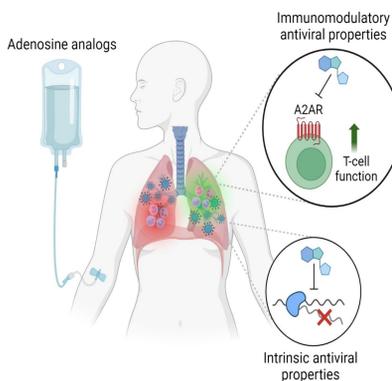


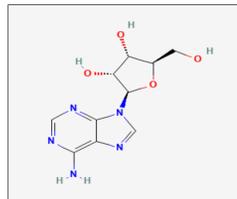
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Introduction



SARS-CoV-2 is a contagious virus that causes COVID-19, a disease that has affected millions of people worldwide and claimed over six million lives. The pandemic has unveiled an urgent need for new antivirals to control emerging infectious diseases and potential future pandemics.

Classic antivirals are currently designed to directly interfere with pathogens. Adenosine analogs are antiviral drugs which resemble the structure of adenosine (1), a metabolite that derives from the breakdown of ATP and can suppress the immune response through activation of the Adenosine A2A Receptor (A2AR) in immune cells.



Adenosine



Forodesine

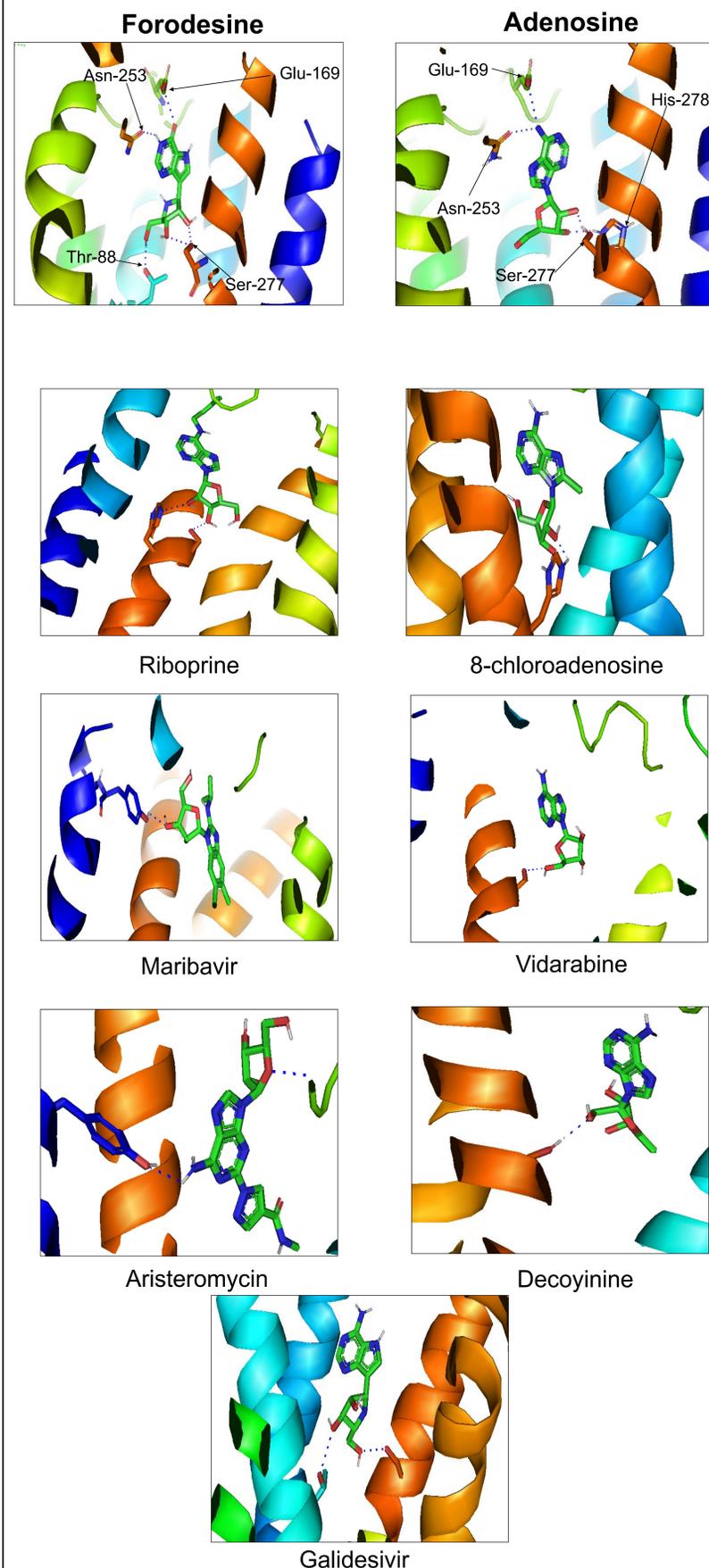
Adenosine is overproduced in some infectious diseases, like COVID-19, where it restricts protective antiviral immune responses. Previous work from our lab showed that adenosine-mediated activation of A2AR causes suppression of CD8+ T-cells effector functions in COVID-19 mouse models (2,3). Importantly, our group showed that adenosine analogs could restore antiviral T-cell responses in COVID-19 models, through blocking the A2AR (2,3). Therefore, we propose that adenosine analogs that interact with A2AR may be endowed with dual – immunomodulatory and intrinsic – antiviral functions. In this study, adenosine analogs were screened using molecular docking to predict whether they would bind to A2AR and CD8+ T-cell proliferation was used as a functional assay to test the action of adenosine analogs on A2AR.

