Sensitivity and specificity of the new Bio-Rad HIV screening test, Access HIV combo V2

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Running title: Sensitivity and specificity of the Access HIV combo V2

Abstract (250/250 mots max):

Background: Diagnosing of Human Immunodeficiency Virus (HIV) types 1 and 2 requires a screening with a highly sensitive and specific enzyme immunoassay and a low detection limit for the HIV-1 p24 antigen to minimize the diagnostic window.

Objectives: To determine the sensitivity, specificity and p24 limit of detection of the Access HIV combo V2 assay.

Study design: Retrospective part of sensitivity: 452 HIV-1 positive samples from 403 chronic (9 different HIV-1 group M subtypes, 22 different HIV-1 group M CRFs, 3 HIV-1 group O), 49 primary HIV-1 infections, 103 HIV-2 positive samples assessed at Pitié-Salpêtrière Hospital, 600 untyped HIV-1, 10 subtype-D and 159 untyped HIV-2 samples assessed in Bio-Rad Laboratories. Prospective part of clinical specificity: all consecutive samples in two blood donor facilities and Pitié-Salpêtrière (6570 patients) tested with Access HIV combo V2 and respectively Prism HIV O Plus (Abbott) or Architect HIV Ag/Ab Combo (Abbott) for Ag/Ab screening, and Procleix Ultrio (Gen Probe) for HIV RNA screening. Limit of detection of p24 antigen was assessed on recombinant virus-like-particles (10 HIV-1 group M subtypes/CRFs, HIV-1 group O).

Results: Sensitivity (95% CI) of Access HIV combo V2 was 100% (99.63-100) for HIV-1 chronic infection, 100% (98.55-100) for HIV-2 chronic infection and 100% (93.00-100) for HIV-1 primary infection. Specificity (95% CI) was 99.98 (99.91-100). Limit of detection for p24 antigen was around 0.43 (IQR [0.38-0.56]) IU/mL, and consistent across the 11 analyzed subtypes/CRFs.

Conclusions: with both high sensitivity and specificity, Access HIV combo V2 is a suitable screening assay for HIV-1/2 infection.

Keywords: Access HIV combo V2, HIV, sensitivity, specificity, accuracy, serology

Introduction

The first target to achieve the 2025 World Health Organization target of ending the Human Immunodeficiency Virus (HIV) pandemic is that at least 95% of people living with HIV knows their status [1]. HIV diagnosis faces three main challenges. Firstly, it has to detect antibodies directed to a remarkable range of diverse antigens from both HIV-1 and HIV-2 [2,3]. Secondly, as a non-negligible part of HIV transmission occurs early after an infection [4–6], diagnostic window should be as reduced as possible. As a consequence, the assay must detect the HIV-1 specific p24 antigen at the lowest limit of detection possible. Lastly, as a false positive HIV diagnosis can have deleterious consequences, a high specificity is needed [7–9].

HIV diagnosis tests have remarkably improved since the beginning of the HIV pandemic, with current recommended ones, or 4th generation assays, able to detect with high sensitivity and specificity all circulating variants with a diagnostic window of about 2 weeks after infection [1,10–12].

The objective of this study was to assess sensitivity and specificity of Bio-Rad's new 4th generation HIV test, Access HIV combo V2 on both real-life settings and well characterized commercial panels. Additionally, we aimed to establish its limit of detection for the p24 antigen using diverse, well-characterized commercial panels.

Materials and methods

Study design – sample collection.

This study included a multicenter retrospective part of clinical sensitivity and a multicenter retrospective and prospective part of clinical specificity.

Two sites were involved in clinical sensitivity. The Service of Virology at the Pitié-Salpêtrière Hospital (Paris, France) provided 452 HIV-1 positive samples from 403 chronic and 49 primary HIV-1 infected patients, and 103 HIV-2 positive samples from chronically infected patients. HIV-1 serum samples from chronically infected patients were selected to account for a large part of HIV-1 diversity (sup table 1) : 9 different HIV-1 group M subtypes, 22 different HIV-1 group M CRFs, 3 HIV-1 group O. Serum samples from HIV-1 primary infection, of which 9 were positive for p24-antigen only (Stage II and III Fiebig [13]), were mostly of HIV-1 subtype B, CRF02 and CRF06 (sup table 1). HIV-antibody and p24 positivity were assessed using Architect HIV Ag/Ab Combo (Abbott®, Rungis, France) and Liaison XL HIV Ab/Ag (Diasorin, Antony, France) as screening assays, with New LAV Blot I and II (Bio-Rad laboratories, Marnes-la-Coquette, France) as confirmatory and differentiation assays. Subtypes and recombinant forms of HIV-1 strains were determined using molecular assay as previously described [14]. Samples were stored frozen at -20°C until use. HIV-positive samples tested at the Bio-Rad Laboratories originated from commercial panels supplied by Seracare, Zeptometrix, and Biomex for seroconversion panels, as detailed in Sup Table 2. Days to first reactive results were compared with the Architect assay. For the Architect assay, we used seroconversion panel data provided by the manufacturer, followed if unavailable by the FDA's published data. If this data was still missing, we extended our review to previously published studies. Chronic HIV samples tested in Bio-Rad Laboratories included a total of 600 ungenotyped HIV-1, 10 HIV-1 subtype D samples and 159 ungenotyped HIV-2 samples.

