

Viscosine as a Potent and Safe Antipyretic Agent Evaluated by Yeast-Induced Pyrexia Model and Molecular Docking Studies

Akhtar Muhammad,^{*,†} Behramand Khan,[†] Zafar Iqbal,[‡] Amir Zada Khan,[‡] Inamullah Khan,[‡] Kashif Khan,[§] Muhammad Alamzeb,^{||} Nasir Ahmad,[†] Khalid Khan,[†] Syed Lal Badshah,[†] Asad Ullah,[†] Sayyar Muhammad,[†] Muhammad Tariq Jan,[†] Said Nadeem,[⊥] and Nurul Kabir[#]

[†]Department of Chemistry, Islamia College University, Peshawar, KPK 25120, Pakistan

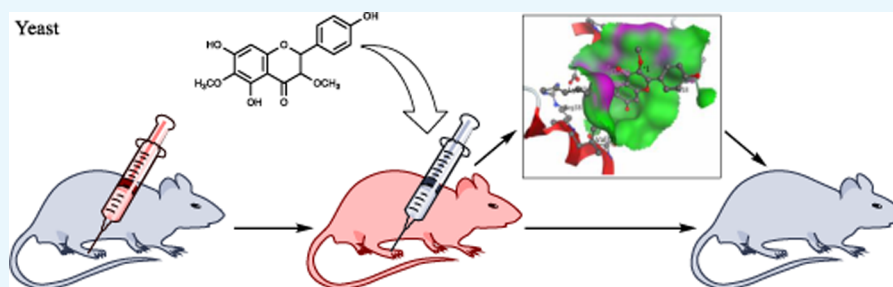
[‡]Department of Pharmacy, University of Peshawar, Peshawar 25120, Pakistan

[§]Department of Chemistry, Sarhad University of Science & Information Technology, Peshawar 25000, Pakistan

^{||}Faculty of Sciences, Department of Chemistry, University of Kotli, Kotli 11100, Azad Jammu and Kashmir, Pakistan

[⊥]Kosk Vocational School of Food Technology, Aydin Adnan Menderes University, Efeler 09010 Aydin, Turkey

[#]Institute of Biological Sciences, Faculty of Science, University of Malaya, 50603 Kuala Lumpur, Malaysia



ABSTRACT: The antipyretic potential of viscosine, a natural product isolated from the medicinal plant *Dodonaea viscosa*, was investigated using yeast-induced pyrexia rat model, and its structure–activity relationship was investigated through molecular docking analyses with the target enzymes cyclooxygenase-1 (COX-1), cyclooxygenase-2 (COX-2), and microsomal prostaglandin E synthase-1 (mPGES-1). The in vivo antipyretic experiments showed a progressive dose-dependent reduction in body temperatures of the hyperthermic test animals when injected with viscosine. Comparison of docking analyses with target enzymes showed strongest bonding interactions (binding energy -17.34 kcal/mol) of viscosine with the active-site pocket of mPGES-1. These findings suggest that viscosine shows antipyretic properties by reducing the concentration of prostaglandin E₂ in brain through its mPGES-1 inhibitory action and make it a potential lead compound for developing effective and safer antipyretic drugs for treating fever and related pathological conditions.

INTRODUCTION

Pyrexia (fever) is associated with many diseases¹ and is an expression of the body's complex immunophysiologic response to infectious or inflammatory stimuli that trigger a cascade of biochemical reactions ultimately producing various endogenous pyrogens.^{2,3} Of these pyrogens, prostaglandin E₂ (PGE₂) is the principal fever mediator in mammals.^{4,5} Although it is body's natural immune response and host-defense mechanism, pyrexia results in general body discomfort and adversely affects the normal functions of various body organs.⁶ Excessive rise of body temperature is controlled by endogenous antipyretics liberated within the brain during fever.⁷ However, the use of antipyretic drugs is many times indispensable.

