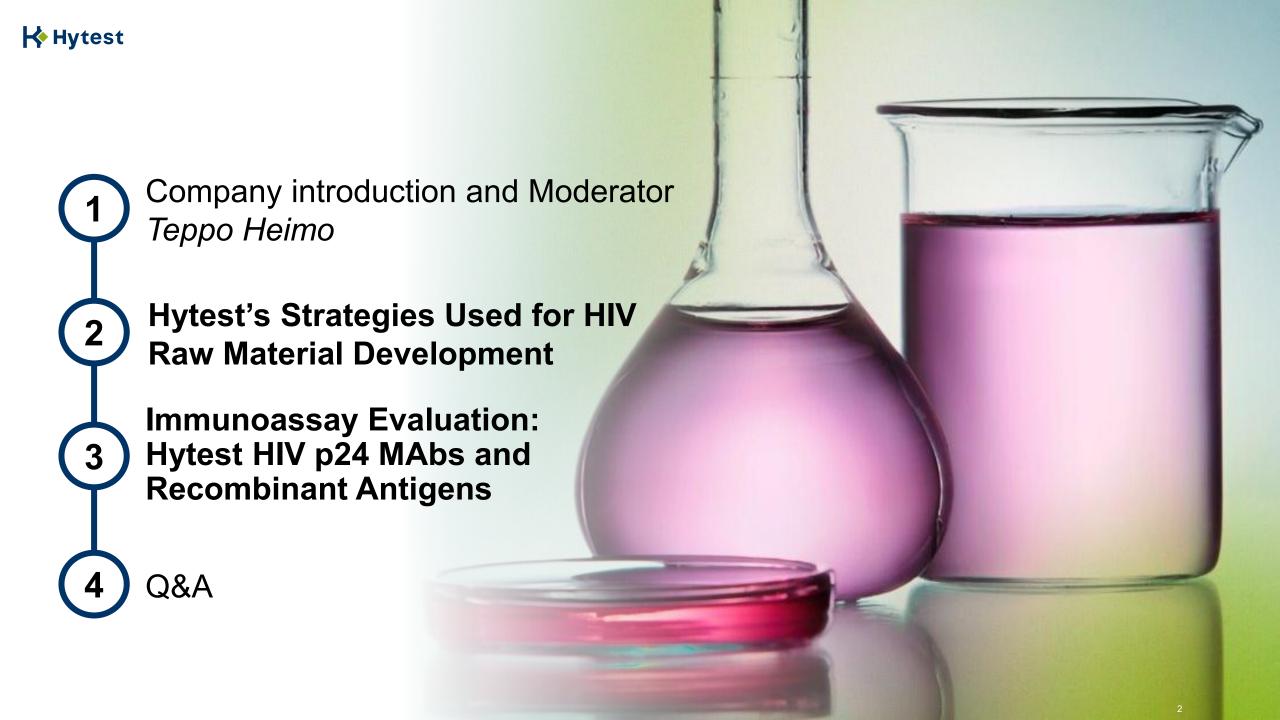


Building Better HIV Testing The Raw Material Journey from Research to Results

November 26th, 2025

Scientific excellence for IVD

1





Who We Are



30+ years of dedication to providing top-tier immunological reagents for the IVD industry and research community



Headquarters in Finland and operations in 3 continents with 150+ employees globally



An extensive product portfolio, featuring selected products as standard reference materials



Products developed based on solid science with 55+ scientific publications



Active participation in IFCC and ADLM standardization committee work



Close collaborations with KOLs



Operations compliant with ISO 9001:2015



Market presence in 55+ countries



We develop and manufacture world-class antibodies and antigens for the IVD industry, leading to better diagnostics







Comprehensive Product Portfolio

Product categories

Monoclonal antibodies Polyclonal antibodies Antigens Sera and Plasma



Clinical Utility

- Screening and monitoring HIV infections in targeted populations to help curb the HIV epidemic
- Early detection and treatment to improve quality of life









HIV monoclonal antibodies for antigen tests

HIV1/2 p24 (Cat# 3H24)

HIV recombinant antigens for serology tests

- HIV-1, gp120-gp41 N-Trx, rec (Cat# 8H11) Available soon!
- HIV-1, gp41-gp120 N-Fc, rec (Cat# 8H12)
- HIV1 gp41 N-HSA, rec (Cat# 8H13)
- HIV-1, gp120 C-Fc, rec (Cat# 8H16)
- HIV-2, gp36 N-HSA, rec (Cat# 8H24) Available soon!
- HIV-2 gp36 C-TnC, rec (Cat# 8H25) Available soon!



Today's Speakers

Natalia Tamm, Ph.D.

Jianwen He, Ph.D.



Natalia Tamm Ph.D.







M.Sc. and Ph.D. degrees from Moscow State University



Co-author of numerous scientific publications and patents



Joined Hytest in 2003 and has been involved in different research and development projects.



Expertise in a broad range of disease areas, including cardiovascular, infectious disease biomarkers and many others.

Deep knowledge in BNP, proBNP, and NT-proBNP biomarkers

Ongoing research focused on HIV biomarkers



Jianwen He Ph.D.











Post-doctorate fellow, University of South California, USA

Doctorate of Medical School of Fudan University (formerly Shanghai Medical University)

Associate Editor of the biblical book Immunoassay Handbook (4th ed. 2013, Publisher Elsevier)

Editor of "Immunoassay: Principle and Application" (2020, Publisher PMPH)

Editor of Chinese-English Bilingual Immunoassay Dictionary (2017, Publisher PMPH)

Published 30 SCI articles

Working and researching on assay principles, technologies & methodologies, and clinical applications of in vitro diagnostics, especially immunodiagnostics, home and abroad

Broad achievements in immunoassay development: biological raw material and reagent design, kinetics and interferences studies, assay and process optimization, clinical application study and troubleshooting



Chief Scientist of IVD BU. Director of IVD Raw Material Business, Mindray



Hytest's Strategies Used for HIV Raw Material Development

Natalia Tamm Ph.D.



Content

- HIV structure and viral genomic diversity
- Development of p24-specific monoclonal antibodies
- Development of HIV-specific recombinant proteins

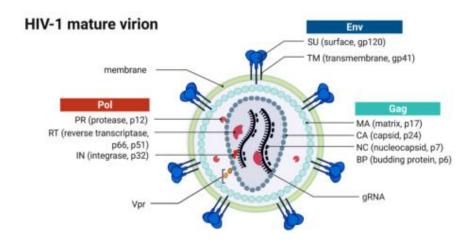


Content

- HIV structure and viral genomic diversity
- Development of p24-specific monoclonal antibodies
- Development of HIV-specific recombinant proteins



HIV: structure



Li G, De Clercq E. 2016. HIV genome-wide protein associations: a review of 30 years of research. Microbiol Mol Biol Rev

