

Review

Discovery and Development of Safe-in-man Broad-Spectrum Antiviral Agents

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Abstract: Viral diseases are one of the leading causes of morbidity and mortality in the world. Broad-spectrum antiviral agents (BSAAs) are key players in control of human viral diseases. Here, we reviewed the discovery and development process of BSAAs, focusing on compounds with available safety profiles in human. In addition, we summarized the information on approved, investigational and experimental safe-in-man BSAAs in freely accessible database at <https://drugvirus.info/>. The number of approved BSAAs will be increased as well as their spectrum of indications will be expanded pending the results of further pre-clinical and clinical studies. This will ultimately reinforce the arsenal of available antiviral options and provide better protection of general population from emerging and re-emerging viral diseases.

Keywords: virus; antiviral drug; drug discovery; drug development; broad-spectrum antiviral agents, BSAAs

1. Introduction

Viruses are one of the major causes of morbidity and mortality in the world (1-4). Antiviral drugs and vaccines are used to fight viral infections in human (5, 6). Previously, there has been a focus on “one drug, one virus” dogma, which relied on targeting virus-specific factors. A counterpoint to this is “one drug, multiple viruses” paradigm, which came with the discovery of broad-spectrum antiviral agents (BSAAs), small-molecules that inhibit a wide range of human viruses (7-12). This paradigm was based on the observation that different viruses utilize similar pathways and host factors to replicate inside a cell (13). Although the concept of BSAAs has been around for almost 50 years, the field received a new impetus recently with recent outbreaks of Ebola, Zika, Dengue, influenza and other viral infections, the discovery of novel host-directed agents as well as development of drug repositioning methodology.

Drug repurposing, also called repositioning, redirecting, reprofiling, is a strategy for generating additional value from an existing drug by targeting disease other than that for which it was originally intended (14, 15). This has significant advantages over new drug discovery since chemical synthesis steps, manufacturing processes, reliable safety, and pharmacokinetic properties in pre-clinical (animal model) and early clinical developmental phases (phase 0, I and IIa) are already available. Therefore, repositioning of launched or even failed drugs from one disease to viral diseases provides unique translational opportunities, including a substantially higher probability of success to market as compared with developing new virus-specific drugs and vaccines, and a significantly reduced cost and timeline to clinical availability (9, 16, 17).

Here, we will describe repositioning of BSAs, focusing on those antivirals, which have been already tested in human as antivirals, antibacterials, antiprotozoals, anthelmintics, etc. Moreover, we will detail the steps of drug development process, from discovery of novel antiviral activities in cell culture to post-market studies (Fig. 1). Finally, we will discuss future perspectives of safe-in-man BSAs and their combinations for treatment of emerging and re-emerging viral infections and co-infections.

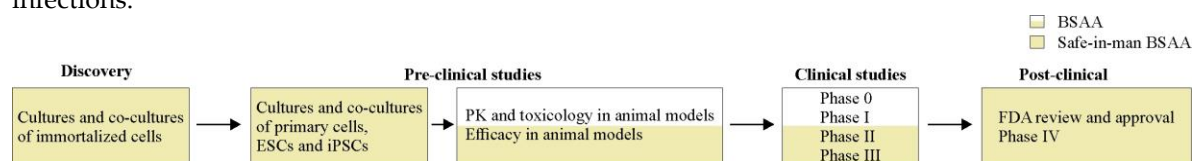


Figure 1. Discovery of novel activities and follow-up development of broad-spectrum antiviral agents (BSAs). Yellow shading indicates a process of discovery and development of safe-in-man BSAs, for which pharmacokinetic (PK) properties in pre-clinical (animal model) and early clinical developmental phases (phase 0-IIa trials) are already available. Abbreviations: ESCs, human embryonic stem cells; iPSCs, human induced pluripotent stem cells (iPSCs).

2. Discovery of novel activities of safe-in-man BSAs in immortalized cell cultures and co-cultures

The discovery of novel activities of BSAs starts with exposing cells to the candidate antiviral agent at different concentrations and infecting the cells with a virus of interest or mock. Immortalized cancerous cell cultures and co-cultures, which express appropriate viral receptors, are most commonly used in this first step. The half-maximal cytotoxic concentrations (CC_{50}) for a compound is calculated based on their dose-response curves obtained on mock-infected cells. The half-maximal effective concentrations (EC_{50}) are calculated based on the analysis of curves obtained on infected cells. Statistical analyses can help to determine if the differences between CC_{50} and EC_{50} are significant, given the inherent variability of the experiment (18). A relative effectiveness of a drug is defined as selectivity index ($SI = CC_{50}/EC_{50}$).

Cell viability assays and cell death assays are commonly used to assess the cytotoxicity and efficacy of BSAs (Fig. 2). Cell viability assays include MTT, MTS, resazurin or similar assays, mitochondrial membrane potential-dependent dyes-based assays, esterase cleaved dye-based assays, ATP-ADP assays, and assays that measure glycolytic flux and oxygen consumption. Other cell death assays include LDH enzyme leakage assays, membrane impermeable dye-based assays, and apoptosis assays, such as Annexin V, TUNEL, and caspase assays (www.abcam.com/kits/cell-health-assays-guide) (19). For example, the Cell Titer Glo assay quantifies ATP, an indicator of metabolically active living cells, whereas Cell Tox Green assay uses fluorescent asymmetric cyanine dye that stains the DNA of dead cells (<https://no.promega.com/products/cell-health-assays/cell-viability-and-cytotoxicity-assays/>) (13, 20-22).

Viral strains or cell lines expressing reporter proteins are also used to assess the efficacy of BSAs in infected cells. For example, TZM-bl cells expressing firefly luciferase under control of HIV-1 LTR promoter allowed quantitation of BSAA action on HIV-1 infection (tat-protein expression by integrated HIV-1 provirus) using firefly luciferase assay (23, 24). RFP-expressing RVFV, nanoLuc-expressing CHIKV and RRV, as well as GFP-expressing FLUAV, HCV and HMPV also allowed identification of novel activities of several BSAs (13, 21, 25-31). In addition, qPCR/RT-qPCR,

RNA/DNA sequencing, RNA/DNA hybridization, CRISPR-CAS immunofluorescence and plaque assays were used to detect inhibitory effects of BSAA on viral replications (32-39).

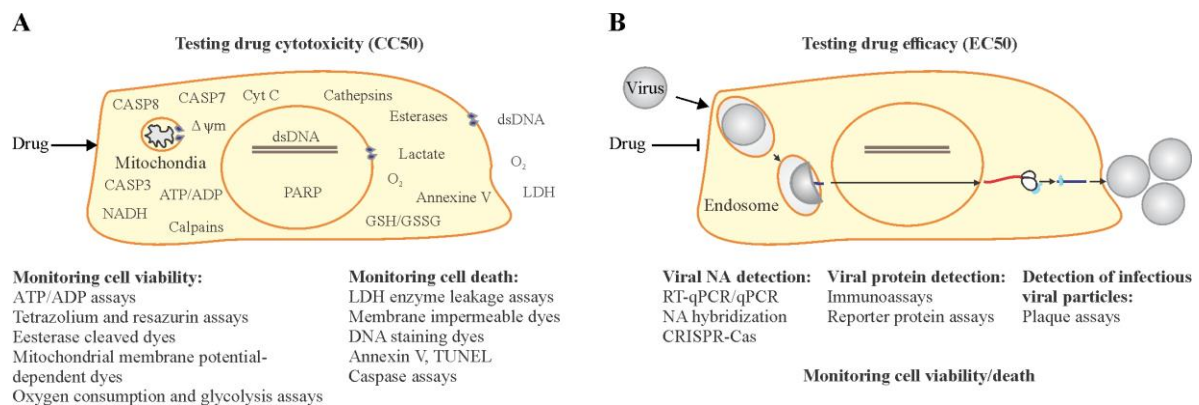


Figure 2. Testing BSAA toxicity (A) and efficacy (B) in immortalized cell cultures and co-cultures.