The commonly used antipyretic and anti-inflammatory agents (i.e., salicylates and nonsteroidal anti-inflammatory drugs (NSAIDs)) stop or lower the formation of the principal fever mediator PEG₂ by inhibiting the cyclooxygenase enzymes cyclooxygenase-1 (COX-1) and cyclooxygenase-2 (COX-2) selectively or nonselectively.⁸ However, COX inhibition can

lead to various health complications. Nonselective COX inhibitors are harmful to the gastrointestinal tract causing gastric distress,⁹ ulceration, and other bleeding disorders.¹⁰ Their prolonged use in children and adolescents may lead to Reye's syndrome (liver damage).¹¹ Prolonged use of selective COX-2 inhibitors, particularly NSAIDs, is linked with higher risks of cardiac attack^{12,13} and stroke, and patients with kidney, heart, or liver conditions are at risk of developing kidney damage.¹⁴ The long-term use of cyclooxygenase inhibitors as antipyretic agents is associated with serious side effects. An alternative antipyretic strategy is the inhibition of prostaglandin E₂ synthases (PGES),^{15,16} especially the microsomal prostaglandin E synthase-1 (mPGES-1). The mPGES-1 catalyzes transformation of PGH₂ to PGE₂ in the last biosynthetic step and is a promisingly safe target^{17,18} of

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antipyretic agents because its inhibition is free of the side effects associated with the COX inhibition.^{19,20} Numerous efforts are being made for the development of next-generation anti-inflammatory and antipyretic drugs based on mPGES-1 inhibitors.^{21–23} Usually for optimized search of lead compounds, structure-based strategy employing ligand–receptor molecular docking²⁴ is adopted for predicting interaction affinities and binding modes of the drug molecules with a particular target receptor.²⁵

We have previously isolated viscosine (Figure 1) from the medicinal plant *Dodonaea viscosa*, which is reported to possess

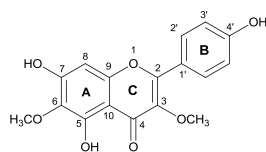


Figure 1. Molecular structure of viscosine.

antinociceptive,²⁶ neuropharmacological,²⁷ lipoxygenase inhibitory,²⁸ hepatoprotective,²⁹ anticholinesterase, and antioxidant properties.³⁰ Since compounds with analgesic and/or anti-inflammatory properties often possess antipyretic properties, we investigated the *in vivo* antipyretic potential of viscosine and its possible mode of action using an *in silico* approach. In this paper, we present the antipyretic potential of viscosine against a yeast-induced pyrexia model in rats. We also report the *in silico* findings showing viscosine as an efficient mPGES-1 inhibitor, suggesting it as a lead compound for developing effective and safe antipyretic drugs.

RESULTS

Yeast induces a biochemical cascade that ultimately leads to the release of biochemical mediators, especially prostaglandin E₂ and interleukins, which in turn upregulate the thermoregulatory center, leading to hyperthermia and pyrexia.³¹ Any compound capable of inhibiting the cascade of pyrexia-inducing mediators will manifest antipyretic activity³² such as paracetamol and other NSAIDs. The antipyretic effects of viscosine on pyrexia are shown in Table 1. Yeast-induced pyrexia model was used to investigate the therapeutic effect of viscosine in relieving fever, which is often associated with inflammation.³³ Upon treatment with viscosine, body temperatures of the test animals were observed to decrease progressively in a dose-dependent manner. Viscosine showed significant antipyretic potential ($P < 0.01–0.001$). In addition to the currently explored antipyretic potential, our previous

studies²⁶ also showed significant analgesic potential of viscosine compared to control, as depicted in its analgesic profile (Figure 2). To investigate the possible mechanism

