPE – From single locus testing to High Throughput Multiplexing in 96 format
Celine Dard[1], Gregory Werner[2] and Mathieu Dewez[3]
[1]EFS Auvergne Rhône Alpes, France; [2]Omixon Biocomputing Ltd, Switzerland; [3]Omixon Biocomputing Ltd, France

P204 | Evaluation of LabScreen PreSorb to proficiently remove pan-reactive DR antibody reactivity
Aisling O'Brien[1], Catherine Owens[1], Patricia Mullany[1], Joseph Kelly[1] and Mary Keogan[1]
[1]National Histocompatibility & Immunogenetics Service for Solid Organ Transplant, Beaumont Hospital, Dublin, Ireland

P205 | Detection and characterization of six novel HLA alleles by next-generation sequencing in a Spanish population during the last year

Amalia Tejeda[1], Jairo Eduardo Niño-Ramírez[1], Daniel Arroyo-Sánchez[1], Antonio Balas[2], Isabel Jiménez Hernaz[1], Pilar Terradillos-Sánchez[1], Ariadna Vicente Parra[1], Beatriz García Martín[1], Yasmín Roldán[1], Ana Balanzategui[1], Miguel Alcoceba[1], Ramón García Sanz[1] and Francisco Javier Gil-Etayo[1]

[1]Laboratorio de HLA-Biología Molecular, Servicio de Hematología, Hospital Universitario de Salamanca, Spain; [2]Histocompatibilidad, Centro de Transfusión de la Comunidad de Madrid, Spain;

P206 | Deciphering alloreactivity: an educational website tailored to teach and learn alloreactivity.

Adèle Dhuyser[1], Cassandra Michel[2] and Alice Aarnink[1]

[1]HLA and Histocompatibility Laboratory, CHRU de Nancy & IMoPA6, UMR7365 CNRS, Université de Lorraine, Nancy, France; [2]Université de Lorraine, Vandoeuvre-les-Nancy, France

SATELLITE SYMPOSIA

WERFEN, MAY 21, 2024, 12:10–12:50, HALL A
Title: Virtual crossmatch in Eurotransplant: A one-year experience

werfen

Cynthia Kramer, PhD
Post-doc, Transplantation Immunology group, Eurotransplant Reference Laboratory, Dept. of Immunology, Leiden University Medical Center, The Netherlands

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CAREDX, MAY 21, 2024, 13:10–13:50, HALL B
Title: Transplantation Excellence: Innovative Solutions for Enhanced Patient Care

CareDx®
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Implementation of NGS-Based HLA typing, AlloSeq Tx Assay, for Donor Registry in Saudi Arabia
Abdullah N. Alsuwaidan, MD FCAP
Dept. of Pathology & Laboratory Medicine, King Faisal Specialist Hospital & Research Center, Riyadh, Saudi Arabia

Long-Term Integrity of Transplanted Stem Cells in Human Recipients
Mirjam Belderbos, MD, PhD
Princess Máxima Center for Pediatric Oncology, Utrecht, The Netherlands

Putting dd-cfDNA into Practise: Clinical Experience from the Hospital Clínic of Barcelona
David Cucchiari, MD, PhD
Renal Transplant Unit, Hospital Clínic, Barcelona, Spain

OMIXON, MAY 21, 2024, 17:30–18:10, HALL A

OMIXON

BIG STEP TOWARDS DIAGNOSTICS WITH NANOPORE SEQUENCING
Dr. Claudia Lehmann
University Hospital Leipzig, Germany, Laboratory for Transplantation

DONOR-DERIVED cfDNA AS DIAGNOSTIC TOOL IN KIDNEY TRANSPLANTATION
Karin Boer, PhD
Erasmus MC Transplant Institute, University Medical Center Rotterdam, Rotterdam, The Netherlands

INTRODUCTION OF NANOTYPE IN THE HLA TYPING ROUTINE ACTIVITY OF A FRENCH HISTOCOMPATIBILITY IMMUNOGENETICS LABORATORY
Romain FERRU-CLEMENT, PhD
Immunogenetics Laboratory, French Blood Centre, Site of Poitiers, France

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HANSA BIOPHARMA, MAY 21, 2024, 17:30–18:10, HALL C
Title: From delisting to organ acceptance for HLAi kidney transplantation – with or without desensitisation

Chair: Dr. Fadi Haidar
Hopitaux Universitaires Geneve, Geneva, Switzerland

Speaker: Dr. Dave Roelen
Leiden University Medical Centre (LUMC), Leiden, The Netherlands

Speaker: Dr. Jean Mihes
Hopital Purpan – CHU de Toulose, Toulose, France

THERMO FISHER SCIENTIFIC, MAY 22, 2024, 12:10–12:50, HALL A
Title: Post-transplant monitoring in the era of dd-cfDNA and single antigen bead analysis

Jakob Nilsson M.D., Ph.D.
Consultant Physician, Director Transplant Immunology, University Hospital Zurich, Switzerland

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GENDX, MAY 22, 2024, 13:10–13:50, HALL A
Title: Timeless Expertise: Innovations in HLA Typing

