# NOVEL DESIGN OF CALANONE DERIVATIVES AS ANTI-LEUKEMIA COMPOUNDS BASED ON QUANTITATIVE STRUCTURE-ACTIVITY RELATIONSHIP ANALYSIS

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Received March 19, 2010; Accepted July 29, 2010

# ABSTRACT

Leukemia drug discovery based on calanone compound was conducted in previous research and produced 6 calanone derivatives. Most of them have lower activities against leukemia cell L1210 than pure calanone. A Quantitative Structure-Activity Relationship (QSAR) analysis is conducted in this work to find more active calanone derivatives. Six compounds were used as the material of the research because they already have anti-leukemia activity data expressed in Inhibitory Concentration of Fifty Percent Cell Lethal (IC<sub>50</sub>, in  $\mu$ g/mL). Calculation of predictors was performed by AM1 semiempirical method. QSAR equation is determined using Principle Component Regression (PCR) analysis, with Log IC<sub>50</sub> as dependent variable. Independent variables (predictors) are atomic net charges, dipole moment ( $\mu$ ), and coefficient partition of n-octanol/water (Log P). This work recommends 3 novel designs of calanone derivatives that may have higher activities (in  $\mu$ g/mL) than those already available, i.e. gemdiol calanone (57.78), 2,4-dinitrophenylhydrazone calanone (30.94) and 2,4,6-trinitrophenylhydrazone calanone (18.96).

Keywords: calanone, QSAR analysis, AM1 semiempirical calculation, leukemia drug

### INTRODUCTION

Calanone is a natural product that is isolated from Calophyllum sp. It has anti-leukemia activity of 59.4  $\mu$ g/mL against leukemia cell L1210. The development of this compound has been conducted. The previous study produced 6 calanone derivatives but only one derivative has higher activity than pure calanone [1]. This problem has to be solved in order to guide novel calanone derivative discovery and to decrease "trial and error" synthesis.

One of computational chemistry applications is Quantitative Structure-Activity Relationship (QSAR) analysis for drug design. QSAR analysis is a combination of computational chemistry and statistical calculation [2]. QSAR analysis was used in some works for designing new compounds such as novel fluoroquinolone derivative as antibacterial compound [3] and antiradical drug based on flavone/flavonol compound [4].

Because of statistical requirement, QSAR analysis commonly needs large number of compounds and predictors especially in Multiple Linear Regression (MLR) calculation. Another regression analysis for small number of compounds with large predictors, called Principle Component Regression (PCR) [5]. Before PCR calculation, Principle Component Analysis (PCA) calculation is conducted to produce principle components. Afterward, regression analysis for obtaining QSAR equation is conducted against the latent variables. QSAR analysis with PCR calculation has been applied on QSAR study of benzalacetone compounds as anti-mutagen drugs [5].

In this paper, QSAR analysis against six experimental data of calanone compounds as antileukemia drugs is performed. PCR statistical calculation is applied in this work. This research is conducted in several steps such as: modeling calanone and its derivatives in three dimensional structures then calculating the theoretical predictors. After that, constructing QSAR equation and designing new calanone compound.

#### **EXPERIMENTAL SECTION**

## Materials

This research is a theoretical study that uses molecular model of calanone compound. All 3Dcartesian coordinates of calanone and its derivatives are

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Table 1. I redictors and now to obtain them						
Number	Symbol	Predictor	Unit	How to obtain		
1	Q	Atomic net charges of the compound	coulomb	Optimizing the structure of the compound by AM1 semiempirical calculation		
2	μ	Dipole moment	debye	Optimizing the structure of the compound by AM1 semiempirical calculation		
3	Log P	Partition coefficient	-	QSAR Properties calculation		

**Table 1.** Predictors and how to obtain them





used in computational chemistry calculation to produce predictors. Six anti-leukemia activities of calanone and its derivatives against leukemia cell L1210 are applied to statistical calculation as dependent variable (see Fig. 1). Anti-leukemia activities data are obtained from previous experimental study [1].

#### Instrumentation

Computational chemistry calculation is conducted at Austrian-Indonesian Centre for Computational Chemistry (AIC), Universitas Gadjah Mada, Indonesia and Physical Chemistry Laboratory, Universitas Jenderal Soedirman, Indonesia. Calculations are performed on 1 personal computer (PC), Intel Pentium IV 2.66 GHz processor, 1 GB RAM, and 40 GB *Harddisk Drive*. Softwares are Hyperchem ver 7.0 (calculating electronic and lipophylic predictor) and SPSS ver 13.0 (statistical analysis).

