

Artificial Intelligence Chemical Report

Cannflavin A

PRESENTED TO

Public

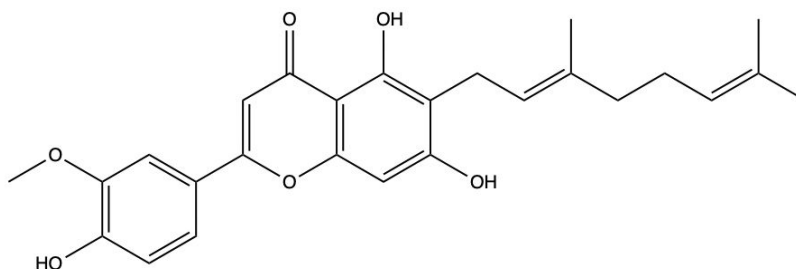
PRESENTED BY

Global Botanical Artificial Intelligence
System (GBAIS)

TABLE OF CONTENTS

Chemical Profile	3
Mass Spectrometry	4
Solubility Solvent Prediction	6
Solvate Prediction	7
Polymorphism Prediction (Coming Soon)	8
Green Retrosynthesis	9
Molecular Dynamic Analysis: 2D NMR Spectra & Dose Response Curves	11
Chemical Disease Pathway	12
Formulation Profile	13
Formulation Risks	14
Excipient Risks	16
Product Market	17
Customer Journals (Coming Soon)	18
References	19

Chemical Profile



Global-Chem Name: Cannflavin A

SMILES:

CC(=CCC/C(=C/CC1=C(C2=C(C=C1O)OC(=CC2=O)C3=CC(=C(C=C3)O)OC)O)/C)C

Molecular Weight: 436.504

Calculated ALogP: 5.82

Calculated LogD_{7.4}: 4.82

LogP (Crippen Method): 5.821

Calculated Acidic PKa: 6.39

Hydrogen Bond Acceptors: 6

Hydrogen Bond Donors: 3

Topological Polar Surface Area: 100.130

Drug-likeness QED Weight: 0.40

Compound Characterization: Digital Twin Mass Spectrometry

Spectra Type: Electrospray Ionization (ESI)

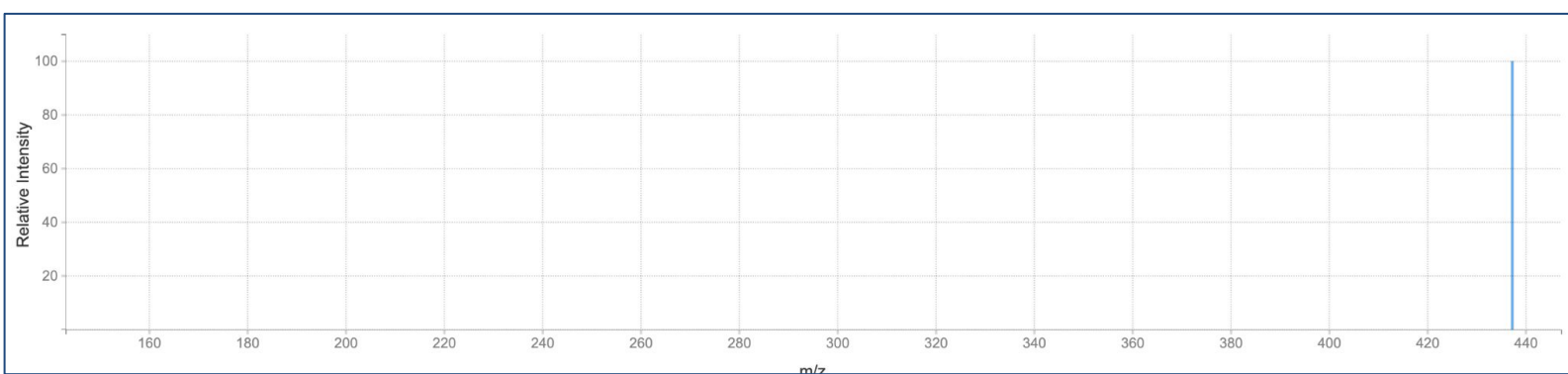
Ion Mode: Positive

Adduct Type: $[M+H]^+$

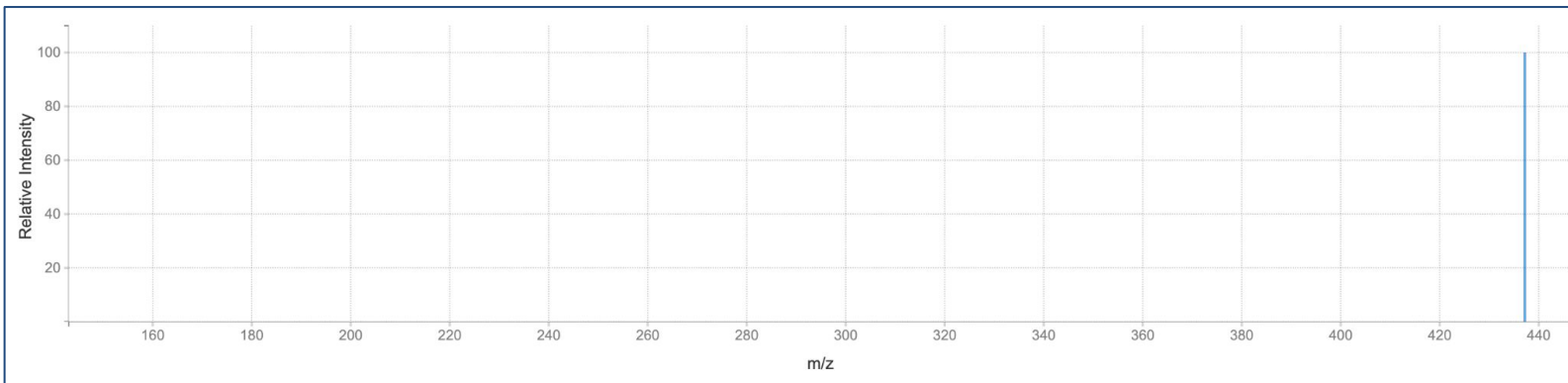
Digital Twin Instrument: Agilent 6510 Q-TOF

Blue lines are the parent mass fragment. **Red** lines are fragments.