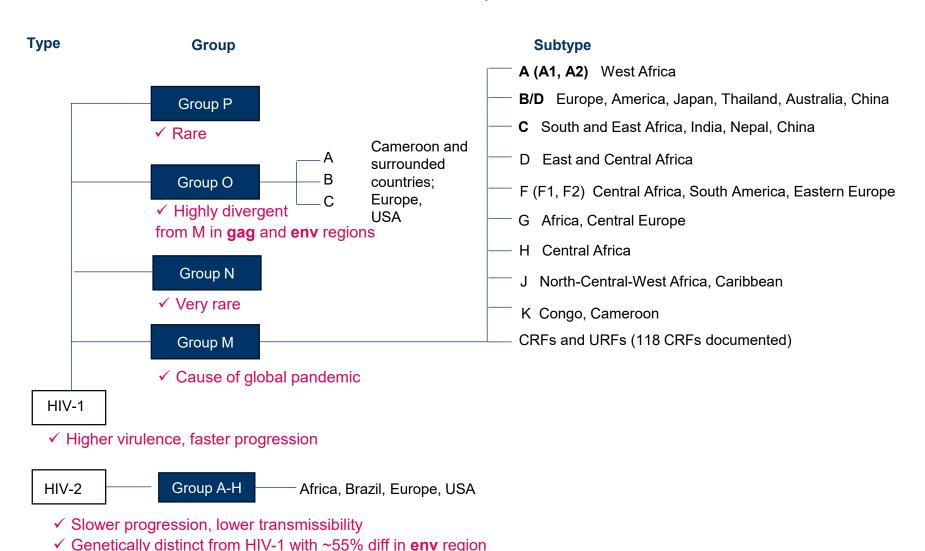
- HIV is a retrovirus causing AIDS (Acquired Immunodeficiency Syndrome).
- Globally, 39.9 million people lived with HIV in 2023
- IVD for detection of HIV infection, not the diagnosis of AIDS

HIV (human immunodeficiency virus)

- A member of the genus *lentivirus* in family *Retroviridae* (Retroviruses)
- Sphere with capsid and envelope
- Two copies of positive-sense single-stranded RNA, 9.7kb
- 16 proteins:
 - Structural proteins:
 - Matrix (p17), capsid (p24), nucleocapsid (p7), and p6
 - O Viral enzymes:
 - Protease (p12), reverse transcriptase (p66/51), and integrase (p32)
 - o Envelope proteins:
 - gp120 and gp41 (HIV-1)/gp36 (HIV-2)
 - Regulatory proteins:
 - Tat and Rev
 - Accessory proteins:
 - Vif, Vpu (HIV-1)/Vpx (HIV-2), Vpr, and Nef



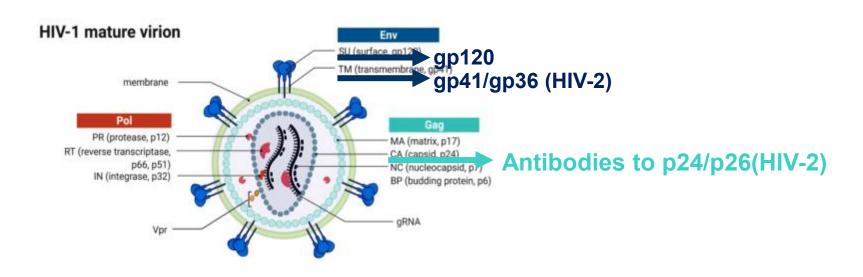
HIV: classification, variation and distribution



13



Specific proteins used for HIV assays development



Immunoassay requirements

- To design carefully due to genetic diversity
- To detect HIV-1 M subtypes/CRFs, O and HIV-2

Generation	4 th Gen
Time	2010
Method	Double- antigen/antibody sandwich
Antigens	Synthetic-peptide or recombinant protein
Antibodies	Monoclonal antibodies to HIV-1 p24 antigen
Analytes	HIV-1 (M and O group) and HIV-2 antibodies, HIV-1 p24 antigen
Feartures	Detection of HIV infection before seroconversion
Problems	Unable to differentiate HIV species
Sereconversion window period	2 weeks



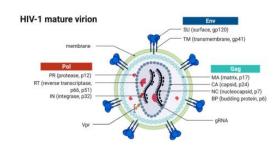
Content

- HIV structure and viral genomic diversity
- Development of p24-specific monoclonal antibodies
- Development of HIV-specific recombinant proteins



Hytest HIV-1/2 immunogens: p24 protein

- Structural protein, a component of the HIV capsid
- p24 monomer 230 aar; possibly phosphorylated (hexamer and pentamer)
- Conserved between types and subtypes (see alignment)
- p24 of HIV-1 Group M, HIV-1 Group O, and HIV-2 are related but not identical

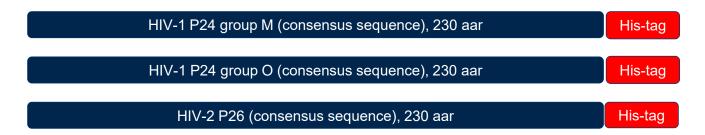


```
HIV-2 B (isolateD205) sp|P15833|P0LHV2D2 - PVQQLAGNYVHLPLSPRTLNAWVKLVEEKKFGAEVVPGFQALSEGCTPYDINQMLNCVGEHQAAMQIIREI
HIV-1 M, B (isolateHXB2) pp 1 V Q N I Q G Q M V H Q A I S P R T L N A W V K V V E E K A F S P E V I P M F S A L S E G A T P Q D L N T M L N T V G G H Q A A M Q M L K E T
HIV-1 0 (isolateMVP5180) 1079666 POLHVIMV PIVTNAQGQMVHQAISPRTLNAWVKAVEEKAFNPEIIPMFMALSEGAVPYDINTMLNAIGGHQGALQVLKEV
                           INEEAADWDQQHP - SPGPMPAGQLRDPRGSDIAGTTSTVEEQIQWMYRAQNPVPVGNIYRRWIQLGLQKCVR
                          INEEAAEWDRVHPVHAGPIAPGQMREPRGSDIAGTTSTLQEQIGWMTNN-PPIPVGEIYKRWIILGLNKIVR
                                                                                                                         143
                spiQ79666|POLHV1MV_INEEAAEWDRTHPPAMGPLPPGQIREPTGSDIAGTTSTQQEQIIWTTRGANSIPVGDIYRKW
                spIP04585IPOLHV1H2 MYSPTSILDIRQGPKEPFRDYVDRFYKTLRAEQASQEVKNWMTETLLVQNANPDCKTILKALGPAATL
                                                                                                                         215
                spiQ79666IPOLHV1MV MYSPVSILDIRQGPKEPFRDYVDRFYKTLRAEQATQEVKNWMTETLLVQNSNPDCKQILKALGPEATLEEMM
                                                                                                                         216
                sp|P15833|POL_HV2D2 TACQGIGGPGQKARLM
                                                                                                                         230
                sp|P04585|POL_HV1H2 TACQGVGGPGHKARVL
                                                                                                                         231
                sp|Q79666|POL_HV1MV VACQGVGGPTHKAKIL
                                                                                                                          232
```

P24 HIV-1 (M-group, subtype B), O-group, p26 HIV-2 alignment. Similarity is highlighted



Hytest HIV-1/2 immunogens: p24 proteins design



- Expression platform: E. coli cells
- Proteins were used for animal immunization and further antibody selection

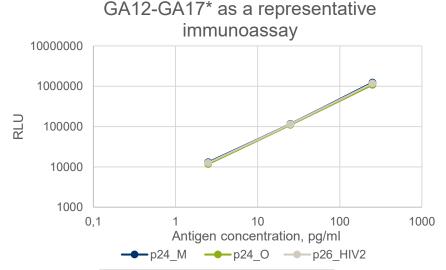


Hytest immunization strategy

• The immunization strategy included p24 **HIV-1 group M**, p24 HIV-1 **group O and p26 HIV-2** antigens to drive the immune response towards recognition of common epitopes within the p24 antigens of both groups of HIV-1 and also HIV-2.



p24 sandwich immunoassay characteristics



HIV-1 genotypes	S/N ratio
D	17,0
B/G	13,9
A/E	13,9
CRF BF	18,9
A1	15,7
B(214)	16,0
A/G	6,7
B (216)	13,7
Н	10,3
С	16,2
CRF G/43_02G	7,9
0	28,7
p26 HIV-2	19,0

The best antibody combinations:

- Recognize p24 HIV-1 gr_M, p24 HIV-1 gr_O, p24 HIV-2 equally effective (the presented antibodies recognize shared epitopes (or conservative fragments) of non-identical p24 proteins
- Recognize the following HIV-1 virus subtypes:
 - A1,B,B,C,D, F1/CRF12_BF/BFrec, G, CRF20_BG, CRF01_AE, CRF02_AG, H, group O (1st WHO International Reference Panel for HIV-1 p24 Antigen NIBSC code: 16/210), and
 - HIV-2 p26 (1st WHO International Reference reagent for HIV-2 p26 Antigen NIBSC code: 16/236)



9 MAbs under Cat.# 3H24 (Hytest)

Cat.#	MAb	Origin	Remarks
3H24	GA12	mouse	CLIA, ELISA, in vitro monoclonal antibodody
	GA15	mouse	CLIA, ELISA, in vitro monoclonal antibodody
	GA17	mouse	CLIA, ELISA, in vitro monoclonal antibodody
	GA18	mouse	CLIA, ELISA, in vitro monoclonal antibodody
	GA32	rabbit	CLIA, ELISA, recombinant monoclonal antibody
	GA34	rabbit	CLIA, ELISA, recombinant monoclonal antibody
	GA38	rabbit	CLIA, ELISA, recombinant monoclonal antibody
	GA39	rabbit	CLIA, ELISA, recombinant monoclonal antibody
	GA54	rat	CLIA, ELISA, recombinant chimeric antibody

According to the obtained data:

- Specificity to p24 from HIV-1 (group M, group O)/ p26 from HIV-2
- Several MAbs combinations can be suggested for the development of HIV p24 immunoassays
- MAbs could be used on CLIA, ELISA platforms
- HIV infection (HIV-1 or HIV-2) can be detected, including group O, CRF_BC, CRF_AE, and other subtypes using only one antibody pair (2 MAbs) instead of a set of monoclonal antibodies (three, four, a special set), used for HIV diagnostics by several IVD companies



Content

- HIV structure and viral genomic diversity
- Development of p24-specific monoclonal antibodies
- Development of HIV-specific recombinant proteins



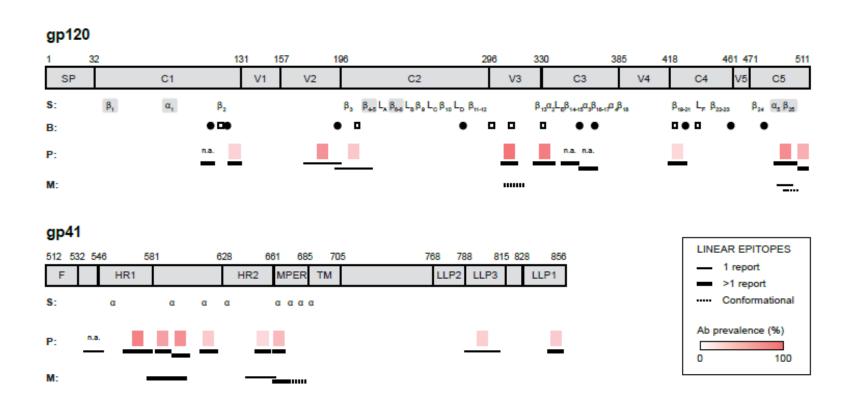
A set of proteins for HIV-1/2 antibody detection

Defining and selecting proteins/epitopes

- High immunogenicity
- Early arising of antibody
- Represent diversity (HIV-1 M/O, subtypes, CRFs; HIV-2)
- Include key immunodominant regions



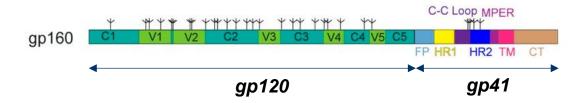
Env proteins represent major molecular targets of HIV-specific antibodies detected in HIV immunoassays



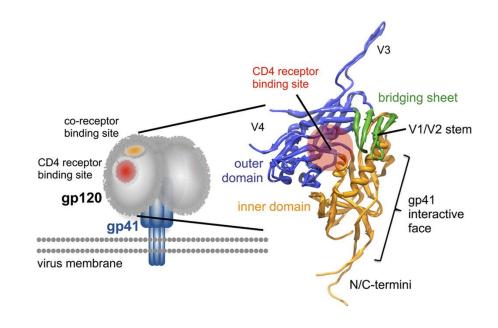
- HIV-1 gp120
- HIV-1 gp41
- HIV-2 gp36



gp120 (env) – one of the proteins used in HIV immunoassays



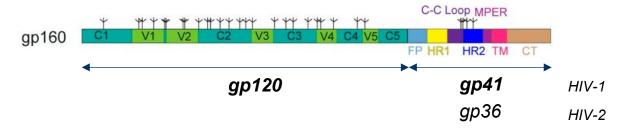
- Envelope glycoprotein
- Trimer
- 481aar HIV-1/503aar HIV-2
- Non-covalently interact with gp41
- V3 and V4 interact with CD4 to trigger viral entry
- Heavily glycosylated (both N- and O-glycans)
- High diversity between gp120 HIV-1/HIV-2



Solution Structure, Conformational Dynamics, and CD4-Induced Activation in Full-Length, Glycosylated, Monomeric HIV gp120; Miklos Guttman et al, Journal of Virology p. 8750–8764 August 2012



gp41 (env) – other protein used in HIV immunoassays



- Transmembrane glycoprotein with several functional regions
- Trimer
- 345aar HIV-1/353aar HIV-2
- Four determined sites of N-glycosylation
- Has several IDRs (immunodominant regions)
- High diversity between gp41 HIV-1 and gp36 HIV-2

```
HIV-1-GP41
               AVGIGALFLGFLGAAGSTMGAASMTLTVQARQLLSGIVQQQNNLLRAIEAQQHLLQLTVW
HIV-2-GP36
               -GVFVLGFLGFLATAGSAMGARSLTLSAQSRTLLAGIVQQQQQLLDVVKRQQEMLRLTVV
                     *****: ***: *** *: **: * **: **** *: : ** . : : **. : *: ****
HIV-1-GP41
HIV-2-GP36
               GTKNLQARVTAIEKYLKHQAQLNSWGCAFRQVCHTTVPW-
               * *: ****: *: *: *** * * * ***: : : * *: ***
HIV-1-GP41
HIV-2-GP36
               EKQVRYLEANISQSLEEAQIQQEKNMYELQKLNSWDILGNWFDLTSWVKYIQYGVHIVVG
               ::::. : 本 . : 本本: 本 水本本本 本本 : 本: 本 本 本本本: : 本 本本 : - : 本本 本
HIV-1-GP41
               LVGLRIVFAVLSIVNRVRQGYSPLS-----FQTHLPTPRGPDRPEGIEEEGGERDRDF
HIV-2-GP36
               IIALRIAIYVVQLLSRFRKGYRPVFSSPPGYLQQIHIHKDRGQPANEGTEEDVGGDSGYD
               HIV-1-GP41
               SIRLVNGSLALIWDDLRSLCLFSYHRLRDLLLIVTRIVELLGR-
HIV-2-GP36
               LWPWPINYVQFLIHLLTRLLIGLYNICRDLLSKNSPTRRLISQSLTAIRD
HIV-1-GP41
               YWSQELKNSAVSLLNATAIAVAEGTDRVIEVVQGACRAIRHIPRRIRQGLERILI
HIV-2-GP36
               YGCEWIQEAFQAFARTTRETLAGAWGWLWEAARRIGRGILAVPRRIRQGAELALL
```

gp41 HIV-1 and gp36 HIV-2 alignment. Similarity is marked with asterix



HIV proteins design; general ideas

Raw materials (proteins)

