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Model systems of human immunodeficiency virus (HIV-1) for *in vitro* efficacy assessment of candidate vaccines and drugs against HIV-1

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Abstract. HIV infection still remains a major challenge for healthcare systems of the world. There are several aspects on counteracting the HIV/AIDS epidemic. The first aspect covers preventive measures including educational campaigns on HIV/AIDS and promotion of a healthy lifestyle, protected sex, and pre-exposure prophylaxis of vulnerable groups. The second aspect is timely HIV testing and the use of antiretroviral therapy when test results come back positive. The third aspect is the scientific research associated with discovering new pharmaceutical agents and developing HIV-1 vaccines. Selecting an adequate tool for quick and accurate *in vitro* efficacy assessment is the key aspect for efficacy assessment of vaccines and chemotherapy drugs. The classical method of virology, which makes it possible to evaluate the neutralizing activity of the sera of animals immunized with experimental vaccines and the efficacy of chemotherapy agents is the method of neutralization using viral isolates and infectious molecular clones, i.e. infectious viral particles obtained via cell transfection with a plasmid vector including the full-length HIV-1 genome coding structural, regulatory, and accessory proteins of the virus required for the cultivation of replication-competent viral particles in cell culture. However, neutralization assessment using viral isolates and infectious molecular clones is demanding in terms of time, effort, and biosafety measures. An alternative eliminating these disadvantages and allowing for rapid screening is the use of pseudoviruses, which are recombinant viral particles, for the analysis of neutralizing activity. Pseudotyped viruses have defective genomes restricting their replication to a single cycle, which renders them harmless compared to infectious viruses. The present review focuses on describing viral model systems for *in vitro* efficacy assessment of vaccines and drugs against HIV-1, which include primary HIV-1 isolates, laboratory-adapted strains, infectious molecular clones, and *env*-pseudoviruses. A brief comparison of the listed models is presented. The HIV-1 *env*-pseudoviruses approach is described in more detail.

Key words: HIV-1; primary isolates; infectious molecular clones; *env*-pseudoviruses; virus neutralization assay.

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Модельные системы вируса иммунодефицита человека (ВИЧ-1), используемые для оценки эффективности кандидатных вакцин и лекарственных препаратов против ВИЧ-1 *in vitro*

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Аннотация. ВИЧ-инфекция по-прежнему остается одной из глобальных проблем здравоохранения во всем мире. Борьба с инфекцией ведется по нескольким направлениям. Во-первых, это профилактические мероприятия, которые включают просвещение населения по проблеме ВИЧ/СПИДа, пропаганду здорового образа жизни, защищенные половые контакты, доконтактную профилактику уязвимых групп населения. Во-вторых, прохождение своевременного тестирования на ВИЧ и применение антиретровирусной терапии в случае его обнаружения. В-третьих, это научные исследования, связанные как с поиском новых лекарственных агентов, так и с разработкой вакцин против ВИЧ-1. Ключевой момент при определении эффективности вакцин и химиотерапевтических препаратов – выбор инструмента, позволяющего быстро и точно оценить их эффективность *in vitro*. Классическим методом вирусологии, позволяющим оценить нейтрализующую активность сывороток животных, иммунизированных экспериментальными вакцинами, и эффективность химиотерапевтических агентов, является метод нейтрализации с использованием вирусных изолятов, а также инфекционных молекулярных клонов, которые представляют собой инфекционные вирусные частицы, полученные путем

трансфекции клеток плазмидным вектором, содержащим полноразмерный геном ВИЧ-1, кодирующий структурные, регуляторные и вспомогательные белки вируса, необходимые для образования репликационно-компетентных вирусных частиц в культуре клеток. При этом метод нейтрализации с использованием вирусных изолятов и инфекционных молекулярных клонов отличается трудоемкостью, продолжительностью и требует повышенных мер биобезопасности. Альтернативным решением, устраняющим указанные недостатки и позволяющим проводить быстрый скрининг, является использование для анализа нейтрализующей активности псевдовирuсов, которые представляют собой рекомбинантные вирусные частицы. В отличие от инфекционных вирусuв, работа с псевдовирuсами безопасна, поскольку геном псевдовирuсов нарушен для того, чтобы их инфекция ограничивалась лишь одним циклом. Данный обзор посвящен описанию модельных вирусных систем, используемых для оценки эффективности вакцин и лекарственных препаратов против ВИЧ-1 *in vitro*: первичных изолятов ВИЧ-1 и лабораторно-адаптированных штаммов, инфекционных молекулярных клонов и *env*-псевдовирuсов. Кратко представлена их сравнительная характеристика. Более подробно описана технология *env*-псевдовирuсов ВИЧ-1.

Ключевые слова: ВИЧ-1; первичные изоляты; инфекционные молекулярные клоны; *env*-псевдовирuсы; анализ нейтрализации вирусuв.

Introduction

The HIV/AIDS pandemic still remains a major problem for healthcare systems of the world with about two million newly infected individuals every year¹. At present, antiretroviral therapy is the most common way to manage HIV infection, as it reduces viral loads and prolongs and improves the quality of life of HIV-infected patients. However, the currently available antiretroviral drugs also have major shortcomings, such as high costs, marked side effects, developing drug resistance, a necessity for regimen changes, and the life-long duration of the therapy (Arts, Hazuda, 2012). Above all that, we are yet to find the cure for HIV infection (Phanuphak, Gulick, 2020). As a result, the development of effective preventive vaccines against HIV/AIDS remains a top priority (Stephenson et al., 2020).

As of today, the RV144 clinical trials performed in Thailand from 2003 to 2009 are considered the most successful. The studied vaccine showed an efficacy of 60 % in 12 months after vaccination and 31.2 % – after a 3.5-year follow-up (Kim et al., 2015). Several years later the RV144 vaccine components were modified to express the antigens of the HIV strains circulating in South Africa. In January 2020, early results of clinical trials showed that the modified vaccine failed to prevent HIV-1 infection in volunteers (Gray et al., 2021). Nowadays, there are still numerous unresolved issues in HIV-1 vaccine development, yet it is clear that it is necessary to use new approaches to its design (Hsu, O'Connell, 2017), hence the intense research for the induction of the protective T and B cell immune response to HIV-1, including broadly neutralizing antibodies (bnAbs) (Shcherbakov et al., 2015; Rudometov et al., 2019b; Jones et al., 2020; Liu et al., 2020; Ng'uni et al., 2020).