For the retrospective part of clinical specificity, 203 frozen serum samples from HIV-negative pregnant women who consulted at Pitié-Salpêtrière Hospital from April 2019 to January 2020 and 10

HTLV+/HIV- serum samples were analyzed. HIV status was assessed using Architect HIV Ag/Ab Combo.

For the prospective part of clinical specificity were included: fresh serum or plasma samples from all consecutive patients with an HIV testing at Pitié-Salpêtrière Hospital from 30/01/2020 to 17/03/2020 (1509 samples), all samples from consecutive blood donors at the Etablissement Français du Sang (EFS) of Bois-Guillaume (France) from 28/01/2020 to 11/02/2020 (2512 samples), and all consecutive blood donors from the EFS of Lille (France) from 10/02/2020 to 13/03/2020 (2549 samples).

Access HIV combo V2 assay

The Access HIV combo V2 assay is a paramagnetic-particle, semi-quantitative chemiluminescent immunoassay (CLIA) designed to detect HIV-1 p24 and HIV-2 p26 antigens, and antibodies to HIV-1 and HIV-2 in human serum or plasma. The test is configured to run on Beckman Coulter's Access immunoassay systems, with a run time of 30 minutes. Results are expressed as S/CO ratio with S/CO <0.9 considered as non-reactive, S/CO ≥1 reactive and S/CO [0.9-<1] gray zone. This test does not distinguish antigen from antibody reactivity.

Sample processing

For the retrospective parts of sensitivity and specificity, all samples were tested with the Access HIV combo V2 according to the manufacturer's recommendations. Test found negative for the study of sensitivity were planned to be repeated once.

For the prospective part of specificity, all samples were tested in parallel with Access HIV combo V2 and either Architect HIV Ag/Ab Combo at the Pitié-Salpêtrière Hospital, or Prism HIV O Plus (Abbott) and Procleix Ultrio (Novartis Diagnostics, Emeryville, CA, USA) at the EFS centers. All gray zone and reactive samples were retested in duplicate, followed, if remaining gray zone or reactive, by a confirmatory assay. Confirmatory assays were conducted using either New LAV Blot I and II at Pitié Salpêtrière Hospital, or INNO-LIA HIV I/II Score (Innogenetics, Gent, Belgium) at the EFS centers.

P24 and p26 antigens analytical sensitivity

Access HIV combo V2 assay sensitivity for p24 and p26 antigens was assessed on the Access 2 platforms using the NIBSC/WHO p24 antigen standards (NIBSC 90/636 and 16/210) and p26 antigen (NIBSC 16/236) with the following dilutions: 1:2, 1:4, 1:8, 1:16, 1:32. Both antigens were diluted in sterile water.

Statistical analysis

Statistical analysis was conducted using R version 4.2.1 software [15]. Categorical variables were expressed as numbers (percentages) and continuous variables as medians (interquartile ranges [IQR]). The 95% confidence intervals (95% CI) were calculated using Wilson confidence interval for proportions [16]. Limit of detection was assessed for p24 antigen using linear regression to calculate the amount of p24 detected for a S/CO of 1. Comparison was done using Chi-squared test for categorical variables, with a significance assigned at a p value < 0.05. Sample size was determined according to the European Commission decision on technical specifications for in vitro diagnostic medical devices [17].

Role of the study sponsor

The sponsor provided reagents and automated equipment used in this study and was responsible for data collection. First and last authors had full access to the study database, generated statistical analyses, prepared first draft of the manuscript and made the decision to submit the manuscript for

publication. Y.C, D.N and M.C are employed by Bio-Rad. V.G, A.GD, C.D, S.G received no personal funding from the study sponsor.

Ethics

The study was conducted according to the principles of the Declaration of Helsinki and in conformity with institutional regulations and guidelines. The evaluated method was performed on sample excess. Patients were systematically notified of any supplementary biological analyses on frozen samples, initially collected as part of routine clinical practice.

