Minimal Requirements for Histocompatibility & Immunogenetics Laboratory

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Medical Director, Blood & Marrow laboratory











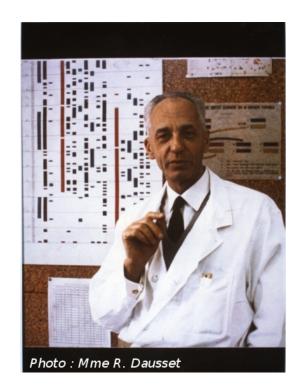
The 4th WBMT Congress and Workshop – Riyadh, KSA - January 15-17, 2017

HLA Discovery, 1958



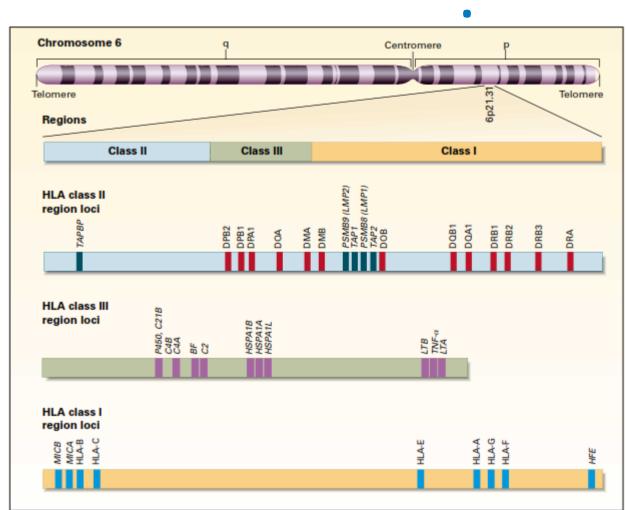
JEAN DAUSSET Prix Nobel de médecine CLIN D'ŒIL À LA VIE LA GRANDE AVENTURE HI A

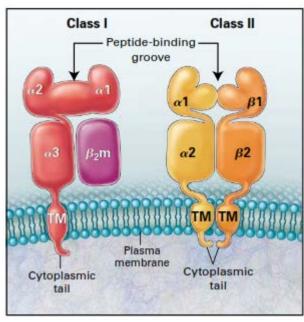
The Nobel Prize in Physiology or Medicine 1980 Baruj Benacerraf, Jean Dausset, George D. Snell



HLA Complex, Chromosome 6

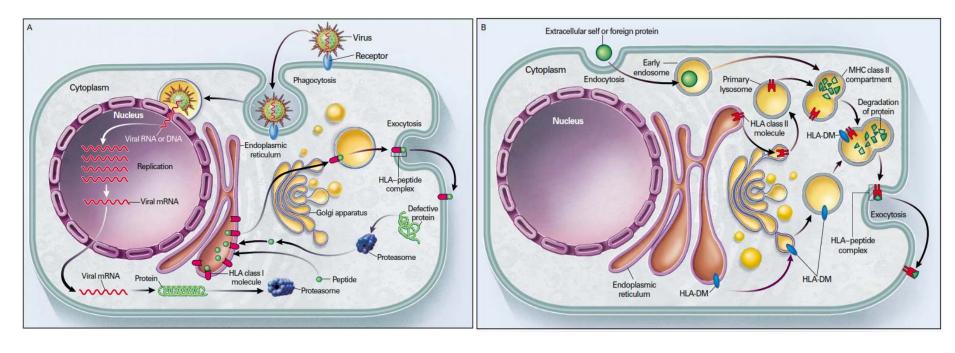






Antigen Processing





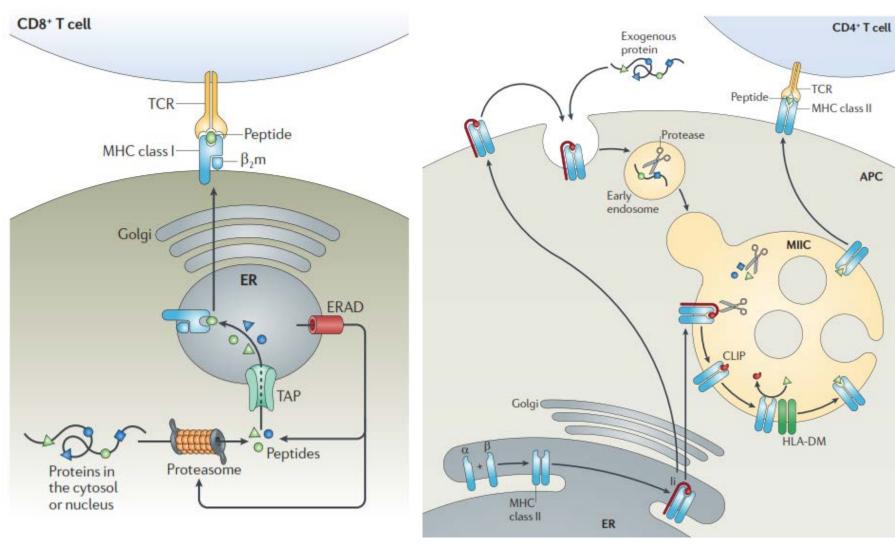
Panel A shows the principal pathways of generating peptides for loading onto HLA class I molecules

Panel B shows the processing of extracellular proteins.