Global IVD companies (2) presumed design

- Based on available info
- Fusions, recombinant and peptides

Antigen design	Format	Expression platform	
HIV-1 gp41_ M	Recombinant protein	E. coli	
HIV-1 gp41_O	Synthetic peptide		
HIV-1 gp41/gp120	Recombinant protein	E. coli	
	Recombinant protein,		
HIV-2 gp36	fused	E. coli	

Hytest design

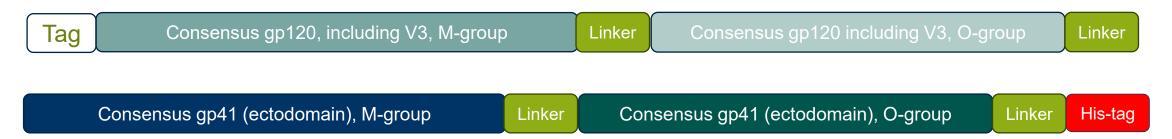
- Maximal diversity
- Consensus sequences
- Different expression platforms for the same protein used for coating and detection
- Sequences based on open-source info

Antigen Design	Format	Expression platform
HIV-1 gp41 _ M	Recombinant protein	Mammalian cells
HIV-2 gp36	Recombinant protein	Mammalian cells
HIV-1 gp120_M,O	Recombinant protein, fused	Mammalian cells
HIV-1 gp120_M,O+gp41_M,O	Recombinant protein, fused	Mammalian cells, <i>E. coli</i>



HIV-specific proteins development strategies

The maximal diversity in one recombinant protein

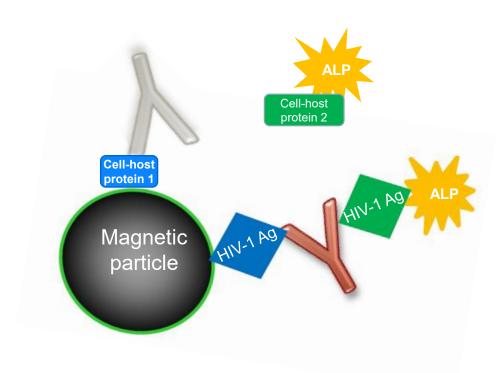


HIV-1 gp120+gp41, fusion protein



HIV-specific proteins development strategies

 The same protein used for double antigen sandwich produced in 2 different expression systems: E. coli and mammalian cells to reduce the risk of interference when used in a double antigen sandwich assay





HIV-1 gp120+gp41, fusion protein expression

E.coli cells

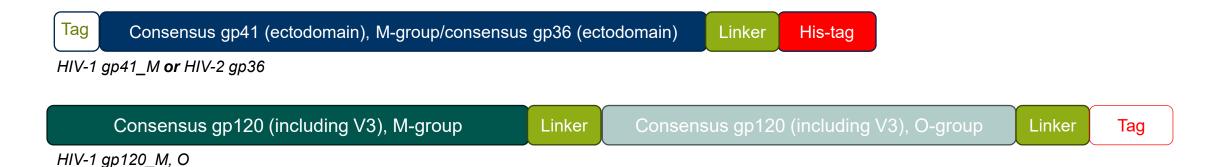
- Recombinant protein gp41/gp120 (fusion) was expressed with protein tag-partner (Thioredoxin)
- No post-translational modifications
- Partly refolded protein
- HIV-1 Trx gp41+gp120 has reactivity with HIV-1 M and O group blood samples (according to the HIV-tests manufactures)
- Acceptable protein stability under different temperature conditions

Mammalian cells

- Recombinant protein gp41/gp120 (fusion) was expressed with protein tag-partner (Fcfragment of human IgG)
- Post-translational modifications (glycosylation) could be similar to HIV proteins circulating in the blood of infected people
- Soluble protein
- HIV-1 gp41+ gp120 Fc has reactivity with HIV-1 M and O group blood samples (according to the HIV-tests manufactures)
- High protein stability under different temperature conditions



HIV-proteins design and expression by mammalian cells



- Recombinant proteins **HIV-1 gp41_M**, **HIV-2 gp36**; **HIV-1 gp120_M**, **O** were expressed with protein tag-partners (human serum albumin (HSA) or human cardiac troponin C (TnC) or Fc- fragment of human IgG)
- Post-translational modifications (glycosylation) could be similar to HIV proteins circulating in the blood of infected people
- All proteins are soluble
- HIV-1 gp41_M, HIV-2 gp36; HIV-1 gp120_M, O have reactivity with blood samples of corresponding group (according to the HIV-tests manufactures)
- High protein stability



HIV-specific proteins recommended for HIV Ab assays development

Cat.#	Product name	Purity-% (SDS-PAGE)	Expression system
8H11	HIV-1, gp120-gp41 N-Trx, recombinant	> 80	Bacterial cells
8H12	HIV-1, gp41-gp120 N-Fc, recombinant	> 80	Mammalian cells
8H13	HIV-1, gp41 N-HSA, recombinant	> 80	Mammalian cells
8H16	HIV-1, gp120 C-Fc, recombinant	> 90	Mammalian cells
8H24	HIV-2, gp36 N-HSA, recombinant	> 90	Mammalian cells
8H25	HIV-2 gp36 C-TnC, recombinant	> 80	Mammalian cells



Thank you

Natalia Tamm





Immunoassay Evolution: Hytest HIV p24 MAbs and Recombinant Antigens

Jianwen He Ph.D.





Content

- HIV immunoassay evolution
- Hytest HIV p24 MAbs evaluation
- Hytest HIV recombinant proteins evaluation





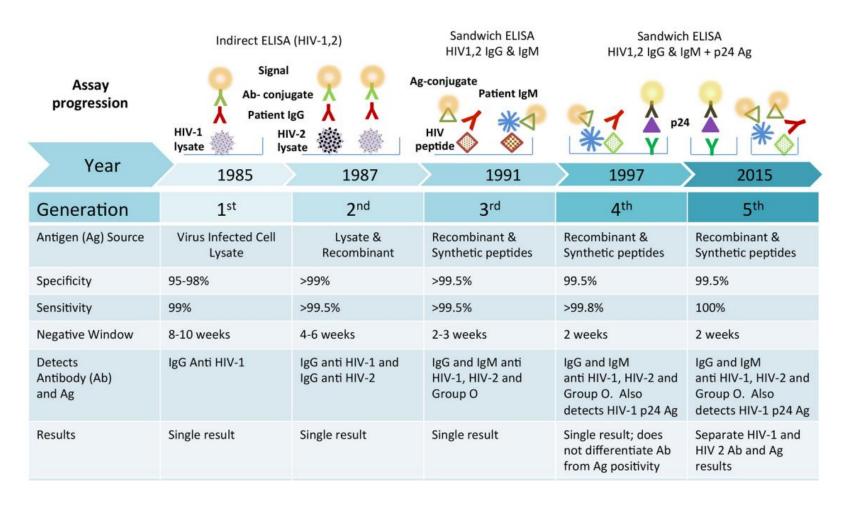
Content

- HIV immunoassay evolution
- Hytest HIV p24 MAbs evaluation
- Hytest HIV recombinant proteins evaluation





HIV Immunoassay Evolution



Becoming more sensitive and specific

- Lysate to recombinant/Peptides
- IgG to IgG+IgM
- HIV antibodies to antigen (gp24)
- Detection to differentiation (types, groups, subtypes)
- Identifies Infection Stage: quick intervention and short diagnostic algorithm

