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# An *In-silico* Pharmacokinetics Study on Cis-Heptadeca-1,9-Diene-4,6-Diyne-3,8-Diol: A Nutraceutical Compound with Anticancer Properties

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#### Authors' contributions

This work was carried out in collaboration among all authors. Authors SME, SIU and PTO designed the study. Authors OCA, OHC and VCN wrote the first draft of the manuscript. Authors PTO, SIU and OAD managed the literature searches. All authors read and approved the final manuscript.

#### Article Information

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# ABSTRACT

The polyacetylenic compound cis-heptadeca-1,9-diene-4,6-diyne-3,8-diol with the trivial name of falcarindiol, has been shown through various studies to exhibit anticancer activities. The beneficial effects of polyacetylenes occur at nontoxic concentrations and thus represent pharmacologically useful properties indicating that polyacetylenes may be important nutraceuticals of vegetables. In

the human diet, carrots are the major dietary source of falcarinol-type polyacetylenes, in particular falcarinol and falcarindiol. The prediction of physicochemical parameters relevant for drug likeness was performed by computational methods. The Lipinski, Ghose and Veber rules were applied to assess drug likeness and to predict whether the compound is likely to be a bioactive compound according to other important parameters such as molecular weight, LogP, number of hydrogen bond donors and hydrogen bond acceptors. The SwissADME tool used vector machine algorithm (SVM) with fastidiously cleaned large datasets of known inhibitors/non-inhibitors as well as substrates/non-substrates for its predictions. Results from the *in silico* pharmacokinetics study on falcarindiol showed that the compound exhibited drug likeness characteristics and as such can serve the purpose of an anticancer agent, being a nutaceutical and bioactive component of falcarindiol.

Keywords: Falcarindiol; polyacetylenes; lipinski; ghose; pharmacokinetics.

#### **1. INTRODUCTION**

Epidemiological studies have provided evidence that a diet high in fruit and vegetables is associated with a reduced risk for the development of certain types of cancer, cardiovascular diseases, diabetes, and other diseases [1]. Compounds associated with the health promoting effects of vegetables are glucosinolates and other organosulfur compounds and their degradation products, carotenoids, phytosterols, polyphenols, vitamins, and dietary fibers [2]. The mechanisms for the protection of these classes of natural products are mostly unknown and may only in part explain the health effects of vegetables. Consequently, in recent years focus has been on other types of potential health promoting compounds [3].

Falcarindiol (FAD) is a natural polyyne found in vegetables and it has been shown to have antiinflammation. antibacterial and anticancer activities, as well as protective effects against hepatotoxicity. Moreover, these beneficial effects occur at non-toxic concentrations and thus represent pharmacologically useful properties [4, 5]. In particular, the anti-proliferative and antitumor functions of FAD were confirmed repeatedly, in colon cancer and breast cancer cells but unfortunately, however, there is little research on the consequence of FAD treatment in the fate determination of stem cells [6,7,8,9].

Falcarindiol has also been isolated from *Aegopodum podraria L.*, (Family: *Apiaceae*) and its antifungal activity towards some other fungi noted [10]. Various other Umbelliferae (family of mostly aromatic flowering plants named after the type genus Apium and commonly known as the celery, carrot or parsley family, or simply as umbellifers), *Apium graveolens L., Falcaria* 

*vulgari Bernh., Oenanthe crocata L. and Opopanax chironium Koch.* also produce falcarindiol [11].

A good number of other acetylenic compounds isolated from different plant families have also been shown to exhibit antifungal activities [12,13]. The few antifungal polyacetylenic compounds that have been investigated appear to be important in disease resistance and since polyacetylenes are widespread in plants, the group as a whole may subsequently be found to be of importance as suggested by Overeem [14].

This study is therefore aimed at predicting the pharmacokinetics parameters of falcarindiol for the confirmation of its drug likeness properties using specific computational biology approach.

