

An Initiative to Exchange Extended RBC Phenotyping Information Using the HL7 FHIR® Open Standard

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Blood donor centers

- Computer systems geared toward effective and safe production of blood products
- Blood establishment computer systems (BECS)

Health system blood banks

- Computer systems geared toward effective and safe transfusion of blood products
- Blood bank laboratory information systems (BBLIS)

Both systems manage:

- Blood products
- Associated metadata

Barcodes provide a minimal digital interface—most of the metadata about the products must be rekeyed.

Need



Donor & Patient Phenotype Reports are Expanding

Positive

Donor Genotype				
Screen Print for DNMDN1	Date: 03/01/2021	Time: 9:11:32		
(DNR/BLD) Donor Phenotypes - U	odate NEXT:	DN		
Id : Blood Type	e: O- CMV: P	(SDBB) Pg 1/1 ositive		
Donor Phenotype	ABS: N	egative	Patient Genoty	pe
S Phenotype Indicator				
- C - Negative	Practitioner Name:	Specimen Type: Whole Bid	Accession Number	4/2020
- C + Positive	Requesting Hospital: Center	Date of Collection: 12/10/2	020 Date fest Fehrmiter, 12/24/20	12020
- E - Negative	Address:	Date Received: 12/11/202		120
- e + Positive				
- V - Negative	Tested Predicted F	henotypes and Results		
- VS - Negative	David Datient Dav	Call Antigan Construct		
k + Positivo	Panel: Patient Rec	i Cell Antigen Genotype		
_ Kpa - Negative	Group: ABO		Group: Duffy	
_ Kpb + Positive	Predicted Phenotynes	Results	Predicted Phenotypes	Results
_ Jsa - Negative		Negative	Fva	Positive
_ Jsb + Positive	<u>^</u>	Positive	Fvb	Negative
_ Fya - Negative	»	Positive	GATA	Positive
_ Fyb + Positive	0	7 03/046		1
_ Jka + Positive _ Jkb + Positive	Group: Rh		Group: Kell	
_ M + Positive	Predicted Phenotypes	Results	Predicted Phenotypes	Results
1 BA 2 QA 3 SB 4 SF 5 HE 6 CL 7 P	D DE D	Positive	к	Negative
	C C	Positive	k	Positive
****	E	Negative	Кра	Negative
	c	Positive	Крв	Positive
	e	Positive	Jsa	Negative
	V	Negative	Jsb	Positive
	VS	Negative		
	hrB	Positive	1	

Manual blood product registration was tolerable with limited metadata. Molecular techniques are rapidly changing this.

- Serologic characterization is expanding.
- Molecular techniques (arrays, targeted sequencing, gene sequencing) are expanding and becoming cost effective for more patients and donors.
- Provenance and molecular characterization information becomes critical when donors become patients.
- Search for potentially compatible blood can become automated and less laborious.

Now is the time to develop an open standard.

- Open standards are developed through a regulated, consensus-driven process.
- Open standards make possible interoperability, creating a common specification.
- Health Level 7 International (HL7®) standards are available under a no-cost licensing agreement.
- HL7 Fast Healthcare Interoperability Resources (FHIR®) is a flexible, scalable standard that streamlines information exchange between healthcare systems.
- HL7 FHIR is endorsed for federal projects under NIH, CDC, CMS, and other HHS agencies.



HL7 FHIR Implementation Guide: How?

- Convene domain stakeholders.
- Represent blood product genotyping and phenotyping information for BECS and BBLIS incorporation.
- Create a common set of resources:
 - for blood product phenotyping.
 - compatible with the FDA biologically derived products effort.*
 - compatible with International Society of Blood Transfusion (ISBT) consensus vocabulary of blood phenotypes.
- Test for feasibility, ballot as national standard, and publish.

RELATIONSHIP BETWEEN FHIR BIOLOGICAL PRODUCT AND A FHIR PATIENT

Process of finding a successful donor for patient blood transfusion match



A standard representation of extended RBC phenotyping promotes:

- Better management of blood product inventory (e.g., preventing accidental release of rare blood phenotypes and streamlining the search for rare phenotypes).
- Safer transfusion practices by
 - Finding the most compatible blood product for testing.
 - Identifying rare units through local, regional, super-regional and national searches
- More efficient public health reporting of adverse reactions, especially when combined with biologically derived product information.

Current activities

- Distributing a white paper and inviting additional support and signatures
- Socializing the effort via HL7 Work Groups–Clinical Genomics, Orders & Observations, Patient Care
- Seeking professional organization endorsement (AABB, CAP, ASCP, ISBT)
- Recruiting vendors and health care organizations (health systems and blood donor centers, software and instrument vendors)
- Full project launch dependent on stakeholder support—endorsement, participation, fund raising for ballot prep, and management

Explore fit and learn more

- Read white paper
- Explore participation:
 - In standards effort
 - In prototyping
 - As a stakeholder

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