3. Pre-clinical evaluation of safe-in-man BSAA

3.1. Evaluation of safe-in-man BSAA in human primary cell cultures

Immortalized cancerous cell cultures/co-cultures and reporter viral strains represent excellent model systems for the discovery of novel activities of safe-in-man BSAA. However, these genetically modified systems have certain limitations (attenuated or incomplete virus replication cycle, accumulation of mutations during repeated cell and virus passaging, defective innate immune responses and viral counter-responses, etc.) (40). Thereby, novel antiviral activities of BSAA should be further validated in primary human cells using different viral strains (including wild-type viruses), different viral loads, different times of compound addition, different endpoint measurements and compound concentration range. Primary cell cultures give more accurate images of drug responses (41-44). They have a low population doubling level and therefore more closely recapitulate the physiological conditions observed in vivo.

Primary cells are cells isolated directly from tissues or blood using enzymatic or mechanical methods. The cells are characterized by their high degrees of specialization, are often fully differentiated and thus require defined culture conditions (serum-free media) in order to preserve their original phenotype. Peripheral blood mononuclear (PBMC), placental, amniotic and fetal primary cultures as well as vaginal/cervical epithelial and male germ cells have been used intensively to validate BSAA activity (42, 45-48). Although primary cell cultures are relevant systems for validation of BSAA, there are technical difficulties limiting their use, such as ethical issues, purity of population of primary cells, and limited shelf life of the cells. In addition, age, race, sex and other genetic and epigenetic factors of donor cells should be considered. For example, common genetic variants in IRF7 and IFITM3 gene loci which is associated with innate immune responses to FLUAV infection in monocyte-derived dendritic cells, could influence on the results of BSAA efficacy experiments (49, 50).

3.2. Evaluation of safe-in-man BSAA in human embryonic stem cell and human induced pluripotent stem cell cultures and co-cultures (organoids)

The obstacles associated with use of human primary cell cultures can be bypassed using human embryonic stem cells (ESCs) and human induced pluripotent stem cells (iPSCs). ESCs are isolated from surplus human embryos, whereas iPSCs are obtained by reprogramming somatic cells. These cells proliferate extensively and retain multi-lineage activity, which allows to generate virtually any cell type of the body. The ESCs- and iPSC-derived cells have been used successfully to investigate the efficacy of several BSAA against HBV, ZIKV, CHIKV and HSV-1 infections (Table S1) (51-56).

iPSCs, ESCs and primary tissue cells can be used to generate complex cultures termed organoids. Organoids are miniature and simplified version of organs. Establishing human airway, gut, skin, cerebral, liver, kidney, breast, retina and brain organoids allowed researchers to study toxicity and

efficacy of several safe-in-man BSAs against coronaviruses, influenza, enteroviruses, rotaviruses and flaviviruses (51, 57-63) (<https://organovir.com/>). However, iPSCs, ESCs and iPSCs/ESCs-derived organoids, have the same disadvantages as human primary cells (genetic differences, line-to-line and organoid batch-to-batch variability). On the other hand, these models allow researchers to predict the behavior of viruses in vivo and therefore to reduce animal use and in cases where animal models are unavailable to initiate clinical trials.

3.2. Evaluation of safe-in-man BSAs in animal models

In vitro and ex vivo models do not fully reflect the complexity and physiology of living organisms. Therefore, several in vivo models have been developed to test novel antiviral activities of BSAs. These include immunocompetent and genetically or chemically immunocompromised mice, guinea pigs, hamsters, ferrets, pigs, macaques and other animals (Fig. 3) (44, 64-68). PK/PD studies determine drug absorption, dosage and half-life of BSAs. Toxicological studies determine if the drugs have any adverse effects on the tissues and organs of the animals and defining the dosage of adverse effects (69-71). Studying the efficacy of BSAs is generally done by treating the animal with the drug or vehicle and infecting it with a virus of interest. Endpoints are usually body weight/mortality (depending on the virus), histopathology, virus titers in organs, presence of clinical signs and development of immunity (72, 73). Although animal models can give the initial characterization of BSA, it is important to keep in mind that they differ significantly from humans, with respect to symptoms, disease manifestation, susceptibility, immune responses, pathogenesis, and pharmacokinetics (74, 75).

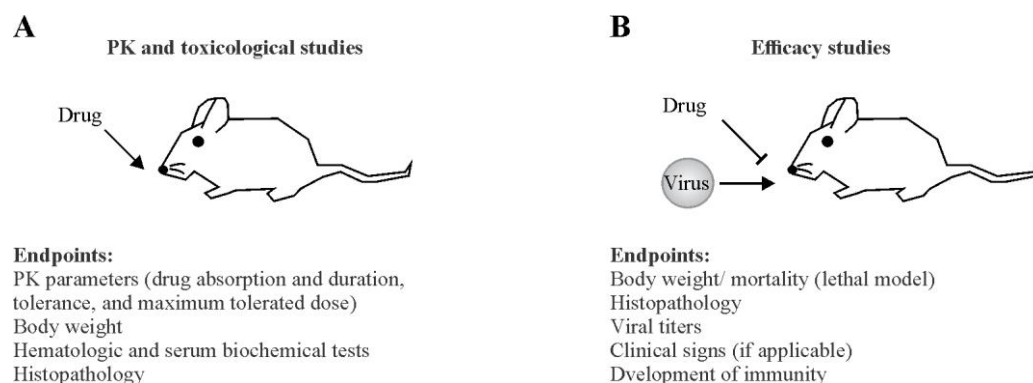


Figure 3. Testing toxicity and efficacy of BSAs in animal models. (A) PK/PD and toxicity studies. (B) Efficacy studies. If BSA is repositioned from another disease (i.e. its PK/PD and toxicity profiles are available for the animal model) it could bypass the safety studies.

4. Clinical trials and post-clinical studies of safe-in-man BSAs

Clinical trials are the most critical and time-consuming step of a drug candidates' journey to being approved (Fig. 4). However, safe-in-man BSAs make this journey relatively short, because they have been already at phase 0, I and sometime at IIa of clinical trials as antibacterial, antiparasitic, anticancer, etc. agent; i.e. they have been administered at sub-therapeutic doses to healthy volunteers to ensure the drugs are not harmful to the participants. Thus, safe-in-man BSAs enter phase II and III, which assess the efficacy, effectiveness, safety and side effects of the drugs in clinic. For this, patients with the viral disease in question are invited to join the study, where they are administered the BSAs at the ideal therapeutic doses. Phase III is the longest of the phases, and include multiple levels of securities to the studies, such as the use of placebos and double-blinded studies, to ensure the data is as unbiased as possible. Upon completing phase III, depending on its performance and efficacy, BSAs may end either being approved or dropped. The U.S Food and Drug Administration (FDA) estimates that only 25-30% BSA candidates which enters phase III are approved for use in the public (76). After approval and marketing of the drug, phase IV may be initiated to follow up on the use of the drug in public, to surveil for rare effects (76, 77).

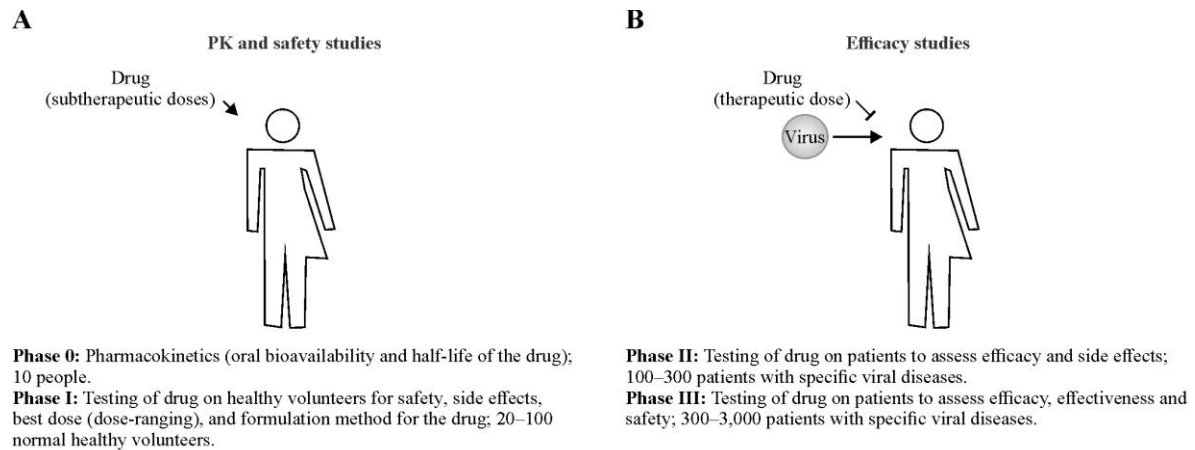


Figure 4. Clinical trials of BSAA. (A) Pharmacokinetics (PK) and safety studies. (B). Efficacy studies. If BSAA is repositioned from another disease (i.e. its safety profile in man is available) it will bypass the PK and safety studies.