Klein and Sato. NEJM, 2000

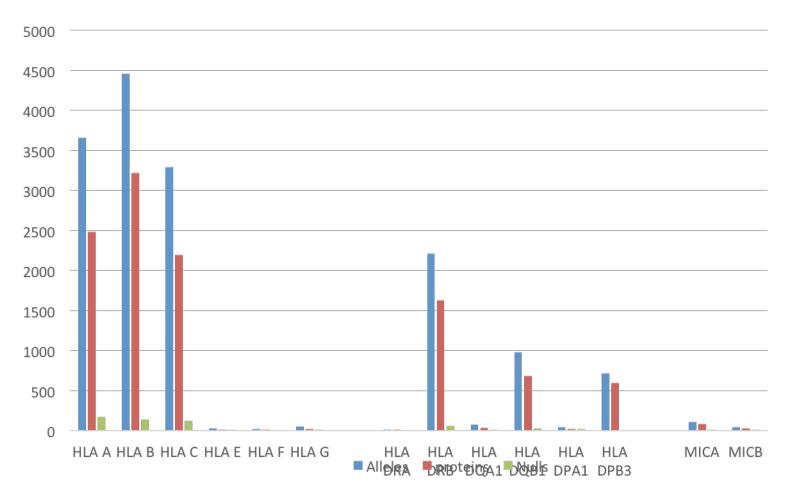
Antigen Presentation





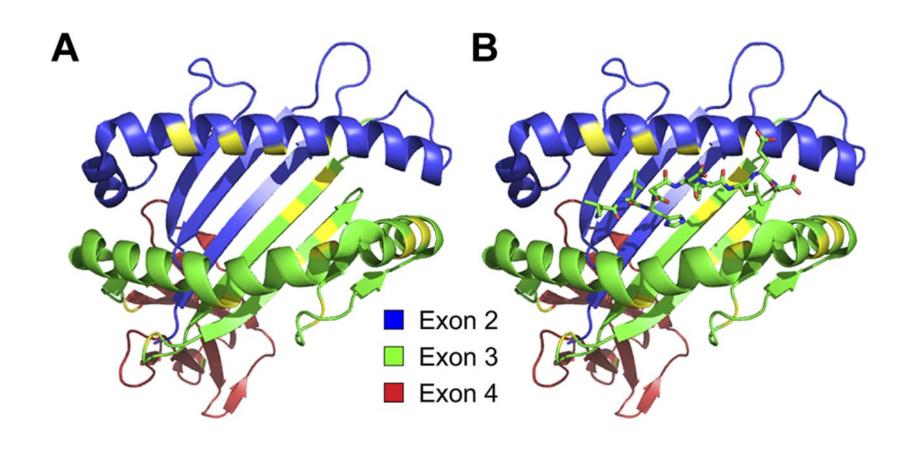
Neefjes et al. Nature Reviews, 2011

The HLA-A, -B, -C, and -DRB1 loci are the most polymorphic genes in the entire human genome



Data were taken from http://www.ebi.ac.uk/imgt/hla/stats.html; accessed January 12, 2017.



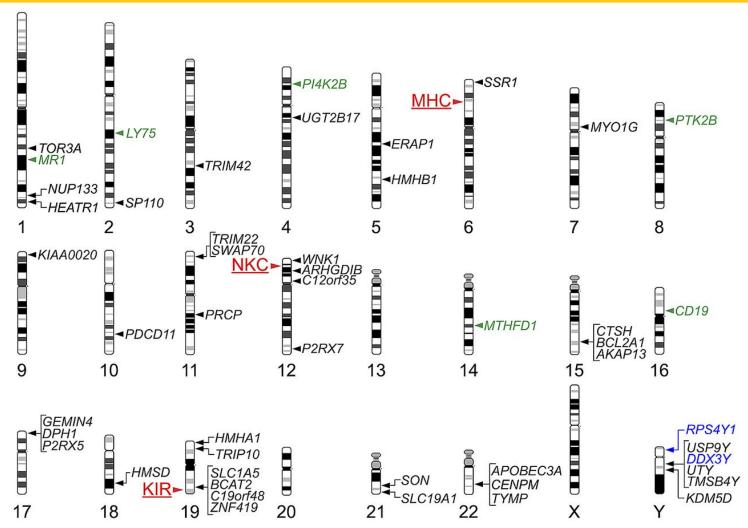


Edus H. Warren et al. Blood 2012;120:2796-2806



Map of genetic loci that can influence histocompatibility in the allogeneic HCT setting





Edus H. Warren et al. Blood 2012;120:2796-2806



Table 1. Diseases Commonly Treated with Hematopoietic Stem-Cell Transplantation.