Results

Sensitivity on chronic HIV infection

All the 403 HIV-1 samples from the retrospective study collected at the Pitié-Salpêtrière Hospital as well as the 25 HIV-1 samples collected during the prospective study of specificity were reactive, yielding a sensitivity of 100% (95%CI [99.11-100]). Each of the 600 ungenotyped and 10 HIV-1 subtype D samples tested in Bio-Rad Laboratories were also reactive, owing a 100% (95% CI [99.37-100]) sensitivity in Bio-Rad Laboratory.

All the 103 retrospective chronically HIV-2 samples collected at Pitié-Salpêtrière Hospital as well as all the 159 HIV-2 samples at Bio-Rad Laboratories were positive, owing a sensitivity of 100% (95% CI [96.40-100]) at Pitié-Salpêtrière and 100% (95% CI [97.64-100]) in Bio-Rad Laboratories.

Pooled estimated sensitivity for the diagnosis of chronic HIV-1 infection was 100% (95% CI [99.63-100]) and for the diagnosis of chronic HIV-2 infection at 100% (95% CI [98.55-100]). These results are summarized fig 1, and S/CO values reported in sup fig 1.

Sensitivity for HIV-1 primary infection

All the 49 retrospective and 2 prospective samples from primary HIV-1 infection collected at Pitié-Salpêtrière Hospital were found reactive on Access HIV combo V2, yielding a sensitivity of 100% (95% CI [93.00-100]). Corresponding S/CO values are reported in sup fig 1.

A total of 415 samples from 41 commercial seroconversion panels were tested in Bio-Rad Laboratories. Results aligned closely with Architect (reference assay). Day to first reactive result was identical for 35 (85.4%) of them, while Access outperformed for 5 (12.2%, with respectively 5, 6, 7, 3 and 5 days later for the Architect) and underperformed for 1 (2.4%, with 4 days earlier for the Architect) of them (sup table 2). Discrepant results between the two assays are summarized table 1.

Specificity

Blood donor samples, routinely tested at EFS Hauts de France–Normandie in Lille (n=2549) and Bois-Guillaume (n=2512) with *Prism HIV O plus* and *Procleix Ultrio* were prospectively analyzed in parallel with Access HIV combo V2. No initial false-reactive sample was identified at the first site, while at the second site, 4 samples were found initial false-reactive (IR). These 4 IR samples were negative after repeating in duplicate. This resulted in an overall IR specificity for blood donors of 99.92% (95% CI [99.80-99.97]) and an overall specificity after repeat testing of 100% (95% CI [99.92-100]). In comparison, Prism HIV O plus assay gave 4 RR false-reactive samples on the blood donor's population. There was no false-reactive sample identified with the Nucleic Acid Amplification Test (NAAT) *Procleix Ultrio*, owing a specificity at 100% (95% CI [99.92-100]). Of note, no sample was identified as true reactive.

At Pitié-Salpêtrière Hospital, among the 1509 samples tested prospectively in parallel with Architect, 27 (1.8%) were found to be true reactive samples (25 chronic and 2 primary infections), while 5 were IR and 1 was repeatedly false-reactive. As so, the hospitalized patient IR specificity assessed at was 99.66% (95% CI [99.74-99.83) and RR specificity 99.93% (95% CI [99.62-99.99]). Architect specificity was identical, with 2 other repeatedly false-reactive samples. We also performed a retrospective exploratory analysis on 203 hospitalized HIV-negative pregnant women. There was no false-reactive sample, owing a specificity of 100% (95% CI [98.1-100]). Of note, RR hospitalized patient specificity was not statistically significantly lower than blood donor specificity (p = 0.51). Also, we performed an exploratory analysis on 10 HTLV-1 positive / HIV-negative samples. One sample tested reactive, with an S/CO ratio at 1.46, repeated at 1.86 and 1.81. Architect was also reactive on this sample. As the HIV Western blots were negative, this sample was considered as a false reactive sample.

As so, pooled (blood donors and hospitalized patients) IR specificity was 99.86% (95% CI [99.74-99.83]) and RR specificity was 99.98% (95% CI [99.91-100%]). Specificity results are summarized fig 2 and S/CO distribution for negative results summarized sup fig 1.

Analytical p24 and p26 antigens sensitivity

The limits of detection for p24 and p26 were assessed using WHO standardized panels (Table 2). The 1st international reference sample for HIV-1 Subtype B had a detection limit of 0.39 IU/mL. The limit of detection for the p24 antigen was consistent across the 11 HIV-1 subtypes, ranging from 0.27 to 0.58 IU/mL with a median of 0.43 (IQR [0.38-0.56]). Since p26 had no assigned unitage, its limit of detection was determined using serial dilution, with samples yielding positive results up to a dilution of 1/8.