Alexander TS. Clin Vaccine Immunol. 2016 4;23(4):249-53

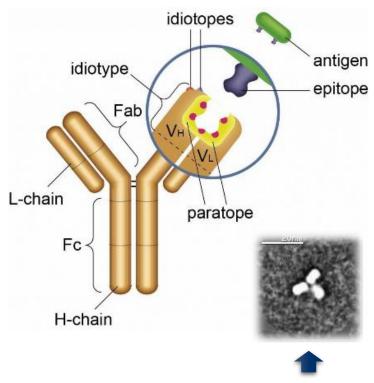




Antibody Is at the Heart of Immunoassay

50% IVD product recalls can be traced to raw materials

IgG Structure and Ag Binding Site



IgG in EM: 3 x 7 x 13 nM

Requirements for Antibody

- High affinity
- High purity
- Stability
- Specificity with defined epitopes
- Batch to batch consistency
- Sustainable supply
- In compliance

"A few good antibodies"





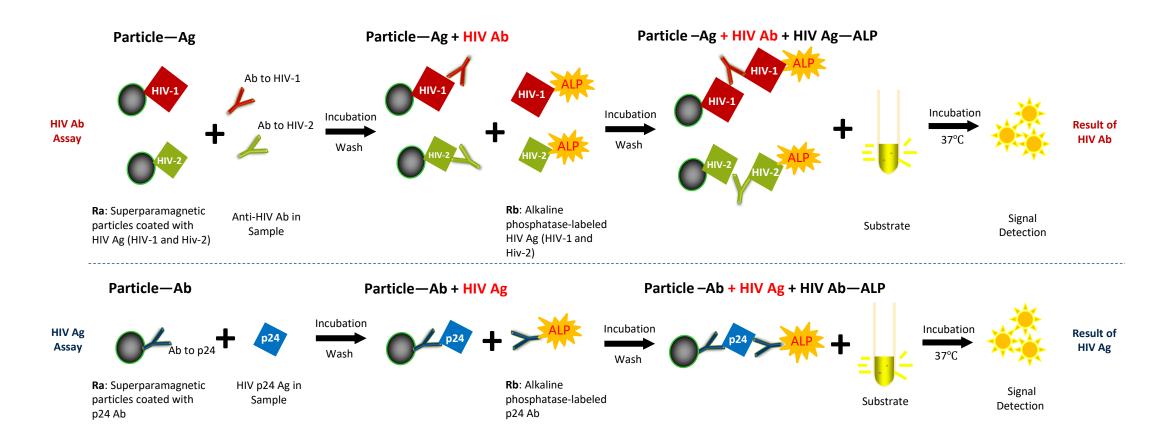
Content

- HIV immunoassay evolution
- Hytest HIV p24 MAbs evaluation
- Hytest HIV recombinant proteins evaluation





HIV Immunoassay Principle: Two-step Sandwich







HIV MAb Evaluation S/N of p24/p26 antigens

Reactivity of Hytest p24 antibody pairs

• MAb pairs demonstrated with high Signal to Noise ratio (S/N) for Hytest HIV-1 M and O group p24 antigens, HIV-2 p26 antigen, and p24 WHO standard (NIBSC code: 90/636).

MAbs evaluated

Cat.#	MAb	Fc
	GA12	Mouse
	GA17	Mouse
	GA32	Rabbit
3H24	GA34	Rabbit
	GA38	Rabbit
	GA39	Rabbit
	GA54	Rat

No.	Capture Mab	Detection Mab	HIV-1 p24 M antigen 10 pg/ml	HIV-1 p24 O antigen 10 pg/ml	HIV-2 p26 antigen 20 pg/ml	p24 WHO standard 2 IU/ml
1	GA12	GA17	10.05	13.52	7.26	8.61
2	GA17	GA54	9.23	14.16	5.89	8.41
3	GA17	GA12	9.31	14.24	7.66	7.70
4	GA32	GA54	9.26	10.22	3.43	7.91
5	GA54	GA17	10.14	13.63	5.78	8.67
6	GA54	GA38	9.30	11.38	5.54	6.74
7	GA12	GA38	7.92	9.48	8.11	7.31
8	GA34	GA54	8.77	11.40	3.48	7.71
9	GA34	GA12	8.12	10.78	7.01	7.33
10	GA39	GA12	8.23	9.58	7.07	7.45
11	GA39	GA54	7.81	9.18	4.11	7.47
12	GA32	GA12	8.24	9.18	6.98	7.29
13	GA38	GA54	7.90	11.00	3.67	6.70
14	GA17	GA34	8.09	9.97	4.89	6.71
15	GA54	GA34	8.35	9.64	4.57	6.59
16	GA17	GA38	8.18	10.59	5.23	6.55
17	GA34	GA38	8.70	9.70	7.91	5.80
18	GA12	GA34	7.29	8.98	6.23	6.96
19	GA34	GA39	7.92	7.62	7.50	6.12
20	GA38	GA12	7.24	8.98	5.86	6.10
21	GA32	GA34	8.45	6.45	5.81	6.06
22	GA38	GA34	8.58	9.01	7.97	5.73
23	GA39	GA34	7.56	6.46	5.86	6.17





HIV MAb Evaluation Limit of Detection (LoD)

- Hytest MAb pairs showed LoD of WHO international standard (NIBSC code: 90/636) about 0.5~0.65 IU/ml, which meets the requirement of COMMISSION IMPLEMENTING REGULATION (EU) 2022/1107 (≤ 2 IU/ml).
- Most of the MAb pairs showed a greater detection capacity for HIV-1 p24 and p26 antigens than results generated in compared groups.

Number of Pair	Capture MAb	Detection MAb	LOD of WHO international standard p24 antigen, IU/ml	HT HIV-1 p24_M, pg/ml	LOD HT HIV-1 p24_O, pg/ml	HT HIV-2 p26, pg/ml
1	GA17	GA54	0.64	4	1.5	15
2	GA17	GA12	0.64	2	2	8
3	GA17	GA38	0.55	1.5	1.5	8
4	GA34	GA54	0.5	2	1.5	20
5	GA38	GA54	0.55	4	1.5	20
6	GA17	GA34	0.64	4	2	15
7	GA39	GA54	0.64	4	4	25
8	GA17	GA32	0.64	2	2	8
9	GA32	GA54	0.55	4	4	30
10	GA17	GA39	0.64	2	2	8
11	GA34	GA39	0.55	2	4	5
12	GA34	GA38	0.5	1	1	3
13	GA38	GA12	0.55	2	1.5	8
14	GA12	GA34	0.55	2	2	8
15	GA12	GA32	0.64	4	4	4
16	GA12	GA39	0.64	4	4	4
17	GA34	GA32	0.55	2	4	8
		Combo1	1.3	5	10	20
	IVD (Combo2	1.0	5	4	Not detected





HIV MAb Evaluation Specificity

• Evaluated with a variety of sample groups, Hytest MAb pairs showed that specificity meets requirement of COMMISSION IMPLEMENTING REGULATION (EU) 2022/1107 (specificity >99.5%).

	Capture MAb etection MAb		GA17 GA12	GA17 GA54	GA17 GA38	GA17 GA34	GA17 GA32	GA17 GA12+GA38
Sample	Category	Sample Number	Number of false reactive (%)					
Collected	Study 1	1000	1 (99.90%)	1	0 (100%)	1 (99.90%)	0 (100%)	1 (99.90%)
Samples	Study 2	1014	0 (100%)	1				
	Study 1	1228	0 (100%)	0 (100%)				
Clinical Samples	Study 2	941	0 (100%)	0 (100%)		/		
Samples	Study 3	1251	2 (99.84%)	2 (99.84%)				
Blood Don	or Samples	989	0 (100%)	0 (100%)				
Total	Samples number	6423	6423	4031	1000	1000	1000	1000
	False Reactive	I	3 (99.95%)	2 (99.95%)	0 (100%)	1 (99.90%)	0 (100%)	1 (99.90%)





HIV MAb Evaluation Reference Panel 1

 Hytest MAb pairs showed higher sensitivity than compared groups in detection of p24 and p26 antigen, respectively.