Selecting an adequate tool for *in vitro* efficacy assessment is an integral part of scientific research aimed at developing vaccines and chemotherapy drugs against viral pathogens, including HIV-1. The neutralizing activity of the sera from the animals immunized with experimental vaccines and the efficacy of chemotherapy agents are conventionally assessed using viral isolates (Jackson et al., 1988). However, this process is demanding in terms of time, effort, and biosafety measures.

An alternative method is to use infectious molecular clones, i.e. infectious viral particles obtained via cell transfection with plasmid vector including the full-length HIV-1 genome coding structural, regulatory, and accessory proteins of the virus required for the cultivation of replication-competent viral particles in cell culture (Peden et al., 1991).

In recent years, many researchers give preference to the pseudotyped virus approach, a safer method suitable for BSL-2 lab settings (Li Q. et al., 2018; Montefiori et al., 2018). Compared to viral isolates and infectious molecular clones, pseudotyped viruses are harmless, because virus replication is restricted to a single cycle due to mutations in coding regions of the genome, which is why pseudotyped viruses are often called single-cycle viruses (Cheresiz et al., 2010; Li Q. et al., 2018).

HIV-1 model systems for *in vitro* efficacy assessment of chemotherapy drugs, bnAbs, and candidate vaccines against HIV-1 will be considered in the present review.

HIV-1 isolates and laboratory-adapted strains

Historically, HIV-1 primary isolates were the first system for analyzing vaccine efficacy and neutralizing the activity of antibodies (Jackson et al., 1988). Viral isolates are obtained via co-cultivation of the peripheral blood mononuclear cells (PBMC) of an HIV-positive patient and the PHA-stimulated PBMC of a healthy donor. Here, the viruses isolated from blood appear as a genetically heterogeneous population due to the quasispecies nature of HIV-1. To eliminate possible selective pressure on viral isolates and ensure optimal preservation of a viral phenotype, the virus is cultivated using primary cell culture, rather than cell lines (Voronin et al., 2007; Van't Wout et al., 2008). The presence of neutralizing antibodies in sera from vaccinated subjects or the efficacy of an antiviral agent is typically identified in a PBMC culture with an added infectious dose of the virus and serial dilutions of immune serum or tested compound. HIV-1 replication suppression is assessed using ELISA by measuring p24 content (structural component of HIV-1 capsid) in the culture medium (Zyryanova et al., 2020a).

However, the use of HIV-1 primary isolates for virus neutralization analysis has several shortcomings, including the use of primary PBMCs for pathogen replication, high biosafety requirements, low repeatability of the results, and therefore

¹ Fact Sheet on HIV/AIDS. World Health Organization, 2020. URL: <https://www.who.int/ru/news-room/fact-sheets/detail/hiv-aids> (Accessed June 2, 2021).

standardization issues (Mascola et al., 1996, 2005). Thus, some HIV-1 strains (IIIB/LAV, MN, SF2) were adapted for replication in immortalized cell lines (H9, CEM) for the sake of simplicity and to ensure repeatability of the experiments in the first years of vaccine development. These were later referred to as laboratory-adapted strains or, more accurately, T cell line adapted strains. Vaccination of volunteers with recombinant trimers based on laboratory-adapted HIV-1 strains induced the antibodies neutralizing these specific laboratory strains. The additional experiments involving HIV-1 primary isolates showed the absence of neutralizing activity against primary isolates, despite intense induction of neutralizing antibodies against the laboratory-adapted strains (Mascola et al., 1996; Montefiori et al., 2018). Apparently, the neutralization analysis performed using laboratory-adapted strains could produce misleading results, and the researchers came back to primary isolates as a more adequate tool for analyzing the virus-neutralizing activity of the antibodies induced as a result of vaccination. Since the method is labor-intensive and does not allow for mass analysis, it began to be used for the concluding stages of research.

HIV-1 infectious molecular clones

Taking into account the cultivation difficulties and significant heterogeneity of HIV-1 primary isolates and laboratory-adapted strains, as well as the variability of donor PBMCs (Polonis et al., 2008), HIV-1 infectious molecular clones (IMCs) were chosen for consistent replication of viral particles. IMCs are obtained via cell transfection with a plasmid vector including a full-length HIV-1 genome to ensure the generation of replication-competent viral particles in a eukaryotic cell culture (Fig. 1). Compared to HIV-1 primary isolates, this approach makes it possible to obtain genetically homogeneous viral particles, since an HIV-1 genome is present in the plasmid vector in the form of DNA (Edmonds et al., 2010; Zyryanova et al., 2020b). To ensure standardization of neutralization analysis using IMCs, modified continuous cell lines with a cell-surface CD4 receptor and CCR5 and CXCR4 co-receptors were genetically engineered (Princen et al., 2004; González et al., 2009). Since IMCs are essentially infectious viral particles, the relevant biosafety requirements are to be fulfilled, similarly to primary isolates and laboratory-adapted strains, and the analysis itself is rather time-consuming.

At the same time, the use of IMCs makes it possible to characterize and study biological properties of genetically different HIV-1 isolates (Ochsenbauer et al., 2012; Baalwa et al., 2013; Wang et al., 2013; Chenine et al., 2018; Zyryanova et al., 2020b), investigate the development mechanisms of drug-resistant HIV-1 strains and the effect of mutations on the biological properties of the virus (Johnston et al., 2005; Pugach et al., 2007; Varghese et al., 2013), and discover new antiretroviral agents (Su et al., 2019; Wagstaff et al., 2019; Mavian et al., 2020).

HIV-1 *env*-pseudoviruses

The use of classical virological methods to work with HIV-1 faces a number of difficulties noted above. *Env*-pseudovirus technology has proved to be a potent tool for quick and adequate assessment of humoral immune response to vaccine

constructs and screening of potential chemotherapeutic agents, specifically entry inhibitors (Montefiori et al., 2018).

