

Favipiravir and Coronavirus

Journal of Bioscience & Biomedical Engineering

Short Communication

Marcos Aurélio Gomes da Silva

Federal University of Juiz de Fora, Brazil

*Correspondence author

Marcos Aurélio Gomes da Silva
Federal University of Juiz de Fora
Brazil

Submitted : 07 April 2020 ; Published : 17 April 2020

Flu is a disease of viral etiology that occurs as annual epidemics in the winter and winter months. pandemic only occasionally. Although it is a mild and self-limited disease, when it affects the population elderly or people with chronic underlying diseases, can present high morbidity and even mortality¹ The best strategy to deal with epidemics annual flu is prevention by applying the adequate influenza vaccine. However, its efficiency and effectiveness is not always what is desired, so that in serious presentations we must use some antiviral treatment.

Favipiravir is an antiviral drug that selectively inhibits the RNA-dependent RNA polymerase of influenza virus. It is phosphoribosylated by cellular enzymes to its active form, favipiravir-ribofuranosyl-5'-triphosphate (RTP). Favipiravir is active against a broad range of influenza viruses, including A (H1N1) pdm09, A (H5N1) and the recently emerged A (H7N9) avian virus. It also inhibits influenza strains resistant to current antiviral drugs, and shows a synergistic effect in combination with oseltamivir, thereby expanding influenza treatment options. Recently, T-705 has been shown to give 100% protection against aerosol Ebola virus E718 infection in immune-deficient mice as well as inhibiting Ebola virus infection in cell culture.

The mechanism of action of favipiravir is inhibition direct replication and transcription of the viral genome type RNA, by blocking RNA polymerase activity RNA-dependent present in influenza viruses and other viruses with RNA genome, including Ebola 10-12 virus. Thus his selectivity index (the ratio of 50% of the concentration cell inhibitory (CC50) / 50% of the inhibitory concentration against influenza (IC50), it is higher than 6,000 in relation to ribavirin, another antiviral inhibitor of RNA polymerases.

The subtypes and strains of known influenza viruses both A, B as C, I include those with demonstrated genetic resistance against neuraminidase inhibitors.

Favipiravir can be administered orally in prodrug form (ribofuranose), being rapidly adsorbed at the intestinal level and converted into the initial active form by cellular nucleosidases⁵⁻⁷. The optimal dose has not yet been definitively established but

could be 1200-1400 mg / first day and 400mg / day for 5-7 days has also proven to be effective in combating coronavirus, patients tested negative for the virus four days after taking the drug. Those infected who did not receive treatment with favipiravir took about 11 days to achieve the same result. In addition, 91% of patients treated with favipiravir showed significant improvements in lung conditions, compared to 62% in those who did not receive the same medication. Japanese doctors are currently using Avigan for clinical studies in patients diagnosed with Covid-19 who have mild or moderate symptoms of the disease. The expectation is that the drug will prevent the virus from multiplying in the body.

We hypothesized that favipiravir would be non-inferior to arbidol in terms of efficacy for moderate symptoms, and improves outcomes clinical recovery of fever, cough, and breathing difficulties compared with antiviral efficacy of arbidol. We therefore assessed the clinical efficacy and safety of favipiravir versus arbidol as treatment for COVID-19.

In moderate COVID-19 patients untreated with antiviral previously, favipiravir can be considered as a preferred treatment because of the higher clinical recovery rate of day 7 and more effectively reduced incidence of fever, cough except manageable antiviral-associated adverse effects.

The time it takes for a patient to be negative in the coronavirus test has been 4 days among those who took favipiravir and 11 days among those who did not. Likewise, improvements in radiographs of the lungs have been observed in 91% of patients treated with favipiravir, compared to 62% of patients in the control group.

Favipiravir is a prodrug that is metabolized in its active form, favipiravir-ribofuranosyl-5'-triphosphate (favipiravir-RTP), available in oral and intravenous formulations.

A mutation in the influenza A virus PB1 gene that confers resistance to favipiravir. The fitness cost of this mutation, resulting from a decrease in polymerase activity, is compensated for by a mutation in the PA gene that can restore polymerase activity and viral growth, while maintaining favipiravir resistance. We use in vitro analyzes and structural modeling to link favipiravir resistance to

a molecular change in the RdRP active site that affects nucleotide incorporation. Our findings suggest that a universal mechanism for favipiravir resistance exists.

This hypermutability favors the rapid adaptation of influenza viruses to environmental and environmental changes, host, as well as immune pressure^{14,15}. Nevertheless in certain occasions this phenomenon can take the virus to the extinction because the mutations determine a decrease in fitness (adaptability) and loss of viability. This concept is the basis of some of the new antivirals, that is to say, they look for drugs that joining the genome and by blocking polymerase determine an increase in the accumulation of mutations and collapse viral progeny.

Together, these observations are important for the use of synthesis in mammalian cells and is not toxic to them.

Among them, influenza viruses have been very sensitive to this new antiviral, including strains with genetic resistance to neuraminidase inhibitors (oseltamivir). Its mechanism of action consists of blocking viral replication and inducing lethal mutagenesis that determines the loss of the infectious activity of influenza viruses. Its activity is intense in the respiratory tract, reducing viral loads to non-infectious levels favipiravir as an antiviral drug against influenza virus infections in the present and future [1-6].

References

1. Perlman S, Netland J (2009) Coronaviruses post-SARS: update on replication and pathogenesis. *Nat. Rev. Microbiol* 7(6): 439-450.
2. Chan JF, To KK, Tse H, Jin DY, Yuen KY (2013) Interspecies transmission and emergence of novel viruses: lessons from bats and birds. *Trends Microbiol* 21(10): 544-555.
3. Lei J, Kusov Y, Hilgenfeld R (2018) Nsp3 of coronaviruses: Structures and functions of a large multi-domain protein. *Antiviral Res* 149: 58-74.
4. Song W, Gui M, Wang X, Xiang Y (2018) Cryo-EM structure of the SARS coronavirus spike glycoprotein in complex with its host cell receptor ACE2. *PLoS Pathog* 14(8): e1007236.
5. Gordon CJ, Tchesnokov EP, Feng JY, Porter DP, Gotte M (2020) The antiviral compound remdesivir potently inhibits RNA-dependent RNA polymerase from Middle East respiratory syndrome coronavirus. *J. Biol. Chem* 295(15): 4773-4779.
6. Monto AS (2003) The role of antivirals in the control of influenza. *Vaccine* 21: 1796-1800.

Copyright: ©2020 Marcos Aurélio Gomes da Silva. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.