T. Olieslagers
Biomedical Scientist at Maastricht University Medical Center

M. Rijkers, PhD
Project Manager R & D GenDx

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3584 CM Utrecht | Utrecht Science Park The Netherlands | +31 (0)30 252 3799 | www.GenDx.com NGS-Pronto® High-Resolution Nanopore HLA typing NGS-Pronto offers amplification and library preparation reagents, complete with 96 barcodes, providing for an easy and efficient workflow. After nanopore sequencing the data can be directly analysed with GenDx software – NGSengine®-Turbo. Learn more at www.gendx.com/product_line/ngs-pronto/ Our products GenDx develops and offers a comprehensive line of reagents and services, analysis software, and education. NGS Chimerism Monitoring experts in transplant diagnostics Education Alexander Numan Building Yalelaan 48 3584 CM Utrecht the Netherlands Phone: +31 30 2523799 Cumberland Metro Office Park 5521 N. Cumberland Avenue Suite 1116 Chicago, IL 60656 USA Phone: +1 312 815 5006 info@gendx.com www.GenDx.com COMING SOON A

Thermo Fisher Scientific

We take pride in our Mission: We enable our customers to make the world healthier, cleaner and safer Customers worldwide trust Thermo Fisher Scientific products and services to help them accelerate innovation and enhance productivity. Together, we are advancing science to make a real dierence. We do that by providing an unmatched combination of innovative technologies, purchasing convenience and comprehensive support through these and other product and service brands: Discover more © 2024 Thermo Fisher Scientific Inc. All rights reserved. All trademarks are the property of Thermo Fisher Scientific and its subsidiaries unless otherwise specified. Information accurate as of posting date, 07-February-2024. R&D investment is calculated based on 2023 figures. Fact sheet >1,000,000 products >120,000 colleagues $1.3B/YR invested in R&D >$40B in revenue The Transplant Diagnostics Division of Thermo Fisher Scientific As part of Thermo Fisher Scientific, we oer products and services to advance the science of transplant diagnostics. Our solutions help transplant labs and clinical teams provide personalizedcare across the patient experience. Dr. Paul Terasaki founded One Lambda™ in 1984 with the goal of developing tests to improve how laboratories match and monitor transplant recipients and donors. Since then, our HLA typing and antibody detection assays have been trusted by transplant laboratories worldwide. Clinicians also rely on data from our patient monitoring tests to make important decisions about post-transplant therapies. Transplantation is life-changing for recipients, and clinical diagnostics play a vital role in supporting all phases of treatment. Through our commitment to scientific innovation, product quality, patient advocacy, and excellent customer service, we are helping the transplant community raise the

Werfen

Powering Patient Care:

Werfen is a growing, family-owned, innovative company founded in 1966 in Barcelona, Spain. We are a worldwide leader in specialized diagnostics in the areas of Hemostasis, Acute Care Diagnostics, Transfusion, Autoimmunity, and Transplant. Through our Original Equipment Manufacturing (OEM) business line, we research, develop, and manufacture customized assays and biomaterials. We operate directly in 30 countries and in more than 100 territories through distributors. Our Headquarters and Technology Centers are located in the US and Europe, and our workforce is more than 7.000 strong.

SILVER SPONSORS

OMIXON BIOCOMPUTING LTD.

Omixon is a global transplantation diagnostic company with a mission to provide histocompatibility laboratories with innovative technologies to improve transplant outcomes. Omixon is headquartered in Budapest, Hungary, with operations in the United States, Brazil and the Netherlands serving more than 100 laboratories worldwide. Building on multidisciplinary competences in bioinformatics, software engineering, molecular biology and regulatory science, Omixon transforms molecular biology innovations into state-ofthe-art products in transplant diagnostics. Omixon was the first to successfully introduce a next generation sequencing (NGS) based HLA genotyping kit and software in 2014 by bringing Holotype HLA product and HLA Twin software to market. HLA Twin delivers a leading software in high-resolution HLA genotyping and is used in more than 100 laboratories worldwide. Omixon was the first to release a high resolution HLA genotyping product NanoTYPE HLA with less than 5 hours turnaround time on the Oxford Nanopore sequencing platform complemented by NanoTYPER analysis software. In 2024 Omixon introduces HoloGRAFT ONE RUO with a promise of frequent and affordable testing of donor-derived cell-free DNA (dd-cfDNA) levels.

Hansa Biopharma

Hansa Company short description – Mar 2024 Hansa Biopharma is based in Lund, Sweden with operations in other European countries and the U.S. Hansa leverages its proprietary antibody-cleaving enzyme technology platform to target pathogenic and disease causing antibodies. Hansa's lead product, imlifidase, is an antibody-cleaving enzyme which enables kidney transplantation in highly sensitized patients. The European Commission conditionally approved Idefirix (imlifidase) for the desensitization of highly sensitized adult kidney transplant patients with a positive crossmatch against an available deceased donor with use reserved for patients unlikely to be transplanted under available kidney allocation systems including prioritisation programmes for highly sensitised patients.

DiagnoSeq

Diagnoseq Genetic Diseases Evaluation Center (GDEC) as one of the largest „Genomic Service Provider" in Turkey and one of the largest „HLA Tissue Typing" laboratories in Europe offers extensive services either as raw data or as clinical report to its customers. Diagnoseq adopts medical ethics, good laboratory practices, information security and confidentiality as its basic principles. Diagnoseq Tissue Typing Laboratory received EFI (European Federation of Immunogenetics) accreditation in 2018, ISO 27001 certification in 2022 and has also successfully completed the CAP Accreditation (College of American Pathologists) in 2023. Diagnoseq with its advanced technology infrastructure provides short turnaround times to deliver the results in as short as 5 days for NGS based HLA Typing and 10 days for Whole Exome Sequencing (WES) and Clinical Exome Sequencing (CES).

BRONZE SPONSORS

SOCIAL NETWORKING EVENTS

WELCOME COCKTAIL

Monday, May 20, 2024, 19:30–21:30
Place: Exhibition Hall
The Welcome Cocktail is open to all conference participants.