HIV-1 *env*-pseudoviruses are recombinant viral particles obtained via eukaryotic cell transfection with the two plasmids referred to as core and envelope. The core plasmid includes genes of structural (Gag and Pol), regulatory (Tat and Rev), and accessory (Vpu, Vpr, Vif, and Nef) HIV-1 proteins necessary for viral particle assembly, as well as sequences required for viral RNA packaging (Ψ). The envelope plasmid carries an envelope glycoprotein gene (*Env*) of certain HIV-1 subtype. As a result of transfection, viral particles with a defective genome incapable of assembling infectious daughter virions are obtained (Li M. et al., 2005; Li Q. et al., 2018). Electron microscopy studies show that the HEK293 cell line transfection with two plasmids produces viral particles morphologically identical to the HIV-1 virions (Zaitsev et al., 2019; Ladinsky et al., 2020).

The determination of the functional activity of *env*-pseudoviruses and analyses neutralization are carried out on a TZM-bl cell line, which is a continuous, genetically modified HeLa cell line with cell-surface CD4 receptors and CCR5 and CXCR4 co-receptors. In addition, firefly luciferase and β -galactosidase *E. coli* reporter genes are integrated into the TZM-bl cell line genome under transcriptional control of HIV-1 long terminal repeat. When a pseudotyped virus enters the target TZM-bl cell, synthesis of a viral Tat protein triggers luciferase reporter gene expression detectable by a luminometer. Here, high luminescence intensity indicates that pseudotyped viral particles have entered target cells, whereas suppressed luminescence indicates that the HIV-1 *env*-pseudoviruses have been neutralized (Platt et al., 1998; Wei et al., 2002). A general work technique of *env*-pseudovirus system is shown in Fig. 2.

An *env*-pseudovirus system has a number of distinct advantages. First, since TZM-bl is a stable continuous cell line, it may be used as a substitute for human primary T cells, thereby reducing the need for individual donor cells. Second, *env*-pseudoviruses are harmless compared to viral isolates and IMCs requiring higher biosafety levels, which makes experimental studies more complicated and expensive. Third, Env protein forms trimer structures at the surface of pseudotyped viral particles, which are identical to those of the natural virus. However, the main advantage of the pseudotyped virus technology is that it makes it possible to obtain the equivalents of the viral particles of various HIV-1 subtypes and strains, thereby providing broad coverage of HIV-1 genetic diversity (Seaman et al., 2010; Montefiori et al., 2018). In addition, the neutralization assessment method using *env*-pseudoviruses favors further optimization and standardization (Wei et al., 2002; Seaman et al., 2010; Sarzotti-Kelsoe et al., 2014). A brief comparison of HIV-1 primary isolates and laboratory-adapted strains, IMCs, and *env*-pseudoviruses is presented in the Table.

It should be noted that the protocols and recommendations for neutralization assessment using *env*-pseudoviruses are available at the website of the Los Alamos National Laboratory (<https://www.hiv.lanl.gov/content/nab-reference-strains/html/home.htm>). In addition, the HIV Reagent Program supported by the National Institute of Allergy and Infectious Diseases and curated by the National Collection of Type Cultures

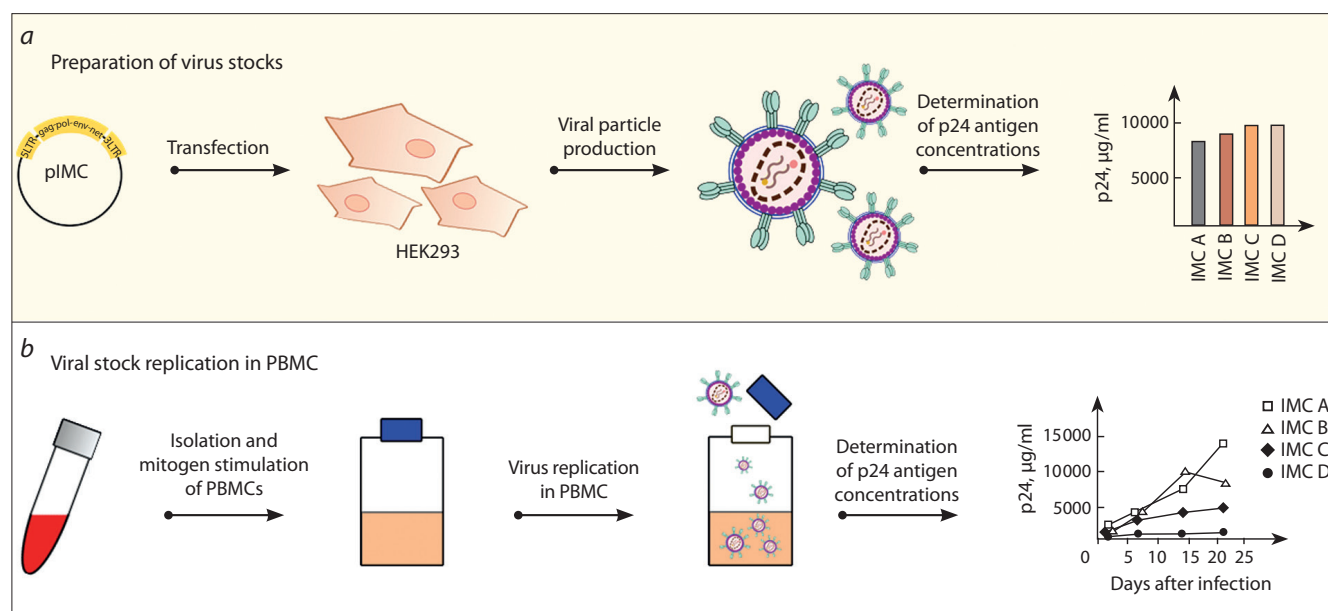


Fig. 1. HIV-1 IMC technology.

Conditionally, HIV-1 IMCs are obtained in two stages. At the first stage (a), viral particles, also referred to as virus stock, are produced via HEK293 cell line transfection. At the second stage (b), the virus stock is further replicated for several weeks using PHA-stimulated PBMCs from a healthy donor. The titers of viral particles are measured at each stage using the ELISA based on p24 antigen content in the culture medium. A replication-competent virus is produced if p24 capsid protein content becomes at least 1000 times as high as its initial content in the culture medium.