Discussion

Overall, Access HIV combo V2 displayed high sensitivity for both chronic HIV-1 samples with a sensitivity of 100% (95% CI [99.63-100]), and for HIV-1 primary infection samples with a sensitivity of 100% (95%CI [93.00-100]). The sensitivity for HIV-2 infection was also 100% (95% CI [98.55-100]). Specificity was also high, at 99.98% (95% CI [99.91-100]). Limit of detection of p24 antigen was low, around 0.43 IU/mL and consistent across the analyzed HIV-1 groups, subtypes/CRFs.

High sensitivity and specificity were expected findings, consistent with previous reports from all other commercial 4th generation assays [11,18–23]. Analytical sensitivity for p24 antigen was around 0.43 IU/mL, consistent across the 11 HIV-1 subtypes/CRFs. This finding contrasts with the previous Access HIV combo version, which had a limit of detection for subtype B around 3 times higher [24] and performed very poorly on non-subtype B samples, with limit of detection often over 10 IU/mL [25]. Compared with published data [26], Access HIV combo V2 (median, [IQR] p24 Ag limit of detection of 0.43 IU/mL, [0.38-0.56 IU/mL]) had a similar limit of detection as ARCHITECT HIV Ag/Ab Combo (0.57 IU/mL, [0.43-0.64 IU/mL]) and BioPlex 2200 HIV Ag-Ab assays (0.27 IU/mL [0.21-0.36]), outperformed Liaison® XL Murex HIV ab/Ag and Elecsys® HIV combi PT assays (0.67 IU/mL, [0.58-0.72]), but underperformed if compared with the Elecsys HIV Duo assay (0.33 IU/mL, [0.30-0.37 IU/mL), as described in sup table 3. However, we were unable to link these gaps in detection thresholds to potential difference in HIV-1 window period, as to date we have not managed to gather any non-reactive 4th generation HIV-1 sample that was positive on HIV-1 NAAT. As this contrast with current guidelines that advise the use of NAAT in this setting [27,28], further studies are needed to address the relevance of this guideline in the setting of increasing p24 sensitivities of 4th generation assays. This study has several limitations. The genotypes of HIV-1 responsible for primary infection at Pitié-Salpêtrière were predominantly HIV-1 group M subtype B and CRF02_AG, reflecting the French and European epidemiologies [2]. Furthermore, since their sera were primarily screened using the

Architect platform, this part of the study couldn't ascertain if Access had a shorter window period. Regarding the commercial seroconversion panel, a similar bias toward subtype B might exist since most blood samples originate from US patients. Furthermore, Architect's results were extracted from manufacturer's data or previously published studies (Sup table 2) instead of being generated from samples stored within the same conditions. The limits of detection for p24 antigen from different HIV-1 subtypes and p26 antigen were derived from commercial recombinant virus-like particles rather than patient sera, to facilitate future comparison. This specification, however, represents a surrogate marker for HIV-1 primary infection and cannot be rigorously translated into window periods. Finally, this assay is designed to detect p26 antigen, to shorter the window period for HIV-2 infection. However, due to a lack of HIV-2 primary infection sample we were unable to validate this hypothesis.

As a conclusion, Access HIV combo V2, with both high sensitivity and specificity is a suitable screening assay for HIV-1 and HIV-2 infections.

References

[1] Consolidated guidelines on HIV prevention, testing, treatment, service delivery and monitoring: recommendations for a public health approach, World Health Organization, Geneva, Switzerland, 2021.

[2] N. Bbosa, P. Kaleebu, D. Ssemwanga, HIV subtype diversity worldwide, Current Opinion in HIV and AIDS 14 (2019) 153–160. https://doi.org/10.1097/COH.0000000000000534.

[3] B. Visseaux, M. Bertine, Q. Le Hingrat, V. Ferré, C. Charpentier, F. Collin, F. Damond, S. Matheron, S. Hué, D. Descamps, HIV-2 diversity displays two clades within group A with distinct geographical distribution and evolution, Virus Evolution 7 (2021) veab024. https://doi.org/10.1093/ve/veab024.

[4] C.D. Pilcher, H.C. Tien, J.J. Eron, Jr., P.L. Vernazza, S. Leu, P.W. Stewart, L. Goh, M.S. Cohen, Quest Study and the Duke-UNC-Emory Acute HIV Consortium, Brief but Efficient: Acute HIV Infection and the Sexual Transmission of HIV, J INFECT DIS 189 (2004) 1785–1792. https://doi.org/10.1086/386333.