NETWORKING EVENT

Wednesday, May 22, 2024, 19:45–01:00
Price: 105 € / per person
Place: Domain Du Grand Malagny, 48-50 Route de Malagny, 1294 Genthod
Transportation: Transportation from the venue and vice versa will be provided by shuttle buses
Dress code: Business casual/smart casual. Comfortable shoes are recommended

GENDX TULIP RUN (5 KM)

Wednesday, May 22, 2024, from 06:40
Price: 25 € / per person
Meeting point: the main entrance of the Palexpo Geneva at 6:15
Detailed schedule: 06:40 – Warm-up
07:00 – Start run
08:00 – Tulip run awards & heading back to the hotel

CareDx
Your Partner in Transplant Care™

EFI GENEVA 2024

Meet us at Booth #2

Join us at our EFI 2024 Symposium

Margot D., Stem Cell Transplant Recipient

Transplantation Excellence: Innovative Solutions for Enhanced Patient Care

Tuesday 21 May, 2024
Time: 13.10 - 13.50
Hall: B

REGISTER

LIVE SYMPOSIUM NOT ENDORSED BY EFI

Save the date to explore insights into the future of pre- and post-transplant excellence

Moderator

Curtis Lind - VP, Head of R&D Products
CareDx

Guest Speakers

Implementation of NGS-Based HLA typing, AlloSeq Tx Assay, for Donor Registry in Saudi Arabia
Abdullah N. Alsuwaidan, MD, FCAP
Department of Pathology & Laboratory Medicine King Faisal Specialist Hospital & Research Center
Riyadh, Saudi Arabia

Long-Term Integrity of Transplanted Stem Cells in Human Recipients
Mirjam Belderbos, MD, PhD
Princess Máxima Center for Pediatric Oncology
Utrecht, the Netherlands

Putting dd-cfDNA into Practice: Clinical Experience from the Hospital Clínic of Barcelona
David Cucchiari, MD, PhD
Renal Transplant Unit, Hospital Clínic
Barcelona, Spain

AlloSeq Tx AlloSeq HCT AlloSeq cfDNA

AlloSeq cfDNA 5 YEARS CareDx

MAR 141 Version 1.0.

CareDx®

AlloSeq Tx

Better Matching. Better Method.

THE NEXT GENERATION IN GENETIC MATCHING WITH INNOVATIVE HYBRID-CAPTURE TECHNOLOGY

Margot D.,
Stem cell transplant recipient

Expandable Gene Content
without Affecting Lab Workflow

Easy Single Tube Workflow
with Early Indexing Step

No Long-range PCR =
No Amplification Inefficiencies

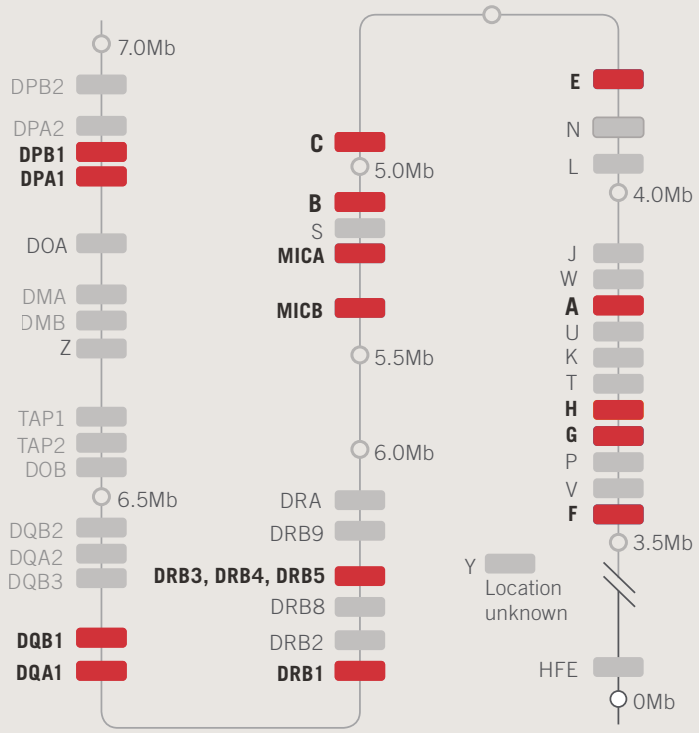
Fast Software Analysis
with Data Upload at 1 min/sample*

*as per internal testing

Availability

AlloSeq Tx9 (HLA-A, -B, -C, -DRB1/3/4/5, DQB1, DPB1) covers classical HLA loci

AlloSeq Tx17 (HLA-A, -B, -C, -E, -F, -G, -H, DRB1/3/4/5, DQA1, DQB1, DPA1, DPB1) moves beyond the traditional transplant related loci to consider more transplant associated genes