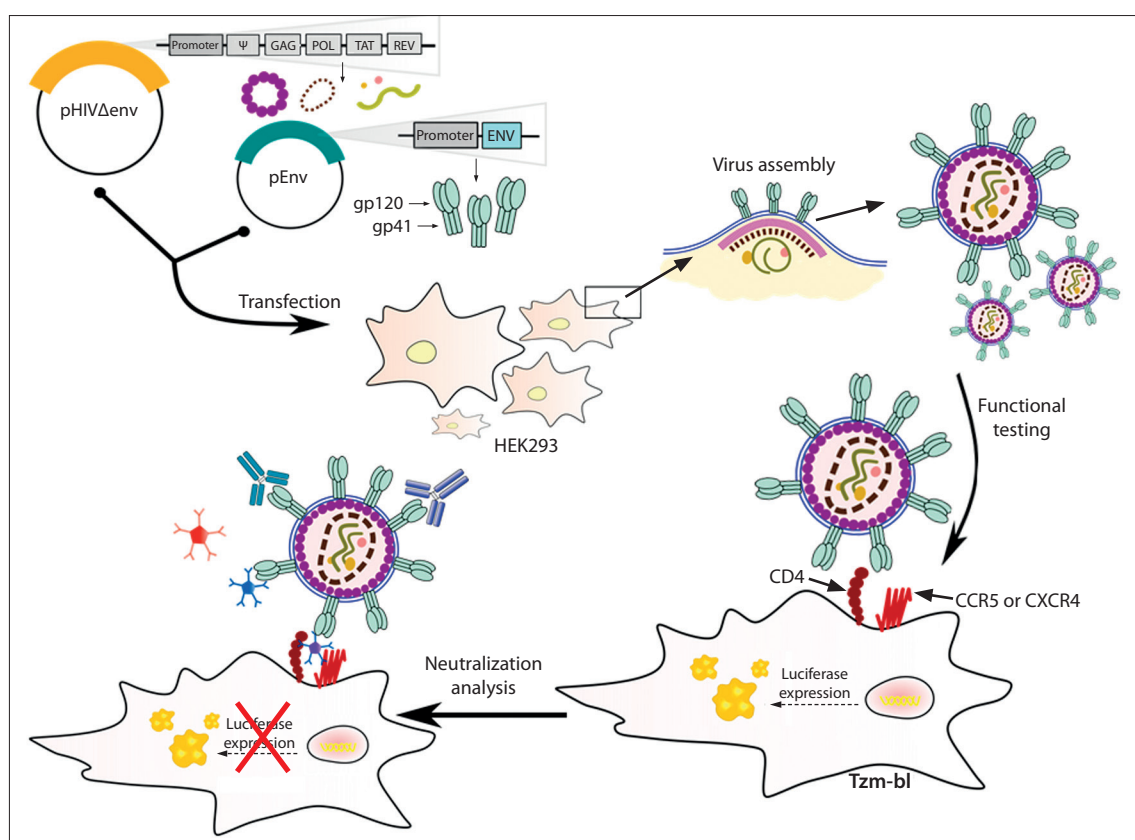


Fig. 2. HIV-1 env-pseudovirus technology.

Experimental study of env-pseudoviruses is done in several stages: Stage 1 includes viral particle assembly via HEK293 cell line transfection with two plasmids, referred to as core and envelope; at Stage 2, one measures the functional activity of pseudotyped viral particles, i.e. their ability to infect target cells and trigger firefly luciferase reporter gene expression; at Stage 3 the neutralization level is analyzed using immune sera or chemotherapeutic agents to measure their ability to block pseudotyped virus entry to target cells.

Comparison of HIV-1 model systems used for *in vitro* efficacy assessment of vaccines and drugs against HIV-1

Parameter	<i>Env</i> -pseudoviruses	Infectious molecular clones	Primary isolates and laboratory-adapted strains
Biosafety requirements (hazard level)	Low	High	High
Analysis speed	High	Low	Low
Standardization	High	High	Low
Investigation of virus properties and virus cycle	Only at the entry stage	All stages of HIV-1 life cycle	All stages of HIV-1 life cycle
Cultivation conditions	Continuous cell lines	Continuous cell lines; PBMCs	PBMCs

makes it possible to obtain all the components (cell lines, plasmids, monoclonal antibodies) required for implementing the technology.

Here are several noteworthy applications of *env*-pseudoviruses panels. Antiviral activity of clinically approved co-receptor antagonist Maraviroc was demonstrated using 160 HIV-1 subtype B *env*-pseudoviruses and 40 *env*-pseudoviruses of other HIV-1 subtypes (Dorr et al., 2005). The activity of Ibalizumab, a monoclonal antibody binding to the CD4 receptor, was demonstrated using 116 *env*-pseudoviruses of subtypes A, B, C, and CRF01_AE (Pace et al., 2013). HIV-1 *env*-pseudoviruses panels were also used to investigate the bnAbs spectrum with respect to various genetic variants of HIV-1. For instance, the neutralization breadth of 98 % for bnAb 10E8 was demonstrated using a panel of 181 *env*-pseudoviruses of subtypes A, B, C, D, G, CRF01_AE, and CRF02_AG (Huang et al., 2012); neutralization breadth of 91 % for bnAb VRC01 was demonstrated using 196 *env*-pseudoviruses (Wu X. et al., 2010); neutralization breadth of 49 % for bnAb VRC34.01 was demonstrated using 179 *env*-pseudoviruses (Kong et al., 2016). It is the introduction of pseudotyped virus panels, including a wide range of genetically diverse HIV-1 variants, that led to a breakthrough in the production and characterization of monoclonal broadly neutralizing antibodies.