[5] E.D.M.B. Kroon, N. Phanuphak, A.J. Shattock, J.L.K. Fletcher, S. Pinyakorn, N. Chomchey, S. Akapirat, M.S. De Souza, M.L. Robb, J.H. Kim, F. Van Griensven, J. Ananworanich, D.P. Wilson, RV254/SEARCH 010 Study Group, Acute HIV infection detection and immediate treatment estimated to reduce transmission by 89% among men who have sex with men in Bangkok, Journal of the International AIDS Society 20 (2017) 21708. https://doi.org/10.7448/IAS.20.1.21708.

[6] C. Verhofstede, V. Mortier, K. Dauwe, S. Callens, J. Deblonde, G. Dessilly, M.-L. Delforge, K. Fransen, A. Sasse, K. Stoffels, D. Van Beckhoven, F. Vanroye, D. Vaira, E. Vancutsem, K. Van Laethem, Exploring HIV-1 Transmission Dynamics by Combining Phylogenetic Analysis and Infection Timing, Viruses 11 (2019) 1096. https://doi.org/10.3390/v11121096.

[7] R. Bhattacharya, S. Barton, J. Catalan, When good news is bad news: psychological impact of false positive diagnosis of HIV, AIDS Care 20 (2008) 560–564. https://doi.org/10.1080/09540120701867206.

[8] S.M. Coleman, N. Gnatienko, C.A. Lloyd-Travaglini, M.R. Winter, C. Bridden, E. Blokhina, D. Lioznov, J. Adong, J.H. Samet, T. Liegler, J.A. Hahn, False-positive HIV diagnoses: lessons from Ugandan and Russian research cohorts, HIV Clinical Trials 19 (2018) 15–22. https://doi.org/10.1080/15284336.2018.1429846.

[9] C.S. Kosack, L. Shanks, G. Beelaert, T. Benson, A. Savane, A. Ng'ang'a, B. Andre, J.-P.B. Zahinda, K. Fransen, A.-L. Page, HIV misdiagnosis in sub-Saharan Africa: performance of diagnostic algorithms at six testing sites, Journal of the International AIDS Society 20 (2017) 21419. https://doi.org/10.7448/IAS.20.1.21419.

[10] Centers for Disease Control and Prevention (U.S.), B. Bernard M., Association of Public Health Laboratorie, O. S. Michele, W. Laura G., B. Berry, W. Barbara G., W. Kelly E., P. Michael A., Laboratory testing for the diagnosis of HIV infection : updated recommendations, Centers for Disease Control and Prevention, 2014. https://doi.org/10.15620/cdc.23447.

[11] V. Lemee, M. Leoz, M. Etienne, F. De Oliveira, J.-C. Plantier, Performance of the Liaison XL Murex HIV Ab/Ag Test on Clinical Samples Representing Current Epidemic HIV Variants, J Clin Microbiol 52 (2014) 3277–3279. https://doi.org/10.1128/JCM.01089-14.

[12] T.D. Ly, A. Ebel, V. Faucher, V. Fihman, S. Laperche, Could the new HIV combined p24 antigen and antibody assays replace p24 antigen specific assays?, Journal of Virological Methods 143 (2007) 86–94. https://doi.org/10.1016/j.jviromet.2007.02.013.

[13] E.W. Fiebig, D.J. Wright, B.D. Rawal, P.E. Garrett, R.T. Schumacher, L. Peddada, C. Heldebrant, R. Smith, A. Conrad, S.H. Kleinman, M.P. Busch, Dynamics of HIV viremia and antibody seroconversion in plasma donors: implications for diagnosis and staging of primary HIV infection, AIDS 17 (2003) 1871–1879. https://doi.org/10.1097/00002030-200309050-00005.

[14] V. Guiraud, J. Bocobza, M. Desmonet, F. Damond, J.-C. Plantier, G. Moreau, M. Wirden, K. Stefic, F. Barin, A. Gautheret-Dejean, Are Confirmatory Assays Reliable for HIV-1/HIV-2 Infection