5. Database of safe-in-man BSAA

We have developed a database for safe-in-man BSAA, which is available at <https://drugvirus.info/>. The drug annotations were obtained from PubChem, DrugBank, DrugCentral, PubMed and clinicaltrials.gov databases (Table S1) (78–80). The information on virus families were exported from Virus Pathogen Database and Analysis Resource (Table S2) (81). The database summarizes activities and developmental stages of BSAA (Fig. 5). The database allows interactive exploration of virus-BSAA interactions. It also includes information on BSA targets. A feedback form is available on the website. The website will be updated upon request or as soon as a new safe-in-man BSAA emerge or novel activity for an existing BSAA is reported.

The database includes 21 BSAA which were approved by FDA, EMA or other agencies. These BSAA altogether target 15 viruses. For example, favipiravir, also known as T-705, was approved against FLUAV in Japan; cidofovir is an injectable antiviral medication used as a treatment for CMV retinitis in people with AIDS; ribavirin, also known as tribavirin, is used for treatment of RSV and HCV infections; pleconaril is used against viruses in the picornaviridae family, including enterovirus and rhinovirus; and valacyclovir is used against CMV, EBV, HBV, HSV-1, HSV-2 and VZV infections. Twenty BSAA are undergoing surveillance studies (phase IV). Azithromycin, chloroquine, cyclosporine, ezetimibe, mycophenolic acid, nitazoxanide and rapamycin progressed to phase IV studies without approvals from national or international authorities (NCT01779570, NCT02058173, NCT02564471, NCT00821587, NCT03360682, NCT02328963, NCT02768545, NCT01624948, NCT01770483, NCT02683291, NCT01624948, NCT01469884, NCT03901001, NCT01412515, NCT02990312).

The database also includes 48 safe-in-man BSAA, which undergo clinical studies as antivirals. There are currently 21 compounds in phase I, 34 agents in phase II and 11 compounds in phase III clinical trials. For example, nitazoxanide, remdesivir and brincidofovir are under clinical investigations against different viral infections (NCT03336619, NCT00302640, NCT03605862, NCT03719586, NCT01276756, NCT03905655, NCT01529073, NCT03395405, NCT03216967, NCT01431326, NCT02087306, NCT01769170). The rest of safe-in-man BSAA are still in pre-clinical or discovery stages. Of the drugs in this group, niclosamide is one of the interesting compounds because it showed the broadest spectrum of activities in vitro and in some cases in vivo (31, 61, 82–89). We believe that emetine and gemcitabine could be also pursued as potential BSAA candidates (31, 42, 90). ABT-263, also known as navitoclax, is another interesting BSAA, which is, by contrast to other compounds, facilitates death of infected cells without affecting non-infected cells (20, 91, 92).

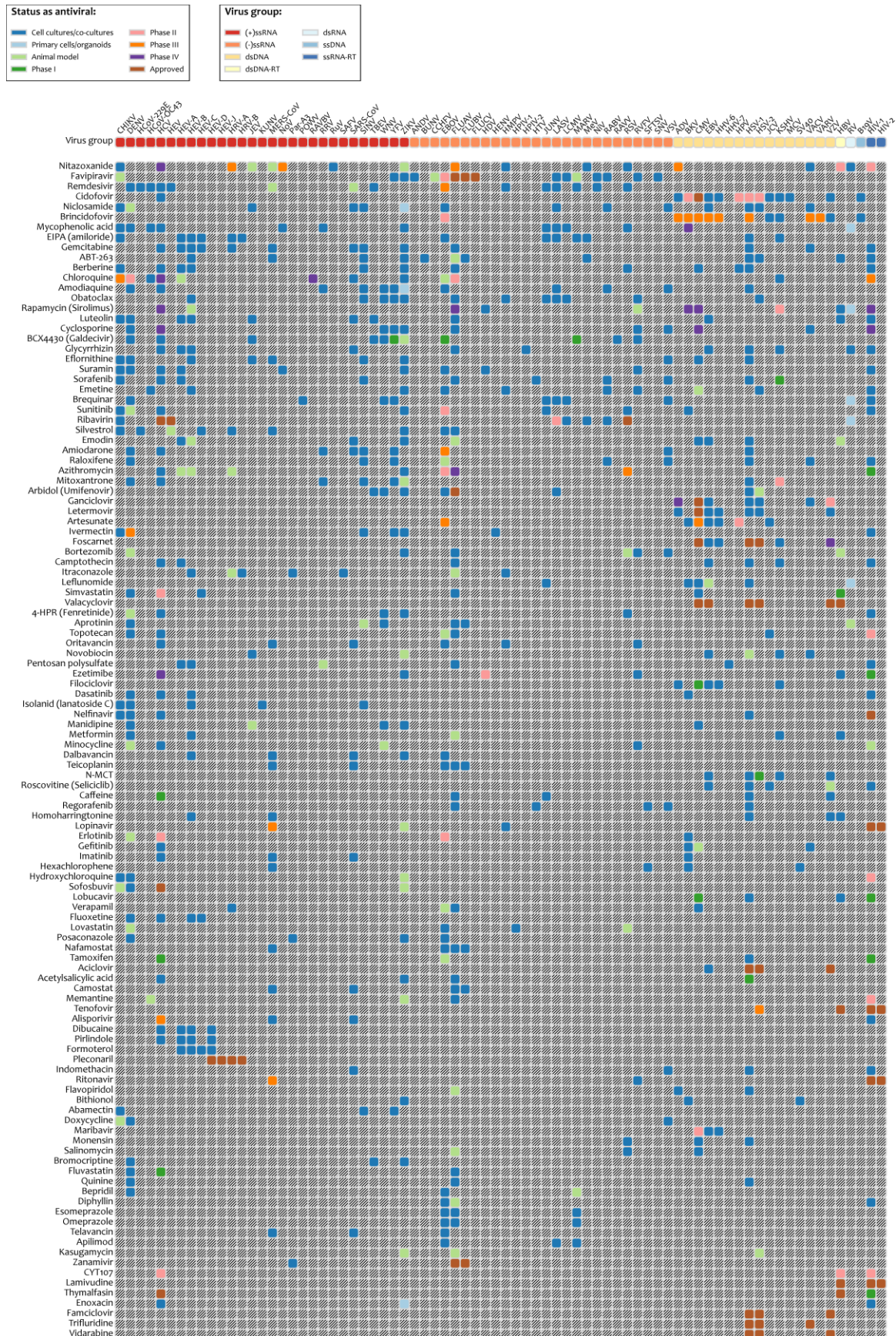


Figure 5. Hundred and nineteen safe-in-man broad-spectrum antiviral agents (BSAAs) and viruses they inhibit. A snapshot is taken from <https://drugvirus.info/> website. Viruses are clustered by virus groups. BSAAs are ranged from the highest to lowest number of targeted viruses. Different shadings indicate different development status of BSAAs. Gray shading indicates that the antiviral activity has not been either studied or reported. Abbreviations: ds, double-stranded; RT, reverse transcriptase; ss, single-stranded.

Altogether, the database contains 119 approved, investigational and experimental safe-in-man BSAAs, which inhibit 83 human viruses, belonging to 25 viral families. The BSAAs inhibit viral, host or both viral and host factors (Table S1). Analysis of BSAA targets and structures (Fig. 6) revealed that the most abundant are nucleotide and nucleoside analogues which inhibit viral RNA and DNA polymerases. Imatinib, erlotinib, gefitinib, and dasatinib that inhibit tyrosine kinases are the most abundant host-directed BSAAs. Most of the host targets (except Bcl-xL protein) are essential for viral replication but redundant for the cell, which is critical for reducing putative toxicities associated with blocking cellular pathways. The limited diversity of the targets and scaffolds could slow down the development of BSAAs.