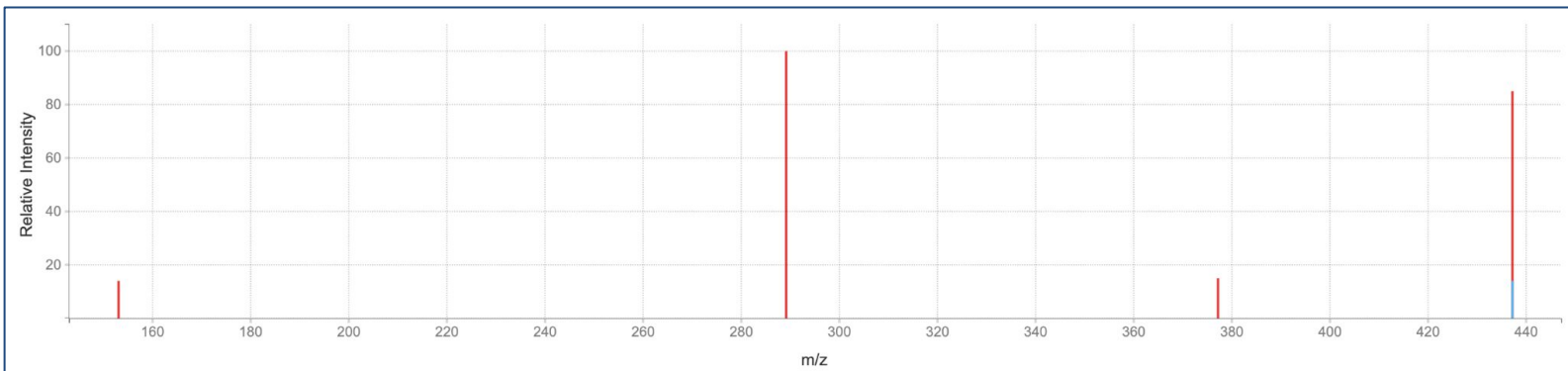
Predicted Low Energy MsMs Spectrum (10V), $[M+H]^+$



Predicted Medium Energy MsMs Spectrum (20V), $[M+H]^+$

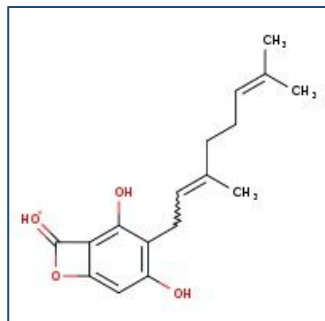


Predicted High Energy MsMs Spectrum (40V), $[M+H]^+$



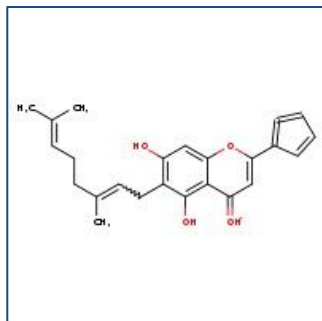
Compound Characterization: Mass Spectrometry (Major/Minor Fragments)

Major

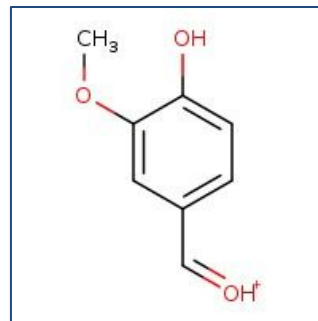


Mass: 289.143

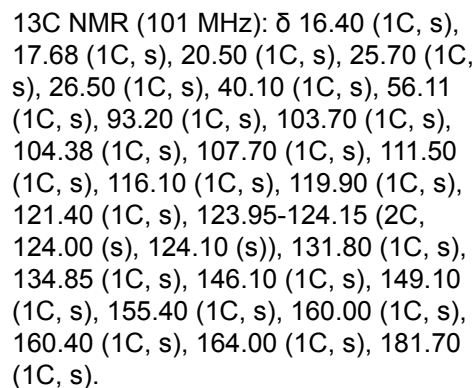
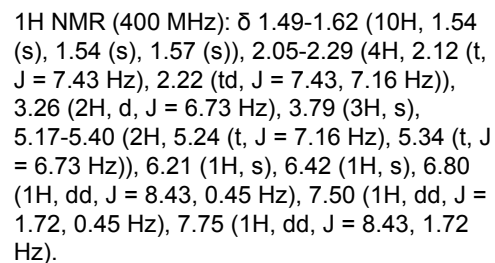
Minor