Env-pseudoviruses panels are extensively used to study the humoral immune response induced by candidate vaccines against HIV-1 at a design stage and during pre-clinical and clinical trials, since the presence of virus-neutralizing antibodies in the vaccinated subjects is among the key indicators of HIV vaccine effectiveness (Rudometov et al., 2019a; Ou et al., 2020). Recent papers by Xu et al., who developed a vaccination regimen based on fusion peptide (FP) of gp41, a key structural component of HIV-1, may be cited as an example. Earlier, they identified the VRC34.01 antibody from an HIV-positive donor, which was aimed at the conservative N-terminal region of HIV-1 FP. Since FP is a short linear peptide, it has low natural immunogenicity, which is why garden snail hemocyanin widely used in biotechnology was used as a carrier protein. Immunization of laboratory animals by an FP bound to garden snail hemocyanin with subsequent boosting by a BG505 trimer resulted in induction of antibodies with neutralization breadth of 31 % demonstrated using a panel of 208 *env*-pseudoviruses of various HIV-1 subtypes (Xu et al., 2018).

In conclusion of this review, it should be mentioned that an HIV-1 pseudotyping system is tolerant to incorporation

of surface proteins of various enveloped viruses. Since most laboratory experiments and studies involving viruses are to be performed in BSL-3 or BSL-4 lab settings, the use of pseudotyped viruses instead of wild-type ones makes it possible for various research groups to study viruses of interest and design antiviral drugs and vaccines against highly dangerous viruses. For example, HIV-1 pseudotyping system was used to obtain the viral particles carrying surface glycoproteins of Ebola virus (Mohan et al., 2015), Marburg virus (Zhang L. et al., 2019), Lassa fever (Zhang X. et al., 2019), Middle East respiratory syndrome coronavirus (Zhao et al., 2013), Rabies virus (Nie et al., 2017), Chikungunya virus (Wu J. et al., 2017), and Nipah virus (Nie et al., 2019). In addition, this technology is extensively used in designing pseudotyped virus platforms for SARS-CoV-2 (Hu et al., 2020; Hyseni et al., 2020; Johnson et al., 2020).

Conclusion

All technologies considered above have their own advantages and shortcomings and most certainly complement each other in integrated studies. Despite the labor-intensity of primary isolate and IMC technologies in neutralization assessments, these models still remain valuable tools for investigating the biological properties of viruses. However, *env*-pseudovirus technology has currently become the base method for efficacy assessment of HIV-1 vaccines and antiviral agents (potential entry inhibitors). Its main advantages include safety, high repeatability of the results, standardization potential, and ability to work with virus particles exposing surface glycoproteins of multiple virus subtypes.

References

Arts E.J., Hazuda D.J. HIV-1 antiretroviral drug therapy. *Cold Spring Harb. Perspect. Med.* 2012;2(4):1-23. DOI 10.1101/cshperspect.a007161.

Baalwa J., Wang S., Parrish N.F., Decker J.M., Keele B.F., Learn G.H., Yue L., Ruzagira E., Ssemwanga D., Kamali A. Molecular identification, cloning and characterization of transmitted/founder HIV-1 subtype A, D and A/D infectious molecular clones. *Virology*. 2013; 436(1):33-48. DOI 10.1016/j.virol.2012.10.009.

Chenine A.L., Merbah M., Wiczorek L., Molnar S., Mann B., Lee J., O’Sullivan A.M., Bose M., Sanders-Buell E., Kijak G.H., Herrera C., McLinden R., O’Connell R., Michael N.L., Robb M.L., Kim J.H., Polonis V.R., Tovanabutra S. Neutralization sensitivity of a novel HIV-1 CRF01_AE panel of infectious molecular clones. *J. Acquir. Immune Defic. Syndrom.* 2018;78(3):348-355. DOI 10.1097/QAI.0000000000001675.