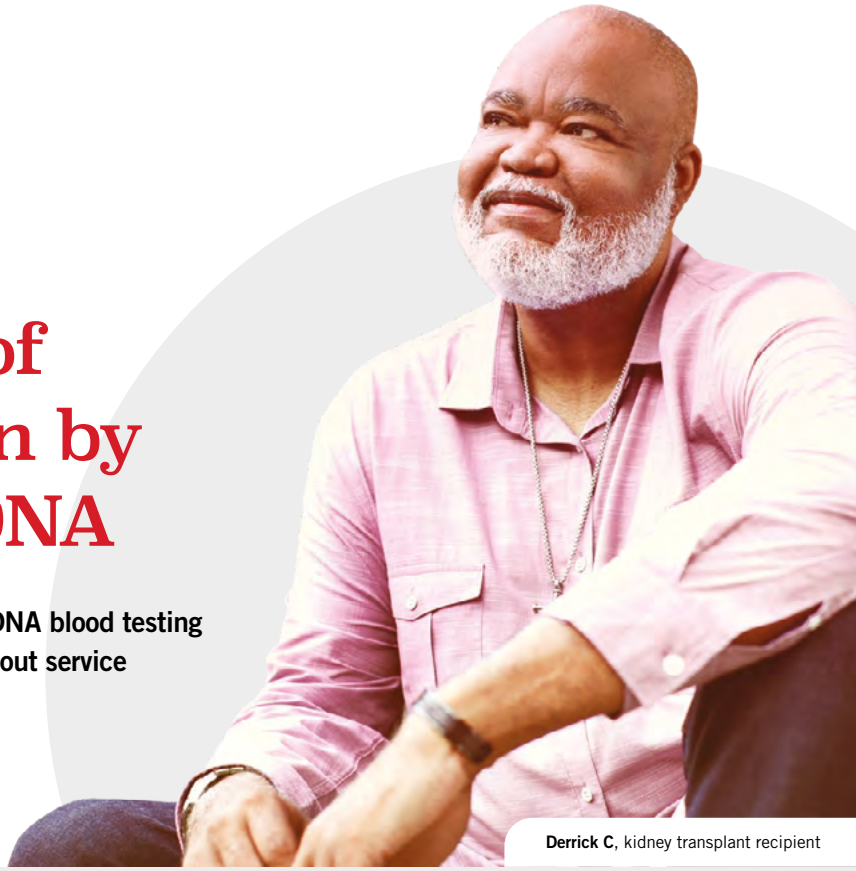
For more information visit **https://CareDx.com/AlloSeqTx** or reach out to your local CareDx representative.

MAR093 Revision 05 Effective 2023-03

AlloSeq cfDNA

Risk Assessment of Allograft Rejection by Measuring dd-cfDNA

An innovative NGS-based solution that enables dd-cfDNA blood testing as a CE-IVD kit for laboratory implementation or send out service



Derrick C, kidney transplant recipient

NGS Based CE-IVD Kit

- + Targets 202 bi-allelic SNPs across 22 autosomes
- + Escalate testing up to 24 samples/run
- + From cfDNA sample to sequencing < 1.5 hrs hand-on time
- + Validated on Illumina MiniSeq and MiSeq
- + No prior genotyping required
- + Multiplexed amplification of targets & sample indexing/barcoding into one single reaction
- + Low cfDNA input of 10ng
- + Ability to monitor more than one donor cfDNA contributor*

Send Out Service

+ Get your results through our high-quality send out service with short turn around time

Collect blood samples in your lab → Send to CareDx → Easy to interpret reports delivered to you

Easy sample collection process | Technical expertise | Full logistics support
No lab setup or NGS equipment required | Cost and resource effective

* Would require recipient genotyping

CareDx®

MAR121 Revision 1 Effective 2023-06

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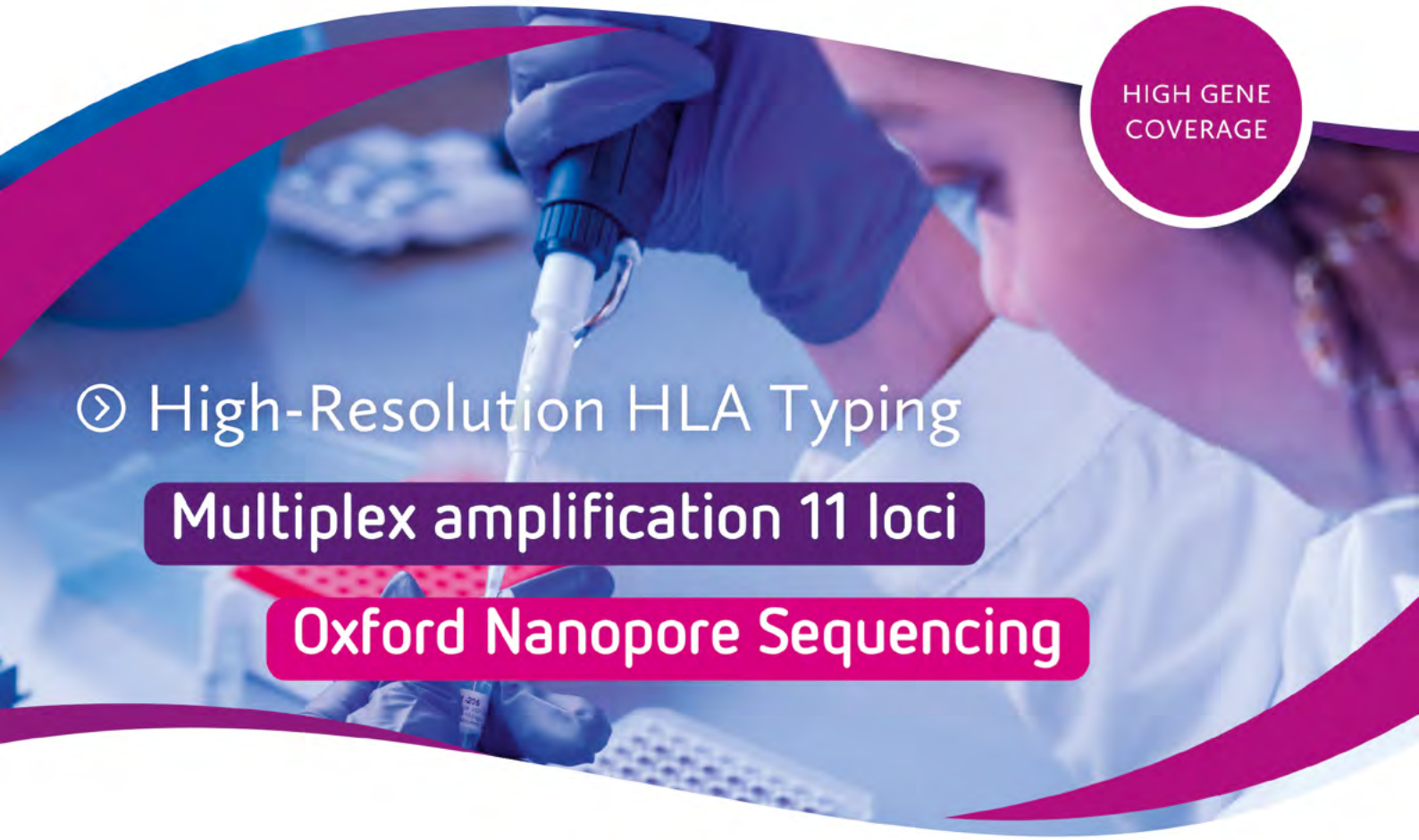
COMING SOON