Assay		Hytest Pair GA17 + GA12	Hytest Pair GA17 + GA54	IVD Combo1	IVD combo2	
Sample	Genotype	Conc.	COI	COI	COI	COI
	A1	1 IU/ml	3.35	2.92	0.881	2.29
	B 16/214	1 IU/ml	4.82	4.16	1.55	2.35
	B 16/216	1 IU/ml	3.62	3.17	1.64	1.26
	С	1 IU/ml	3.58	3.26	1.23	1.96
	D	1 IU/ml	4.16	3.57	1.1	2.51
p24 WHO Reference	F1	1 IU/ml	3.39	3.04	1.83	1.26
Panel 16/210	G	1 IU/ml	2.55	2.12	1.4	2.21
	BG	1 IU/ml	3.68	3.28	1.57	1.43
	AE	1 IU/ml	3.26	2.68	1.31	2.5
	AG	1 IU/ml	3.6	2.93	1.62	1.86
	Н	1 IU/ml	4.49	3.92	1.26	2.12
	Group O	1 IU/ml	6.48	5.41	2.06	3.56
	p24_M	2pg/ml	1.22	0.9	0.678	0.46
HyToot Antigon	p24_O	2pg/ml	1.85	1.38	0.54	0.96
HyTest Antigen	HIV-2 p26	20pg/ml	3.02	1.54	5.95	0.38
	HIV-2 p26	10pg/ml	1.8	0.91	0.653	0.11





HIV MAb Evaluation Reference Panel 2

 Hytest MAb pairs showed higher sensitivity than compared groups in detection of p24 and p26 antigen, respectively.

	Assay		Hytest Pair GA17 + GA12	Hytest Pair GA17 + GA54	IVD Combo1	IVD combo2
Sample	Genotype	Conc.	COI	COI	COI	COI
·	В	5pg/ml	1.40	1.10	1.56	1.29
	CRF-07-BC	5pg/ml	1.38	1.01	1.16	0.11
	CRF-07-BC	5pg/ml	1.46	1.33	1.16	0.85
	CRF-01-AE	5pg/ml	1.61	1.13	1.42	1.82
	CRF-01-AE	5pg/ml	1.20	1.09	2	1.71
	CRF-08-BC	5pg/ml	1.70	1.23	1.16	1.38
	CRF-08-BC	10pg/ml	1.97	1.46	1.49	0.23
	CRF-55-01B	5pg/ml	1.54	1.41	1.54	1.88
	CRF-55-01B	5pg/ml	2.13	1.58	1.93	2.59
04/ 00 D (A1	15pg/ml	1.49	1.17	2.01	1.21
p24/p26 Reference	В	15pg/ml	3.75	1.47	1.33	0.97
Panel From A Third	В	10pg/ml	1.82	1.53	2.04	0.14
Party	С	5pg/ml	1.56	1.35	1.12	0.37
	D	10pg/ml	1.64	1.42	0.989	0.21
	F1/CRF12-BF/BFrec	5pg/ml	1.09	0.97	1.29	0.1
	G	30pg/ml	1.57	1.54	1.91	0.4
	CRF20-BG	5pg/ml	1.22	1.10	1.34	0.14
	CRF01-AE	5pg/ml	1.43	1.19	1.11	1.22
	CRF02-AG	5pg/ml	1.40	1.21	1.41	0.68
	Н	5pg/ml	1.50	1.46	1.06	0.36
	HIV-1 O	10pg/ml	1.79	1.28	0.394	1.88
	HIV-2 A	250pg/ml	2.77	1.11	4.86	0.25





Content

- HIV immunoassay evolution
- Hytest HIV p24 MAbs evaluation
- Hytest HIV recombinant proteins evaluation





HIV Antigen Evaluation- Sensitivity

 11 undiluted positive HIV-1 samples were evaluated, and the Hytest HIV-1 antigen pairs demonstrated sensitivity of 100%. 11 different diluted HIV-2 samples were tested, and the Hytest HIV-2 antigen pairs demonstrated sensitivity of 100%.

Recombinant Antigens Evaluated

No.	Cat. #	Antigen	Expression System
1	8H11	HIV-1, gp120-gp41 N-Trx, rec	E. coli
2	8H12	HIV-1, gp41-gp120 N-Fc, rec	Mammalian cells
3	8H13	HIV-1, gp41 N-HSA, rec	Mammalian cells
4	8H24	HIV-2, gp36 N-HSA, rec	Mammalian cells
5	8H25	HIV-2, gp36 C-TnC, rec	Mammalian cells

HIV-1 Positive Samples

Capture Antigen	8H11	8H11	IVD Combo1	IVD Combo2
Detection Antigen	8H12	8H13		
No.1	1.00	3.23	59.228	3.53
No.2	1.49	2.34	6.621	3.69
No.3	4.13	5.34	18.691	9.88
No.4	4.77	51.76	9.308	8.13
No.5	18.07	46.44	25.498	49.74
No.6	23.16	47.99	24.068	102.2
No.7	23.86	61.58	38.319	138.2
No.8	39.49	77.6	45.984	294.4
No.9	52.72	81.08	29.218	161.9
No.10	56.84	107.28	93.722	217
No.11	188.54	235.45	93.499	190.4

HIV-2 Positive Samples

Capture Antigen	8H24	8H24	8H25	IVD Combo1
Detection Antigen	8H24	8H25	8H24	
No.1	1.74	1.90	1.86	1.76
No.2	3.62	2.41	4.45	7.51
No.3	12.49	8.53	20.41	15.92
No.4	14.88	12	26.5	22.02
No.5	4.12	2.93	9.91	1.48
No.6	19.72	14.68	49.99	26.83
No.7	17.5	14.89	44.1	34.67
No.8	10.42	10.72	20.45	5.58
No.9	16.81	14.47	39.01	32.13
No.10	24.64	20.92	66.67	45.01
No.11	20.46	18.4	54.66	37.18





HIV Antigen Evaluation- HIV-1 Specificity

• Evaluated using various types of interference samples and clinical specimens, 2 Hytest HIV-1 antigen pairs demonstrated high specificity, satisfying the required performance criteria of COMMISSION IMPLEMENTING REGULATION (EU) 2022/1107 (specificity >99.5%).

	Particle Antigen	8H11	8H11	
	Conjugate Antigen		8H12	8H13
Type of Inte	erference	Sample Number	False R	Reactive
Interference	e Sample	78	1	3
ANA Positiv	e Sample	88	0	0
RF Sai	mple	139	0	1
Total	Sample Number	305	305	305
Total	False reactive	1	1	4
Sample C	ategory	Sample Number	False Reactive	
Collected Samples	Study 1	856	3 (99.65%)	3 (99.65%)
Collected Samples	Study 2	1014	1	3 (99.70%)
	Study 1	850	0 (100%)	0 (100%)
Clinical Samples	Study 2	941	1	1 (99.89%)
	Study 3	1251	1 (99.92%)	1 (99.92%)
Blood Dono	Blood Donor Samples		1	2 (99.80%)
Total	Sample Number	5901	2957	5901
iotai	False reactive	1	4 (99.86%)	10 (99.83%)





HIV Antigen Evaluation- HIV-2 Specificity

• Evaluated with different types of interference samples and clinical samples, 4 Hytest HIV-2 antigen pairs had high specificity, which met the requirement (specificity >99.5%).