Mass: 377.174



Mass: 153.054



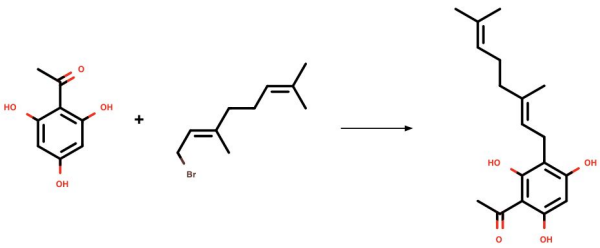
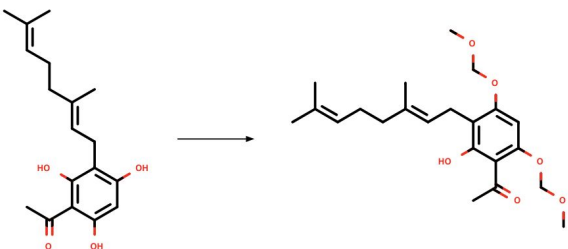
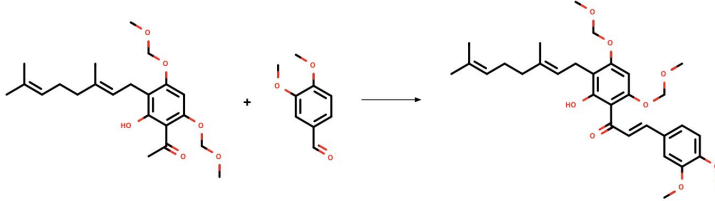
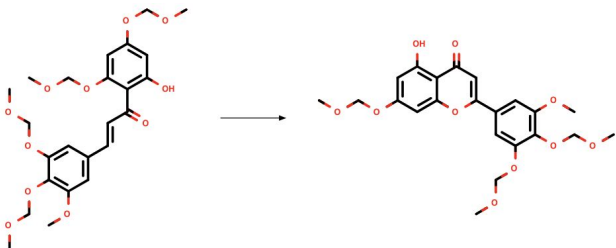
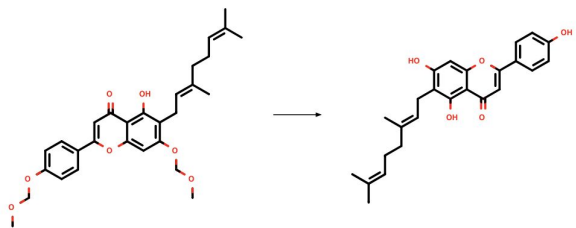
Solubility Solvent Prediction

Solvent	Temperature	LogS	AI Choice
Ethyl Alcohol	298	0.244	Yes
Methyl Alcohol	298	0.300	Yes
Dichloromethane	298	0.456	Yes
n-hexane	298	-4.538	No
Ethyl Acetate	298	0.631	Partially
Trichloromethane	298	0.058	Partially
Ethyl Ether	298	-0.769	No
H2O	298	-6.456	No
Dimethylformamide	298	2.210	Yes
DMSO	298	1.988	Yes

Solvate Prediction

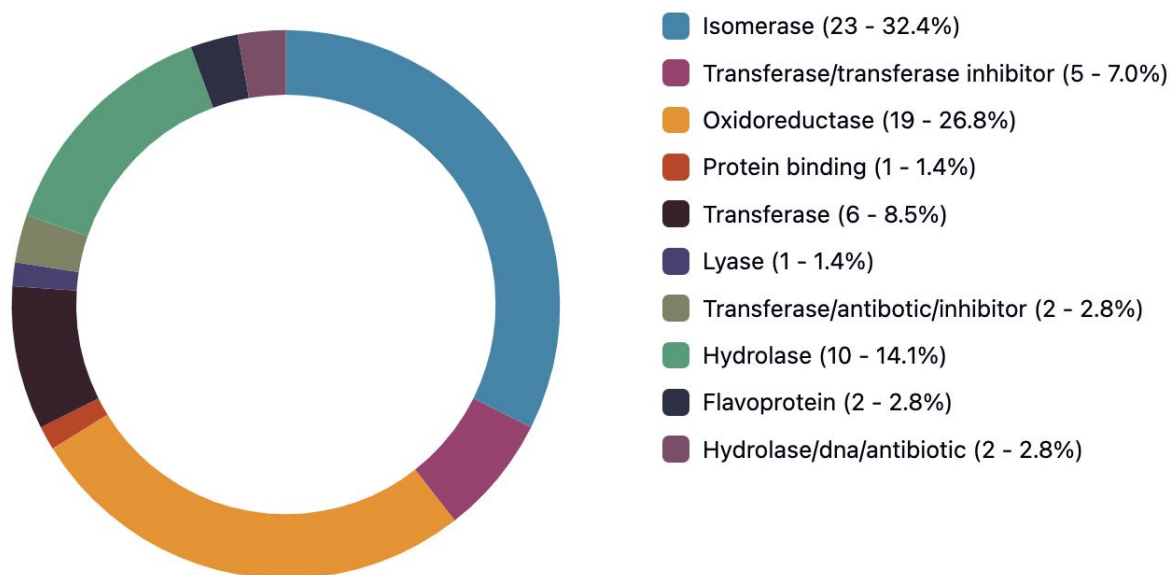
Chemical	Solvent	Probability	AI Choice
Cannaflavin A	Water	0.9647	Yes
Cannaflavin A	Ethyl Alcohol	0.7124	Yes
Cannaflavin A	Methyl alcohol	0.9616	Yes
Cannaflavin A	Dichloromethane	1.0000	Yes
Cannaflavin A	n-Hexane	0.0782	No
Cannaflavin A	Ethyl acetate	0.9530	Yes
Cannaflavin A	Trichloromethane	1.0000	Yes
Cannaflavin A	Ethyl ether	0.7905	Yes
Cannaflavin A	Dimethylformamide	0.9827	Yes
Cannaflavin A	DMSO	0.9851	Yes
Cannaflavin A	Oxolane	0.9743	Yes