- Cheresiz S.V., Grigoryev I.V., Semenova E.A., Pustyniyak V.O., Vlasov V.V., Pokrovsky A.G. A pseudovirus system for the testing of antiviral activity of compounds in different cell lines. *Doklady Biochemistry and Biophysics*. 2010;435:295-298. DOI 10.1134/S1607672910060049.
- Dorr P., Westby M., Dobbs S., Griffin P., Irvine B., Macartney M., Mori J., Rickett G., Smith-Burchnell C., Napier C., Webster R., Armour D., Price D., Stammen B., Wood A., Perros M. Maraviroc (UK-427,857), a potent, orally bioavailable, and selective small-molecule inhibitor of chemokine receptor CCR5 with broad-spectrum anti-human immunodeficiency virus type 1 activity. *Antimicrob. Agents Chemother.* 2005;49(11):4721-4732. DOI 10.1128/AAC.49.11.4721-4732.2005.
- Edmonds T.G., Ding H., Yuan X., Wei Q., Smith K.S., Conway J.A., Wiczorek L., Brown B., Polonis V., West J.T., Montefiori D.C., Kappes J.C., Ochsenbauer Ch. Replication competent molecular clones of HIV-1 expressing *Renilla* luciferase facilitate the analysis of antibody inhibition in PBMC. *Virology*. 2010;408(1):1-13. DOI 10.1016/j.virol.2010.08.028.
- González N., Pérez-Olmeda M., Mateos E., Cascajero A., Alvarez A., Spijkers S., García-Pérez J., Sánchez-Palomino S., Ruiz-Mateos E., Leal M., Alami J. A sensitive phenotypic assay for the determination of human immunodeficiency virus type 1 tropism. *J. Antimicrob. Chemother.* 2010;65(12):2493-2501. DOI 10.1093/jac/dkq379.
- Gray G.E., Bekker L.G., Laher F., Malahleha M., Allen M., Moodie Z., Grunenberg N., Huang Yu., Grove D., Prigmore B. Vaccine efficacy of ALVAC-HIV and bivalent subtype C gp120-MF59 in adults. *New Eng. J. Med.* 2021;384(12):1089-1100. DOI 10.1056/NEJMoa2031499.
- Hsu D.C., O'Connell R.J. Progress in HIV vaccine development. *Hum. Vaccines Immunother.* 2017;13(5):1018-1030. DOI 10.1080/21645515.2016.1276138.
- Hu J., Gao Q., He C., Huang A., Tang N., Wang K. Development of cell-based pseudovirus entry assay to identify potential viral entry inhibitors and neutralizing antibodies against SARS-CoV-2. *Genes Dis.* 2020;7(4):551-557. DOI 10.1016/j.gendis.2020.07.006.
- Huang J., Ofek G., Laub L., Louder M.K., Doria-Rose N.A., Longo N.S., Imamichi H., Bailer R.T., Chakrabarti B., Sharma S.K., Munir Alam S., Wang T., Yang Y., Zhang B., Migueles S.A., Wyatt R., Haynes B.F., Kwong P.D., Mascola J.R., Connors M. Broad and potent neutralization of HIV-1 by a gp41-specific human antibody. *Nature*. 2012;491(7424):406-412. DOI 10.1038/nature11544.
- Hyseni I., Molesti E., Benincasa L., Piu P., Casa E., Temperton N.J., Manenti A., Montomoli E. Characterisation of SARS-CoV-2 lentiviral pseudotypes and correlation between pseudotype-based neutralisation assays and live virus-based micro neutralisation assays. *Viruses*. 2020;12(9):1-18. DOI 10.3390/v12091011.
- Jackson J.B., Coombs R.W., Sannerud K., Rhame F.S., Balfour H.H., Jr. Rapid and sensitive viral culture method for human immunodeficiency virus type 1. *J. Clin. Microbiol.* 1988;26(7):1416-1418. PMID 3165981.
- Johnson M.C., Lyddon T.D., Suarez R., Salcedo B., LePique M., Graham M., Ricana C.L., Robinson C.A., Ritter D.G. Optimized pseudotyping conditions for the SARS-COV-2 spike glycoprotein. *J. Virol.* 2020;94(21):1-10. DOI 10.1128/JVI.01062-20.
- Johnston E., Dupnik K.M., Gonzales M.J., Winters M.A., Rhee S.Y., Imamichi T., Shafer R.W. Panel of prototypical infectious molecular HIV-1 clones containing multiple nucleoside reverse transcriptase inhibitor resistance mutations. *AIDS*. 2005;19(7):731-733. DOI 10.1097/01.aids.0000166098.54564.0c.
- Jones L.D., Moody M.A., Thompson A.B. Innovations in HIV-1 vaccine design. *Clin. Ther.* 2020;42(3):499-514. DOI 10.1016/j.clinthera.2020.01.009.
- Kim J.H., Excler J.L., Michael N.L. Lessons from the RV144 Thai phase III HIV-1 vaccine trial and the search for correlates of protection. *Annu. Rev. Med.* 2015;66:423-437. DOI 10.1146/annurev-med-052912-123749.
- Kong R., Xu K., Zhou T., Acharya P., Lemmin T., Liu K., Ozorowski G., Soto C., Taft J., Bailer R., Cale E.M., Chen L., Choi C.W., Chuang G., Doria-Rose N.A., Druz A., Georgiev I.S., Gorman J., Huang J., Gordon Joyce M., Louder M.K., Ma X., McKee K., O'Dell S., Pancera M., Yang Y., Blanchard S.C., Mothes W., Burton D.R., Koff W.C., Connors M., Ward A.B., Kwong P.D., Mascola J.R. Fusion peptide of HIV-1 as a site of vulnerability to neutralizing antibody. *Science*. 2016;352(6287):828-833. DOI 10.1126/science.aac0474.
- Ladinsky M.S., Gnanapragasam P.N., Yang Z., West A.P., Kay M.S., Bjorkman P.J. Electron tomography visualization of HIV-1 fusion with target cells using fusion inhibitors to trap the pre-hairpin intermediate. *eLife*. 2020;9. DOI 10.7554/eLife.58411.
- Li M., Gao F., Mascola J.R., Stamatatos L., Polonis V.R., Koutsoukos M., Voss G., Goepfert P., Gilbert P., Greene K.M., Bilska M., Kothe D.L., Salazar-Gonzalez J.F., Wei X., Decker J.M., Hahn B.H., Montefiori D.C. Human immunodeficiency virus type 1 *env* clones from acute and early subtype B infections for standardized assessments of vaccine-elicited neutralizing antibodies. *J. Virol.* 2005;79(16):10108-10125. DOI 10.1128/JVI.79.16.10108-10125.2005.
- Li Q., Liu Q., Huang W., Li X., Wang Y. Current status on the development of pseudoviruses for enveloped viruses. *Rev. Med. Virol.* 2018;28(1):1-10. DOI 10.1002/rmv.1963.
- Liu Y., Cao W., Sun M., Li T. Broadly neutralizing antibodies for HIV-1: efficacies, challenges and opportunities. *Emerg. Microbes Infect.* 2020;9(1):194-206. DOI 10.1080/22221751.2020.1713707.
- Mascola J.R., Snyder S.W., Weislow O.S., Belay S.M., Belshe R.B., Schwartz D.H., Clements M.L., Dolin R., Graham B.S., Gorse G.J. Immunization with envelope subunit vaccine products elicits neutralizing antibodies against laboratory-adapted but not primary isolates of human immunodeficiency virus type 1. *J. Infect. Dis.* 1996; 173(2):340-348. DOI 10.1093/infdis/173.2.340.
- Mascola J.R., D'Souza P., Gilbert P., Hahn B.H., Haigwood N.L., Morris L., Petropoulos C.J., Polonis V.R., Sarzotti M., Montefiori D.C. Recommendations for the design and use of standard virus panels to assess neutralizing antibody responses elicited by candidate human immunodeficiency virus type 1 vaccines. *J. Virol.* 2005;79(16): 10103-10107. DOI 10.1128/JVI.79.16.10103-10107.2005.
- Mavian C., Coman R.M., Zhang X., Pomeroy S., Ostrov D.A., Dunn B.M., Sleasman J.W., Goodenow M.M. Molecular docking-based screening for novel inhibitors of the human immunodeficiency virus type 1 protease that effectively reduce the viral replication in human cells. *bioRxiv*. 2020;1-11. DOI 10.1101/2020.11.14.382895.
- Mohan G.S., Ye L., Li W., Monteiro A., Lin X., Sapkota B., Polack B.P., Compans R.W., Yang C. Less is more: Ebola virus surface glycoprotein expression levels regulate virus production and infectivity. *J. Virol.* 2015;89(2):1205-1217. DOI 10.1128/JVI.01810-14.
- Montefiori D.C., Roederer M., Morris L., Seaman M.S. Neutralization tiers of HIV-1. *Curr. Opin. HIV AIDS*. 2018;13(2):1-9. DOI 10.1097/COH.0000000000000442.
- Ng'uni T., Chasara C., Ndhlovu Z.M. Major scientific hurdles in HIV vaccine development: historical perspective and future directions. *Front. Immun.* 2020;11:1-17. DOI 10.3389/fimmu.2020.590780.
- Nie J., Liu L., Wang Q., Chen R., Ning T., Liu Q., Huang W., Wang Y. Nipah pseudovirus system enables evaluation of vaccines *in vitro* and *in vivo* using non-BSL-4 facilities. *Emerg. Microbes Infect.* 2019;8(1):272-281. DOI 10.1080/22221751.2019.1571871.
- Nie J., Wu X., Ma J., Cao S., Huang W., Liu Q., Li X., Li Y., Wang Y. Development of *in vitro* and *in vivo* rabies virus neutralization as-

