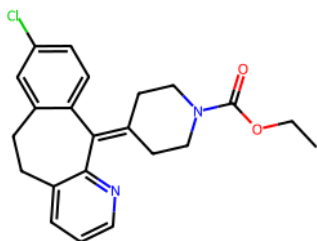


Astinova AI Platform — ADMET Profiling Report

Subject: Loratadine (public reference compound) · in silico ADMET across absorption, distribution, metabolism, excretion, toxicity

Model-based predictions, expert-reviewed. Illustrative sample on a public molecule.



MW (Da)	382.9
logP	4.89
TPSA (Å²)	42.4
HBD / HBA	0 / 3
QED	0.70
Lipinski violations	4

SMILES: CCOC(=O)N1CCC(=C2c3ccc(Cl)cc3CCc3ccnc32)CC1

Absorption

Endpoint	Value	Interpretation
Human intestinal absorption (prob)	1.000	High
Caco-2 permeability (log cm/s)	-4.59	Permeable
Oral bioavailability (prob)	0.685	Favourable
Aqueous solubility (log mol/L)	-4.85	Low solubility

Distribution

Endpoint	Value	Interpretation
BBB permeability (prob)	0.955	Predicted BBB-permeable
P-gp substrate (prob)	0.700	Low efflux
Plasma protein binding (%)	96.9	Highly bound

Metabolism (CYP)

Endpoint	Value	Interpretation
CYP3A4 inhibitor (prob)	0.663	Lower risk
CYP2C9 inhibitor (prob)	0.861	DDI risk
CYP2D6 inhibitor (prob)	0.355	Lower risk

Excretion & toxicity

Endpoint	Value	Interpretation
Half-life (Obach, h)	47.6	Model estimate
hERG inhibition (prob)	0.684	Moderate
DILI (prob)	0.747	Hepatotoxicity flag
AMES mutagenicity (prob)	0.080	Low
Clinical toxicity (prob)	0.089	Low
Carcinogenicity (prob)	0.089	Low
LD50 (log mol/kg)	2.90	Moderate acute tox.

Predictions are in silico estimates generated by the Astinova AI Platform and reviewed by a chemist. Probabilities are model outputs, not experimental values; hERG, DILI, and CYP flags should be confirmed with the corresponding in vitro assays before decision-making.