Polymorphism Predictions (Coming Soon)

Step	Reference Reaction	DOI
1		Sun, Qiu, et al. "Synthesis of Natural-like Acylphloroglucinols with Anti-Proliferative, Anti-Oxidative and Tube-Formation Inhibitory Activity." European Journal of Medicinal Chemistry, vol. 85, Oct. 2014, pp. 621–28
2		Jung, Doo Hwan, et al. "New Synthetic Routes to Biologically Interesting Geranylated Flavanones and Geranylated Chalcones: First Total Synthesis of (±)-Prostratol F, Xanthoangelol, and (±)-Lespeol." Helvetica Chimica Acta, vol. 93, no. 4, Apr. 2010, pp. 635–47.
3		Jung, Doo Hwan, et al. "New Synthetic Routes to Biologically Interesting Geranylated Flavanones and Geranylated Chalcones: First Total Synthesis of (±)-Prostratol F, Xanthoangelol, and (±)-Lespeol." Helvetica Chimica Acta, vol. 93, no. 4, Apr. 2010, pp. 635–47.
4		Guz, Nathan R., and Frank R. Stermitz. "Synthesis and Structures of Regioisomeric Hydnocarpin-Type Flavonolignans." Journal of Natural Products, vol. 63, no. 8, Aug. 2000, pp. 1140–45.
5		Heterocycles; vol. 32; 3; (1991); p. 499 - 510

Molecular Dynamic Analysis: 2D NMR Spectra & Dose Response Curves

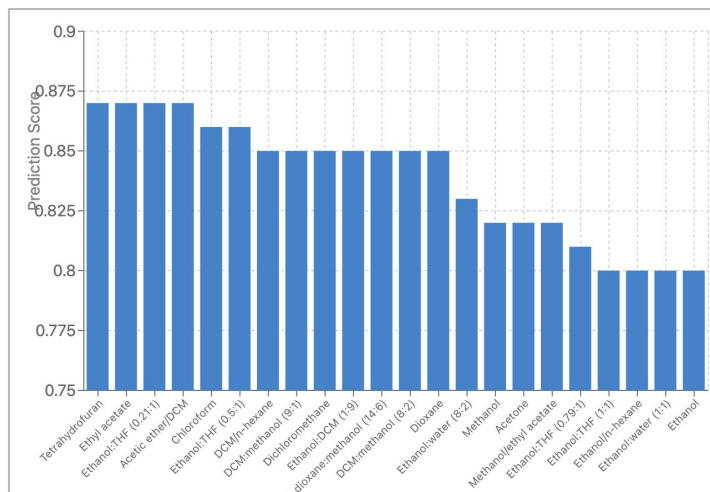
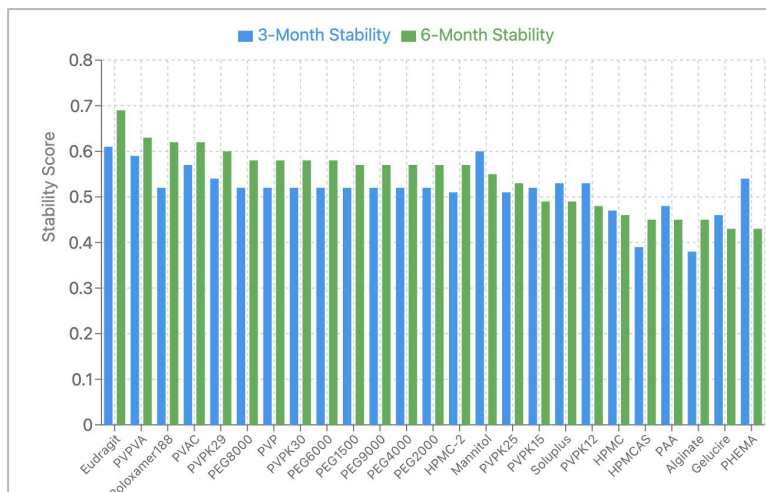
Chemical Disease Pathway



Target Distribution of Protein Families

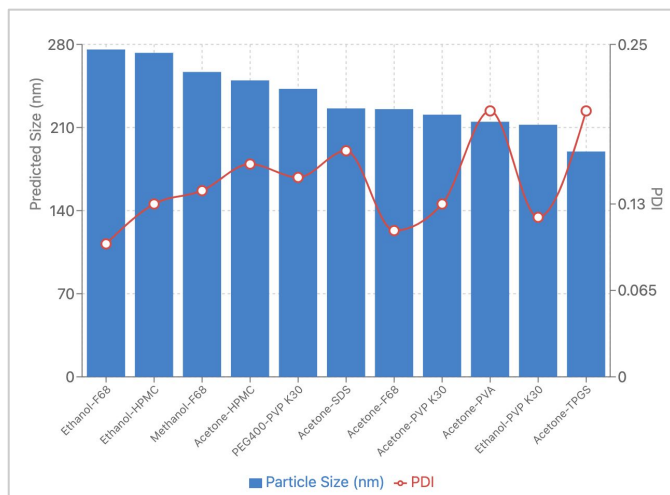
Major Target Identified	PDB ID	Physiological Effects	Risks	Drug Before	Adverse Effects
PIM1	4XH6	Anti-inflammatory, Diabetes Mellitus, Obesity	Yes	Thioridazine	Hospitalization (initial or prolonged):24 Death: 22 Life-Threatening condition:6 Disability:4
RSK2	4GUE	Modulate cell proliferation, transformation, Cell Cycle transition	Yes	N/A	N/A