Autologous transplantation*

Cancers

Multiple myeloma

Non-Hodgkin's lymphoma

Hodgkin's disease

Acute myeloid leukemia

Neuroblastoma

Ovarian cancer

Germ-cell tumors

Other diseases

Autoimmune disorders

Amyloidosis

Allogeneic transplantation†

Cancers

Acute myeloid leukemia

Acute lymphoblastic leukemia

Chronic myeloid leukemia

Myelodysplastic syndromes

Myeloproliferative disorders

Non-Hodgkin's lymphoma

Hodgkin's disease

Chronic lymphocytic leukemia

Multiple myeloma

Juvenile chronic myeloid leukemia

Other diseases

Aplastic anemia

Paroxysmal nocturnal hemoglobinuria

Fanconi's anemia

Blackfan-Diamond anemia

Thalassemia major

Sickle cell anemia

Severe combined immunodeficiency

Wiskott-Aldrich syndrome

Inborn errors of metabolism

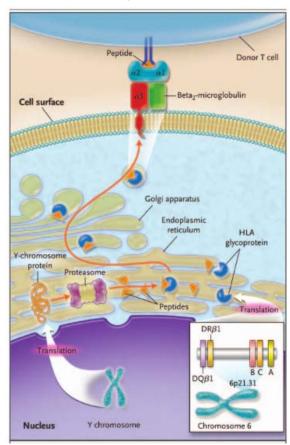
The NEW ENGLAND JOURNAL of MEDICINE

REVIEW ARTICLE

MEDICAL PROGRESS

Hematopoietic Stem-Cell Transplantation

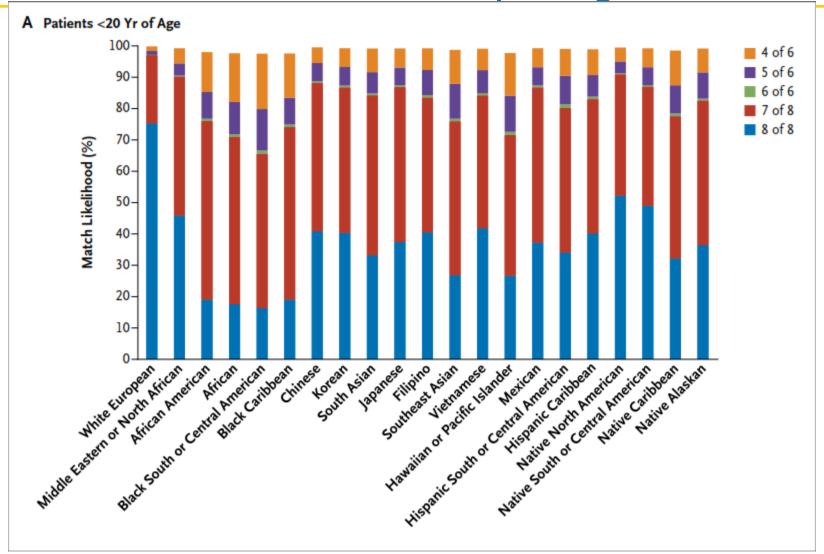
Edward A. Copelan, M.D.



Copelan. NEJM, 2006

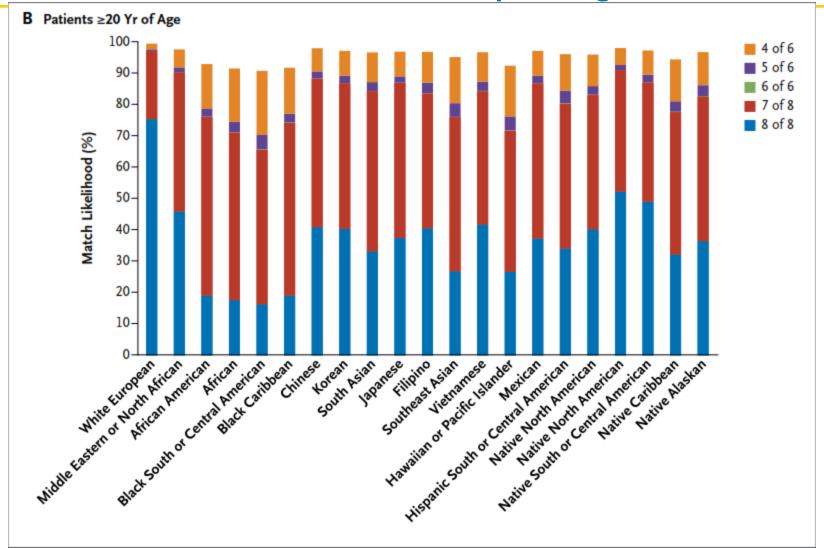
Match Likelihoods According to Racial and Ethnic Group and Age





Match Likelihoods According to Racial and Ethnic Group and Age









Personnel Qualifications

Quality Assurance

Testing Methods

Laboratory Accreditation



Personnel Qualifications

- The laboratory must employ one or more individuals who meet the qualifications and fulfill the responsibilities of the Director, and Technical Supervisor
- The number of staff must be large enough to carry out the volume and variety of tests required



- Facilities
- Specimen submission and requisition
- Laboratory Procedure Manual
- Proficiency Testing and Competency Evaluation
- Records and test reports
- Testing referred to other laboratories