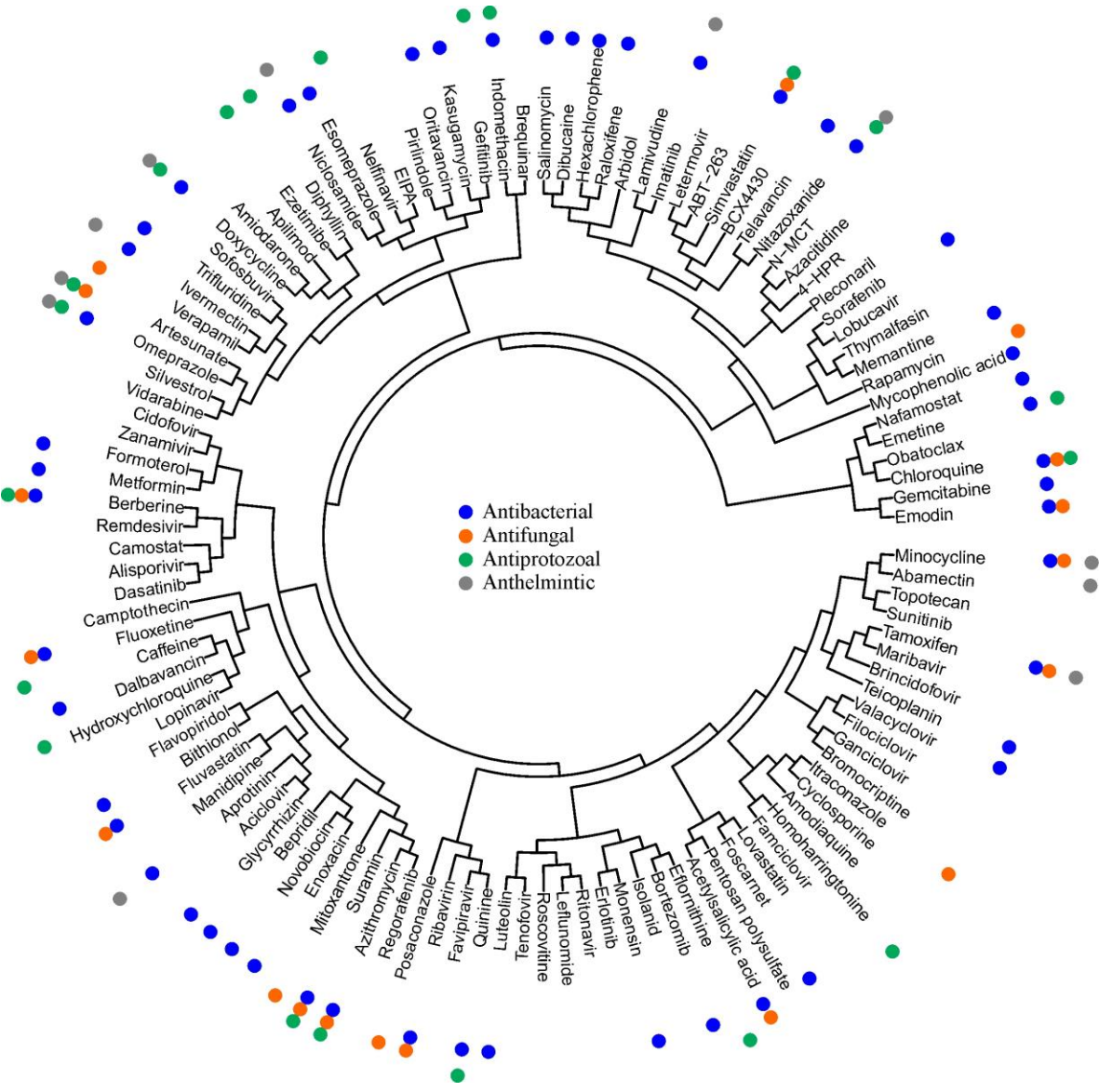


Figure 6. Structure-activity relationship of safe-in-man broad-spectrum antiviral agents.

6. Conclusions and future perspectives

Here, we reviewed the discovery and development processes of safe-in-man BSAs. In addition, we developed a database which consists of 119 BSAs. These BSAs block viral replication completely, reduce the viral burden to a level at which host immune responses can deal with it or facilitate apoptosis of infected cells. The database will be updated as soon as a new safe-in-man BSA emerge or novel activity for an existing BSA is reported.

Emerging BSAs, such as 5,6-dimethoxyindan-1-one, saliphenylhalamide, and GS-5734 (22, 42, 90, 93-95), whose safety profiles in humans are not yet available, could serve as valuable antivirals in the future, pending the results of further pre-clinical and clinical investigations. The follow-up studies as well as the results of on-going, finalized or terminated clinical trials should be made publicly available to allow posterization and translation of emerging and existing BSAs into clinical practice.

BSAs could be combined with other antiviral agents to obtain synergistic or additive effects against certain viruses (17, 96). For example, it was reported that obatoclax and saliphenylhalamide, as well as gemcitabine and pimodivir (JNJ872) possessed synergistic effects against ZIKV and FLUAV infections in vitro, respectively (90, 97). Moreover, many combination therapies, which include BSAs, became a standard for the treatment of HIV and HCV infections. These include abacavir/dolutegravir/lamivudine (Triumeq), darunavir/cobicistat/emtricitabine/tenofovir (Symtuza), lopinavir/ritonavir (Kaletra), ledipasvir/sofosbuvir and sofosbuvir/velpatasvir (98-100).

By contrast to individual drugs, combinations of 2-3 BSAs could be used to target even broader range of viruses (17, 101). Such combinations could serve as front line therapeutics against poorly characterized emerging viruses or re-emerging drug-resistant viral strains. For example, a cocktail of nitazoxanide, favipiravir, and niclosamide could be developed for the treatment of infections of viruses belonging to 11 families.

Fifty BSAs possess not only antiviral but also antibacterial activity (Fig. 6; Table S1) (102). Moreover, 13 of the 50 agents are approved as antibiotics (2 withdrawn). These agents with dual activity could be used for treatment of viral and bacterial co-infections or for protection of patients from the secondary infections. For example, azithromycin could be used against FLUAV and *Chlamydomphila pneumoniae*, *Haemophilus influenzae*, *Mycoplasma pneumoniae* or *Streptococcus pneumoniae* infections (NCT01779570) (103).

In addition, BSAs showed activity against a wide range of other medically important human pathogens, including fungi, protozoa and parasites (Table S1) (104), pointing out that some pathogens utilize common mechanisms to infect hosts. Moreover, structure-activity relationship analysis of BSAs suggest that some agents, such as doxycycline, artesunate, omeprazole, nitazoxanide, suramin, azithromycin, minocycline and chloroquine, could have novel antibacterial, antiprotozoal, antifungal or anthelmintic activities (Fig. 6). If confirmed, this could lead to development of broad-spectrum anti-infective drugs.

BSAs could also serve as treatment of other co-morbidities simplifying the therapy and lowering its cost (Table S1). For example, the concomitant actions of ezetimibe and statins could be beneficial for treatment of both hypertension and several viral infections in patients with these co-morbidities (NCT00908011, NCT00099684, NCT00843661, NCT03490097, NCT00994773, NCT00441493).

In conclusion, BSAs could play a pivotal role in the battle against emerging and re-emerging viral diseases. Discovery of novel BSAs as well as repositioning existing safe-in-man BSAs may shorten time and resources, needed for development of virus-specific drugs and vaccines. In the future, BSAs will have global impact by decreasing morbidity and mortality from viral and other diseases, maximizing the number of healthy life years, improving the quality of life of infected patients and decreasing the costs of patient care.

Supplementary Materials: The following are available online at <http://www.mdpi.com/XXXX>. Table S1: Safe-in-man broad-spectrum antiviral agents; Table S2: Human viruses and associated diseases.