Formulation Profile



Solid Dispersions Stability
Technique: 1-Hot Melt Extrusion
Storage Temperature: 25.0°C
Preparation Temperature: 60.0°C
Relative Humidity: 75.0%

Phospholipid Complex Combination
Reaction Temperature: 50.0°C
Drug Phospholipid Ratio: 0.50
Reaction Time: 2.0h



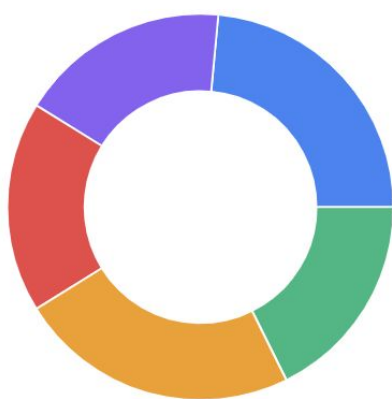
Nanosize Particle Predictions
Method: Anti-solvent
Concentration S1: 0.5
Volume Ratio: 0.1
Volume of Water: 2
Ultrasonication Time: 1.5
Stirring Speed: 1000
Ultrasonication Power: 300
Concentration API: 10

Characterization Results

Size 141.10 nm	Zeta Potential -17.68 mV	PDI 0.13	Encapsulation 72.78 %
Lipid Composition 46.7% 33.3% 20% ● DSPC ● Cholesterol ● DSPG			
Drug/Lipid Ratio 21.29%		Solvent Chloroform	
Preparation Method Thin Film Hydration		Temperature 60.0°C	
Preparation Time 60.0 min		Filtration Yes (200 nm)	
Sonication No			

Formulation Risks

Excipient Hazard Flags



● **Alkaline excipients**

Bentonite, Calcium Carbonate,
Magnesium Oxide, Sodium Bicarbonate

● **Metal ion impurities**

Calcium Alginate, Calcium Phosphate,
Zinc Stearate

● **Oxidizing excipients**

Acacia, Attapulgite, Calcium Oxide

● **Aldehyde impurities**

HPMC, Lactose, PEG, Starch

● **Peroxide impurities**

Lanolin, PVP, PVPP

The excipient risk analysis identifies five major risk categories that require careful consideration in formulation. The risk profile includes potential interactions with alkaline excipients (such as calcium carbonate and sodium bicarbonate), metal ion-containing excipients (like calcium alginate and zinc stearate), and oxidizing excipients (including acacia and calcium oxide). Additional concerns arise from excipients containing aldehyde impurities (such as HPMC, PEG, and starch) and peroxide impurities (found in materials like PVP and lanolin). These chemical interactions could affect drug stability and efficacy, necessitating careful excipient selection during formulation development.