#### Procedure

#### Molecular Modeling of Calanone Derivatives

Each compound is drawn into a three dimensional (3D) model of compound. Then calculations of parameters are set up for each model.

#### **Generating Molecular Predictors**

Predictors can be generated from experimental data or theoretical data. The predictors in this research are theoretical predictors that are generated by computational chemistry calculation. They consist of the physicochemical properties of calanone and its derivatives that are used as independent variables in statistical analysis. They are assumed as an important factor on reaction mechanism and physical properties of drug such as atomic charges, dipole moment, and partition coefficient. Table 1 is the details of theoretical molecular predictors. Dependent variable in this study is biological activity of the compounds.

In the case of atomic charge calculation, only atoms in the main structure of Calanone are used as predictors (see Fig. 2). Additional parameters of computational calculation are RHF spin pairing, lowest state and 0.001 kcal.mole<sup>-1</sup> of RMS gradient.

# Constructing QSAR Equation and Designing the New Compound

Determination of QSAR equation is started with PCA calculation against the theoretical predictors. After that, PCR calculation is conducted to PCA results where the Log  $IC_{50}$  as dependent variable and all of



Fig 2. Numbering of atom in the main structure of calanone [9]

predictors as independent variables. By looking to the quality of statistical parameters, the selected QSAR equation is chosen. The general form of QSAR equation is described below:

$$A = C + b_1 x_1 + b_2 x_2 + b_3 x_3 + \dots$$

Where: A = compound's activities

C = a constant $x_n = n^{th}$  variable =  $n^{th}$  predictor  $b_n = a$  coefficient of  $n^{th}$  predictor

After obtaining QSAR equation, the design of new calanone derivative was then carried out on the basis of some following considerations such as: the most influenced predictor to the activity and the ease of synthesis procedure for the new compound. So, with this consideration, the more active new calanone compound can be produced.

## **RESULT AND DISCUSSION**

## **Molecular Modeling**

Molecular modeling is important for describing the chemical system. The detailed checking of 3D structure of the model is also important. If the structure is not correctly represented, the calculation of predictors would also give wrong results. Modeling chemical system is one of important things besides choosing calculation method in computational chemistry study [2]. The result from modeling step is a 3D coordinate of the compound. This coordinate is then applied in predictor calculation.

#### **Generating Molecular Predictors**

Predictors are generated by optimization of calanone structure based on AM1 semiempirical calculation. The advantages using of AM1 semiempirical method are that the calculation is not time consuming, electronic structure data can be easily obtained and the method is suitable for bulky structure of organic compound such as calanone [2]. AM1 semiempirical calculation method is already applied in QSAR analysis of antibacterial fluoroquinolone compounds and gives a good compromising of guality QSAR equation and the need of computer specification [3]. The selected predictors are the physicochemical properties which influence the anti-leukemia activity of calanone compounds. Table 2 shows the predictors data obtained from calculation.

Table 2 shows the theoretical characteristics of calanone compounds. Atomic net charges and dipole moment describes the electronic properties, where as partition coefficient reflects the solubility property of compounds. In the atomic net charges value, the most significant difference is observed in the 19<sup>th</sup> C element (C<sub>19</sub>). These significant differences in atomic net charge are due to the difference in functional group of each calanone compound. For instance, atomic net charges of C<sub>19</sub> in oxym calanone and isobutyl calanone are -0.005 and -0.002 Coulomb, respectively. This fact is caused by the existence of functional group on their  $C_{19}$  that pushed the electron density to the  $C_{19}$ , such as alkyl functional group [6]. In contrast, other compounds have positive value of C<sub>19</sub> atomic net charges, because they have functional groups that pull electron density from C<sub>19</sub>. All of the obtained predictors are applied for constructing the QSAR equation.

Dipole moment values are also different among the compounds. The highest value is observed in benzoylphenylalanine ester calanone which has a lot of electronegative atom in C<sub>19</sub> functional group. Dipole moment value is influenced by the difference in electronegativity among the atoms and the geometry of certain functional group in the structure [6].