Screening of adenosine analogs

Adenosine Analog	ΔG	Interacting residues
Forodesine	-7.3	Glu-169, Asn-253, Ser-277, Thr-88
Riboprine	-8.2	His-278, Ser-277
8-chloroadenosine	-7.2	His-278
Maribavir	-7.2	Tyr-9
Vidarabine	-7.0	Ser-277
Aristermocin	-7.0	Glu-169, Asn-253, Ser-277
Decoyinine	-6.4	Ser-277
Galidesivir	-7.4	Ser-277, Thr-88
Adenosine	-7.3	Glu-169, Asn-253, Ser-277, His-278

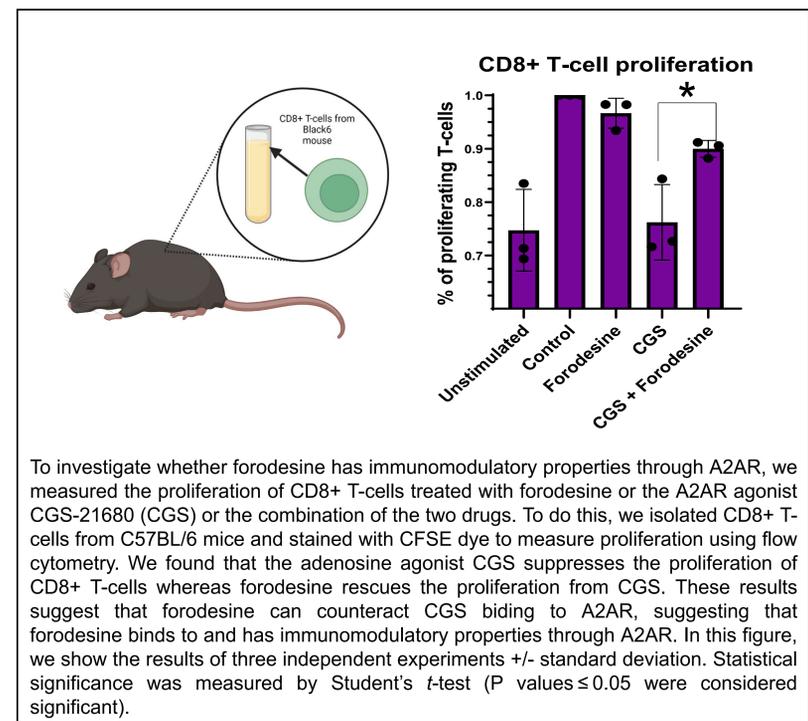
We selected eight adenosine analogs to screen for binding to A2AR. These analogs were known or predicted to inhibit the replication of SARS-CoV-2 and were used in clinical settings (1). As shown in the table, using molecular docking - AutodockVina software, we calculated the affinity for binding to A2AR (ΔG) and the residues of interaction between A2AR and the analogs. Forodesine, Riboprine and 8-chloroadenosine showed the lowest ΔG and similar interacting residues as adenosine, suggesting they likely bind to A2AR.

Molecular Docking of the adenosine analogs



We displayed molecular docking results using Pymol and analyzed the interaction between the analogs and A2AR compared with adenosine. Among all analyzed analogs, forodesine was ultimately selected because it had the most residues in common with adenosine-A2AR complex and a low ΔG , thus is predicted to bind A2AR.

Forodesine shows immunomodulatory properties



Conclusion

We found that forodesine treatment restored T-cell proliferation from the A2AR agonist CGS in CD8+ T-cells, suggesting that forodesine binds and blocks A2AR. Our results indicate that adenosine analogs, like forodesine, may be endowed with immunomodulatory properties through A2AR, in addition to their intrinsic antiviral properties. In conclusion, adenosine analogs, like forodesine, could be used as novel dual – immunomodulatory and intrinsic – antiviral drugs. These compounds could represent game-changing therapies, not only to control COVID-19, but also other emerging viral diseases and future pandemics.

Future work: Future work will focus on: characterizing forodesine using *in vitro* and *in vivo* functional assays, including testing the drug *in vivo* in non-infected and SARS-CoV-2 infected mice; testing the other adenosine analogs *in vitro* and *in vivo*, like we did for forodesine.

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