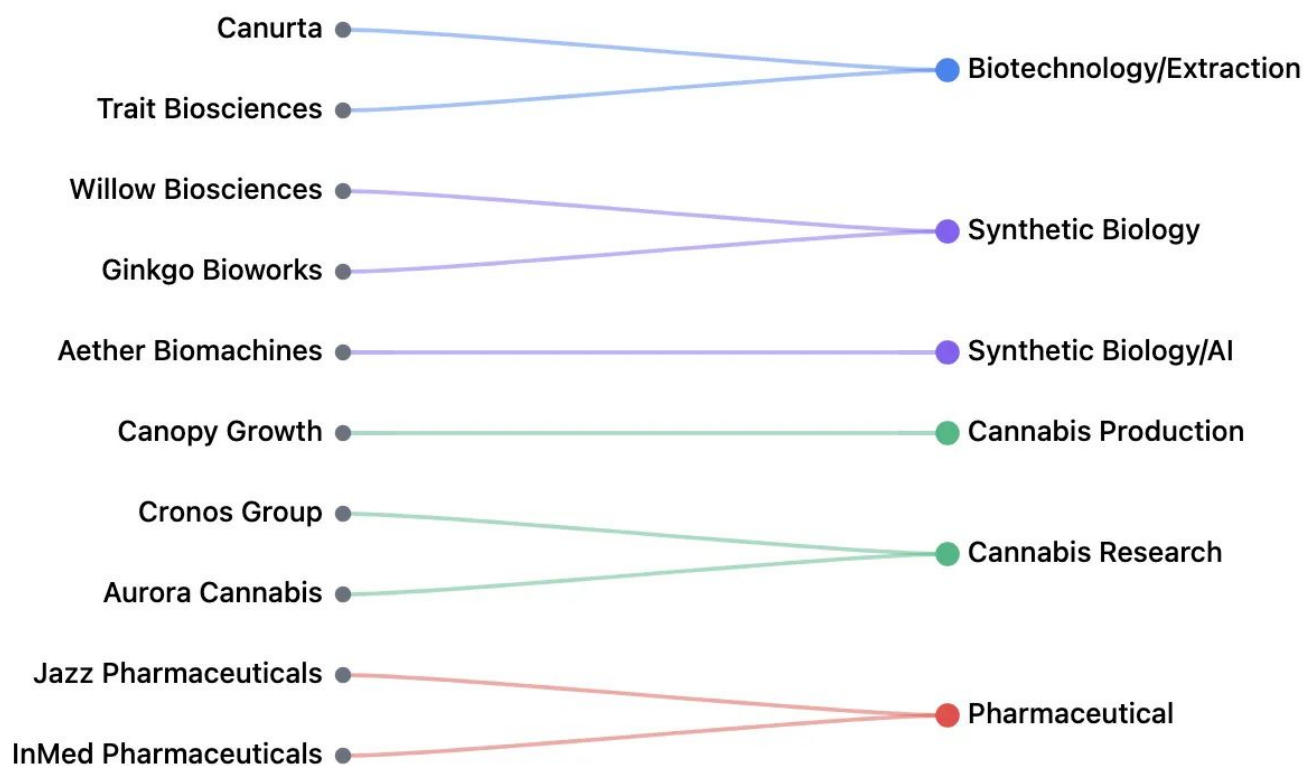
Toxicology Endpoints

Pulmonary embolism, pulmonary hypertensio	No	Pulmonary edema-KRFP	No
Drug-induced rhabdomyolysis	No	Bronchitis-KRFP	Yes
Acute oral toxicity	Yes	Mutagenicity	Yes
Toxicity on avian species	No		
Honey bee toxicity	No		
Reproductive and development toxicity	No		
Nongenotoxic carcinogenicity	No		
Inhalation toxicity	No		
Drug-induced ototoxicity	No		
Endocrine disruption-estrogen receptor	No		
Eye irritation	No		
Chemical-induced hematotoxicity	No		
Endocrine disruption-pregnane X receptor	No		
Mitochondrial toxicity	No		
Bioconcentration	No		
Chemical aquatic toxicity: Daphnia magna	Yes		
Nephrotoxicity	Yes		
Asthma-KRFP	No		
Endocrine disruption-androgen receptor	No		
Endocrine disruption	No		
Drug-induced neurotoxicity	No		
Drug-induced autoimmune diseases	No		
Mitochondrial Toxicity	No		
Skin sensitization	Yes		
hERG inhibition	Yes		
Pulmonary fibrosis-KRFP	No		
Hepatotoxicity	Yes		
Respiratory infection-KRFP	No		
Pneumonia-KRFP	No		
Chemical aquatic toxicity: Fathead minnow	No		
Idiosyncratic adverse drug reaction	Yes		
Bronchospasm-KRFP	Yes		
Chemical aquatic toxicity: Tetrahymena pyrifo	Yes		

The toxicological assessment reveals 11 positive endpoints out of 36 total endpoints tested. Key concerns include aquatic toxicity (affecting *Daphnia magna* and *Tetrahymena*), cardiac safety (hERG inhibition), and organ-specific toxicity (hepatotoxicity and nephrotoxicity). The compound also shows potential for acute oral toxicity, skin sensitization, and mutagenicity. Several respiratory-related effects are indicated, including bronchitis and bronchospasm. However, the compound shows no significant risks for many critical endpoints including reproductive toxicity, endocrine disruption, neurotoxicity, or carcinogenicity. The majority of KRFP (Knowledge-based Respiratory Failure Prediction) endpoints are negative.

Product Market

In the pursuit of Cannflavin A development, each industry sector brings unique approaches. Canurta leads specialized extraction with proprietary methods specifically for cannflavins, while Trait Biosciences focuses on water-soluble cannabinoid technology. In synthetic biology, Willow Biosciences and Ginkgo Bioworks are developing biosynthetic production methods for rare cannabinoids including Cannflavin A. Aether Biomachines applies AI to optimize biosynthesis processes. Traditional cannabis companies like Canopy Growth and Cronos Group incorporate Cannflavin A research into broader cannabinoid programs, while pharmaceutical companies Jazz and InMed explore medical applications of cannabis-derived compounds including flavonoids.



Customer Journals (Coming Soon)