Solubility property is described by the Log P The highest Log P value is found in value. calanyloctanoic (see Table 2). It is an ester compound with long hydrocarbon chain that can contribute non polar property to whole structure [6]. Hydrocarbon compounds have non polar character and easily dissolved in non polar solvents such as n-hexane and carbon tetrachloride [6].

Compound		Name		qC <sub>1</sub>	qC <sub>2</sub>	qC₃	qC <sub>4</sub>	qC₅
1	Calanone			-0.134	-0.106	-0.052	-0.089	-0.138
2	Calanol			-0.139	-0.084	-0.053	-0.107	-0.136
3	Calanyloc	Calanyloctanoic			-0.085	-0.056	-0.106	-0.135
4	Oxym cala	Oxym calanone			-0.085	-0.052	-0.107	-0.135
5	Benzoylph	Benzoylphenylalanine ester			-0.107	-0.056	-0.085	-0.138
6	Isobutvlcalanone			-0.138	-0.089	-0.050	-0.106	-0.134
Compound	$qC_6$	qC7	qC <sub>8</sub>	qC <sub>9</sub>	<b>qC</b> <sub>10</sub>	<b>qC</b> 11	qC <sub>12</sub>	qC <sub>13</sub>
1	-0.121	0.114	-0.216	0.219	0.343	-0.264	0.229	-0.215
2	-0.120	0.121	-0.207	0.192	0.346	-0.272	0.214	-0.207
3	-0.119	0.122	-0.209	0.190	0.345	-0.272	0.219	-0.207
4	-0.121	0.113	-0.207	0.203	0.341	-0.266	0.216	-0.203
5	-0.119	0.124	-0.206	0.198	0.348	-0.273	0.213	-0.196
6	-0.122	0.113	-0.201	0.186	0.341	-0.265	0.202	-0.192
Compound	qC <sub>14</sub>	qC <sub>15</sub>	qC <sub>16</sub>	qC <sub>17</sub>	qC <sub>18</sub>	qC <sub>19</sub>	qC <sub>20</sub>	qC <sub>21</sub>
1	0.234	-0.305	0.124	-0.221	-0.064	0.342	-0.136	-0.066
2	0.228	-0.228	0.124	-0.224	-0.062	0.121	-0.124	-0.107
3	0.232	-0.276	0.124	-0.221	-0.064	0.127	-0.100	-0.115
4	0.209	-0.194	0.124	-0.221	-0.065	-0.005	-0.084	-0.110
5	0.198	-0.230	0.123	-0.217	-0.067	0.126	-0.121	-0.127
6	0.175	-0.191	0.123	-0.219	-0.067	-0.002	-0.077	-0.149
Compound	qC <sub>22</sub>	qC <sub>23</sub>	$qC_{24}$	qC <sub>25</sub>	$qC_{26}$	qC <sub>27</sub>	qO <sub>1</sub>	qO <sub>2</sub>
1	-0.141	-0.098	-0.143	-0.082	-0.225	-0.220	-0.191	-0.194
2	-0.127	-0.124	-0.128	-0.131	-0.218	-0.226	-0.209	-0.198
3	-0.129	-0.122	-0.129	-0.112	-0.219	-0.226	-0.217	-0.196
4	-0.135	-0.116	-0.134	-0.092	-0.219	-0.225	-0.185	-0.196
5	-0.130	-0.118	-0.133	-0.092	-0.226	-0.218	-0.169	-0.196
6	-0.130	-0.130	-0.128	-0.109	-0.217	-0.227	-0.199	-0.196
				lan D				
Compound	qO <sub>3</sub>	qO <sub>4</sub>	μ	log P	Notes: q = atomic net charges (coulomb) µ = dipole moment (debye) log P = partition coefficient			
1	-0.282	-0.244	5.856	0.43				liomb)
2	-0.297	-0.244	4.066	0.42				
3	-0.292	-0.236	5.558	2.32				
4	-0.288	-0.238	5.258	1.08				
5	-0.337	-0.242	8.789	1.24				
6	-0 296	-0 255	4 962	1.84				

Table 2. Theoretical predictors calculated by AM1 semiempirical method

# Construction of QSAR Equation and Design of the New Compound

# PCA and PCR

The predictors that are used for constructing QSAR equation are quite a lot. In contrast, the experimental data e.g. activity of the compound are too few. To solve this problem PCR calculation method is applied. Before PCR calculation was done, PCA calculation was

conducted on predictor data in order to reduce the amount of data and produce principle components. The PCA result shows that first 4 principle components (t) can be applied in regression analysis. The PCA result and the calculation of 1<sup>st</sup> until 4<sup>th</sup> latent variables for each calanone compound are given in Table 3 and 4.