#### 2. MATERIALS AND METHODS

#### 2.1 In silico ADMET Screening

The ADMET properties including aqueous solubility, blood brain barrier (BBB), plasma protein binding, CYP2D6 binding, gastrointestinal absorption and hepatotoxicity were evaluated for these molecules within human. The models used to predict the ADMET properties in this protocol are derived from a variety of experimental data sources and are catalogued in the product documentation [15].

## 2.2 Screening for Pharmacokinetics and Drug-likeness

Pharmacokinetics and drug-likeness prediction for falcarindiol was also performed using an online tool SwissADME [16] of the Swiss Institute of Bioinformatics (http://www.sib.swiss). This was used to evaluate individual ADME behaviors of the compound [17].

In the Swiss ADME program package, the topological polar surface area (TPSA) was calculated according to Ertl *et al* [18]. Furthermore, the lipophilicity was predicted according to Wildman et al. [19], the solubility was predicted according to Ali et al. [20], and violations of the Pfzer flter for oral bioavailability were assessed according to Lipinski et al. [21]. Results are indicated in Fig. 3.

The analysis task was done to check whether falcarindiol was an inhibitor of isoforms of Cytochrome P450 (CYP) family such as CYP1A2 and CYP2D6. In addition to other predicted pharmacokinetics (such as gastro intestinal absorption, P-glycoprotein and Blood brain barrier) and drug-likeness prediction such as Ghose and Veber rules and the bioavailability score [22,23].

#### 2.3 2D Structure Modeling and File Conversion

2D structural models were drawn in MarvinSketch (ChemAxon Software) and the falcarindiol SMILES was translated into a mol2 file by online SMILES translator and structure file generator found in the OpenBabel software [24].

# 2.4 Ligand Minimization

Falcarindiol 3D structure was minimized using the UCSF Chimera software [25]. UCSF Chimera is an extensible program for analyzing and interactively visualizing molecular structures and related data which include supramolecular assemblies. density maps, alignment of sequences. results from molecular and conformational docking trajectories ensembles [26].

# 2.5 Visualization of Atoms

Atoms making up falcarindiol was visualized using the Pymol molecular visualizer [27]. PyMOL is an open-source tool for model visualization and it is made available for utilization in structural biology [28]. The *Py* aspect of the name of the software is a reference pointer that it is extensible and extends by the python programming language [29].

## 3. RESULTS AND DISCUSSION

In this study, we subjected the designed falcarindiol chemical structure to in silico ADMET screening, using the SwissADME online software, to predict the overall absorption. distribution, metabolism, excretion, and toxicity risks. In the in silico evaluation, the analysis of different descriptors (calculated octanol/water partition coefficient, molecular weight, molecular volume, number of hydrogen bond donors and acceptor groups) of all compounds revealed that they are highly hydrophobic to penetrate the biological membranes, as determined by the Lipinski "rule-of-five" which states that the absorption or permeation of a molecule is more likely when the molecular weight is under 500 g/mol, the value of log P is lower than 5, and the molecule has utmost 5 H-donor and 10 Hacceptor atoms (cLogP<5, MW<500, HBD<5 and HBA<10) [21, 30] as shown in Fig. 3. The Lipinski (Pfzer) filter is the pioneer rule-of-five [31], therefore, this result suggests that falcarindiol have a high theoretical oral bioavailability.

According to the pharmacokinetic properties, falcarindiol showed a high gastrointestinal absorption property which gives credence to it oral bioavailability score. Falcarindiol according to the *in silico* pharmacokinetics report possess the blood brain barrier permeability attribute but it however showed no inhibition to the Cytochrome P450 isomers and the P-glycoprotein with the exemption of the CYP2C9 isomer. The druglikeness prediction was also conducted depending on the selected Ghose, Veber rules and bioavailability score.