- says based on a high-titer pseudovirus system. *Sci. Rep.* 2017;7(1): 1-12. DOI 10.1038/srep42769.
- Ochsenbauer C., Edmonds T.G., Ding H., Keele B.F., Decker J., Salazar M.G., Salazar-Gonzalez J.F., Shattock R., Haynes B.F., Shaw G.M., Hahn B.H., Kappes J.C. Generation of transmitted/founder HIV-1 infectious molecular clones and characterization of their replication capacity in CD4 T lymphocytes and monocyte-derived macrophages. *J. Virol.* 2012;86(5):2715-2728. DOI 10.1128/JVI.06157-11.
- Ou L., Kong W.P., Chuang G.Y., Ghosh M., Gulla K., O'Dell S., Variale J., Barefoot N., Changela A., Chao C.W., Cheng Ch., Druz A., Kong R., McKee K., Rawi R., Sarfo E., Schön A., Shaddeau A., Tsybovsky Ya., Verardi R., Wang Sh. Preclinical development of a fusion peptide conjugate as an HIV vaccine immunogen. *Sci. Rep.* 2020;10(1):1-13. DOI 10.1038/s41598-020-59711-y.
- Pace C.S., Fordyce M.W., Franco D., Kao C.Y., Seaman M.S., Ho D.D. Anti-CD4 monoclonal antibody ibalizumab exhibits breadth and potency against HIV-1, with natural resistance mediated by the loss of a V5 glycan in envelope. *J. Acquir. Immune Defic. Syndr.* 2013;62(1):1-9. DOI 10.1097/QAI.0b013e3182732746.
- Peden K., Emerman M., Montagnier L. Changes in growth properties on passage in tissue culture of viruses derived from infectious molecular clones of HIV-1LA1, HIV-1MAL, and HIV-1ELI. *Virology.* 1991;185:661-672. DOI 10.1016/0042-6822(91)90537-L.
- Phanuphak N., Gulick R.M. HIV treatment and prevention 2019: current standards of care. *Curr. Opin. HIV AIDS.* 2020;15(1):4-12. DOI 10.1097/COH.0000000000000588.
- Platt E.J., Wehrly K., Kuhmann S.E., Chesebro B., Kabat D. Effects of CCR5 and CD4 cell surface concentrations on infections by macrophagetropic isolates of human immunodeficiency virus type 1. *J. Virol.* 1998;72:2855-2864. DOI 10.1128/JVI.72.4.2855-2864.1998.
- Polonis V.R., Brown B.K., Borges A.R., Zolla-Pazner S., Dimitrov D.S., Zhang M.Y., Barnett S.W., Ruprecht R.M., Scarlatti G., Fenyo E., Montefiori D.C., McCutchan F.E., Michael N.L. Recent advances in the characterization of HIV-1 neutralization assays for standardized evaluation of the antibody response to infection and vaccination. *Virology.* 2008;375(2):315-320. DOI 10.1016/j.virol.2008.02.007.
- Princen K., Hatse S., Vermeire K., De Clercq E., Schols D. Establishment of a novel CCR5 and CXCR4 expressing CD4+ cell line which is highly sensitive to HIV and suitable for high-throughput evaluation of CCR5 and CXCR4 antagonists. *Retrovirology.* 2004;1(1): 1-13. DOI 10.1186/1742-4690-1-2.
- Pugach P., Marozsan A.J., Ketas T.J., Landes E.L., Moore J.P., Kuhmann S.E. HIV-1 clones resistant to a small molecule CCR5 inhibitor use the inhibitor-bound form of CCR5 for entry. *Virology.* 2007;361(1):212-228. DOI 10.1016/j.virol.2006.11.004.
- Rudometov A.P., Chikae A.N., Rudometova N.B., Antonets D.V., Lomzov A.A., Kaplina O.N., Ilyichev A.A., Karpenko L.I. Artificial anti-HIV-1 immunogen comprising epitopes of broadly neutralizing antibodies 2F5, 10E8, and a peptide mimic of VRC01 discontinuous epitope. *Vaccines.* 2019a;7(3):1-18. DOI 10.3390/vaccines 7030083.
- Rudometov A.P., Rudometova N.B., Shcherbakov D.N., Lomzov A.A., Kaplina O.N., Shcherbakova N.S., Ilyichev A.A., Bakulina A.Yu., Karpenko L.I. The structural and immunological properties of chimeric proteins containing HIV-1 MPER sites. *Acta Naturae.* 2019b; 11(3):56-65. DOI 10.32607/20758251-2019-11-3-56-65.
- Sarzotti-Kelsoe M., Bailer R.T., Turk E., Lin C.L., Bilks M., Greene K.M., Gao H., Todd C.A., Ozaki D., Seaman M.S., Mascola J.R., Montefiori D.C. Optimization and validation of the TZM-bl assay for standardized assessments of neutralizing antibodies against HIV-1. *J. Immunol. Methods.* 2014;409:131-146. DOI 10.1016/j.jim.2013.11.022.
- Seaman M.S., Janes H., Hawkins N., Grandpre L.E., Devoy C., Giri A., Coffey R.T., Harris L., Wood B., Daniels M.G. Tiered categorization of a diverse panel of HIV-1 Env pseudoviruses for assessment of neutralizing antibodies. *J. Virol.* 2010;84(3):1439-1452. DOI 10.1128/JVI.02108-09.
- Shcherbakov D.N., Bakulina A.Y., Karpenko L.I., Ilyichev A.A. Broadly neutralizing antibodies against HIV-1 as a novel aspect of the immune response. *Acta Naturae.* 2015;7(4):11-21. PMID 26798488.
- Stephenson K.E., Wagh K., Korber B., Barouch D.H. Vaccines and broadly neutralizing antibodies for HIV-1 prevention. *Annu. Rev. Immunol.* 2020;38:673-703. DOI 10.1146/annurev-immunol-080219-023629.
- Su S., Rasquinha G., Du L., Wang Q., Xu W., Li W., Lu L., Jiang S. A peptide-based HIV-1 fusion inhibitor with two tail-anchors and palmitic acid exhibits substantially improved *in vitro* and *ex vivo* anti-HIV-1 activity and prolonged *in vivo* half-life. *Molecules.* 2019;24(6):1-13. DOI 10.3390/molecules24061134.
- Van't Wout A.B., Schuitemaker H., Kootstra N.A. Isolation and propagation of HIV-1 on peripheral blood mononuclear cells. *Nat. Protoc.* 2008;3(3):363-370. DOI 10.1038/nprot.2008.3.
- Varghese V., Mitsuya Y., Fessel W.J., Liu T.F., Melikian G.L., Katzenstein D.A., Schiffer C.A., Holmes S.P., Shafer R.W. Prototypal recombinant multi-protease-inhibitor-resistant infectious molecular clones of human immunodeficiency virus type 1. *Antimicrob. Agents Chemother.* 2013;57(9):4290-4299. DOI 10.1128/AAC.00614-13.
- Voronin Y., Chohan B., Emerman M., Overbaugh J. Primary isolates of human immunodeficiency virus type 1 are usually dominated by the major variants found in blood. *J. Virol.* 2007;81(19):10232-10241. DOI 10.1128/JVI.01035-07.
- Wagstaff K.M., Headey S., Telwate S., Tyssen D., Hearps A.C., Thomas D.R., Tachedjian G., Jans D.A. Molecular dissection of an inhibitor targeting the HIV integrase dependent preintegration complex nuclear import. *Cell. Microbiol.* 2019;21(1):1-13. DOI 10.1111/cmi.12953.
- Wang Z., Hong K., Zhang J., Zhang L., Li D., Ren L., Liang H., Shao Y. Construction and characterization of highly infectious full-length molecular clones of a HIV-1 CRF07_BC isolate from Xinjiang, China. *PLoS One.* 2013;8(11):1-9. DOI 10.1371/journal.pone.0079177.
- Wei X., Decker J.M., Liu H., Zhang Z., Arani R.B., Kilby J.M., Saag M.S., Wu X., Shaw G.M., Kappes J.C. Emergence of resistant human immunodeficiency virus type 1 in patients receiving fusion inhibitor (T-20) monotherapy. *Antimicrob. Agents Chemother.* 2002; 46:1896-1905. DOI 10.1128/aac.46.6.1896-1905.2002.
- Wu J., Zhao C., Liu Q., Huang W., Wang Y. Development and application of a bioluminescent imaging mouse model for Chikungunya virus based on pseudovirus system. *Vaccine.* 2017;35(47):6387-6394. DOI 10.1016/j.vaccine.2017.10.007.
- Wu X., Yang Z.Y., Li Y., Hogerkerp C.M., Schief W.R., Seaman M.S., Zhou T., Schmidt S.D., Wu L., Xu L. Rational design of envelope identifies broadly neutralizing human monoclonal antibodies to HIV-1. *Science.* 2010;329(5993):856-861. DOI 10.1126/science.1187659.
- Xu K., Acharya P., Kong R., Cheng C., Chuang G.Y., Liu K., Louder M.K., O'Dell S., Rawi R., Sastry M. Epitope-based vaccine design yields fusion peptide-directed antibodies that neutralize diverse strains of HIV-1. *Nat. Med.* 2018;24(6):857-867. DOI 10.1038/s41591-018-0042-6.
- Zaitsev B.N., Taranov O.S., Rudometova N.B., Shcherbakova N.S., Ilyichev A.A., Karpenko L.I. An optimized method for counting viral particles using electron microscopy. *Vavilovskii Zhurnal Genetiki i Selektii = Vavilov Journal of Genetics and Breeding.* 2019;23: 337-342. DOI 10.18699/VJ19.498.