Facilities

- Sufficient space to all procedures carried out
- Proper maintenance of equipments, instruments and test systems (with ample records)
- Laboratories performing amplification of nucleic acids must use physical and/or biochemical barriers to prevent DNA contamination
- Adequate lighting and ventilations



Specimens issues

- The laboratory must have available and follow written policies and procedures regarding specimen collection
- The laboratory must assure adequate information contained in the requisition form
- The laboratory must maintain a system to ensure reliable specimen identification



Laboratory Procedure Manual

- All procedures in use in the laboratory must be detailed in a procedure manual, and immediately available
- Each procedure must be reviewed periodically (according to standards) by the Director and written evidence of this review must be in the manual



Proficiency Testing

 The laboratory must participate in External proficiency Testing (EPT) program (s) to cover all the accredited laboratory applications (HLA typing, antibody screening and identification, crosshatching, etc.)



Competency Evaluation

 The laboratory Director and the technical staff must participate in pertinent continuing education



Records and test reports

 The laboratory must maintain records of subjects tested for a period dictated by national laws and accreditation standards



Records and test reports

- The report must contain:
 - a. Name of the individual tested or unique identifier(s) and relationship to the patient if applicable.
 - b. Date(s) of collection of sample(s)
 - c. Date of the report



Records and test reports

- The report must contain:
 - d. Test results.
 - e. Techniques used.
 - f. Appropriate interpretations and the signature of the Laboratory Director, or, in his/her absence, by a designee



Testing referral

 An accredited laboratory may engage another laboratory to perform testing not done by the primary laboratory

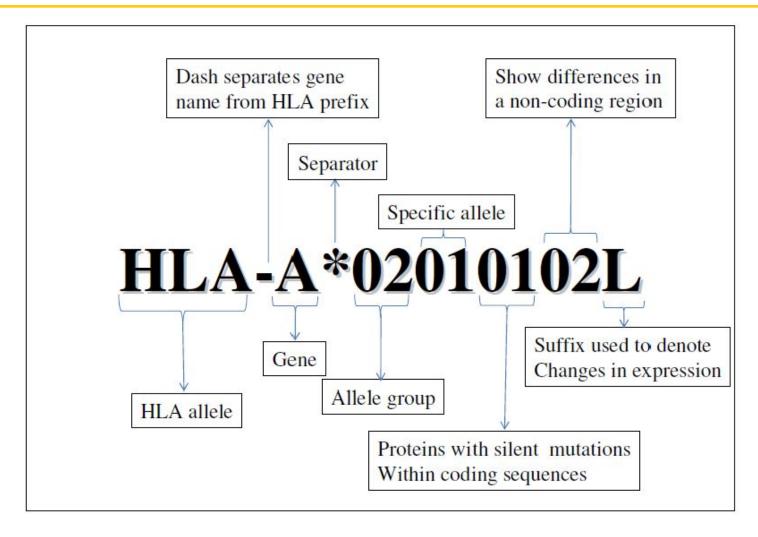


Testing Methods

- Serological vs. Molecular-based HLA typing
 - HLA Nomenclature
 - Resolution level
 - Haplotype assignment
- Chimerism studies

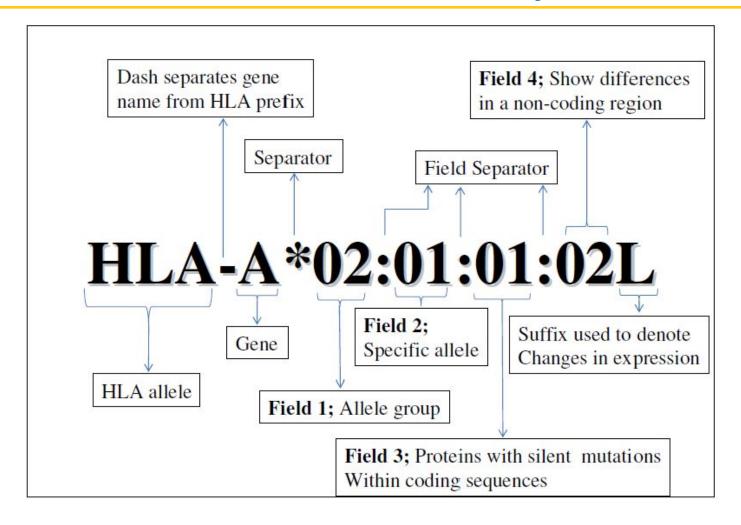
HLA Nomenclature, old





HLA Nomenclature, new





HLA Typing Methods



1950 s	Discovery of HLA system
	 Serological typing
1980 S	 First HLA genes cloned, sequenced
1990 S	 DNA/PCR based HLA typing
	• Sequence entire MHC (HGP)
2000	Database of all HLA alleles
2000 S	• SBT, Luminex SSO
	Next Generation Sequencing

Levels of Resolution



Low resolution (2 digit)

 same serotype group (e.g. A*02)

Intermediate resolution

(allele string)