Author Contributions: Conceptualization, D.E.K.; Investigation and Validation, P.I.A., A.I., H.L. and D.E.K.; Data Curation & Visualization, A.I. and D.E.K.; Writing & Editing, all authors; Project Administration and Funding Acquisition, D.E.K.

Funding: This research was funded by Estonian Research Council Mobilitas pluss top researcher grant (contract No. MOBTT3).

Acknowledgments: This study was supported by the European Regional Development Fund, the Mobilitas Pluss Project MOBTT39 (to D.K.).

Conflicts of Interest: The authors declare no conflict of interest.

References

1. Disease GBD, Injury I, Prevalence C. 2018. Global, regional, and national incidence, prevalence, and years lived with disability for 354 diseases and injuries for 195 countries and territories, 1990-2017: a systematic analysis for the Global Burden of Disease Study 2017. *Lancet* 392:1789-1858.
2. DALYs GBD, Collaborators H. 2018. Global, regional, and national disability-adjusted life-years (DALYs) for 359 diseases and injuries and healthy life expectancy (HALE) for 195 countries and territories, 1990-2017: a systematic analysis for the Global Burden of Disease Study 2017. *Lancet* 392:1859-1922.
3. WHO. 2015. WHO publishes list of top emerging diseases likely to cause major epidemics. www.who.int/medicines/ebola-treatment/WHO-list-of-top-emerging-diseases/en/.
4. Howard CR, Fletcher NF. 2012. Emerging virus diseases: can we ever expect the unexpected? *Emerg Microbes Infect* 1:e46.
5. Marston HD, Folkers GK, Morens DM, Fauci AS. 2014. Emerging viral diseases: confronting threats with new technologies. *Sci Transl Med* 6:253ps10.
6. De Clercq E, Li G. 2016. Approved Antiviral Drugs over the Past 50 Years. *Clin Microbiol Rev* 29:695-747.
7. Bekerman E, Einav S. 2015. Infectious disease. Combating emerging viral threats. *Science* 348:282-3.
8. Debing Y, Neyts J, Delang L. 2015. The future of antivirals: broad-spectrum inhibitors. *Curr Opin Infect Dis* 28:596-602.
9. Ianevski A, Andersen PI, Merits A, Bjorås M, Kainov D. 2019. Expanding the activity spectrum of antiviral agents. *Drug Discov Today* 24:1224-1228.
10. de Clercq E, Montgomery JA. 1983. Broad-spectrum antiviral activity of the carbocyclic analog of 3-deazaadenosine. *Antiviral Res* 3:17-24.
11. Rada B, Dragun M. 1977. Antiviral action and selectivity of 6-azauridine. *Ann N Y Acad Sci* 284:410-7.
12. Sidwell RW, Huffman JH, Khare GP, Allen LB, Witkowski JT, Robins RK. 1972. Broad-spectrum antiviral activity of Virazole: 1-beta-D-ribofuranosyl-1,2,4-triazole-3-carboxamide. *Science* 177:705-6.
13. Bosl K, Ianevski A, Than TT, Andersen PI, Kuivanen S, Teppor M, Zusinaite E, Dumpis U, Vitkauskienė A, Cox RJ, Kallio-Kokko H, Bergqvist A, Tenson T, Merits A, Oksenych V, Bjorås M, Anthonsen M, Shum D, Kaarbo M, Vapalahti O, Windisch

- MP, Superti-Furga G, Snijder B, Kainov D, Kandasamy RK. 2019. Common Nodes of Virus–Host Interaction Revealed Through an Integrated Network Analysis. *Front Immunology* 4:2186.
14. Nishimura Y, Hara H. 2018. Editorial: Drug Repositioning: Current Advances and Future Perspectives. *Front Pharmacol* 9:1068.
 15. Pushpakom S, Iorio F, Eyers PA, Escott KJ, Hopper S, Wells A, Doig A, Guillems T, Latimer J, McNamee C, Norris A, Sanseau P, Cavalla D, Pirmohamed M. 2019. Drug repurposing: progress, challenges and recommendations. *Nat Rev Drug Discov* 18:41-58.
 16. Pizzorno A, Padey B, Terrier O, Rosa-Calatrava M. 2019. Drug Repurposing Approaches for the Treatment of Influenza Viral Infection: Reviving Old Drugs to Fight Against a Long-Lived Enemy. 10.
 17. Zheng W, Sun W, Simeonov A. 2018. Drug repurposing screens and synergistic drug-combinations for infectious diseases. *Br J Pharmacol* 175:181-191.
 18. Meneghini KA, Hamasaki DI. 1967. The electroretinogram of the iguana and Tokay gecko. *Vision Res* 7:243-51.
 19. Shen L, Niu J, Wang C, Huang B, Wang W, Zhu N, Deng Y, Wang H, Ye F, Cen S, Tan W. 2019. High-Throughput Screening and Identification of Potent Broad-Spectrum Inhibitors of Coronaviruses. *J Virol* 93.
 20. Bulanova D, Ianevski A, Bugai A, Akimov Y, Kuivanen S, Paavilainen H, Kakkola L, Nandania J, Turunen L, Ohman T, Ala-Hongisto H, Pesonen HM, Kuisma MS, Honkimaa A, Walton EL, Oksenych V, Lorey MB, Guschin D, Shim J, Kim J, Than TT, Chang SY, Hukkanen V, Kuleskiy E, Marjomaki VS, Julkunen I, Nyman TA, Matikainen S, Saarela JS, Sane F, Hober D, Gabriel G, De Brabander JK, Martikainen M, Windisch MP, Min JY, Bruzzone R, Aittokallio T, Vaha-Koskela M, Vapalahti O, Pulk A, Velagapudi V, Kainov DE. 2017. Antiviral Properties of Chemical Inhibitors of Cellular Anti-Apoptotic Bcl-2 Proteins. *Viruses* 9.
 21. Ianevski A, Zusinaite E, Kuivanen S, Strand M, Lysvand H, Teppor M, Kakkola L, Paavilainen H, Laajala M, Kallio-Kokko H, Valkonen M, Kantele A, Telling K, Lutsar I, Letjuka P, Metelitsa N, Oksenych V, Bjoras M, Nordbo SA, Dumpis U, Vitkauskienė A, Ohrmalm C, Bondeson K, Bergqvist A, Aittokallio T, Cox RJ, Evander M, Hukkanen V, Marjomaki V, Julkunen I, Vapalahti O, Tenson T, Merits A, Kainov D. 2018. Novel activities of safe-in-human broad-spectrum antiviral agents. *Antiviral Res* 154:174-182.
 22. Muller KH, Spoden GA, Scheffer KD, Brunnhofer R, De Brabander JK, Maier ME, Florin L, Muller CP. 2014. Inhibition by cellular vacuolar ATPase impairs human papillomavirus uncoating and infection. *Antimicrob Agents Chemother* 58:2905-11.
 23. Sarzotti-Kelsoe M, Bailer RT, Turk E, Lin CL, Bilska M, Greene KM, Gao H, Todd CA, Ozaki DA, Seaman MS, Mascola JR, Montefiori DC. 2014. Optimization and validation of the TZM-bl assay for standardized assessments of neutralizing antibodies against HIV-1. *J Immunol Methods* 409:131-46.
 24. Xing L, Wang S, Hu Q, Li J, Zeng Y. 2016. Comparison of three quantification methods for the TZM-bl pseudovirus assay for screening of anti-HIV-1 agents. *J Virol Methods* 233:56-61.

