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STUDY BIOINFORMATICS WHITE TURMERIC (CURCUMA ZEDOARIA) AGAINST BREAST CANCER

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ABSTRACT

In Indonesia, almost 70% of people with this disease have experienced cancer, which is found in an advanced stage. This cancer occurs because of an unhealthy lifestyle, so that the immune system decreases. Cancer is usually experienced by many women, one of which is breast cancer. This breast cancer arises from the swelling of the woman's breast area or it can be called a tumor. White turmeric (Curcuma zedoria) is a traditional medicine in Indonesia. This white turmeric contains 11 active compounds, there are 5 main active compounds, namely 1,8-cineole, curcumin, D-alpha-pinene, D-borneol and D-champene. The methods used are modeling and computation using phytochem, chemdraw ultra 12.0 and chem 3D Pro 12.0, Switzerland. The conclusion from the results of bioinformatic research on white turmeric extract (curcuma zedoria.) Can be used as an anti-breast cancer drug.

Key words: White turmeric; breast cancer

1. INTRODUCTION

This research is about the many medicinal properties of herbs that have not been utilized by researchers until now [1,2]. The method used in this study was ethnobotany, which consisted of using herbal medicines used by certain ethnicities, proven by experimental pharmacological activity and continued with the isolation of compounds that had biological activity [3,4,5,6].

One of the herbal remedies that can be used for medicine, namely white turmeric (curcuma zedori) is a plant originating from the Himalayas, India and countries scattered in

Asia[7,8,9]. White turmeric is also found in Indonesia which is used as a traditional medicine in various regions, namely Sumatra, West Java, and Central Java, where the altitude reaches 1000 asl. White turmeric is a rhizomatous or herbaceous type of plant which is an annual plant that grows to form grass with a height of approximately 1 meter, its stems are upright and is a rimbang plant (tubers and underground branches)[10,11]. The chemical content of white turmeric is curcumin, demethoxicumin and essential oil containing monoterpenes (1,8-cinoele) and other ingredients such as ethyl-pmetoxycinamate and D-borneol[12,13,14].

Chemical reactions are very important in life[15]. Where chemical reactions can occur in the stomach, eyes, mouth and other organs. Chemical reactions occur in living things, namely humans, animals and plants and the universe. A chemical reaction will produce new products and new properties, chemical reactions can occur spontaneously or not, there are enormous benefits of chemical reactions in our lives[16].

In white turmeric, which can be used as a cancer drug is its extraction[17]. This extract from white turmeric will react with abnormal body cells, thus helping to stabilize problematic cells. It is suspected that white turmeric contains D-borneol which can attach to abnormal body cells, so that problem cells in the body that should develop are stopped due to the activity of D-borneol sticking[18,19].

2. LITERATURE REVIEW

2.1 Bioinformatics tools used in circRNA research

Since 2012, a large number of bioinformatics tools have been developed for circRNA study[20]. Although the functions of circRNA tools are diverse, they can be classified into three main categories that include circRNA identification tools[21]. A few tools such as disease-associated circRNA databases are technically databases but do not contain primary data. We group tools with similar functions together. Tools that primarily recognize circRNAs are classified as circRNA identification tools[22,23]]. Databases that provide basic annotation information are classified as circRNA annotation databases. Finally, stand-alone tools or online services are classified as other tools.

2.2 CircRNA identification tools

Prior to identification, RNA library construction and sequencing provide the raw reads needed. Several library preparation methods can be used for RNA sequencing and subsequent circRNA identification[24]. In theory, polyA+ selected RNAs should not contain circRNAs. Nonetheless, libraries sequenced from this selection of RNA may still contain a tiny

proportion of circRNA, because selection is not absolutely accurate. Despite this minor imperfection, polyA+ selected datasets can be used as negative controls. Another factor in identification is specificity[25]. To improve this aspect of identification, the false-positive rate of back spliced junction (BSJ) reads can be reduced in various ways. For example, BSJ reads can be supported by its presence in different samples, by a strict cut-off for read counts or by determining whether the circRNA enrichment step is adopted in the library preparation[26]. Most identification tools prefer a circRNA-enriched RNA-Seq dataset as an input. Another strategy is to use paired-end read sequencing to improve the identification of decoy reads that are typically discarded as alignment artifacts. This can be helpful for filtering out false-positive circRNAs.

Notably, most tools have an essential step termed remapping for discovering BSJ reads. Except for segemehl, k-mer based tools and machine-learning-based tools, all identification tools depend on external aligners, with Bowtie, BWA and STAR as the common choices. Most circRNA identification tools belong to the stand-alone category. Nearly all tools release their source code via GitHub and the most popular program language is Python, with R and Perl also being popular languages and Shell as a common choice for building a pipeline. Most pipelines run under Linux or a UNIX-like system. Some tools need to be compiled from source code and some tools need to be manually installed with the dependency environment. Currently, it is more convenient to install tools in Linux, with many tools supporting easy installation methods, such as Conda (bioconda), Docker,

Python package index (PyPI) and BiocManager (Bioconductor)[27]. Most have well-written documents and tutorials to help users install and run their tools. Although installation can be straightforward, some computer science skills may be needed (e.g. programming, Linux system knowledge) to perform an analysis[28]. Therefore, a well-designed, free, online interface or stand-alone tools with a user-friendly interface is generally required for users without advanced computational training. Based on implementation form, circRNA identification tools can be divided into three categories that include BSJ-based, integrated-based and machine learning-based.

