

Yt Blood Group System

Number of antigens 2

High prevalence Yt^a
 Polymorphic Yt^b

Terminology

ISBT symbol (number) YT (011)
 Obsolete name Cartwright
 History Named after the high prevalence antigen, Yt^a ; became a system in 1964 after discovery of the antithetical antigen.

Expression

Other blood cells Not on lymphocytes, granulocytes or monocytes
 Tissues Brain, muscle, nerves

Gene

Chromosome 7q22.1
 Name *YT (ACHE)*
 Organization 6 exons distributed over 2.2 kbp of gDNA (exons 5 and 6 are alternatively spliced)
 Product Acetylcholinesterase (AChE)



Database accession numbers

GenBank M55040 (mRNA); L42812 (DNA)
 Entrez Gene ID 43

Molecular bases of Yt phenotypes

The reference allele, *YT*01* or *YT*A* (Accession number M55040) encodes Yt^a (YT1). Nucleotide differences from this reference allele, and the amino acids affected, are given.

Allele encodes	Allele name	Exon	Nucleotide	Amino acid	Ethnicity (prevalence)
Yt(a-b+) or YT:-1,2	<i>YT*02.01</i> or <i>YT*B</i>	2	1057C>A	His353Asn	Israeli Jews, Israeli Arabs, Druse > Blacks > Europeans (Common)
Yt(a-b+) or YT:-1,2	<i>YT*02.02</i> or <i>YT*B</i>	2 5	1057C>A 1775C>G	His353Asn Pro592Arg	Israeli Arabs, Druse, and Jews

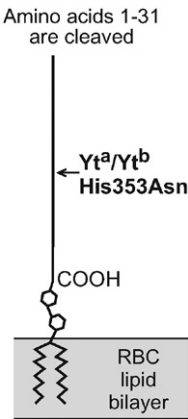
Amino acid sequence¹

MRPPQCLLHT	PSLASPLLLL	LLWLLGGGVG	AEGREDAELL	VTVRGGRLRG	50
IRLKTPGGPV	SAFLGIPFAE	PPMGPRRFLP	PEPKQPWSGV	VDATTFQSVC	100
YQYVDTLYPG	FEGTEMWNP	RELSEDCLYL	NVWTPYPRPT	SPTPVLVWIY	150
GGGFYSGASS	LDVYDGRFLV	QAERTVLVSM	NYRVGAFGFL	ALPGSREAPG	200
NVGLLDQRLA	LQWVQENVAA	FGGDPTSVTL	FGESAGAASV	GMHLLSPPSR	250
GLFHRAVLQS	GAPNGPWATV	GMGEARRRAT	QLAHLVGCPP	GGTGGNDTEL	300
VACLKTRPAQ	VLVNHEWHVL	PQESVFRFSF	VPVVDGDFLS	DTPEALINAG	350
DFHGLQVLVG	VVKDEGSYFL	VYGAPGFSKD	NESLISRAEF	LAGVRVGVPQ	400
VSDLAEEAVV	LHYTDWLHPE	DPARLREALS	DVVGDNHNVV	PVAQLAGRLA	450
AQGARVYAYV	FEHRASTLSW	PLWMGVPHGY	EIEFIFGIPL	DPSRNYTAE	500
LIFAQRLMRY	WANFARTGDP	NEPRDPKAPQ	WPPYTAGAQQ	YVSLDLRPLE	550
VRRGLRAQAC	AFWNRFLPKL	LSATASEAPS	TCPGFTHGEA	APRGLPLPL	600
LLLHQLLLLF	LSHLRRL				617

YT encodes a signal peptide of 31 amino acids, which is cleaved from the membrane bound protein. The carboxyl-terminal 29 amino acids are cleaved from the RBC GPI-linked form.

Carrier molecule

GPI-linked glycoprotein that probably exists as a dimer in the RBC membrane.



<i>M_r</i> (SDS-PAGE)	160,000 (72,000 as monomer under reducing conditions)
CHO: N-glycan	3 sites
CHO: O-glycan	Present
Cysteine residues	8
Copies per RBC	7,000–10,000 (or 3,500–5,000 dimers)

Function

AChE terminates nerve impulse transmission. AChE is in many tissues in various forms as a result of alternative splicing and post-translational modification. Function in RBC unknown.

Disease association

PNH III RBCs are deficient in AChE. Levels are reduced in myelodysplasias associated with chromosome 7 abnormalities and in some cases of SLE.