DISCOVER NGS-Pronto®



High-Resolution HLA Typing

Multiplex amplification 11 loci

Oxford Nanopore Sequencing

Only 15 min. sequencing/sample

Run up to 96 samples at a time

Full phasing No fragmentation

Dedicated software NGSengine®-Turbo

Interested?

Contact us for more information at support@gendx.com

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ORDER NOW

NGS-Turbo®
HIGH-RESOLUTION HLA TYPING WITHIN 3 HOURS

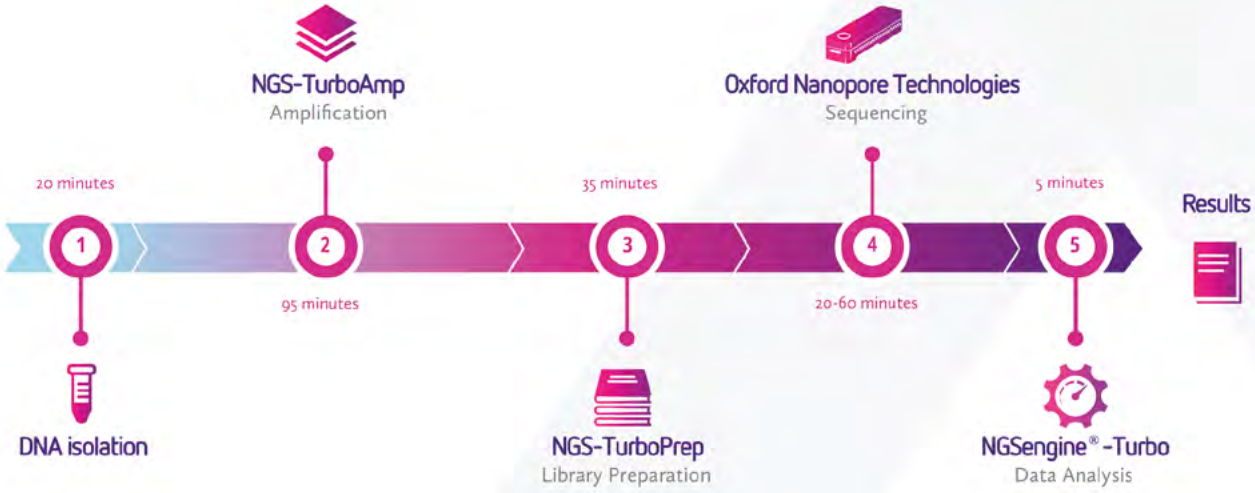
OUR SOLUTION FOR NANOPORE HLA SEQUENCING

Fast and easy

With a turnaround time of less than three hours and an easy and robust workflow, NGS-Turbo® is designed for HLA typing in situations where time is a critical factor. In combination with Oxford Nanopore Technology sequencing devices, accurate sequencing-based HLA typing results are generated at incredible speed.

Workflow simplicity

The NGS-TurboAmp kit offers 11 HLA loci using a whole-gene approach. The extremely quick 1-tube library preparation with NGS-TurboPrep requires no fragmentation and can be safely paused at any time. After nanopore sequencing the data can be directly analyzed with NGSengine®-Turbo.