3. EXPERIMENTAL

This study uses a systematic literacy review (SLR) method to review studies, assess and interpret and collect information about sedswa contained in white turmeric (curcuma zedoria) and its role in helping to cure breast cancer [29, 30]. To determine the content, phytochem, chemdraw ultra 12.0, chem 3D Pro 12.0, AND Swiss were used to construct the structure of the compounds that were already in the previous https://pubchem.ncbi.nlm.nig.gov/ and determine the parameters (coordinate and cartesian tables) in addition to it is also used http://swisstargetprediction.ch/ to predict the compound is more active to cure any disease.

4. RESULTS AND DISCUSSION

4.1 Chemical content in White Turmeric (Curcuma Zedoaria)

Table 1. Data of White Turmeric (Curcuma Zedoaria)

Activity	Chemical	Part	Low PPM	High PPM	StdDev
67	1,8-Cineole	Rhizoma	902.0	1920.0	-0,59
5	Bis-	Rhizoma	-	1000.0	-1.0
	Desmethoxycurcumin				
135	Curcumin	Rhizoma	-	1000.0	-0.98
0	D-Alpha-Pinene	Rhizoma	141.0	300.0	
4	4 D-Borneol		141.0	300.0	-1.0
0	D-Camphene	Rhizoma	329.0	700.0	1.0
0	D-Cmohor	Rhizoma	395.0	840.0	-1.05
0	EO	Rhizoma	10000.0	1500	
0	Sequiterpene-	Rhizoma	4800.0	9600.0	
	Alcohols				
0	Sesquiterpenes	Rhizoma	1000.0	2000.000	

(source: courtesy of Sindy Afrianti, et al.)

After Optimization

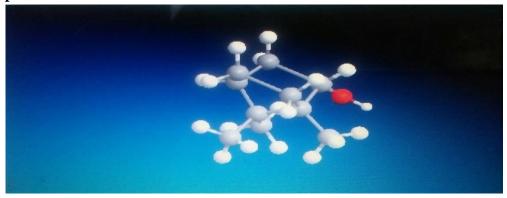


Figure 1. -----MM2----- (source : courtesy of Sindy Afrianti, *et al.*)

Table 2. Cartesian Table

Atom	X(A)	Y(A)	Z(A)			
C(1)	-0.7961	11.244	0.3227			
C(2)	-0.8382	0.6692	-16.222			
C(3)	-21.572	-0.0921	-16.222			
O(11)	0.3072	0.5598	0.9781			
H(12)	-0.7006	22.045	0.5737			
H(13)	-0.8728	15.751	-22.678			
H(14)	0.0462	0.1222	-20.192			

(source: courtesy of Sindy Afrianti, et al.)

Table 3. Internal Coordinate Table

Atom					2nd		2nd
	Atom	Bond	Angle		Angle	2nd	angle
	Bond	Length	Atom	Angle	atom	Angle	Type
C(1)							
C(2)	C(1)	19.979					
C(3)	C(2)	15.230	C(1)	975.915			
H(13)	C(2)	11.130	C(1)	1.123.277	C(3)	1.123.282	Pro-S
H(14)	C(2)	11.130	C(1)	1.162.630	C(3)	1.162.633	Pro-R
C(4)	C(3)	15.230	C(2)	1.094.700	C(1)	791.358	Dihedral
C(7)	C(3)	15.230	C(2)	1.040.001	C(4)	1.039.999	Pro-R
H(17)	C(4)	11.130	C(3)	1.162.631	C(5)	1.162.629	Pro-S
O(11)	C(1)	14.020	C(2)	1.123.281	C(6)	1.123.281	Pro-S

(source: courtesy of Sindy Afrianti, et al.)

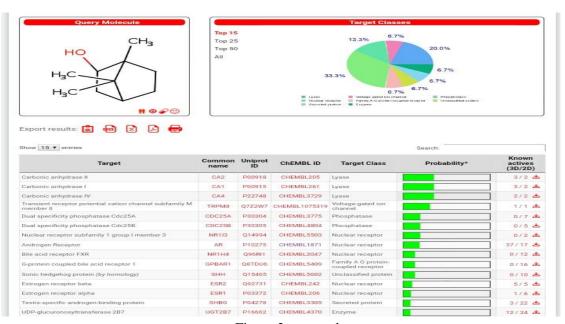


Figure 2. target classes (source : courtesy of Sindy Afrianti, *et al.*)

5. CONCLUSION

The conclusion from the results of bioinformatic research on white turmeric extract (curcuma zedoria.) Can be used as an anti-breast cancer drug.

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