The Ghose filter (Amgen) [22] defines druglikeness constraints as follows: calculated log P is between -0.4 and 5.6. MW is between 160 and 480, molar refractivity is between 40 and 130, and the total number of atoms is between 20 and 70. Veber (GSK) [23], rule defines drug likeness constraints as Rotatable bond count  $\leq$  10 and polar surface area (PSA)  $\leq$  140. Just as the screening process with Lipinski Rule of Five showed that there were no compound violations, suggestions according to the screening process with Ghose rules show that falcarindiol can be accepted as it showed no violations (Fig. 3). Also, the screening process with Veber rules showed that falcarindiol met the criteria for drug likeness assessment. Using the above models also indicate that falcarindiol violates none of the standards defined by each drug likeness prediction model.

The Bioavailability score was implemented without changes from Martin et al., and it is similar but seeks to predict the probability of a compound to have at least 10% oral bioavailability in rat or measurable Caco-2 permeability. The synthetic accessibility score of falcarindiol has also shown that it falls within the range of compounds that can be easily synthesized. We therefore recommend the laboratory synthesis of falcarindiol for further studies such as the molecular docking with structural modifications and preclinical trials [32].

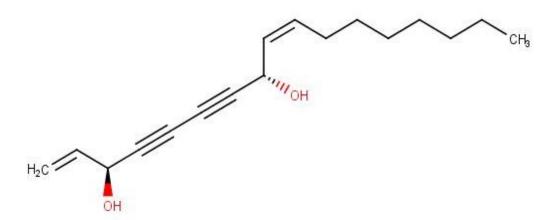


Fig. 1. 2D structure of falcarindiol

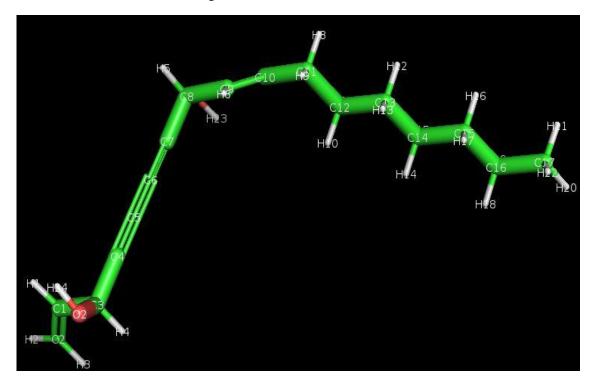


Fig. 2. Minimized 3D structure of falcarindiol with labeled atoms

Molecule 1			
H @ @			Water Solubility
	LIPO	Log S (ESOL) 8	-3.69
		Solubility	5.37e-02 mg/ml ; 2.06e-04 mol/l
	CH,	Class 0	Soluble
	Y	Log S (Ali) 😣	-4.95
		Solubility	2.95e-03 mg/ml; 1.13e-05 mol/l
		Class 🥯	Moderately soluble
		Log S (SILICOS-IT)	-2.35
	NSATU POLAR	Solubility	1.17e+00 mg/ml ; 4.49e-03 mol/l
HC AN	Not Contract of Co	Class 0	Soluble
			Pharmacokinetics
	INSOLU	GI absorption 9	High
SMILES CCCCCCC/C=CV	C@@H](C#CC#C[C@H](C=C)0)0	BBB permeant	Yes
Physicochemical Properties		P-gp substrate 9	No
Formula	C17H24O2	CYP1A2 inhibitor 🥹	No
Molecular weight	260.37 g/mol	CYP2C19 inhibitor 😣	No
Num, heavy atoms	19	CYP2C9 inhibitor 9	Yes
Num. arom. heavy atoms	0	CYP2D6 inhibitor 9	No
Fraction Csp3	0.53	CYP3A4 inhibitor 9	No
Num, rotatable bonds	8	Log Ko (skin permeation)	-4.78 cm/s
Num, H-bond acceptors	2		Druglikeness
Num. H-bond donors	2	Lipinski 🥹	Yes: 0 violation
Molar Refractivity	81.53	Ghose 9	Yes
TPSA 🥹	40.46 Å*	Veber 😑	Yes
	Lipophilicity	Egan 🥹	Yes
Log Poly (iLOGP)	4.02	Muegge 9	Yes
Log P <sub>olw</sub> (XLOGP3) 🥯	4.38	Bioavailability Score	0.55
Log P <sub>olw</sub> (WLOGP) 🤨	2.98		Medicinal Chemistry
Log P <sub>olw</sub> (MLOGP) 🥯	3.33	PAINS 0	0 alert
Log Poly (SILICOS-IT) 🥹	4.35	Brenk 🥹	2 alerts: isolated_alkene, triple_bond 9
Consensus Log Poly	3.81	Leadlikeness 🧐	No; 2 violations: Rotors>7, XLOGP3>3.5
CIW .	ar i ar a	Synthetic accessibility	4.65