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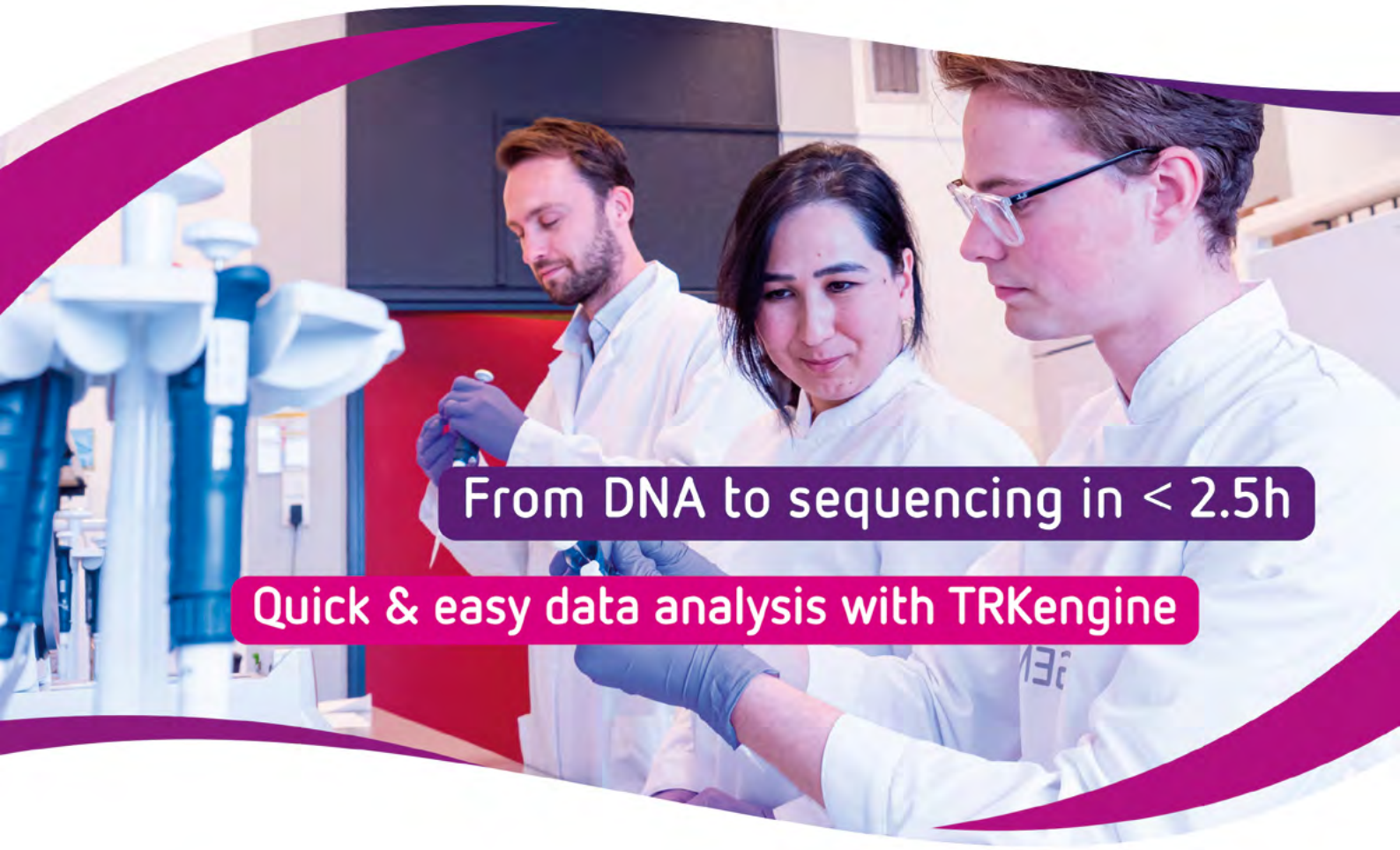
Chimerism monitoring by NGS

NGStrack® & TRKengine®



- Reliable at all chimerism %
- Up to 0.1% sensitivity
- Only 60 ng DNA needed per sample

Interested?

Contact us for more information at support@gendx.com
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Join us at booth #3

We'll be showcasing our assays, protocols and software as our advancements continue to reshape the landscape of HLA genotyping and antibody detection technologies.

Join the Community

Join us at our industry symposia

Wednesday, 22nd May 12.10 – 12.50 | Hall A

Post-transplant monitoring in the era of dd-cfDNA and single antigen bead analysis

Jakob Nilsson M.D., Ph.D.
Consultant Physician, Director Transplant Immunology
University Hospital Zurich

Redefining HLA antibody isolation, bead by bead

This innovative assay isolates individual HLA antibodies from human serum in a targeted manner using adsorption and elution via antigen-coated magnetic beads.

One Lambda™ MagSort™ utilizes a comprehensive panel of 59 HLA antigens across 6 loci, individually coupled to magnetic beads. The ability to elute specific antigens of interest provides a more focused dataset and provides users with a robust and standardized approach.

Capabilities

Targeted technology

- Individual antigen-coated magnetic beads
- Elute antibodies that bind to a specific antigen
- Results in a more focused HLA antibody assessment

Customizable panels

- 59 antigens across 6 locus specific kits
- Flexibility to focus only on antigens of interest
- Coverage across well-documented antigens

Efficient workflow

- Up to 45 minutes of hands-on time and as little as 4 hours to workflow completion
- Uses equipment already found in most laboratories
- Immediate antigen availability.

For more information on this new product, visit the Thermo Fisher Scientific booth

ADAPT.

Are you ready to make a change?

Change can be intimidating, but without change there is limited growth. Differences in MFI values between vendors can be challenging to explain, but they should not be seen as a determent. There are many ways to help overcome MFI differences between vendors and with the right partner and tools you can overcome this barrier. As Darwin said, "It is not the strongest that survive, nor the most intelligent, but the ones that are most adaptable to change."

CLICK HERE to find out what your peers experienced by adapting to change.

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GROWTH.

Looking for ways to advance your HLA antibody analysis knowledge?

Antibody analysis can be complex and overwhelming. With staff turnover it can be difficult to train new employees on analysis creating a greater burden on existing staff. Immucor's MATCH IT!® Antibody software is specifically designed for simplifying the complexities of HLA analysis. Adaptive, effective and powerful, MATCH IT!® Antibody software will turn any tech into an antibody analysis expert.