Particle Antigen		8H24	8H24	8H25	8H25	
Conjugate Antigen			8H24	8H25	8H24	8H25
Type of I	nterference	Number		False Read	tive	
Interferer	nce Sample	78	0	0	0	0
ANA Posi	tive Sample	88	/	0	/	/
RF S	ample	102	/	0	/	/
Total	Sample Number	268	78	268	78	78
Total	False Reactive	/	0	0	0	0
Sample	Category	Number		False Read	tive	
Collected Samples	Study 1	1638	0 (100%)	0 (100%)	0 (100%)	1 (99.87%)
Collected Samples	Study 2	1014	0 (100%)	0 (100%)	/	/
	Study 1	850	/	0 (100%)	/	/
Clinical Samples	Study 2	941	/	0 (100%)	/	/
	Study 3	1251	/	0 (100%)	/	/
Blood Doi	nor Samples	989	/	0 (100%)	/	/
Total	Sample Number		2652	6683	1638	750
	False Reactive	/	0 (100%)	0 (100%)	0 (100%)	1 (99.87%)





HIV Ab/Ag Combo Evaluation-Genotype

Hytest HIV Ab/Ag combo assays were able to detect all different genotypes similar to compared groups.

	Assay Pairs		HIV Combo1 HIV-1: 8H11+8H13 HIV-2: 8H24+8H25 p24: GA17+GA12	HIV Combo2 HIV-1: 8H11+8H12 HIV-2: 8H24+8H25 p24: GA17+GA54	IVD Combo1	IVD Combo2
No.	HIV Subtype	HIV-1 Viral Load*	COI	COI	COI	COI
1	Α	262,186	13.73	18.01	25.5	16.17
2	В	46	145.41	246.52	821	1038.44
3	В	1,361,613	349.45	673.57	1625	566.84
4	В	761,620	432.19	781.81	274	940.22
5	С	> 10,000,000	6.20	7.48	5.66	3.33
6	C / CRF_BC	1,427,474	25.84	23.03	66.7	69.84
7	C / CRF_BC	> 10,000,000	11.58	10.38	7.09	6.7
8	C / CRF_BC	> 10,000,000	53.14	57.85	36.8	17.19
9	C / CRF08_BC	98,128	4.17	4.61	44.1	30.92
10	C / CRF08_BC	1,076	34.85	37.51	188	53.09
11	C / CRF08_BC	90,675	268.82	429.63	280	452.14
12	C / CRF31_BC	10,594	232.53	451.57	2788	1157.45
13	CRF_BG	2,711	220.97	396.42	137	620.13
14	CRF01_AE	12,688	100.72	162.14	251	257.72
15	CRF02_AG	> 10,000,000	13.27	14.14	10.3	6.44
16	CRF03_AB	163	44.32	47.49	43.1	9.04
17	CRF22_01A1	548,193	170.99	304.56	41.4	189.41
18	D	243,267	4.01	4.53	4.16	0.55
19	F2	28,294	348.48	680.02	404	279.91
20	G	9,629	387.05	711.00	1055	774.07
21	1	> 10,000,000	6.95	9.32	7.37	4.63
22	1	> 10,000,000	2.42	2.30	1.96	1.95

^{*}All tested samples are diluted 100 times





HIV Ab/Ag Combo Evaluation-Seroconversion Panel

 Hytest HIV Ab/Ag combo assays demonstrated similar sensitivity as compared group.

	Assay Pairs		HIV Combo1 HIV-1: 8H11+8H13 HIV-2: 8H24+8H25 p24: GA17+GA12	HIV Combo2 HIV-1: 8H11+8H12 HIV-2: 8H24+8H25 p24: GA17+GA54	IVD Combo1
No.	Seroconversion Panel	Sample Size	Number of HIV reactive samples		
1	HIV9021	17	4	4	4
2	HIV9018	9	3	3	3
3	HIV9030	16	3	3	3
4	HIV9011	11	2	2	2
5	HIV9028	7	2	2	2
6	HIV 9012	8	3	4	3
7	HIV 9077	23	12	12	12
8	SCP-HIV-002	20	11	12	11
9	SCP-HIV-003	7	4	4	4
10	SCP-HIV-004	7	3	3	3
11	SCP-HIV-005	25	21	21	21
12	SCP-HIV-007	9	4	4	4
13	0600-0240	5	2	2	2
14	0600-0245	7	2	2	2
15	0600-0248	10	3	3	3
16	0600-0251	10	3	4	4
17	0600-0256	4	2	2	2
18	0600-0258	4	2	2	2
19	0600-0261	4	3	3	2
20	0600-0270	4	3	3	2
21	0600-0272	6	3	3	3
22	0600-0243	9	2	2	2
	Seroconversion Panel Teste	d	22	22	22
	Total Tested Samples		222	222	222
	HIV Reactive Samples	97	100	96	





HIV Ab/Ag Combo Evaluation-Specificity

- A total of 4272 different types of interference samples and samples from different populations were evaluated with HIV Ab/Ag combo assays.
- Hytest HIV Ab/Ag combo assays showed high specificity (>99.9%).

HIV Ab/Ag Combo Evaluation

Assay	HIV Combo1	HIV Combo2
HIV-1 pair	8H11 + 8H13	8H11 + 8H12
HIV-2 pair	8H24 + 8H25	8H24 + 8H25
p24 pair	GA17 + GA12	GA17 + GA54
Sample number	4272	4272
Number of false reactive	3	4
Specificity	99.93%	99.91%

Sources of Evaluated Samples

Assay		HIV Combo1	HIV Combo2
Sample Category	Number Tested	Number of F	False Positive
Hospital 1	1690	2	2
Hospital 2	40	0	0
Hospital 3	739	0	2
Hospital 4	213	0	0
Triglyceride-rich Samples	189	0	0
Hemolyzed Samples	189	0	0
RF Samples	69	0	0
ANA Positive Samples	299	1	0
IgM-rich Samples	2	0	0
Patients with Diabetes	95	0	0
Patients with Uremia	172	0	0
Patients with Dialysis	8	0	0
Old People	124	0	0
Pregnant women	75	0	0
Newborn	100	0	0
Children	84	0	0
Cardiovascular Diseases	184	0	0
Total	4272	3	4





Conclusions

The evaluated HIV p24 monoclonal antibodies using Mindray CLIA System exhibit high sensitivity and specificity in the detection of both p24 and p26 antigens.

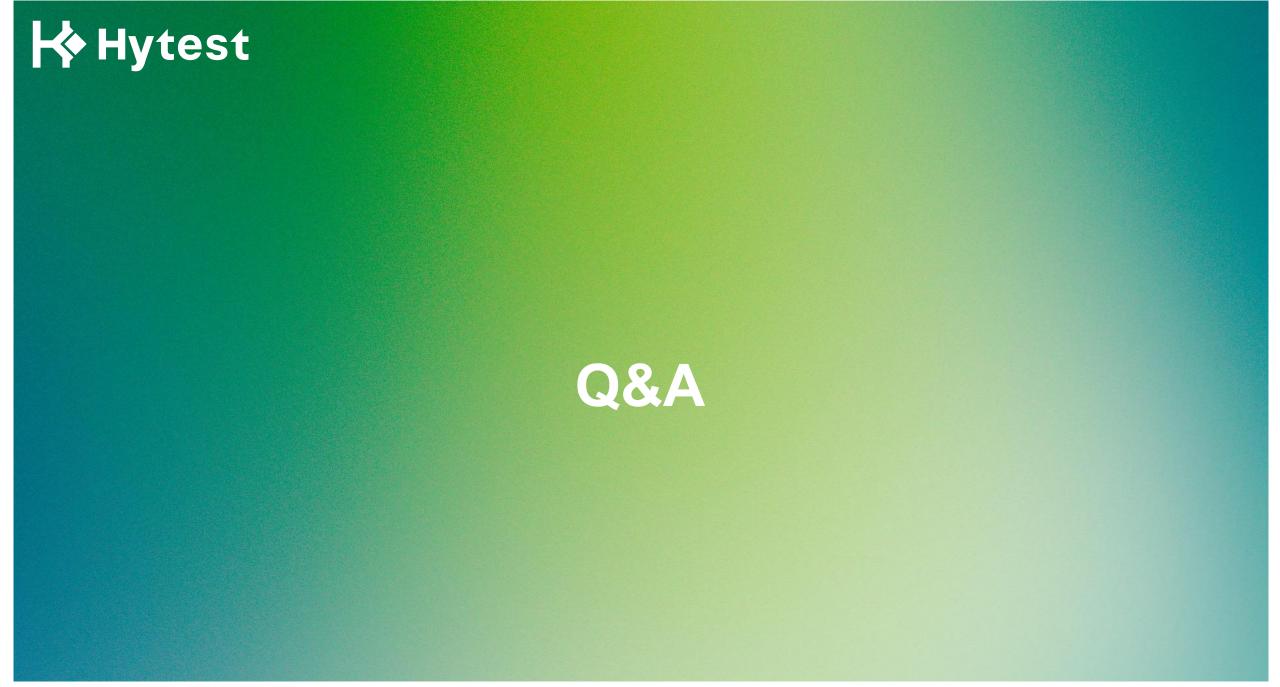
The assessed HIV recombinant antigens using Mindray CLIA System demonstrate high sensitivity and specificity.