References

- Gaulton A., Bellis L. J., Bento A. P., Chambers J., Davies M., Hersey A., et al. (2011). ChEMBL: a Large-Scale Bioactivity Database for Drug Discovery. *Nucleic Acids Res.* 40 (D1), D1100–D1107. 10.1093/nar/gkr777 [DOI] [PMC free article] [PubMed] [Google Scholar]
- Godden J. W., Xue L., Bajorath J. (2000). Combinatorial Preferences Affect Molecular Similarity/diversity Calculations Using Binary Fingerprints and Tanimoto Coefficients. *J. Chem. Inf. Comput. Sci.* 40 (1), 163–166. 10.1021/ci990316u [DOI] [PubMed] [Google Scholar]
- Gold L. S., Sawyer C. B., Magaw R., Backman G. M., De Veciana M., Levinson R., et al. (1984). A Carcinogenic Potency Database of the Standardized Results of Animal Bioassays. *Environ. Health Perspect.* 58, 9–319. 10.1289/ehp.84589 [DOI] [PMC free article] [PubMed] [Google Scholar]
- Hanson R. M. (2016). Jmol SMILES and Jmol SMARTS: Specifications and Applications. *J. Cheminform* 8 (1), 50. 10.1186/s13321-016-0160-4 [DOI] [PMC free article] [PubMed] [Google Scholar]
- Hua Y., Shi Y., Cui X., Li X. (2021). In Silico prediction of Chemical-Induced Hematotoxicity with Machine Learning and Deep Learning Methods. *Mol. Divers* 25 (3), 1585–1596. 10.1007/s11030-021-10255-x [DOI] [PubMed] [Google Scholar]
- Huang X., Tang F., Hua Y., Li X. (2021). In Silico prediction of Drug-induced Ototoxicity Using Machine Learning and Deep Learning Methods. *Chem. Biol. Drug Des.* 98 (2), 248–257. 10.1111/cbdd.13894 [DOI] [PubMed] [Google Scholar]
- Jiang C., Yang H., Di P., Li W., Tang Y., Liu G. (2019). In Silico prediction of Chemical Reproductive Toxicity Using Machine Learning. *J. Appl. Toxicol.* 39 (6), 844–854. 10.1002/jat.3772 [DOI] [PubMed] [Google Scholar]
- Jiang C., Zhao P., Li W., Tang Y., Liu G. (2020). In Silico prediction of Chemical Neurotoxicity Using Machine Learning. *Toxicol. Res.* 9 (3), 164–172. 10.1093/toxres/taaa016 [DOI] [PMC free article] [PubMed] [Google Scholar]
- Kalgutkar A. S. (2020). Designing Around Structural Alerts in Drug Discovery. *J. Med. Chem.* 63 (12), 6276–6302. 10.1021/acs.jmedchem.9b00917 [DOI] [PubMed] [Google Scholar]
- Kühne R., Ebert R.-U., Schüürmann G. (2006). Model Selection Based on Structural Similarity–Method Description and Application to Water Solubility Prediction. *J. Chem. Inf. Model.* 46 (2), 636–641. 10.1021/ci0503762 [DOI] [PubMed] [Google Scholar]
- Li X., Chen L., Cheng F., Wu Z., Bian H., Xu C., et al. (2014). In Silico prediction of Chemical Acute Oral Toxicity Using Multi-Classification Methods. *J. Chem. Inf. Model.* 54 (4), 1061–1069. 10.1021/ci5000467 [DOI] [PubMed] [Google Scholar]
- Li X., Chen Y., Song X., Zhang Y., Li H., Zhao Y. (2018). The Development and Application of In Silico Models for Drug Induced Liver Injury. *RSC Adv.* 8 (15), 8101–8111. 10.1039/C7RA12957B [DOI] [PMC free article] [PubMed] [Google Scholar]
- Li X., Zhang Y., Chen H., Li H., Zhao Y. (2017a). In Silico prediction of Chronic Toxicity with Chemical Category Approaches. *RSC Adv.* 7 (66), 41330–41338. 10.1039/C7RA08415C [DOI] [Google Scholar]
- Li X., Zhang Y., Chen H., Li H., Zhao Y. (2017b). Insights into the Molecular Basis of the Acute Contact Toxicity of Diverse Organic Chemicals in the Honey Bee. *J. Chem. Inf. Model.* 57 (12), 2948–2957. 10.1021/acs.jcim.7b00476 [DOI] [PubMed] [Google Scholar]
- Li X., Zhang Y., Li H., Zhao Y. (2017c). Modeling of the hERG K⁺ Channel Blockage Using Online Chemical Database and Modeling Environment (OCHEM). *Mol. Inf.* 36 (12), 1700074. 10.1002/minf.201700074 [DOI] [PubMed] [Google Scholar]
- Limban C., Nuță D. C., Chiriță C., Negreș S., Arsene A. L., Goumenou M., et al. (2018). The Use of Structural Alerts to Avoid the Toxicity of Pharmaceuticals. *Toxicol. Rep.* 5, 943–953. 10.1016/j.toxrep.2018.08.017 [DOI] [PMC free article] [PubMed] [Google Scholar]
- Lovrić M., Molero J. M., Kern R. (2019). PySpark and RDKit: Moving towards Big Data in Cheminformatics. *Mol. Inf.* 38 (6), 1800082. 10.1002/minf.201800082 [DOI] [PubMed] [Google Scholar]
- Nelms M. D., Mellor C. L., Cronin M. T. D., Madden J. C., Enoch S. J. (2015). Development of an In Silico Profiler for Mitochondrial Toxicity. *Chem. Res. Toxicol.* 28 (10), 1891–1902. 10.1021/acs.chemrestox.5b00275 [DOI] [PubMed] [Google Scholar]
- O’Boyle N. M. (2012). Towards a Universal Smiles Representation - a Standard Method to Generate Canonical Smiles Based on the InChI. *J. Cheminform* 4 (1), 22. 10.1186/1758-2946-4-22 [DOI] [PMC free article] [PubMed] [Google Scholar]
- Patlewicz G., Jeliakova N., Safford R. J., Worth A. P., Aleksiev B. (2008). An Evaluation of the Implementation of the Cramer Classification Scheme in the ToxTree Software. *SAR QSAR Environ. Res.* 19 (5-6), 495–524. 10.1080/10629360802083871 [DOI] [PubMed] [Google Scholar]