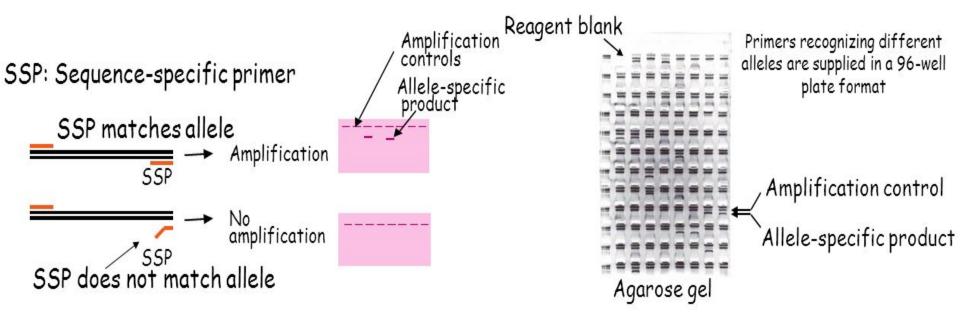
 identifies alleles that have common sequence (e.g. A*02:05/08/22)

High resolution (minimum 4 digit)

identifies single allele e.g.
 A*02:05

HLA TYPING DNA-BASED TYPING METHODS

SSP-PCR: Sequence-specific PCR (allele-specific primers)



Very rapid test that can be performed in 3-4 hours from the time a sample is received. PCR-SSP is used for typing deceased organ donors when speed is an important consideration.

HLA TYPING DNA-BASED TYPING METHODS

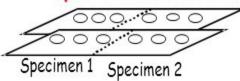
PCR-SSOPH: Sequence-specific oligonucleotide probe hybridization

Specimen 1 (Type A*0203)

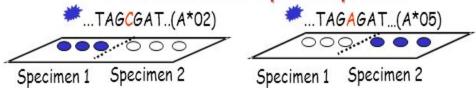
TAG C GAT TAG AGAT TAG

- Intermediate resolution
- Sreening test to identify potential donors or individuals who may later require higher resolution testing
- · High volume, relatively low cost

Amplify, denature, and spot onto membranes



Probe with allele-specific probes



xMAP® Technology, Multi - Analyte Profiling



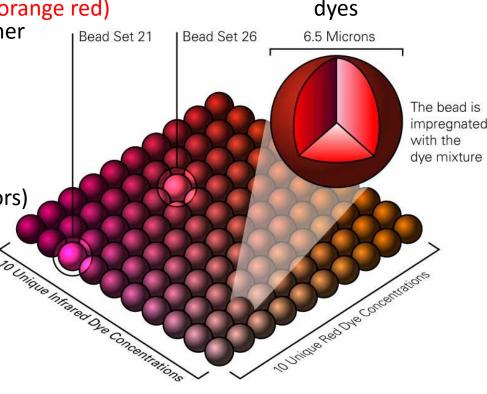
- 5.6 micron (non magnetic) or 6.5 micron (magnetic) beads
- Up to 100 (500) different beads/well
 - Bead color = spectral address

A combination of red and infrared (& orange red)

distinguishes one bead set from another

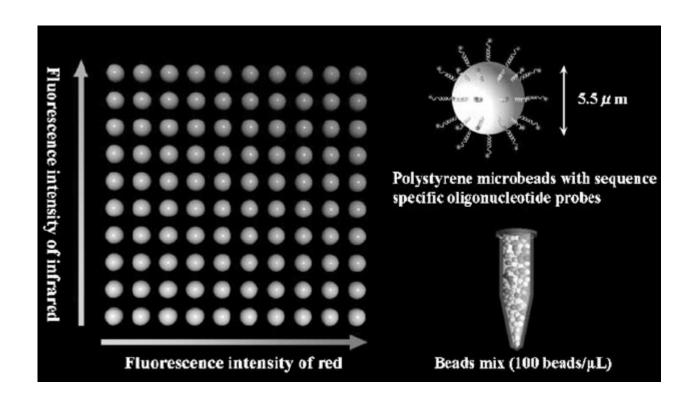
Small bead size allows:

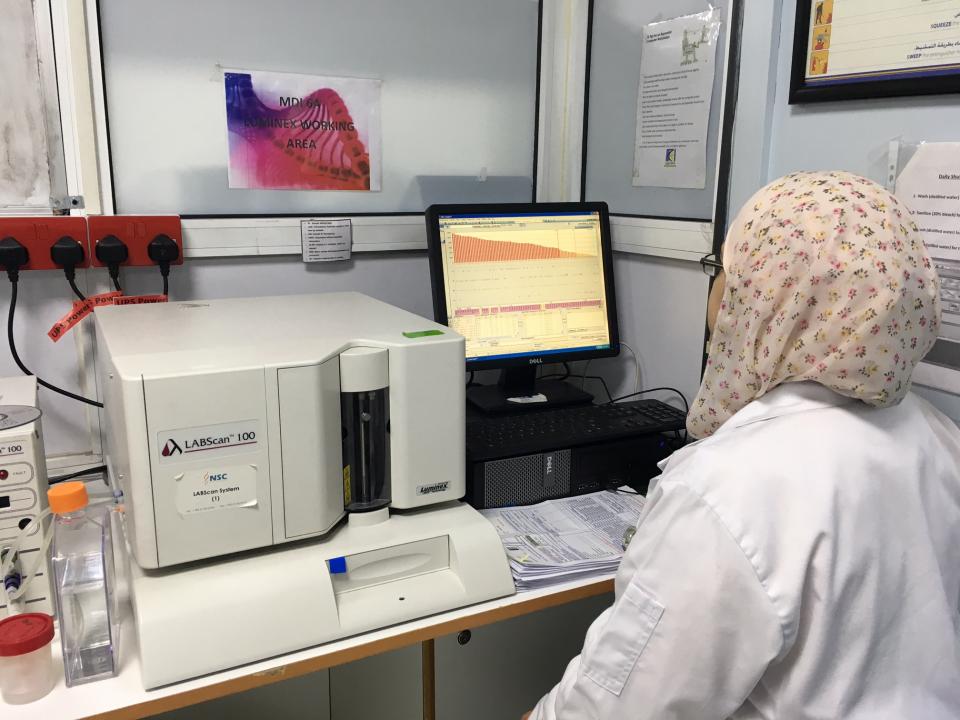
- Liquid suspension assay with
 - a high surface-to-volume-ratio
 - fast kinetics (liquid-phase behaviors)