25. Habjan M, Penski N, Spiegel M, Weber F. 2008. T7 RNA polymerase-dependent and -independent systems for cDNA-based rescue of Rift Valley fever virus. *J Gen Virol* 89:2157-66.
26. Lee M, Yang J, Jo E, Lee JY, Kim HY, Bartenschlager R, Shin EC, Bae YS, Windisch MP. 2017. A Novel Inhibitor IDPP Interferes with Entry and Egress of HCV by Targeting Glycoprotein E1 in a Genotype-Specific Manner. *Sci Rep* 7:44676.
27. Kittel C, Sereinig S, Ferko B, Stasakova J, Romanova J, Wolkerstorfer A, Katinger H, Egorov A. 2004. Rescue of influenza virus expressing GFP from the NS1 reading frame. *Virology* 324:67-73.
28. de Graaf M, Herfst S, Schrauwen EJ, van den Hoogen BG, Osterhaus AD, Fouchier RA. 2007. An improved plaque reduction virus neutralization assay for human metapneumovirus. *J Virol Methods* 143:169-74.
29. Utt A, Quirin T, Saul S, Hellstrom K, Ahola T, Merits A. 2016. Versatile Trans-Replication Systems for Chikungunya Virus Allow Functional Analysis and Tagging of Every Replicase Protein. *PLoS One* 11:e0151616.
30. Jupille HJ, Oko L, Stoermer KA, Heise MT, Mahalingam S, Gunn BM, Morrison TE. 2011. Mutations in nsP1 and PE2 are critical determinants of Ross River virus-induced musculoskeletal inflammatory disease in a mouse model. *Virology* 410:216-27.
31. Andersen PI, Krpina K, Ianevski A, Shtaida N, Jo E, Yang J, Koit S, Tenson T, Hukkonen V, Anthonsen MW, Bjoras M, Evander M, Windisch MP, Zusinaite E, Kainov DE. 2019. Novel Antiviral Activities of Obatoclax, Emetine, Niclosamide, Brequinar, and Homoharringtonine. *Viruses* 11:pii: E964.
32. Konig A, Yang J, Jo E, Park KHP, Kim H, Than TT, Song X, Qi X, Dai X, Park S, Shum D, Ryu WS, Kim JH, Yoon SK, Park JY, Ahn SH, Han KH, Gerlich WH, Windisch MP. 2019. Efficient long-term amplification of hepatitis B virus isolates after infection of slow proliferating HepG2-NTCP cells. *J Hepatol* 71:289-300.
33. Fischer C, Torres MC, Patel P, Moreira-Soto A, Gould EA, Charrel RN, de Lamballerie X, Nogueira RMR, Sequeira PC, Rodrigues CDS, Kümmerer BM, Drosten C, Landt O, Bispo de Filippis AM, Drexler JF. 2017. Lineage-Specific Real-Time RT-PCR for Yellow Fever Virus Outbreak Surveillance, Brazil. *Emerg Infect Dis* 23:1867-71.
34. Laamiri N, Aouini R, Marnissi B, Ghram A, Hmila I. 2018. A multiplex real-time RT-PCR for simultaneous detection of four most common avian respiratory viruses. *Virology* 515:29-37.
35. Landry ML. 1990. Nucleic acid hybridization in viral diagnosis. *Clinical Biochemistry* 23:267-277.
36. Boonham N, Kreuze J, Winter S, van der Vlugt R, Bergervoet J, Tomlinson J, Mumford R. 2014. Methods in virus diagnostics: from ELISA to next generation sequencing. *Virus Res* 186:20-31.
37. Perez JT, Garcia-Sastre A, Manicassamy B. 2013. Insertion of a GFP reporter gene in influenza virus. *Curr Protoc Microbiol* Chapter 15:Unit 15G.4.
38. Sashital DG. 2018. Pathogen detection in the CRISPR-Cas era. *Genome medicine* 10:32-32.

39. Zhou L, Peng R, Zhang R, Li J. 2018. The applications of CRISPR/Cas system in molecular detection. *Journal of cellular and molecular medicine* 22:5807-5815.
40. Carter M, Shieh J. 2015. Chapter 14 - Cell Culture Techniques, p 295-310. *In* Carter M, Shieh J (ed), *Guide to Research Techniques in Neuroscience* (Second Edition) doi:<https://doi.org/10.1016/B978-0-12-800511-8.00014-9>. Academic Press, San Diego.
41. Postnikova E, Cong Y, DeWald LE, Dyall J, Yu S, Hart BJ, Zhou H, Gross R, Logue J, Cai Y, Deilulis N, Michelotti J, Honko AN, Bennett RS, Holbrook MR, Olinger GG, Hensley LE, Jahrling PB. 2018. Testing therapeutics in cell-based assays: Factors that influence the apparent potency of drugs. *PLoS One* 13:e0194880.
42. Denisova OV, Kakkola L, Feng L, Stenman J, Nagaraj A, Lampe J, Yadav B, Aittokallio T, Kaukinen P, Ahola T, Kuivanen S, Vapalahti O, Kantele A, Tynell J, Julkunen I, Kallio-Kokko H, Paavilainen H, Hukkanen V, Elliott RM, De Brabander JK, Saelens X, Kainov DE. 2012. Obatoclax, saliphenylhalamide, and gemcitabine inhibit influenza A virus infection. *J Biol Chem* 287:35324-32.
43. Koban R, Neumann M, Dausgs A, Bloch O, Nitsche A, Langhammer S, Ellerbrok H. 2018. A novel three-dimensional cell culture method enhances antiviral drug screening in primary human cells. *Antiviral Res* 150:20-29.
44. Alves MP, Vielle NJ, Thiel V, Pfaender S. 2018. Research Models and Tools for the Identification of Antivirals and Therapeutics against Zika Virus Infection. *Viruses* 10.
45. Fink SL, Vojtech L, Wagoner J, Slivinski NSJ, Jackson KJ, Wang R, Khadka S, Luthra P, Basler CF, Polyak SJ. 2018. The Antiviral Drug Arbidol Inhibits Zika Virus. *Sci Rep* 8:8989.
46. Robinson CL, Chong ACN, Ashbrook AW, Jeng G, Jin J, Chen H, Tang EI, Martin LA, Kim RS, Kenyon RM, Do E, Luna JM, Saeed M, Zeltser L, Ralph H, Dudley VL, Goldstein M, Rice CM, Cheng CY, Seandel M, Chen S. 2018. Male germ cells support long-term propagation of Zika virus. *Nat Commun* 9:2090.
47. Rausch K, Hackett BA, Weinbren NL, Reeder SM, Sadovsky Y, Hunter CA, Schultz DC, Coyne CB, Cherry S. 2017. Screening Bioactives Reveals Nanchangmycin as a Broad Spectrum Antiviral Active against Zika Virus. *Cell Rep* 18:804-815.
48. Barrows NJ, Campos RK, Powell ST, Prasanth KR, Schott-Lerner G, Soto-Acosta R, Galarza-Munoz G, McGrath EL, Urrabaz-Garza R, Gao J, Wu P, Menon R, Saade G, Fernandez-Salas I, Rossi SL, Vasilakis N, Routh A, Bradrick SS, Garcia-Blanco MA. 2016. A Screen of FDA-Approved Drugs for Inhibitors of Zika Virus Infection. *Cell Host Microbe* 20:259-70.
49. Lee MN, Ye C, Villani AC, Raj T, Li W, Eisenhaure TM, Imboywa SH, Chipendo PI, Ran FA, Slowikowski K, Ward LD, Raddassi K, McCabe C, Lee MH, Frohlich IY, Hafler DA, Kellis M, Raychaudhuri S, Zhang F, Stranger BE, Benoist CO, De Jager PL, Regev A, Hacohen N. 2014. Common genetic variants modulate pathogen-sensing responses in human dendritic cells. *Science* 343:1246980.
50. Zhang YH, Zhao Y, Li N, Peng YC, Giannoulitou E, Jin RH, Yan HP, Wu H, Liu JH, Liu N, Wang DY, Shu YL, Ho LP, Kellam P, McMichael A, Dong T. 2013. Interferon-induced transmembrane protein-3 genetic variant rs12252-C is associated with severe influenza in Chinese individuals. *Nat Commun* 4:1418.