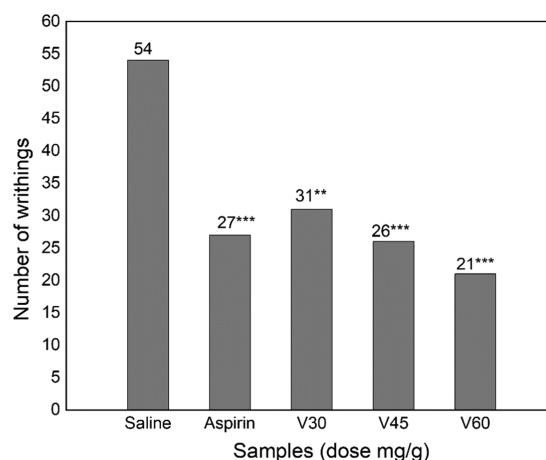


Figure 2. Analgesic profile of viscosine on acetic acid-induced writhing model: Saline, 10 mL/kg; aspirin, 150 mg/kg; V30, viscosine 30 mg/kg; V45, viscosine 45 mg/kg; V60, viscosine 60 mg/kg.

behind its antipyretic action, viscosine was docked with the possible target enzymes COX-1, COX-2, and mPGES-1, and their structure–activity relationships were investigated. The docking results are presented in Figures 3–5.

DISCUSSION

In antipyretic studies, both viscosine (V60) and aspirin showed significant ($P < 0.01$) antipyretic activity after 2 h of their injection. The antipyretic potential of viscosine increased with passage of time. Both the low dosages of viscosine (V30 and V60) showed highly significant antipyretic potential at the later stages comparable to the reference drug, which showed significant effect at much higher dosage.

Results from the analgesic activities showed that viscosine as well as aspirin significantly suppressed the writhing response in a dose-dependent manner. It is apparent, however, that viscosine is significantly effective at lower doses compared to aspirin. In conclusion, viscosine revealed promising antipyretic potential during *in vivo* analyses and deserved further exploration of the molecular mechanisms involved behind its significant analgesic and antipyretic effects.

Docking analysis with COX-1 showed (Figure 3) a total of four interactions: a single interaction between hydroxyl group at position 5 of ring A with residue Gly45, two interactions

Table 1. Antipyretic Action of Viscosine against Yeast-Induced Pyrexia in Mice^a

time (h)	dosage (mg/kg)			
	saline	V30	V60	aspirin (100)
	body temperature			
0	35.03 ± 0.26	34.77 ± 0.13	34.78 ± 0.12	35.02 ± 0.09
0.5	36.84 ± 0.19	37.11 ± 0.18	36.85 ± 0.14	36.07 ± 0.14
1	36.96 ± 0.12	36.83 ± 0.14	36.65 ± 0.17	36.59 ± 0.11
2	36.89 ± 0.09	36.54 ± 0.19	36.33 ± 0.09**	36.26 ± 0.29**
3	36.72 ± 0.11	36.04 ± 0.16**	36.12 ± 0.07**	35.81 ± 0.04***
4	36.59 ± 0.09	35.59 ± 0.12***	35.67 ± 0.09***	35.45 ± 0.02***
5	36.57 ± 0.06	35.26 ± 0.19***	35.28 ± 0.11***	35.24 ± 0.08***

^aData are expressed as mean ± standard error of the mean. Significant at ** $P < 0.01$, *** $P < 0.001$ compared to control.

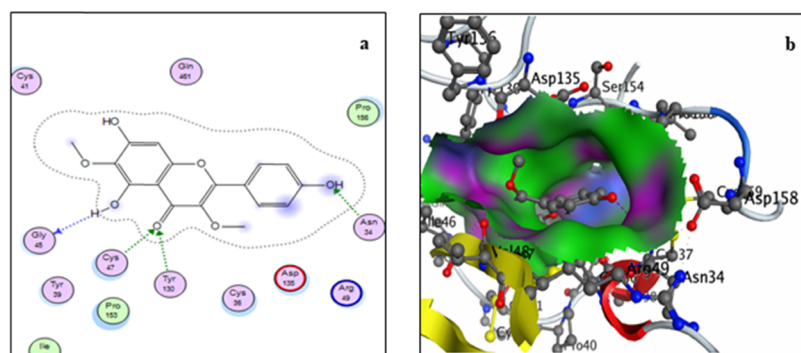


Figure 3. Two-dimensional (a) and three-dimensional (3D) (b) binding-site interaction models of COX-1 with viscosine: The blue highlight represents ligand exposure; H-bond lengths, 2.77–3.0 Å.