Fig. 3. Swiss ADME in silico pharmacokinetics result for falcarindiol

#### 4. CONCLUSION

The computational analysis on the pharmacokinetics and drug likeness parameters of falcarindol has confirmed the safety of the compound for oral administration. Haven shown the ability to cross the blood brain barrier as seen in the above results, falcarindiol can only be administered as a neuroactive drug to avoid side effects as regarding the central nervous system.

## **COMPETING INTERESTS**

Authors have declared that no competing interests exist.

#### REFERENCES

1. Maynard M, Gunnell D, Emmett P, Frankel S, Smith GD. Fruit, vegetables, and

antioxidants in childhood and risk of adult cancer: the Boyd Orr cohort. J. Epidemiol. Community Health. 2003;57:218–225.

- Trichopoulou A, Naska A, Antoniou A, Friel S, Trygg K, Turrini A. Vegetable and fruit: The evidence in their favour and the public health perspective. Int. J. Vitam. Nutr. Res. 2003;73:63–69.
- Brandt K, Christensen LP, Hansen-Moeller J, Hansen SL, Haraldsdottir J, Jespersen L, Purup S, Kharazmi A, Barkholdt V, Froekiær H, Kobæk-Larsen M. Health promoting compounds in vegetables and fruits: A systematic approach for identifying plant components with impact on human health. Trends Food Sci. Technol. 2004;15:384–393.
- Metzger BT, Barnes DM, Reed JD. Purple carrot (*Daucus carota* L.) polyacetylenes decrease lipopolysaccharide-induced expression of inflammatory proteins in

macrophage and endothelial cells. J Agric Food Chem. 2008;56:3554–3560.

- 5. Brennan CA, Garrett WS. Gut microbiota, infammation, and colorectal cancer. Annu Rev Microbiol. 2016;70:395–411.
- Deng S, Wang Y, Inui T, Chen SN, Farnsworth NR, Cho S, et al. Anti-TB polyynes from the roots of Angelica sinensis. Phytother Res. 2008;22:878–882.
- Stavri M, Gibbons S. The antimycobacterial constituents of dill (*Anethum graveolens*). Phytother Res. 2005;19:938–941.
- 8. Lu T, Gu M, Zhao Y, Zheng X, Xing C. Autophagy contributes to falcarindiolinduced cell death in breast cancer cells with enhanced endoplasmic reticulum stress. PLoS ONE. 2017;12:e0176348.
- Jin HR, Zhao J, Zhang Z, Liao Y, Wang CZ, Huang WH, Li SP, He TC, Yuan CS, Du W. The antitumor natural compound falcarindiol promotes cancer cell death by inducing endoplasmic reticulum stress. Cell Death Dis. 2012;3:e376.
- 10. Kemp MS. Falcarindiol: An antifungal polyacetylen from *Aegopodum podararia*. Phytochemistry. 1978;17:102.
- Bohlmann F. Polyacetyle verbindungen, CXXXIX. Notiz Uber die Inhaltsto:fe von Petersilieund Sellerie-Wurzeln. Chemisches Berihte, 1967;10:3454. Bohlmann, F. (1971). In Th Biolog and Chestry of th Umbellferae (Ed. by V. H. Heywood), p. 279. Academic Press, London, New York.
- 12. Allen HE, Thomas CA. Trans-tans-3, 11tridecadiee-5, 7, 9-triyene-1,2-diol, a antifungal polyacetylene from diseased safflower (*Carthamus tinctorius*). Phytochemistry. 1971a;16:1579.
- Allen HE, Thomas CA. A second antifungal polyacetylene compound from Phytophthora-infected safflower. Phytopathology. 1971b;61:1107.
- Overeem J. In: Biochemical Aspects of Plant-Parasite Relationships (Ed, by J, Friend & D.R. Trelfall). Academic Press, London, New York. 1976;198.
- 15. Discovery Studio. 2.1 (2008) San Diego, CA, USA.
- Zoete V, Daina A, Bovigny C, Michielin O. Swiss similarity: A web tool for low to ultra high throughput ligand-based virtual screening. J Chem Inf Model. 2016;56: 1399-1404.
- 17. Daina A, Michielin O, Zoete V. Swissadme. A free web tool to evaluate