51. Zhou T, Tan L, Cederquist GY, Fan Y, Hartley BJ, Mukherjee S, Tomishima M, Brennand KJ, Zhang Q, Schwartz RE, Evans T, Studer L, Chen S. 2017. High-Content Screening in hPSC-Neural Progenitors Identifies Drug Candidates that Inhibit Zika Virus Infection in Fetal-like Organoids and Adult Brain. *Cell Stem Cell* 21:274-283 e5.
52. Lanko K, Eggermont K, Patel A, Kaptein S, Delang L, Verfaillie CM, Neyts J. 2017. Replication of the Zika virus in different iPSC-derived neuronal cells and implications to assess efficacy of antivirals. *Antiviral Res* 145:82-86.
53. Iwasawa C, Tamura R, Sugiura Y, Suzuki S, Kuzumaki N, Narita M, Suematsu M, Nakamura M, Yoshida K, Toda M, Okano H, Miyoshi H. 2019. Increased Cytotoxicity of Herpes Simplex Virus Thymidine Kinase Expression in Human Induced Pluripotent Stem Cells. *Int J Mol Sci* 20.
54. Simonin Y, Erkilic N, Damodar K, Cle M, Desmetz C, Bollere K, Taleb M, Torriano S, Barthelemy J, Dubois G, Lajoix AD, Foulongne V, Tuaillon E, Van de Perre P, Kalatzis V, Salinas S. 2019. Zika virus induces strong inflammatory responses and impairs homeostasis and function of the human retinal pigment epithelium. *EBioMedicine* 39:315-331.
55. Ferreira AC, Reis PA, de Freitas CS, Sacramento CQ, Villas Boas Hoelz L, Bastos MM, Mattos M, Rocha N, Gomes de Azevedo Quintanilha I, da Silva Gouveia Pedrosa C, Rocha Quintino Souza L, Correia Loiola E, Trindade P, Rangel Vieira Y, Barbosa-Lima G, de Castro Faria Neto HC, Boechat N, Rehen SK, Bruning K, Bozza FA, Bozza PT, Souza TML. 2019. Beyond Members of the Flaviviridae Family, Sofosbuvir Also Inhibits Chikungunya Virus Replication. *Antimicrob Agents Chemother* 63.
56. Xia Y, Carpentier A, Cheng X, Block PD, Zhao Y, Zhang Z, Protzer U, Liang TJ. 2017. Human stem cell-derived hepatocytes as a model for hepatitis B virus infection, spreading and virus-host interactions. *J Hepatol* 66:494-503.
57. Yin Y, Chen S, Hakim MS, Wang W, Xu L, Dang W, Qu C, Verhaar AP, Su J, Fuhler GM, Peppelenbosch MP, Pan Q. 2018. 6-Thioguanine inhibits rotavirus replication through suppression of Rac1 GDP/GTP cycling. *Antiviral Res* 156:92-101.
58. Watanabe M, Buth JE, Vishlaghi N, de la Torre-Ubieta L, Taxidis J, Khakh BS, Coppola G, Pearson CA, Yamauchi K, Gong D, Dai X, Damoiseaux R, Aliyari R, Liebscher S, Schenke-Layland K, Caneda C, Huang EJ, Zhang Y, Cheng G, Geschwind DH, Golshani P, Sun R, Novitsch BG. 2017. Self-Organized Cerebral Organoids with Human-Specific Features Predict Effective Drugs to Combat Zika Virus Infection. *Cell Rep* 21:517-532.
59. Li C, Deng YQ, Wang S, Ma F, Aliyari R, Huang XY, Zhang NN, Watanabe M, Dong HL, Liu P, Li XF, Ye Q, Tian M, Hong S, Fan J, Zhao H, Li L, Vishlaghi N, Buth JE, Au C, Liu Y, Lu N, Du P, Qin FX, Zhang B, Gong D, Dai X, Sun R, Novitsch BG, Xu Z, Qin CF, Cheng G. 2017. 25-Hydroxycholesterol Protects Host against Zika Virus Infection and Its Associated Microcephaly in a Mouse Model. *Immunity* 46:446-456.
60. Sacramento CQ, de Melo GR, de Freitas CS, Rocha N, Hoelz LV, Miranda M, Fintelman-Rodrigues N, Marttorelli A, Ferreira AC, Barbosa-Lima G, Abrantes JL,

- Vieira YR, Bastos MM, de Mello Volotao E, Nunes EP, Tschoeke DA, Leomil L, Loiola EC, Trindade P, Rehen SK, Bozza FA, Bozza PT, Boechat N, Thompson FL, de Filippis AM, Bruning K, Souza TM. 2017. The clinically approved antiviral drug sofosbuvir inhibits Zika virus replication. *Sci Rep* 7:40920.
61. Xu M, Lee EM, Wen Z, Cheng Y, Huang WK, Qian X, Tcw J, Kouznetsova J, Ogden SC, Hammack C, Jacob F, Nguyen HN, Itkin M, Hanna C, Shinn P, Allen C, Michael SG, Simeonov A, Huang W, Christian KM, Goate A, Brennand KJ, Huang R, Xia M, Ming GL, Zheng W, Song H, Tang H. 2016. Identification of small-molecule inhibitors of Zika virus infection and induced neural cell death via a drug repurposing screen. *Nat Med* 22:1101-1107.
 62. Yin Y, Wang Y, Dang W, Xu L, Su J, Zhou X, Wang W, Felczak K, van der Laan LJ, Pankiewicz KW, van der Eijk AA, Bijvelds M, Sprengers D, de Jonge H, Koopmans MP, Metselaar HJ, Peppelenbosch MP, Pan Q. 2016. Mycophenolic acid potently inhibits rotavirus infection with a high barrier to resistance development. *Antiviral Res* 133:41-9.
 63. Yin Y, Bijvelds M, Dang W, Xu L, van der Eijk AA, Knipping K, Tuysuz N, Dekkers JF, Wang Y, de Jonge J, Sprengers D, van der Laan LJ, Beekman JM, Ten Berge D, Metselaar HJ, de Jonge H, Koopmans MP, Peppelenbosch MP, Pan Q. 2015. Modeling rotavirus infection and antiviral therapy using primary intestinal organoids. *Antiviral Res* 123:120-31.
 64. Morrison TE, Diamond MS. 2017. Animal Models of Zika Virus Infection, Pathogenesis, and Immunity. *J Virol* 91.
 65. Haese NN, Broeckel RM, Hawman DW, Heise MT, Morrison TE, Streblow DN. 2016. Animal Models of Chikungunya Virus Infection and Disease. *J Infect Dis* 214:S482-S487.
 66. Taylor G. 2017. Animal models of respiratory syncytial virus infection. *Vaccine* 35:469-480.
 67. Thangavel RR, Bouvier NM. 2014. Animal models for influenza virus pathogenesis, transmission, and immunology. *J Immunol Methods* 410:60-79.
 68. Louz D, Bergmans HE, Loos BP, Hoeben RC. 2013. Animal models in virus research: their utility and limitations. *Crit Rev Microbiol* 39:325-61.
 69. Alabaster V., In Vivo Pharmacology Training Group. 2002. The fall and rise of in vivo pharmacology. *Trends Pharmacol Sci* 23:13-8.
 70. Rizk ML, Zou L, Savic RM, Dooley KE. 2017. Importance of Drug Pharmacokinetics at the Site of Action. *Clinical and translational science* 10:133-142.
 71. Parasuraman S. 2011. Toxicological screening. *Journal of pharmacology & pharmacotherapeutics* 2:74-79.
 72. Smee DF, Barnard DL. 2013. Methods for evaluation of antiviral efficacy against influenza virus infections in animal models. *Methods Mol Biol* 1030:407-25.
 73. Oh DY, Hurt AC. 2016. Using the Ferret as an Animal Model for Investigating Influenza Antiviral Effectiveness. *Frontiers in Microbiology* 7.
 74. Barré-Sinoussi F, Montagutelli X. 2015. Animal models are essential to biological research: issues and perspectives. *Future science OA* 1:FSO63-FSO63.