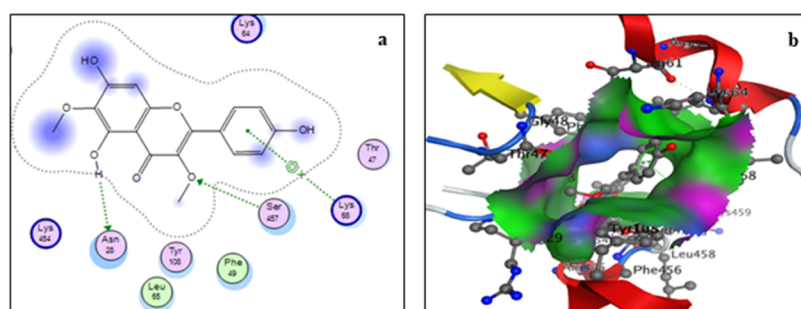


Figure 4. Two-dimensional (a) and three-dimensional (b) binding-site interaction models of COX-2 with viscosine: The blue highlight represents ligand exposure; H-bond lengths, 2.77–3.0 Å.

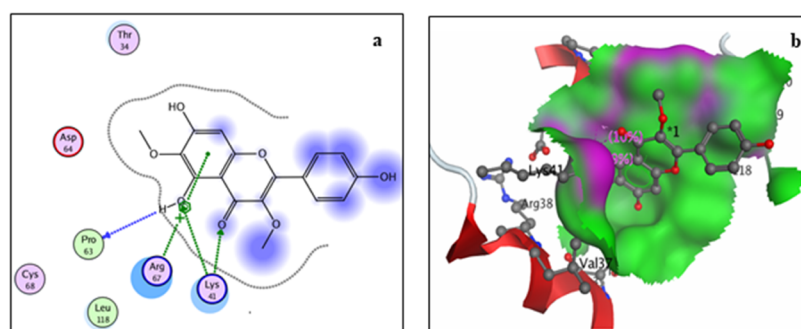


Figure 5. Two-dimensional (a) and three-dimensional (b) binding-site interaction models of mPGES-1 with viscosine: The blue highlight represents ligand exposure; H-bond lengths, 2.77–3.0 Å.

involving the carbonyl group at position 4 of the ring C with residues Cys47 and Tyr130, and a single interaction between the hydroxyl group at position 4' of the ring B with residue Asn34 of the receptor. Both the methoxy and aromatic rings showed no interactions with the receptor. All of the interactions are through hydrogen bonding and, thermodynamically, the inhibition is moderate with a binding energy of -13.34 kcal/mol.

Docking studies of viscosine with COX-2 also showed moderate ligand–receptor interactions (Figure 4). Overall, three interactions with an overall binding energy of -10.46 kcal/mol were observed, including an arene–cation interaction between ring B and residue Lys68 and two hydrogen-bonding interactions between hydroxyl moiety at position 5 with residue Asn28 and between 3-methoxy group and residue Ser457.

Docking with mPGES-1 receptor also showed four interactions (Figure 5). Of these, ring A showed two arene–

cation-type interactions with residues Lys41 and Arg67, while the 4-carbonyl and 5-hydroxyl groups of the ligand showed hydrogen-bonding interactions with Lys41 and Pro63, respectively. Residue Lys41 of the receptor showed a hydrogen-bonding interaction with carbonyl group of ring C and an arene–cation interaction with ring A of the ligand. It is evident from the data that viscosine interacts strongly with mPGES-1 with a predicted overall binding energy of -17.34 kcal/mol.