pharmacokinetics, drug-likeness and medicinal chemistry friendliness of small molecules. Sci Rep. 2017;7:427-517.

- Ertl P, Rohde B, Selzer P. Fast calculation of molecular polar surface area as a sum of fragment-based contributions and its application to the prediction of drug transport properties. J Med Chem. 2000;43:3714–7.
- 19. Wildman SA, Crippen GM. Prediction of physicochemical parameters by atomic contributions. J Chem Inf Comput Sci. 1999;39:868–73.
- Ali J, Camilleri P, Brown MB, Hutt AJ, Kirton SB. Revisiting the general solubility equation: *In silico* prediction of aqueous solubility incorporating the effect of topographical polar surface area. J Chem Inf Model. 2012;52:420–8.
- 21. Lipinski CA, Lombardo F, Dominy BW, Feeney PJ. Experimental and computational approaches to estimate solubility and permeability in drug discovery and development settings. Adv Drug Deliv Rev. 2001;46:3–26.
- 22. Ghose AK, Viswanadhan VN, Wendoloski JJ. A knowledge-based approach in designing combinatorial or medicinal chemistry libraries for drug discovery. A qualitative and quantitative characterization of known drug databases. J Comb Chem. 1999;1:55-68.
- 23. Veber DF, Johnson SR, Cheng HY, Smith BR, Ward KW, et al. Molecular properties that influence the oral bioavailability of drug candidates. J Med Chem. 2002;45:2615-2623.
- Noel MO, Michael B, Craig AJ, Chris M, Tim V, Geoffrey R. Hutchison Open Babel: An open chemical toolbox. Journal of Cheminformatics. 2011;3:33.
- Pettersen EF, Goddard TD, Huang CC, Couch GS, Greenblatt DM, Meng EC, Ferrin TE. UCSF Chimera--a visualization system for exploratory research and analysis. J Computational Chemistry. 2004;25(13):1605–1612.
- 26. Savjani KT, Gajjar AK, Savjani JK. Drug solubility: importance and enhancement techniques. ISRN Pharm. 2012;195727.
- Zhu K, Day T, Warshaviak D, Murrett C, Friesner R, Pearlman D. Antibody structure determination using a combination of homology modeling, energy-based refinement, and loop prediction. Proteins. 2014;82(8):1646-1655.

Okoli et al.; AFSJ, 7(3): 1-7, 2019; Article no.AFSJ.47204

- Salam NK, Adzhigirey M, Sherman W, Pearlman DA. Structure-based approach to the prediction of disulfide bonds in proteins. Protein Engineering, Design and Selection. 2014;27(10):365-374.
- 29. Beard H, Cholleti A, Pearlman D, Sherman W, Loving KA. Applying physics-based scoring to calculate free energies of binding for single amino acid mutations in protein-protein complexes, PLoS ONE. 2013;8(12):828-849.
- Lipinski CA. Lead-and drug-like compounds: The rule-of-five revolution. Drug Discov Today Technol. 2004;1:337-341.
- Lipinski CA, Lombardo F, Dominy BW, Feeney PJ. Experimental and computational approaches to estimate solubility and permeability in drug discovery and development settings. Adv Drug Deliv Rev. 2012;64:4-17.
- 32. Martin YC. A bioavailability score. J Med Chem. 2005;48:3164-3170.

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