75. Shanks N, Greek R, Greek J. 2009. Are animal models predictive for humans? *Philosophy, ethics, and humanities in medicine : PEHM* 4:2-2.
76. U.S Food and Drug Administration. 2018. The Drug Development Process: Step 3. <https://www.fda.gov/patients/drug-development-process/step-3-clinical-research>. Accessed
77. Umscheid CA, Margolis DJ, Grossman CE. 2011. Key concepts of clinical trials: a narrative review. *Postgraduate medicine* 123:194-204.
78. Ursu O, Holmes J, Bologna CG, Yang JJ, Mathias SL, Stathias V, Nguyen DT, Schurer S, Oprea T. 2019. DrugCentral 2018: an update. *Nucleic Acids Res* 47:D963-D970.
79. Kim S, Chen J, Cheng T, Gindulyte A, He J, He S, Li Q, Shoemaker BA, Thiessen PA, Yu B, Zaslavsky L, Zhang J, Bolton EE. 2019. PubChem 2019 update: improved access to chemical data. *Nucleic Acids Res* 47:D1102-D1109.
80. Wishart DS, Feunang YD, Guo AC, Lo EJ, Marcu A, Grant JR, Sajed T, Johnson D, Li C, Sayeeda Z, Assempour N, Iynkkaran I, Liu Y, Maciejewski A, Gale N, Wilson A, Chin L, Cummings R, Le D, Pon A, Knox C, Wilson M. 2018. DrugBank 5.0: a major update to the DrugBank database for 2018. *Nucleic Acids Res* 46:D1074-D1082.
81. Pickett BE, Greer DS, Zhang Y, Stewart L, Zhou L, Sun G, Gu Z, Kumar S, Zaremba S, Larsen CN, Jen W, Klem EB, Scheuermann RH. 2012. Virus pathogen database and analysis resource (ViPR): a comprehensive bioinformatics database and analysis resource for the coronavirus research community. *Viruses* 4:3209-26.
82. Mazzon M, Ortega-Prieto AM, Imrie D, Luft C, Hess L, Czieso S, Grove J, Skelton JK, Farleigh L, Bugert JJ, Wright E, Temperton N, Angell R, Oxenford S, Jacobs M, Ketteler R, Dorner M, Marsh M. 2019. Identification of Broad-Spectrum Antiviral Compounds by Targeting Viral Entry. *Viruses* 11.
83. Hulseberg CE, Feneant L, Szymanska-de Wijs KM, Kessler NP, Nelson EA, Shoemaker CJ, Schmaljohn CS, Polyak SJ, White JM. 2019. Arbidol and Other Low-Molecular-Weight Drugs That Inhibit Lassa and Ebola Viruses. *J Virol* 93.
84. Kao JC, HuangFu WC, Tsai TT, Ho MR, Jhan MK, Shen TJ, Tseng PC, Wang YT, Lin CF. 2018. The antiparasitic drug niclosamide inhibits dengue virus infection by interfering with endosomal acidification independent of mTOR. *PLoS Negl Trop Dis* 12:e0006715.
85. Cairns DM, Boorgu D, Levin M, Kaplan DL. 2018. Niclosamide rescues microcephaly in a humanized in vivo model of Zika infection using human induced neural stem cells. *Biol Open* 7.
86. Huang L, Yang M, Yuan Y, Li X, Kuang E. 2017. Niclosamide inhibits lytic replication of Epstein-Barr virus by disrupting mTOR activation. *Antiviral Res* 138:68-78.
87. Wang YM, Lu JW, Lin CC, Chin YF, Wu TY, Lin LI, Lai ZZ, Kuo SC, Ho YJ. 2016. Antiviral activities of niclosamide and nitazoxanide against chikungunya virus entry and transmission. *Antiviral Res* 135:81-90.
88. Fang J, Sun L, Peng G, Xu J, Zhou R, Cao S, Chen H, Song Y. 2013. Identification of three antiviral inhibitors against Japanese encephalitis virus from library of pharmacologically active compounds 1280. *PLoS One* 8:e78425.

89. Wu CJ, Jan JT, Chen CM, Hsieh HP, Hwang DR, Liu HW, Liu CY, Huang HW, Chen SC, Hong CF, Lin RK, Chao YS, Hsu JT. 2004. Inhibition of severe acute respiratory syndrome coronavirus replication by niclosamide. *Antimicrob Agents Chemother* 48:2693-6.
90. Kuivanen S, Bessalov MM, Nandania J, Ianevski A, Velagapudi V, De Brabander JK, Kainov DE, Vapalahti O. 2017. Obatoclax, saliphenylhalamide and gemcitabine inhibit Zika virus infection in vitro and differentially affect cellular signaling, transcription and metabolism. *Antiviral Res* 139:117-128.
91. Kakkola L, Denisova OV, Tynell J, Viiliainen J, Ysenbaert T, Matos RC, Nagaraj A, Ohman T, Kuivanen S, Paavilainen H, Feng L, Yadav B, Julkunen I, Vapalahti O, Hukkanen V, Stenman J, Aittokallio T, Verschuren EW, Ojala PM, Nyman T, Saelens X, Dzek K, Kainov DE. 2013. Anticancer compound ABT-263 accelerates apoptosis in virus-infected cells and imbalances cytokine production and lowers survival rates of infected mice. *Cell Death Dis* 4:e742.
92. Shim JM, Kim J, Tenson T, Min JY, Kainov DE. 2017. Influenza Virus Infection, Interferon Response, Viral Counter-Response, and Apoptosis. *Viruses* 9.
93. Sheahan TP, Sims AC, Graham RL, Menachery VD, Gralinski LE, Case JB, Leist SR, Pyrc K, Feng JY, Trantcheva I, Bannister R, Park Y, Babusis D, Clarke MO, Mackman RL, Spahn JE, Palmiotti CA, Siegel D, Ray AS, Cihlar T, Jordan R, Denison MR, Baric RS. 2017. Broad-spectrum antiviral GS-5734 inhibits both epidemic and zoonotic coronaviruses. *Sci Transl Med* 9.
94. Patil SA, Patil V, Patil R, Beaman K, Patil SA. 2017. Identification of Novel 5,6-Dimethoxyindan-1-one Derivatives as Antiviral Agents. *Med Chem* 13:787-795.
95. Muller KH, Kainov DE, El Bakkouri K, Saelens X, De Brabander JK, Kittel C, Samm E, Muller CP. 2011. The proton translocation domain of cellular vacuolar ATPase provides a target for the treatment of influenza A virus infections. *Br J Pharmacol* 164:344-57.
96. Cheng YS, Williamson PR, Zheng W. 2019. Improving therapy of severe infections through drug repurposing of synergistic combinations. *Curr Opin Pharmacol* 48:92-98.
97. Fu Y, Gaelings L, Soderholm S, Belanov S, Nandania J, Nyman TA, Matikainen S, Anders S, Velagapudi V, Kainov DE. 2016. JNJ872 inhibits influenza A virus replication without altering cellular antiviral responses. *Antiviral Res* 133:23-31.
98. Moreno S, Perno CF, Mallon PW, Behrens G, Corbeau P, Routy JP, Darcis G. 2019. Two-drug vs. three-drug combinations for HIV-1: Do we have enough data to make the switch? *HIV Med* 20 Suppl 4:2-12.
99. Isakov V, Paduta D, Viani RM, Enejosa JV, Pasechnikov V, Znoyko O, Ogurtsov P, Bogomolov PO, Maevskaya MV, Chen X, Shulman NS. 2018. Ombitasvir/paritaprevir/ritonavir+dasabuvir+ribavirin for chronic hepatitis C virus genotype 1b-infected cirrhotics (TURQUOISE-IV). *Eur J Gastroenterol Hepatol* 30:1073-1076.
100. De Clercq E. 2019. Fifty Years in Search of Selective Antiviral Drugs. *J Med Chem* 62:7322-7339.

101. Foucquier J, Guedj M. 2015. Analysis of drug combinations: current methodological landscape. *Pharmacol Res Perspect* 3:e00149.
102. Schor S, Einav S. 2018. Combating Intracellular Pathogens with Repurposed Host-Targeted Drugs. *ACS Infect Dis* 4:88-92.
103. Mandell LA, Wunderink RG, Anzueto A, Bartlett JG, Campbell GD, Dean NC, Dowell SF, File TM, Jr., Musher DM, Niederman MS, Torres A, Whitney CG, Infectious Diseases Society of A, American Thoracic S. 2007. Infectious Diseases Society of America/American Thoracic Society consensus guidelines on the management of community-acquired pneumonia in adults. *Clin Infect Dis* 44 Suppl 2:S27-72.
104. Montoya MC, Krysan DJ. 2018. Repurposing Estrogen Receptor Antagonists for the Treatment of Infectious Disease. *MBio* 9.