- Pires D. E. V., Blundell T. L., Ascher D. B. (2015). pkCSM: Predicting Small-Molecule Pharmacokinetic and Toxicity Properties Using Graph-Based Signatures. *J. Med. Chem.* 58 (9), 4066–4072. 10.1021/acs.jmedchem.5b00104 [DOI] [PMC free article] [PubMed] [Google Scholar]
- Schyman P., Liu R., Desai V., Wallqvist A. (2017). vNN Web Server for ADMET Predictions. *Front. Pharmacol.* 8, 889. 10.3389/fphar.2017.00889 [DOI] [PMC free article] [PubMed] [Google Scholar]
- Shi Y., Hua Y., Wang B., Zhang R., Li X. (2022). In Silico Prediction and Insights into the Structural Basis of Drug Induced Nephrotoxicity. *Front. Pharmacol.* 12, 793332. 10.3389/fphar.2021.793332 [DOI] [PMC free article] [PubMed] [Google Scholar]
- Sun L., Zhang C., Chen Y., Li X., Zhuang S., Li W., et al. (2015). In Silico prediction of Chemical Aquatic Toxicity with Chemical Category Approaches and Substructural Alerts. *Toxicol. Res.* 4 (2), 452–463. 10.1039/C4TX00174E [DOI] [Google Scholar]
- Sushko I., Salmina E., Potemkin V. A., Poda G., Tetko I. V. (2012). ToxAlerts: a Web Server of Structural Alerts for Toxic Chemicals and Compounds with Potential Adverse Reactions. *J. Chem. Inf. Model.* 52 (8), 2310–2316. 10.1021/ci300245q [DOI] [PMC free article] [PubMed] [Google Scholar]
- Tomasulo P. (2002). ChemIDplus-super Source for Chemical and Drug Information. *Med. Ref. Serv. Q.* 21 (1), 53–59. 10.1300/J115v21n01_04 [DOI] [PubMed] [Google Scholar]
- Wang Q., Li X., Yang H., Cai Y., Wang Y., Wang Z., et al. (2017). In Silico prediction of Serious Eye Irritation or Corrosion Potential of Chemicals. *RSC Adv.* 7 (11), 6697–6703. 10.1039/C6RA25267B [DOI] [Google Scholar]
- Wang Y., Lu J., Wang F., Shen Q., Zheng M., Luo X., et al. (2012). Estimation of Carcinogenicity Using Molecular Fragments Tree. *J. Chem. Inf. Model.* 52 (8), 1994–2003. 10.1021/ci300266p [DOI] [PubMed] [Google Scholar]
- Wang Z., Chen J., Hong H. (2021). Developing QSAR Models with Defined Applicability Domains on PPAR γ Binding Affinity Using Large Data Sets and Machine Learning Algorithms. *Environ. Sci. Technol.* 55 (10), 6857–6866. 10.1021/acs.est.0c07040 [DOI] [PubMed] [Google Scholar]
- Wishart D. S., Feunang Y. D., Guo A. C., Lo E. J., Marcu A., Grant J. R., et al. (2017). DrugBank 5.0: a Major Update to the DrugBank Database for 2018. *Nucleic Acids Res.* 46 (D1), D1074–D1082. 10.1093/nar/gkx1037 [DOI] [PMC free article] [PubMed] [Google Scholar]
- Wu Y., Zhu J., Fu P., Tong W., Hong H., Chen M. (2021). Machine Learning for Predicting Risk of Drug-Induced Autoimmune Diseases by Structural Alerts and Daily Dose. *Ijerp* 18 (13), 7139. 10.3390/ijerp18137139 [DOI] [PMC free article] [PubMed] [Google Scholar]
- Xiong G., Wu Z., Yi J., Fu L., Yang Z., Hsieh C., et al. (2021). ADMETlab 2.0: an Integrated Online Platform for Accurate and Comprehensive Predictions of ADMET Properties. *Nucleic Acids Res.* 49 (W1), W5–W14. 10.1093/nar/gkab255 [DOI] [PMC free article] [PubMed] [Google Scholar]
- Yang H., Li J., Wu Z., Li W., Liu G., Tang Y. (2017). Evaluation of Different Methods for Identification of Structural Alerts Using Chemical Ames Mutagenicity Data Set as a Benchmark. *Chem. Res. Toxicol.* 30 (6), 1355–1364. 10.1021/acs.chemrestox.7b00083 [DOI] [PubMed] [Google Scholar]
- Yang H., Lou C., Li W., Liu G., Tang Y. (2020). Computational Approaches to Identify Structural Alerts and Their Applications in Environmental Toxicology and Drug Discovery. *Chem. Res. Toxicol.* 33 (6), 1312–1322. 10.1021/acs.chemrestox.0c00006 [DOI] [PubMed] [Google Scholar]
- Yang H., Lou C., Sun L., Li J., Cai Y., Wang Z., et al. (2018a). admetSAR 2.0: Web-Service for Prediction and Optimization of Chemical ADMET Properties. *Bioinformatics* 35 (6), 1067–1069. 10.1093/bioinformatics/bty707 [DOI] [PubMed] [Google Scholar]
- Yang H., Sun L., Li W., Liu G., Tang Y. (2018b). Identification of Nontoxic Substructures: a New Strategy to Avoid Potential Toxicity Risk. *Toxicol. Sci.* 165 (2), 396–407. 10.1093/toxsci/ky146 [DOI] [PubMed] [Google Scholar]
- Yang H., Sun L., Li W., Liu G., Tang Y. (2018c). In Silico prediction of Chemical Toxicity for Drug Design Using Machine Learning Methods and Structural Alerts. *Front. Chem.* 6, 30. 10.3389/fchem.2018.00030 [DOI] [PMC free article] [PubMed] [Google Scholar]
- Yap C. W. (2011). PaDEL-descriptor: An Open Source Software to Calculate Molecular Descriptors and Fingerprints. *J. Comput. Chem.* 32 (7), 1466–1474. 10.1002/jcc.21707 [DOI] [PubMed] [Google Scholar]
- Zhang C., Cheng F., Sun L., Zhuang S., Li W., Liu G., et al. (2015). In Silico prediction of Chemical Toxicity on Avian Species Using Chemical Category Approaches. *Chemosphere* 122, 280–287. 10.1016/j.chemosphere.2014.12.001 [DOI] [PubMed] [Google Scholar]