xMAP® Technology, Multi - Analyte Profiling







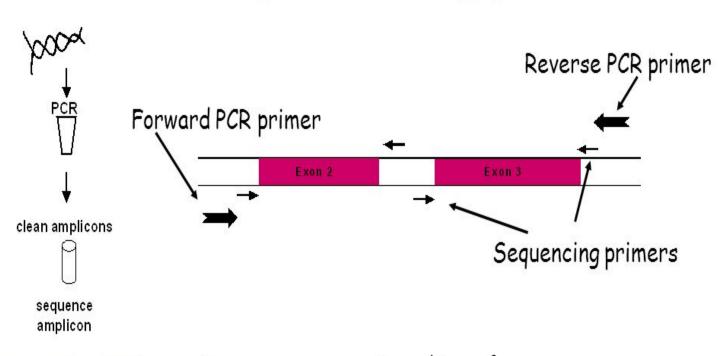
HLA TYPING

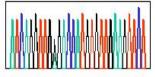
DNA-BASED TYPING METHODS

SBT: Sequence-based typing

Polymorphic regions are amplified by PCR and then sequenced

The highest resolution HLA typing

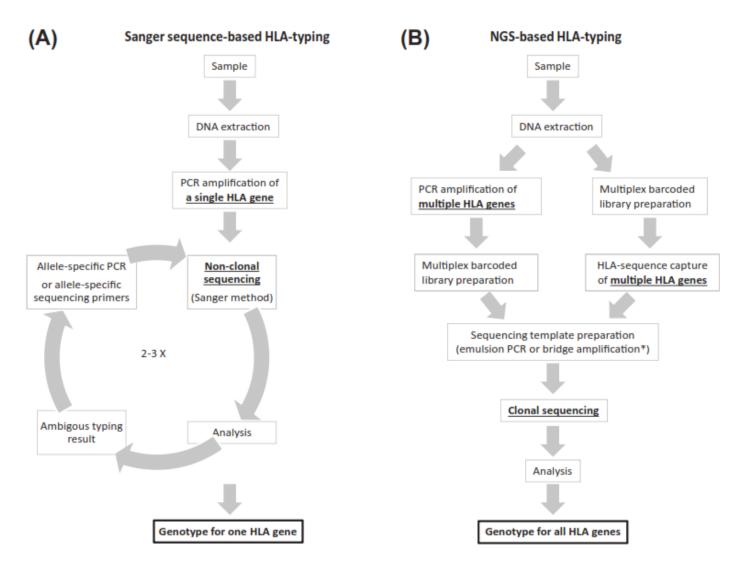




Isolate DNA

Sequences are compared to reference → sequences for previously assigned alleles

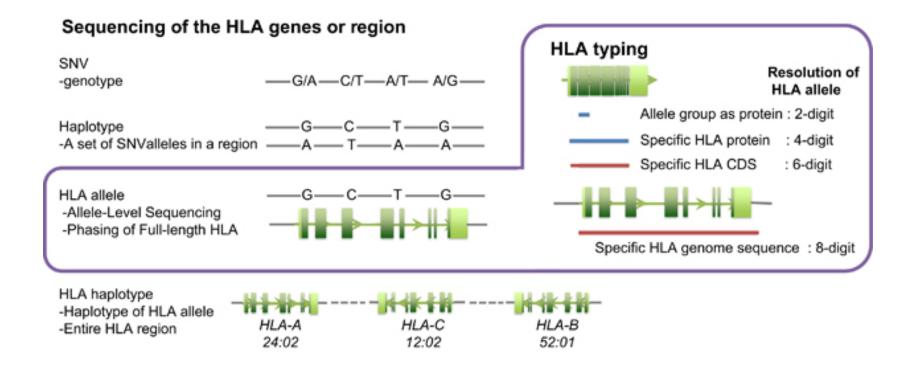




R. Carapito et al. / Human Immunology 77 (2016) 1016–1023

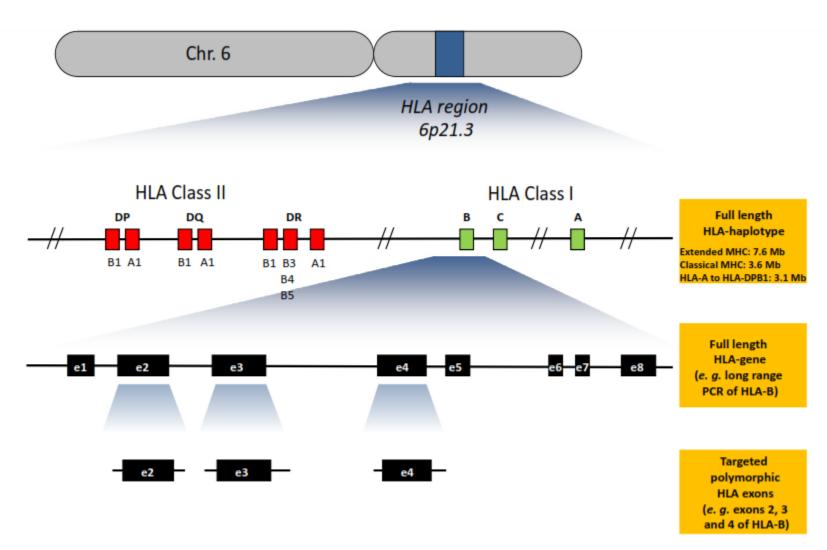
Sequence-based Typing





Targeted regions of the HLA by NGS approaches





R. Carapito et al. / Human Immunology 77 (2016) 1016–1023



Contents lists available at ScienceDirect







journal homepage: www.elsevier.com/locate/humimm

Invited Review

Clinical validation of NGS technology for HLA: An early adopter's perspective



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UNC Hospitals, McLendon Clinical Laboratories, United States
University of North Carolina at Chapel Hill School of Medicine, Department of Pathology and Laboratory Medicine, Chapel Hill, NC 27514, United States

ARTICLE INFO

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ABSTRACT

Clinical validation of NGS for HLA typing has been a topic of interest with many laboratories investigating the merits. NGS has proven effective at reducing ambiguities and costs while providing more detailed information on HLA genes not previously sequenced. The ability of NGS to multiplex many patients within a single run presents unique challenges and sequencing new regions of HLA genes requires application of our knowledge of genetics to accurately determine HLA typing. This review represents my lab-

- NGS can reduce the ambiguity rate, cost, and TAT for HLA typing.
- Validation of NGS for clinical HLA typing is challenging due to the numerous issues, including: sample types, complexity of the HLA genes, reliance on software for accurate HLA typing, and many more.

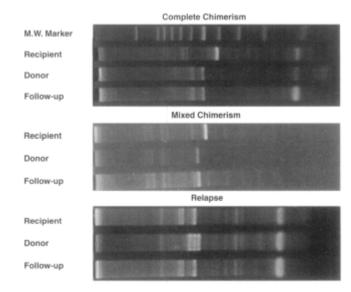


Original Article



The Use of Amplified Variable Number of Tandem Repeats (VNTR) in the Detection of Chimerism Following Bone Marrow Transplantation A Comparison With Restriction Fragment Length Polymorphism (RFLP) by Southern Blotting

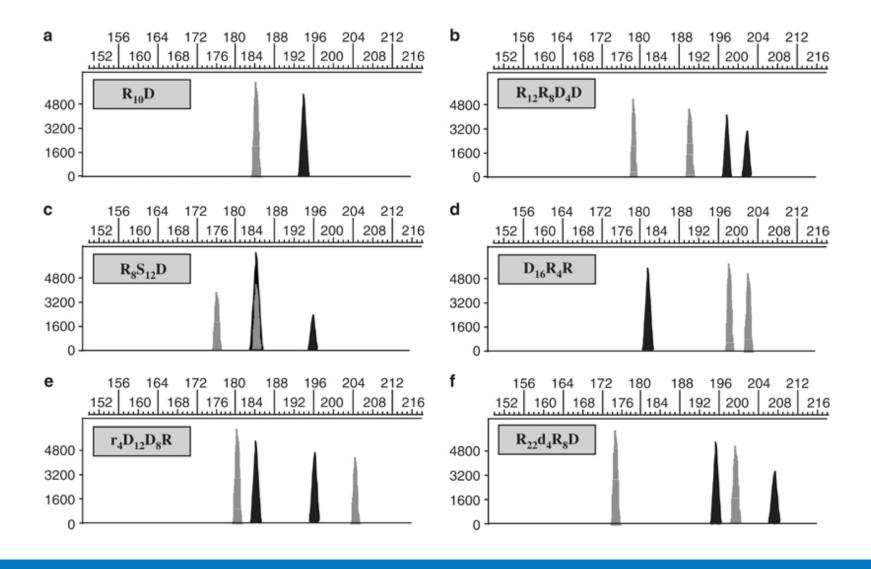
JOSEPH J. SREENAN, MD, JAMES D. PETTAY, MT(ASCP), ABDELGHANI TBAKHI, MD, GRIGORIOS TOTOS, MD, LINDA M. SANDHAUS, MD, MICHAEL L. MILLER, DO, BRIAN BOLWELL MD, AND RAYMOND R. TUBBS, DO



Am J Clin Pathol 1997;107:292-298.