The first 4 latent variables are applied because they can describe over 95% the variation in the predictors [7]. All of data in Table 4 are used in PCR

Table 3. Results of PCA calculation						
Component -	Initial Eigenvalues					
Component	Total	% of Variance	Cumulative %			
1	16.7	50.5	50.5			
2	8.9	26.9	77.4			
3	4.0	12.2	89.6			
4	2.1	6.3	95.9			
5	1.3	4.1	100.0			

Table 4. Latent variables as input of regression analysis

Compound	t <sub>1</sub>	t <sub>2</sub>	t <sub>3</sub>	t4	log IC <sub>50</sub>
1	3.5833	4.1100	3.0538	1.1619	1.7738
2	2.3393	2.7112	2.4796	0.7910	1.8451
3	1.9821	4.3197	3.2661	2.4024	1.8762
4	2.3717	3.9066	2.8880	1.5401	1.9222
5	3.9638	6.8895	4.2145	2.3246	1.7959
6	1.6918	3.8990	2.8962	1.9611	2.0233



Fig 3. The structure of new calanone derivatives. Their theoretical activities are (1) = 57.78  $\mu$ g/mL, (2) = 30.94  $\mu$ g/mL and (3) = 18.96  $\mu$ g/mL [9].

analysis for constructing the QSAR equation.

## Selecting the best QSAR equation

Selection of the best QSAR equation is conducted with evaluation of the statistical parameters such as r(correlation coefficient),  $r^2$  (determination coefficient), F(variance analysis), SE (standard error) and PRESS (Prediction Residual Sum of Squares). The QSAR equation from the PCR analysis is:

Log IC<sub>50</sub> =  $2.8055 - 0.1135 t_1 + 0.2678 t_2 - 0.5696 t_3$ 

Some characteristics of the equation are as follows: *n* (the number of data) = 6, *r* = 0.991,  $r^2 = 0.982$ , SE = 0.0193, and  $F_{calc}/F_{table} = 1.987$  and *PRESS* = 7.49 10<sup>-4</sup>. The symbol  $t_1 - t_3$  are latent variables.

This QSAR equation has good correlation (*r* value) between activities and latent variables and can explain over 98% variations of data (see  $r^2$  value). Small error is also showed in this QSAR equation (see *SE* value). It is also accepted at 95% confidence level because the  $F_{calc}/F_{table}$  is more than 1.00.

## Designing the new compounds

There are 3 factors that are used in the selection of the new calanone derivative candidates, i.e. theoretical activity obtained from QSAR equation calculation, the possibility in synthesizing the new compound and the ease to get the reactant for synthesis [9]. Based on the prediction calculation of their activity, some new compounds which may have better anti-leukemia activity are proposed e.g. gemdiol calanone (1), 2,4-dinitrophenylhydrazone calanone (2) and 2,4,6-trinitrophenylhydrazone calanone (3) (see Fig. 3).

These three candidates could guide the synthesis procedures of new candidates of calanone derivatives become more focus. All the synthesis works can be started by using pure calanone that produced at Chemistry Research Center, The Indonesian Institute of Sciences (LIPI), Serpong, Indonesia.

# CONCLUSION

QSAR analysis can be conducted for the small number of compounds but possessing large predictors using PCR method as regression analysis. Designing 3 more active candidates based on QSAR equation could focus the synthesis and prevent "trial and error" process. Therefore, high cost synthesis of novel calanone derivative could be avoided. More study is needed in order to synthesis the recommended compounds and to increase the theory level of calculation method in obtaining predictor data.

## ACKNOWLEDGEMENT

This work was supported by Basic Research Incentive Program, The Ministry of Research and Technology Republic of Indonesia, Chemistry Research Center of the Indonesian Institute of Sciences (LIPI) and also Austrian-Indonesian Centre for Computational Chemistry (AIC), Chemistry Department, Universitas Gadjah Mada.

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