- Zhang L., Lei S., Xie H., Li Q., Liu S., Liu Q., Huang W., Xiao X., Wang Y. Screening and identification of Marburg virus entry inhibitors using approved drugs. *Virol. Sin.* 2020;35:235-239. DOI 10.1007/s12250-019-00184-3.
- Zhang X., Yan F., Tang K., Chen Q., Guo J., Zhu W., He S., Banadyga L., Qiu X., Guo Y. Identification of a clinical compound losmapimod that blocks Lassa virus entry. *Antiviral Res.* 2019;167:68-77. DOI 10.1016/j.antiviral.2019.03.014.
- Zhao G., Du L., Ma C., Li Y., Li L., Poon V.K., Wang L., Yu F., Zheng B.J., Jiang S., Zhou Y. A safe and convenient pseudovirus-based inhibition assay to detect neutralizing antibodies and screen for viral entry inhibitors against the novel human coronavirus MERS-CoV. *Virol. J.* 2013;10(1):1-8. DOI 10.1186/1743-422X-10-266.
- Zyryanova D.P., Bogacheva N.V., Totmenin A.V., Gashnikova N.M. HIV-1 CRF63_02A6 models as a tool for evaluating efficacy of developing antiretroviral drugs. *Infektsiya i Immunitet = Russian Journal of Infection and Immunity.* 2020;10(4):769-774. DOI 10.15789/2220-7619-MHC-1261. (in Russian)
- Zyryanova D.P., Totmenin A.V., Bogacheva N.V., Gashnikova N.M. Construction and characterization of infectious molecular clones of HIV-1 CRF63_02A6. *AIDS Res. Hum. Retrovir.* 2020;36(3):227-233. DOI 10.1089/aid.2019.0177.

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