CLICK HERE to see how you can become an expert.

werfen.com

Rev1-MAR2024

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EVOLVE.

Looking for more solutions to help aid in your decision making process?

How could more tools in your toolbox better aid you in your decision making process for critical transplant patients? When you have more information at your disposal, you can optimize your decision making process for better efficiency and delivery of results. Make the best use of your valuable time by adding LIFECODES Single Antigen Class I and Class II to your workflow.

CLICK HERE to see how a fellow member of the HLA community has accomplished this.

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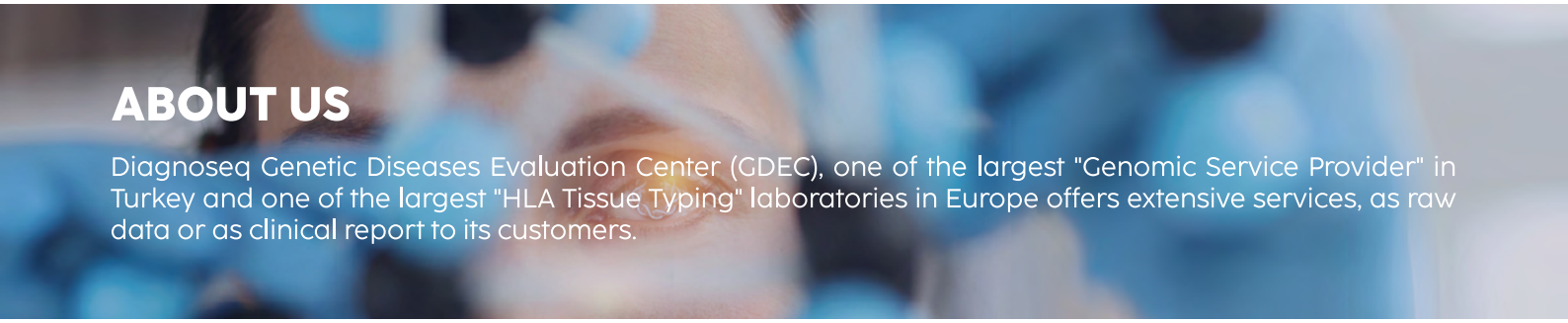
DiagnoSeq
Life of Gene



ABOUT US

Diagnoseq Genetic Diseases Evaluation Center (GDEC), one of the largest "Genomic Service Provider" in Turkey and one of the largest "HLA Tissue Typing" laboratories in Europe offers extensive services, as raw data or as clinical report to its customers.

SERVICES

We provide the highest quality HLA Typing and Genomic services. Our mission, as a high throughput NGS laboratory is to provide the most efficient solutions to our customers for Tissue Typing and other genomics services (WES, WGS, Molecular Karyotyping, Oncology)

We welcome opportunities to cooperate in the areas where you can benefit from our services.

CERTIFICATES

EFI ACCREDITATION

The European Federation for Immunogenetics (EFI) awards the EFI certificate to laborories that the quality requirements set by EFI (Standards). EFI is a European organisation that focuses on immunogenetics, tissue typing and transplantation. The EFI certificate is required by a number of organisations operating in the field of stem cell and solid organ transplantation including JACIE, NMDP and the Eurotransplant foundation.

CAP ACCREDITATION

The CAP accredits laboratories performing testing on specimens from human beings or animals, using methodologies and clinical application within the expertise of the program. CAP Laboratory Accreditation helps laboratories: Maintain accuracy of test results and ensure accurate patient diagnosis. Meet required standards from CLIA, FDA and OSHA. CAP requirements commonly exceed the standards, bolstering patient care and safety.

ISO 15189

ISO 15189 standard; Based on ISO/IEC 17025 and ISO 9001, it aims to ensure safety in medical laboratories, the accuracy and required by patients and clinical staff directly responsible for treatment.

ISO 27001

Internationally recognized ISO/IEC 27001 is an excellent framework for managing and protecting our information assets to keep them safe and secure. In order to protect personal data and sensitive information, ISO/IEC 27001 standard enables implementation of a robust approach to manage information security in our organization.