Based on the strongly supportive values of the binding energies obtained in our *in silico* investigations, it can be safely deduced that viscosine possesses favorable structural features to effectively interact and inhibit mPGES-1. This suggests viscosine to be a safe antipyretic agent and should be considered as a lead compound with promising potential for developing safe antipyretic drugs.

CONCLUSIONS

The in vivo studies performed on pyrexia-induced rats showed that viscosine possesses strong antipyretic actions. Low dosages of viscosine showed high antipyretic potential at later phases compared to aspirin. Viscosine also significantly suppressed the writhing response in a dose-dependent manner. Molecular docking studies suggest that viscosine showed stronger interactions with microsomal prostaglandin E synthase-1 than the cyclooxygenases and support the hypothesis that febrile response is reduced through mPGES-1 inhibition. Comparison of the binding energies of viscosine to that of other reported compounds³² suggests the need for further studies to confirm that viscosine strongly inhibits mPGES-1 prior to its consideration for developing safe antipyretic drugs.

EXPERIMENTAL SECTION

Materials and Methods. Experiments were conducted with adult wistar rats (weighing 180–260 g) and Swiss albino mice (weighing 18–25 g). All of the animals were housed individually at 24 ± 1 °C with abundant access to water and food until the experiment day, when only water was made available. Animal experiments performed in the manuscript were conducted in compliance with institutional guidelines,³³ and permission for the experiments was previously granted by the local ethics committee for research on laboratory animals.

Yeast-Induced Pyrexia Model. The yeast-induced pyrexia model, as reported by Al-Ghamdi,³⁴ was selected for evaluating the antipyretic properties of viscosine. Pyrexia was induced into the test animals by subcutaneously injecting a 15% aqueous yeast solution at a dosage of 10 mL/kg. Rectal temperatures of test animals were recorded 24 h before and after injecting yeast through a digital thermometer (Hartmann, Germany). The rats that did not show a minimum elevation of 0.5 °C in rectal temperatures after the yeast administration were excluded from experiment. The selected animals were divided into five groups (each group comprising six animals) and were separately treated with normal saline water, viscosine, and the reference aspirin. Rectal temperature of each animal was subsequently recorded for up to 5 h initially at 30 min and afterward at 1 h time intervals.

Molecular Docking Analysis. Viscosine was taken as a ligand for molecular docking. The ligand file was designed and optimized using the ChemBioDraw software (v. 14) and then converted to a 3D format through the “translate molecular files” tools available in the SYBYL-X 2.0 platform. Crystallographic structures of the targets COX-1 (PDB ID: 3KK6), COX-2 (PDB ID: 3LN1), and mPGES-1 (PDB ID: 3DWW) were taken from the archive of Protein Data Bank (RSCB PDB). The ligand was docked with targets using the MOE software.

ASSOCIATED CONTENT

Accession Codes

Chemical Purity: Viscosine was purified through reverse-phase high-performance liquid chromatography, recrystallized, and characterized through single X-ray diffraction and NMR spectroscopy. The crystallographic data are available at the Cambridge Crystallographic Data Centre (CCDC 1041249).

AUTHOR INFORMATION

Corresponding Author

*E-mail: dr.akhtarmuhammad@icp.edu.pk. Phone: +92 3009 007 393.

ORCID

Akhtar Muhammad: 0000-0002-5296-216X

Notes

The authors declare no competing financial interest.

ABBREVIATIONS USED

COX-1, cyclooxygenase 1; COX-2, cyclooxygenase 2; mPGES-1, microsomal prostaglandin E synthase-1; NSAIDs, nonsteroidal anti-inflammatory drugs; SAR, structure–activity relationship; V30, viscosine 30 mg/kg; V45, viscosine 45 mg/kg; V60, viscosine 60 mg/kg

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