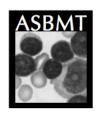
STR-PCR-based chimerism testing (







Biology of Blood and Marrow Transplantation 7:473-485 (2001)
© 2001 American Society for Blood and Marrow Transplantation



Review

Establishment of Complete and Mixed Donor Chimerism After Allogeneic Lymphohematopoietic Transplantation: Recommendations From a Workshop at the 2001 Tandem Meetings

Joseph H. Antin, ¹ Richard Childs, ² Alexandra H. Filipovich, ³ Sergio Giralt, ⁴ Stephen Mackinnon, ⁵ Thomas Spitzer, ⁶ Daniel Weisdorf ^{7*}

¹Dana-Farber Cancer Institute, Boston, Massachusetts; ²National Heart, Lung, and Blood Institute, Bethesda, Maryland; ³Children's Hospital Medical Center, Cincinnati, Ohio; ⁴M. D. Anderson Cancer Center, Houston, Texas; ⁵University College London, London, United Kingdom; ⁶Massachusetts General Hospital, Boston, Massachusetts; ⁷University of Minnesota Medical School, Minneapolis, Minnesota

Correspondence and reprint requests: Joseph H. Antin, MD, Dana-Farber Cancer Institute, 44 Binney St, Boston, MA

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www.nature.com/leu

ORIGINAL ARTICLE

The EuroChimerism concept for a standardized approach to chimerism analysis after allogeneic stem cell transplantation

T Lion¹, F Watzinger¹, S Preuner¹, H Kreyenberg², M Tilanus³, R de Weger⁴, J van Loon⁴, L de Vries⁴, H Cavé⁵, C Acquaviva⁵, M Lawler⁶, M Crampe⁶, A Serra⁷, B Saglio⁷, F Colnaghi⁸, A Biondi⁸, JJM van Dongen⁹, M van der Burg⁹, M Gonzalez¹⁰, M Alcoceba¹⁰, G Barbany¹¹, M Hermanson¹¹, E Roosnek¹², C Steward¹³, J Harvey¹⁴, F Frommlet¹⁵ and P Bader² on behalf of the EuroChimerism Consortium (EU-Project number: QLG1-CT-2002-01485)

Hematopoietic stem cell transplantation is becoming an increasingly important approach to treatment of different malignant and non-malignant disorders. There is thus growing demand for diagnostic assays permitting the surveillance of donor/recipient chimerism posttransplant. Current techniques are heterogeneous, rendering uniform evaluation and comparison of diagnostic results between centers difficult. Leading laboratories from 10 European countries have therefore performed a collaborative study supported by a European grant, the EuroChimerism Concerted Action, with the aim to develop a standardized diagnostic methodology for the detection and monitoring of chimerism in patients undergoing allogeneic stem cell transplantation. Following extensive analysis of a large set of microsatellite/short tandem repeat (STR) loci, the EuroChimerism (EUC) panel comprising 13 STR markers was established with the aim to optimally meet the specific requirements of quantitative chimerism analysis. Based on highly stringent selection criteria, the EUC panel provides multiple informative markers in any transplant setting. The standardized STR-PCR tests permit detection of donor- or recipient-derived cells at a sensitivity ranging between 0.8 and 1.6%. Moreover, the EUC assay facilitates accurate and reproducible quantification













Hindawi Publishing Corporation Biotechnology Research International Volume 2016, Article ID 8589270, 6 pages http://dx.doi.org/10.1155/2016/8589270



Research Article

Chimerism Analysis of Cell-Free DNA in Patients Treated with Hematopoietic Stem Cell Transplantation May Predict Early Relapse in Patients with Hematologic Malignancies

Mahmoud Aljurf, Hala Abalkhail, Amal Alseraihy, Said Y. Mohamed, 1 Mouhab Ayas,³ Fahad Alsharif,¹ Hazza Alzahrani,¹ Abdullah Al-Jefri,³ Ghuzayel Aldawsari, Ali Al-Ahmari, Asim F. Belgaumi, Claudia Ulrike Walter, Claudia Ulrike Walter, Hassan El-Solh, Walid Rasheed, and Maher Albitar

¹Oncology Centre, King Faisal Specialist Hospital and Research Centre, Riyadh 11211, Saudi Arabia

²Pathology and Laboratory Medicine, King Faisal Specialist Hospital and Research Centre, Riyadh 11211, Saudi Arabia

Minimal Requirements for Histocompatibility & Immunogenetics laboratory

Laboratory Accreditation

- To certify that the laboratory is meeting the minimum requirements as per given sets of standards
 - ASHI
 - EFI
 - CAP
 - ISO 15189





Thank Mou