Thank you

Jianwen He





Q: Can you explain your quality control process that you are using for your (Hytest's) products?

A: Hytest's quality control process can be described like this:

- -We use raw materials/chemical reagents only from suppliers who are approved on our list of acceptable suppliers (based on quality, reliability etc) and we follow performance of our suppliers continuously
- -In-coming inspection is performed on the materials (this is in many case a document review to confirm acceptable specifications etc.)
- -Only raw material from product specific list of reagents can be used in manufacturing and QC, the reagents on the list are carefully selected and qualified by R&D before product launch
- -All product lots are QC-tested to confirm that lots correspond to internal specifications and pass predetermined QC acceptance criteria. Hytest uses a broad variety of QC tests for produced proteins, such as appearance control, purity determined by means of SDS-PAGE or capillary electrophoresis, proteins' immunoreactivity determined by means of sandwich immunoassays which use protein-specific monoclonal or polyclonal antibodies.



Q: Why develop 5th-gen assays, when 4th-gen is "good enough"?

A: 4th-generation assays are indeed effective and widely utilized; however, they yield only qualitative "positive/negative" results for antibodies and do not provide information regarding the stage of infection. In contrast, 5th-generation assays offer significant additional advantages. First, they enable differentiation among HIV-1 p24 antigen, HIV-1 antibodies, and HIV-2 antibodies. This enhanced analytical resolution facilitates earlier detection of acute infections and reduces the necessity for confirmatory testing—potentially lowering associated testing costs by up to 40%. In markets such as those in Europe and the United States, where molecular diagnostic testing is particularly costly, this capability represents a compelling value proposition.



Q: How critical is recombinant viral protein stability in 4th/5th-gen reagents?

A: Recombinant viral protein stability is of critical importance for both fourth- and fifth-generation HIV assays.

From the raw material supplier's perspective, the stability of HIV-specific proteins represents a key determinant in the performance of HIV-specific reagents. The use of mammalian expression systems enables the production of proteins with high intrinsic stability during manufacturing, storage, and handling. At Hytest, we systematically optimize protein expression, purification, and storage conditions to maximize protein stability. Furthermore, proper posttranslational modifications and the strategic inclusion of protein tags contribute significantly to the enhanced stability of HIV proteins produced in mammalian cells.

From the assay developer's standpoint, even highly stable recombinant proteins may remain susceptible to aggregation and, in some cases, degradation after incorporation into the assay format—whether immobilized as coated antigens or used as conjugated detection reagents. Therefore, careful selection of raw materials is essential, and stability must be actively maintained through thoughtful reagent design and formulation. This involves optimizing buffer composition, detergents, stabilizers, and overall process parameters to ensure consistent assay performance, particularly given the high sensitivity required in HIV testing.

Reliable performance in fourth- and fifth-generation HIV assays is thus achieved through a combination of robust upstream protein production and meticulous downstream assay formulation.



Q: Can synthetic peptides replace recombinant antigens?

A: Synthetic peptides can't fully replace recombinant proteins. HIV proteins have several epitopes or antigenic determinants for antibody production on their surface. Epitopes can be either continuous or linear, in which several amino acids, usually from six to eight, are present in the sequence; or discontinuous, in which the amino acids that form the epitope are brought together by the three-dimensional folding of the protein. Even though an epitope constitutes only a relatively few amino acids, its reactivity with the antibody is influenced by amino acids not included in the epitope, or by the presence of post translational modifications, which may also be located outside the epitope.

Moreover, it's easier to take into account the genetic diversity while using a recombinant protein, not the single peptide. More epitopes you have in your protein sequence, the more likely you will be able to determine the presence of specific antibodies.

In some cases, to ensure higher sensitivity in the context of recognizing a wide variety of strains, a specific peptide can be added to the test system, but it's not possible to replace the test system with peptides completely. However, as we know there are approaches in the market, which combine the recombinant proteins with the gp41-specific peptides.



Q: Is it necessary and if so, how often and in what way should IVD companies (test system manufacturers) evaluate/verify the current HIV assay for its ability to detect newly emerging HIV strains?

A: HIV is a highly genetically variable virus and circulating recombinant forms (CRFs) continue to emerge and spread globally. In vitro diagnostic (IVD) manufacturers should routinely evaluate their HIV immunoassay reagents for the ability to detect emerging HIV strains and CRFs based on a risk-based approach. While there is no universally defined interval for such evaluations, the following strategies are recommended:

- 1. Maintain an updated inventory of newly identified HIV-1 group M subtypes, novel CRFs, and mutations that may impact antigen—antibody binding. Risk assessments should be periodically revised based on new findings to enable early identification of potential assay vulnerabilities.
- 2. Continuously monitor field complaints, proficiency testing outcomes, regulatory alerts, and scientific literature. If data trends indicate reduced assay sensitivity in specific geographic regions or sample types, this should prompt immediate reassessment.
- 3. Conduct formal re-validation when high-risk viral variants are detected. Such evaluations should employ test panels containing recent clinical isolates and newly characterized CRFs to verify that the assay continues to meet its claimed sensitivity and specificity performance.
- 4. Perform routine lot-to-lot verification testing using HIV reference panels as required by regulatory authorities to ensure manufacturing consistency. Any modifications to antigens, reagents, or production processes should also trigger targeted reevaluation for strain coverage.

Implementing these measures provides a robust framework for ensuring reliable HIV detection across evolving viral strains and emerging CRFs, thereby supporting long-term assay performance and public health surveillance.



Q: Do you have experience in the use of anti-p24 monoclonal antibodies in lateral flow assays?

A: At this stage, we have not conducted internal testing specifically on the lateral flow (LF) platform. However, several customers are currently evaluating our anti-p24 antibodies in their own LF assay development, and these projects are ongoing. While we cannot yet share finalized data, the initial feedback from these evaluations has been positive. Because lateral flow systems can vary significantly in design, materials, and assay configuration, performance is often highly platform-dependent. For this reason, we strongly encourage customers to test the antibodies directly in their own LF setup to obtain the most accurate, application-specific results.



Q: Thank you for the excellent talk! Can you speak to stability testing done for the recombinant HIV proteins?

A: During the product development phase, we conduct both accelerated stability studies and freeze—thaw stability evaluations on our recombinant HIV proteins. When studying stability, the change in protein immunoreactivity is assessed using a sandwich immunofluorescence assay using polyclonal antibodies specific to the protein. Protein resistance to freeze/thaw and to the storage under different temperature conditions is also assessed using SDS-electrophoresis. These tests allow us to confirm that the proteins remain stable under various stress conditions.



Thank you!

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