Differentiation? A Multicenter Study, J Clin Microbiol (2023) e00619-23. https://doi.org/10.1128/jcm.00619-23.
[15] R Core Team, R: A Language and Environment for Statistical Computing, (n.d.).
[16] R.G. Newcombe, Two-sided confidence intervals for the single proportion: comparison of seven methods, Statist. Med. 17 (1998) 857–872. https://doi.org/10.1002/(SICI)1097-0258(19980430)17:8<857::AID-SIM777>3.0.CO;2-E.
[17] European Commission, Commission Implementing Decision (EU) 2019/1244 of 1 July 2019 amending Decision 2002/364/EC as regards requirements for HIV and HCV antigen and antibody combined tests and as regards requirements for nucleic acid amplification techniques with respect to reference materials and qualitative HIV assays, 2019. https://eur-lex.europa.eu/eli/dec_impl/2019/1244/oj.
[18] H. Kutvonen, H. Jarva, M. Lappalainen, S. Kurkela, Comparative evaluation of four commercial analyzers for the serological screening of hepatitis A, B, C and HIV, Journal of Clinical Virology 153 (2022) 105219. https://doi.org/10.1016/j.jcv.2022.105219.
[19] M. Bhatta, S. Banerjee, S. Nandi, S. Dutta, M.K. Saha, Performance of commercially available HIV in vitro diagnostic assays: A systematic review and meta-analysis, Journal of Clinical Virology 146 (2022) 105047. https://doi.org/10.1016/j.jcv.2021.105047.
[20] K.T.D. Thai, H. Götz, B.C.G.C. Slingerland, J. Klaasse, M. Schutten, C.H. GeurtsvanKessel, An analysis of the predictive value of the HIV Ag/Ab screening assay within the performance characteristics of the DiaSorin LIAISON XL for the detection of blood-borne viruses, Journal of Clinical Virology 102 (2018) 95–100. https://doi.org/10.1016/j.jcv.2018.02.018.
[21] D. Wiredja, T.A. Ritchie, G. Tam, C.A. Hogan, B. Pinsky, R.Z. Shi, Performance evaluation and optimized reporting workflow for HIV diagnostic screening and confirmatory tests in a low prevalence setting, Journal of Clinical Virology 145 (2021) 105020. https://doi.org/10.1016/j.jcv.2021.105020.
[22] S. Crowe, B. Bennett, S. Fordan, Impact of the 2014 CDC HIV testing guidelines on detection of acute HIV infections, Journal of Clinical Virology 146 (2022) 105058. https://doi.org/10.1016/j.jcv.2021.105058.
[23] T. Sano, M. Kondo, Y. Yoshimura, N. Tachikawa, H. Sagara, I. Itoda, K. Yamanaka, K. Sudo, S. Kato, M. Imai, Evaluation of a New Vesion of the Human Immunodeficiency Virus Antigen and Antibody Combination Assay with Improved Sensitivity in HIV-1 p24 Antigen Detection, J. J. A. Inf. D 87 (2013) 415–423. https://doi.org/10.11150/kansenshogakuzasshi.87.415.
[24] T.D. Ly, J.C. Plantier, L. Leballais, S. Gonzalo, V. Lemée, S. Laperche, The variable sensitivity of HIV Ag/Ab combination assays in the detection of p24Ag according to genotype could compromise the diagnosis of early HIV infection, Journal of Clinical Virology 55 (2012) 121–127. https://doi.org/10.1016/j.jcv.2012.06.012.
[25] B.N. Vetter, V. Orlowski, K. Fransen, C. Niederhauser, V. Aubert, M. Brandenberger, D. Ciardo, G. Dollenmaier, T. Klimkait, S. Regenass, P. Schmid, V. Schottstedt, F. Suter-Riniker, S. Yerly, C. Shah, J. Böni, J. Schüpbach, Generation of a Recombinant Gag Virus-Like-Particle Panel for the Evaluation of p24 Antigen Detection by Diagnostic HIV Tests, PLoS ONE 9 (2014) e111552. https://doi.org/10.1371/journal.pone.0111552.
[26] X. Qiu, L. Sokoll, T. Duong Ly, C. Coignard, S.H. Eshleman, P. Mohr, C. Huizenga, P. Swanson, G. Cloherty, J. Hackett Jr., An improved HIV antigen/antibody prototype assay for earlier detection of acute HIV infection, Journal of Clinical Virology 145 (2021) 105022. https://doi.org/10.1016/j.jcv.2021.105022.
[27] Morlat Philippe, Prise en charge médicale des personnes vivant avec le VIH - Recommandations du groupe d'experts, in: France, 2018. https://cns.sante.fr/wp-content/uploads/2018/04/experts-vih_prevention-depistage.pdf (accessed February 28, 2024).
[28] E.A. DiNenno, J. Prejean, K. Irwin, K.P. Delaney, K. Bowles, T. Martin, A. Tailor, G. Dumitru, M.M. Mullins, A.B. Hutchinson, A. Lansky, Recommendations for HIV Screening of Gay, Bisexual, and Other Men Who Have Sex with Men — United States, 2017, MMWR Morb. Mortal. Wkly. Rep. 66 (2017) 830–832. https://doi.org/10.15585/mmwr.mm6631a3.

345 Table 1: Commercial seroconversion panels with conflicting results between Access HIV combo V2
346 and Abbott's Architect.