REFERENCES

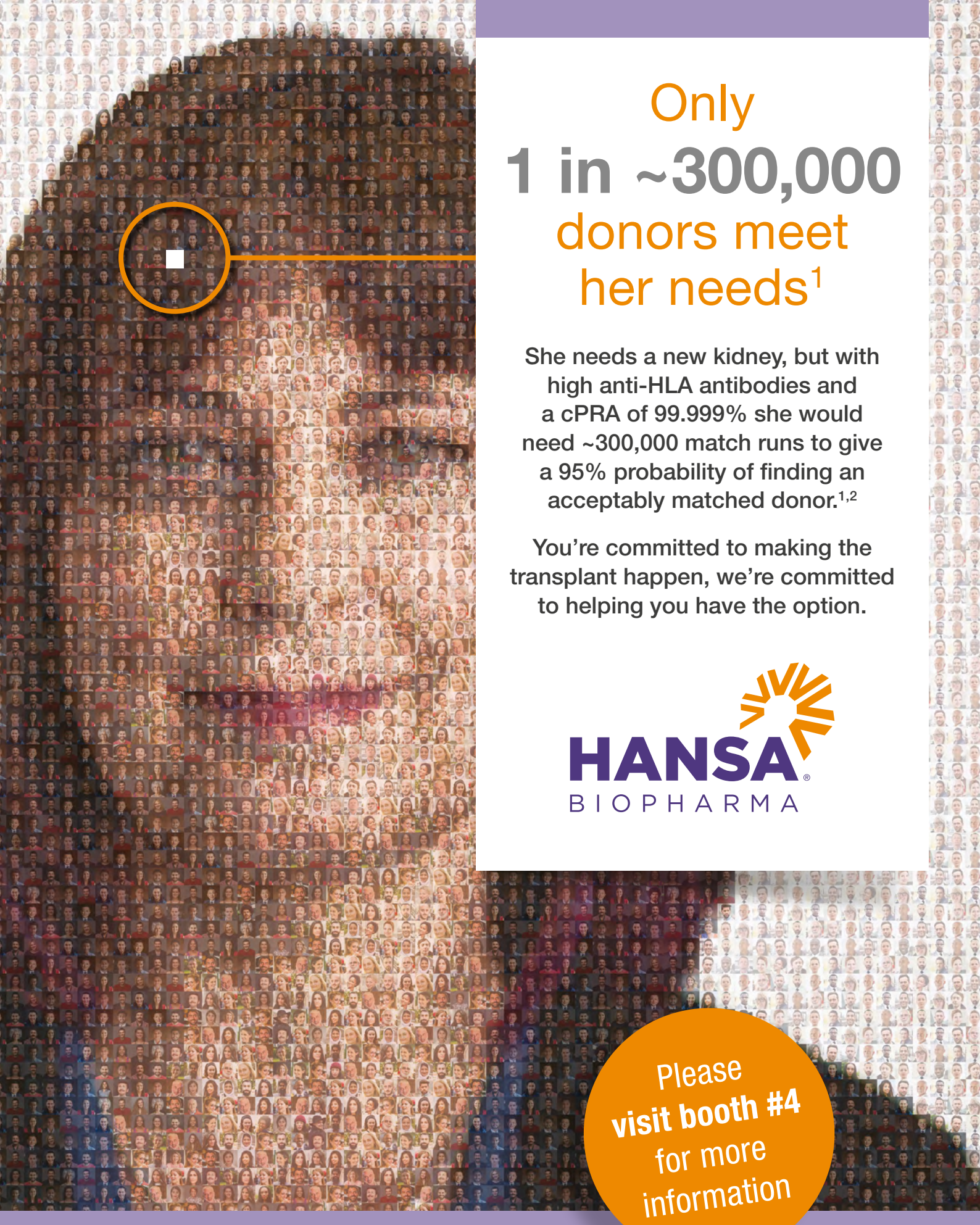
We are proud to provide services to the Donor Registry of Turkiye (TURKOK) and performed more than 1 million HLA Typing.

CONTACT US

DiagnoSeq Genetic Diseases Assessment Center

www.diagnoseq.com.tr **info@diagnoseq.com.tr**

DiagnoSeq



Only 1 in ~300,000 donors meet her needs[1]

She needs a new kidney, but with high anti-HLA antibodies and a cPRA of 99.999% she would need ~300,000 match runs to give a 95% probability of finding an acceptably matched donor.[1,2]

You're committed to making the transplant happen, we're committed to helping you have the option.

HANSA® BIOPHARMA

Please **visit booth #4** for more information

1. Keith DS, et al. Clin J Am Soc Nephrol. 2016;11(4):684–693.
2. Kuppachi S, Axelrod DA. Transpl Int. 2020;33(3):251–259.

This photo has been reproduced with patient approval.

HANSA-PR.GLO.24.001-001 | April 2024

NANOTYPE™ HOLOGRAFT™ omixon

LIVE

NANOTYPE DEMO AT BOOTH #9

CHECK THE DEMO SCHEDULE AT THE OMIXON BOOTH

OMIXON SYMPOSIUM

MAY 21, 2024, 17:30 | HALL A

Big step towards diagnostics with Nanopore Sequencing

Dr. Claudia Lehmann
University Hospital Leipzig, Germany
Laboratory for Transplantation Immunology

Donor-derived cfDNA as diagnostic tool in kidney transplantation

Karin Boer, PhD
Erasmus MC Transplant Institute
University Medical Center Rotterdam
Rotterdam, the Netherlands

Introduction of NanoTYPE in the HLA typing routine activity of a French Histocompatibility and Immunogenetics Laboratory

Romain FERRU-CLEMENT, PhD
Histocompatibility and Immunogenetics Laboratory
French Blood Centre, Site of Poitiers

EFI GENEVA 2024

37TH EFI Conference
May 20-23, 2024

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BAG DIAGNOSTICS

ENJOY THE SYMPHONY OF ANTIBODY DIAGNOSTICS

Be a maestro as you orchestrate your antibody identification, using HISTO SPOT® HLA AB. From diluted serum to results generation, all steps are performed in a **fully automated** instrument; always finely tuned. No need for extra calibration protocols, **saving time, money** and **reagents.** HISTO MATCH software delivers **simple analysis** and **highly specific results** in a virtuoso performance.

HISTO SPOT® HLA AB

- HIGHLY REPRODUCIBLE
- FLEXIBLE THROUGHPUT (1-96 SAMPLES)
- OPTIMIZED SAMPLE BUFFER REDUCES UNSPECIFIC REACTIONS TO A MINIMUM
- CLASS I AND CLASS II - ALL CLINICALLY RELEVANT SEROLOGICAL SPECIFICITIES COVERED

FIND OUT MORE NOW

SAVE THE DATE

38th European Immunogenetics & Histocompatibility Conference

14-17 May 2025, Prague, Czech Republic
Prague Congress Centre

www.efi-conference.org

EFI 2025 Conference, Prague, Czech Republic
38th European Immunogenetics & Histocompatibility Conference