Phenotypes (% occurrence)

Phenotype	Most populations	Israelis
Yt(a+b-)	91.9	74.4
Yt(a+b+)	7.8	23.7
Yt(a-b+)	0.3	1.9
Null: Inherited Yt(a-b-) phenotype not found		
Unusual: One example of transient Yt(a-b-) RBCs reported ²		

References

- ¹ Bartels, C.F., et al., 1993. Mutation at codon 322 in the human acetylcholinesterase (ACHE) gene accounts for YT blood group polymorphism. *Am J Hum Genet* 52, 928–936.
- ² Rao, N., et al., 1993. Human erythrocyte acetylcholinesterase bears the Yt^a blood group antigen and is reduced or absent in the Yt(a–b–) phenotype. *Blood* 81, 815–819.

Yt^a Antigen

Terminology

ISBT symbol (number)	YT1 (011001 or 11.1)
Obsolete name	Cartwright
History	In 1956 when the antibody to this high-prevalence antigen was found, most letters in the patient's name (Cartwright) had been taken by other antigens. The authors thought “why not T?” but to avoid confusion with T polyagglutination, they then said “why T” or “Yt”. The Yt ^a antigen achieved system status in 1964 after discovery of the antithetical antigen.

Occurrence

Most populations	>99.8%
Israeli Jews	98.6%
Israeli Arabs	97.6%
Israeli Druse	97.4%

Antithetical antigen

Yt^b (YT2)

Expression

Cord RBCs	Weak
Altered	Weak or absent from PNH III RBCs

Molecular basis associated with Yt^a antigen¹

Amino acid	His353
Nucleotide	C at bp 1057 in exon 2

Effect of enzymes and chemicals on Yt^a antigen on intact RBCs

Ficin/Papain	Sensitive (variable)
Trypsin	Resistant
α-Chymotrypsin	Sensitive
DTT 200 mM/50 mM	Sensitive/weakened (thus sensitive to WARM™ and ZZAP)
Acid	Resistant

In vitro characteristics of alloanti-Yt^a

Immunoglobulin class	IgG (some are IgG4)
Optimal technique	IAT
Complement binding	Some

Clinical significance of alloanti-Yt^a

Transfusion reaction	No to moderate (rare)/delayed
HDFN	No

Comments

A report of an apparent alloanti-Yt^a in a Yt(a+) person suggests the possibility of heterogeneity of the Yt^a antigen². Experts agree that anti-Yt^a are often benign and antigen-negative blood may not need to be transfused.

References

¹ Bartels, C.F., et al., 1993. Mutation at codon 322 in the human acetylcholinesterase (ACHE) gene accounts for YT blood group polymorphism. *Am J Hum Genet* 52, 928–936.
² Mazzi, G., et al., 1994. Presence of anti-Yt^a antibody in a Yt(a+) patient. *Vox Sang* 66, 130–132.

Yt^b Antigen

Terminology

ISBT symbol (number)	YT2 (011002 or 11.2)
History	Identified in 1964 and named when its antithetical relationship to Yt ^a was recognized.

Occurrence

Europeans	8%
Israeli Jews	21.3%
Israeli Arabs	23.5%
Israeli Druse	26%

Not found in Japanese.

Antithetical antigen

Yt^a (YT1)

Expression

Cord RBCs	Weak
Altered	Weak or absent from PNH III RBCs

Molecular basis associated with Yt^b antigen¹

Amino acid	Asn353
Nucleotide	A at bp 1057 in exon 2

Effect of enzymes and chemicals on Yt^b antigen on intact RBCs

Ficin/Papain	Sensitive (variable)
Trypsin	Resistant
α-Chymotrypsin	Sensitive
DTT 200 mM/50 mM	Sensitive/weakened (thus sensitive to WARM™ and ZZAP)

In vitro characteristics of alloanti-Yt^b

Immunoglobulin class	IgG
Optimal technique	IAT

Clinical significance of alloanti-Yt^b

Transfusion reaction	No
HDFN	No

Comments

Anti-Yt^b is rare and usually occurs in sera with other antibodies. The second example of anti-Yt^b was made by a patient with PNH.

Reference

¹ Bartels, C.F., et al., 1993. Mutation at codon 322 in the human acetylcholinesterase (ACHE) gene accounts for YT blood group polymorphism. Am J Hum Genet 52, 928–936.