Vendor	Sample ID	Days to first reactive result		Difference of first reactive result (Days)*	Source for Architect results
		Access Combo V2	Architect		
Seracare / BBI	PRB944	2	7	5	FDA notice
	PRB945	7	13	6	T. Sano et al [23]
	PRB953	7	3	-4	Manufacturer
	PRB957	16	23	7	FDA notice
	SC9018	25	28	3	Manufacturer
	SC12008	23	28	5	Manufacturer

347 * a positive number indicates that Access Combo V2 is reactive before Architect, while a negative
348 result indicates that Architect is reactive before Access Combo V2.

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Table 2: Analytical sensitivity of the Access HIV combo V2 assay on the Access platform for HIV-1 p24 antigen according to group, subtypes and CRFs

HIV-1 Subtype	Analytical Sensitivity (IU/mL)
B[a]	0.39
A1[b]	0.56
B[b] (16/214)	0.27
B[b] (16/216)	0.35
C[b]	0.47
D[b]	0.53
F1/CRF12_BF/BFrec[b]	0.43
G[b]	0.56
CRF20_BG[b]	0.38
CRF01_AE[b]	0.56
CRF02_AG[b]	0.36
H[b]	0.42
Group O[b]	0.58

[a] WHO reference panel 90/636

[b] WHO reference panel 16/210

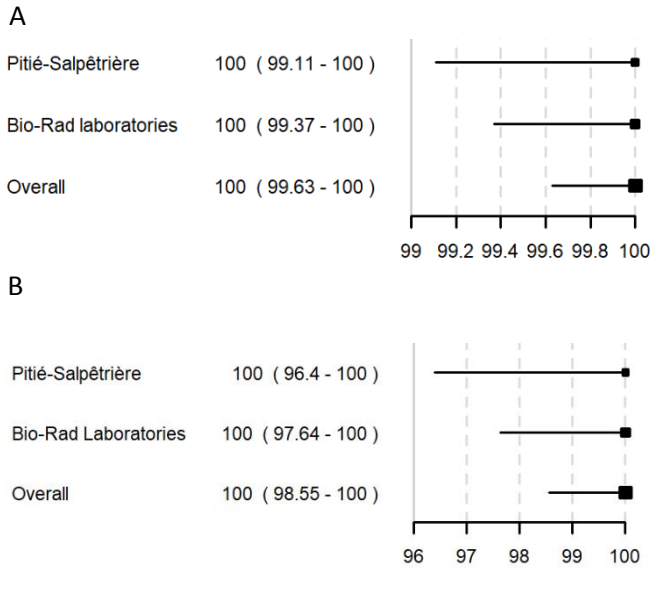
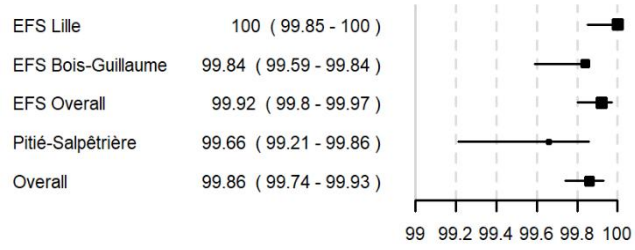


Figure 1: Sensitivity of the Access HIV combo V2 for HIV-1 (A) and HIV-2 (B) chronic infection

A



B

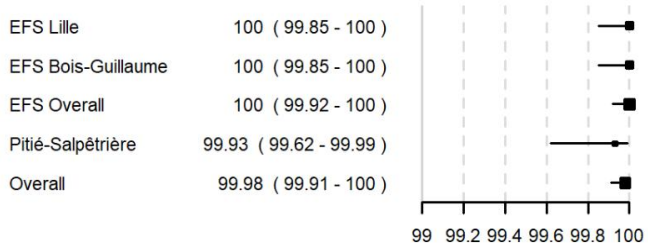


Figure 2: Specificity of the Access HIV combo V2 on first result (A) and after repeat (B)

Supplementary table 1: Subtypes and recombinant forms of retrospective HIV-1 samples used at Pitié-Salpêtrière Hospital.

Subtype/ group/CRF	Stage	No Specimens
Subtype A	Chronic	34
Subtype B	Chronic	94
Subtype C	Chronic	20
Subtype D	Chronic	14
Subtype F	Chronic	17
Subtype G	Chronic	29
Subtype H	Chronic	6
Subtype J	Chronic	1
Subtype K	Chronic	2
Group O	Chronic	3
CRF01	Chronic	22
CRF02	Chronic	97
CRF06	Chronic	15
CRF08	Chronic	1
CRF09	Chronic	7
CRF10	Chronic	1
CRF11	Chronic	8
CRF13	Chronic	6
CRF14	Chronic	6
CRF15	Chronic	2
CRF18	Chronic	3
CRF19	Chronic	2
CRF20	Chronic	1
CRF22	Chronic	2
CRF25	Chronic	1
CRF30	Chronic	1
CRF36	Chronic	2
CRF37	Chronic	1
CRF42	Chronic	1
CRF44	Chronic	1
CRF45	Chronic	1
CRF60	Chronic	2
Subtype A	Primary	1
Subtype B	Primary	20
Subtype C	Primary	2
Subtype D	Primary	1
CRF01/CRF15[a]	Primary	1
CRF02	Primary	10
CRF06	Primary	4
CRF18	Primary	2
Unknown	Primary	8

[a] Genotyping was unable to distinguish between CRF01 and CRF15

393 Supplementary table 2: Commercial seroconversion panels used for the present study with results
394 for Access HIV combo V2 and Abbott's Architect.

Vendor	Sample ID	Days to first reactive result		Source for Architect results
		Access Combo V2	Architect	
Seracare / BBI	PRB944	2	7	FDA notice [1]
	PRB945	7	13	T. Sano et al [2]
	PRB949	18	18	Manufacturer
	PRB950	18	18	Manufacturer
	PRB953	7	3	Manufacturer
	PRB954	17	17	Manufacturer
	PRB955	3	3	Manufacturer
	PRB957	16	23	FDA notice [1]
	PRB958	7	7	FDA notice [1]
	PRB964	22	22	Manufacturer
	PRB966	44	44	Manufacturer
	PRB969	63	63	Manufacturer
	PRB970	0	0	Manufacturer
	PRB973	7	7	Manufacturer
	PRB975	14	14	Manufacturer
	SC-0600-0270	30	30	Manufacturer
	SC-0600-0271	7	7	Manufacturer
	SC-0600-0272	18	18	Manufacturer
Zeptometrix	SC9011	36	36	Manufacturer
	SC9012	16	16	Manufacturer
	SC9013	25	25	Manufacturer
	SC9016	30	30	Manufacturer
	SC9018	25	28	Manufacturer
	SC9020	90	90	Manufacturer
	SC9021	47	47	Manufacturer
	SC9023	78	78	Manufacturer
	SC9024	53	53	Manufacturer
	SC9025	85	85	Manufacturer
	SC9026	44	44	Manufacturer
	SC9030	47	47	Manufacturer
	SC9031	146	146	Manufacturer
	SC9033	82	82	Manufacturer
	SC9089	16	16	Manufacturer
	SC6244	28	28	Manufacturer
	SC12008	23	28	Manufacturer
Biomex	SCP-HIV-002	63	63	Manufacturer
	SCP-HIV-003	17	17	Manufacturer
	SCP-HIV-004	56	56	Manufacturer
	SCP-HIV-005	16	16	Manufacturer
	SCP-HIV-006	15	15	Manufacturer
	SCP-HIV-007	12	12	Manufacturer

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Supplementary table 3: Summary of the p24 antigen limit of detection (IU/mL) on the WHO panel for six 4th generation assays. Data for comparative assays were extracted from Qiu et al. [3].

	Access HIV combo V2	ARCHITECT HIV Ag/Ab Combo	Liaison® XL murex HIV ab/Ag HT	Elecsys HIV Duo	Elecsys® HIV combi PT	BioPlex 2200 HIV Ag-Ab
Median (IQR)[1,2]	0.43 (0.38-0.56)	0.57 (0.43-0.64)	0.67 (0.58-0.72)	0.33 (0.30-0.37)	0.89 (0.74-1.04)	0.27 (0.21-0.36)
P-value for comparison with Access[3]	NA[4]	0.24	**0.0012**	**0.02**	**0.0005**	0.13

1: Inter Quartile Range

2: Results are expressed as IU/mL

3: Based on Wilcoxon's test for paired samples

4: Not applicable

Bibliography for Supplementary tables 2 and 3:

[1] ARCHITECT HIV Ag/Ab Combo package insert, 2010. https://www.fda.gov/media/116836/download.

[2] T. Sano, M. Kondo, Y. Yoshimura, N. Tachikawa, H. Sagara, I. Itoda, K. Yamanaka, K. Sudo, S. Kato, M. Imai, Evaluation of a New Vesion of the Human Immunodeficiency Virus Antigen and Antibody Combination Assay with Improved Sensitivity in HIV-1 p24 Antigen Detection, J. J. A. Inf. D. 87 (2013) 415–423. https://doi.org/10.11150/kansenshogakuzasshi.87.415.

[3] X. Qiu, L. Sokoll, T. Duong Ly, C. Coignard, S.H. Eshleman, P. Mohr, C. Huizenga, P. Swanson, G. Cloherty, J. Hackett Jr., An improved HIV antigen/antibody prototype assay for earlier detection of acute HIV infection, Journal of Clinical Virology. 145 (2021) 105022. https://doi.org/10.1016/j.jcv.2021.105022.