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## THEORETICAL & COMPUTATIONAL CHEMISTRY | RESEARCH ARTICLE

# Prediction of anticancer activities of cynaroside and quercetin in leaf of plants *Cynara scolymus* L and *Artocarpus incisa* L using structure–activity relationship

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**Abstract:** Natural products from plants are an alternative resource in the search for anti-cancer drugs and can have a direct impact on eliminating cancer cells and also reduce cancer side effects. Recently, we have isolated a few flavonoid quercetin and cynaroside from leaf of *cynara scolymus* L and *artocarpus incisa* L in Vietnam, with cytotoxic activity relatively strong in Hela cancer cells. The flavonoid compound is a search target, research and development of anti-cancer agents in clinical use. To clarify the important nature of the activity, the subject QSAR studies on cancer Hela cell line use the multiple linear regression (MLR) gradually, partial least square regression (PLS) and artificial neural network. The MLR and PLS models showed good correlation values of  $R^2 = 0.938$ ,  $R^2_{\text{pred}} = 0.903$ , and  $R^2 = 0.943$ ,  $R^2_{\text{pred}} = 0.912$ , respectively. The MLR model shows the level of importance of atomic charge descriptors. Also, artificial neural network architecture I(6)-HL(4)-O(1) is built with RMSE = 0.00345,  $R^2 = 0.993$ ,  $R^2_{\text{pred}} = 0.971$  using the atomic charge descriptors selected in the MLR model such as neurons of input layer and the anti-cancer activity such as neuron of output layer. The anti-cancer activities of the flavonoids and isoflavonoids in the test group and

### ABOUT THE AUTHORS

Scientific research for us is important and necessary in order to improve and expand the knowledge. Since 1990s, our role of scientific research had become to be important in an university, so we had some projects in QSAR field as:

Studying the process of extracting and purifying the total alkaloid extract from *Dichroa febrifuga* Lour and the screening process for alkaloid from the extract of leaves of *Dichroa febrifuga* Lour. Using artificial neural network for screening process of alkaloids from *Dichroa febrifuga* Lour. During this duration, we had a number of scientific works in the field of computer applications in chemistry. For example, the QSAR in design for new drugs.

In 1990s, we had a project of Ministry of Education and Training in the field of computer applications in the chemistry with the topic: Study of quantitative relationship between the structure and activity of the group antimalarial compounds, anti-cancer, anti-HIV, anti-fungal, and anti-bacterial.

### PUBLIC INTEREST STATEMENT

The studies for *Cynara scolymus* L showed that its ingredients from flowers, leaves, stems, roots are very effective in healing and for food. The flavonoid compounds were extracted from *Cynara scolymus* L working in the treatment of some diseases such as liver, bile, cardio, antioxidants and reduce cholesterol in blood, especially HIV anti-virus (Loi, 2006).

In Vietnam, *Artocarpus Incisa* L scattered only has been planted in the orchard of the Vietnamese family. *Artocarpus Incisa* L is the kind of big trees. The compound groups in leaves *Artocarpus Incisa* L were determined quantitatively; the results showed that the leaves of *Artocarpus Incisa* L contain substances: flavonoids, saponins, anthranoid, tannin, reducing sugar, acid amines and polysaccharides. The water extract from the leaves of *Artocarpus Incisa* L showed that the blood pressure is lowered and decreased heart rate in mice. The water extract from leaves of *Artocarpus Incisa* L effects on cancer cells of the pancreas (Loi, 2006).



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compounds *quercetin* and *cynaroside* isolated from *cynara scolymus* L and *artocarpus incisa* L are compared with experimental data and those from references.

**Subjects: Computational and Theoretical Chemistry; Medicinal & Pharmaceutical Chemistry; Organic Chemistry**

**Keywords: QSAR<sub>MLR</sub> and QSAR<sub>PLS</sub> model; neural network QSAR<sub>ANN</sub> model; anticancer activities HeLa**

## 1. Introduction

Natural products from plants are of interest in searching for new anti-cancer drugs and can have a direct effect on HeLa cancer cells and reduce side effects. Recently, we have isolated a few flavonoids from leaf of *cynara scolymus* L and *artocarpus incisa* L (Loi, 2006) and tested their *in vitro* activities pointed out the relatively strong impacts for cancer cells HeLa (Singh, Kaur, & Silakari, 2014). These flavonoids from leaf of *cynara scolymus* L (Apóstolo, Brutti, & Llorente, 2005; Fritsche, Beindorff, Dachtler, Zhang, & Lammers, 2002; Zhu, Zhang, & Lo, 2004) and *artocarpus incisa* L (El Senousy, Farag, Al-Mahdy, & Wessjohann, 2014) were also tested biologically in the treatment of some diseases such as liver, bile, cardio, antioxidants and reduce cholesterol in blood, especially HIV anti-virus (Abbasi & Samadi, 2014; Moreira, Castelo-Branco, Monteiro, Tavares, & Beltramini, 1998). Flavonoids are polyphenolic compounds in most plants (Mahapatra, Bharti, & Asati, 2015; Priyadarsini et al., 2010; Ziberna, Fornasaro, Čvorović, Tramer, & Passamonti, 2014). The flavonoids have been shown their activities and role of food within flavonoids in the cancer inhibition are widely studied (Gavin & Durako, 2012; Lee, Boyce, & Breadmore, 2012; Pawlikowska-Pawłęga et al., 2014).

In recent years, the methods of quantum chemistry calculations are widely applied to the study of chemical properties and seeking new drugs. The field of new drug design by computer tools has become an important tool nowadays. Study on quantitative relationships between structure and activity (QSAR) of natural compounds is of concern for new drug researchers and pharmaceutical manufacturing facilitators. In Vietnam, there are a number of works of scientists from universities and institutions published in the journal (Phuong Thuy & Tat, 2012a, 2012b). The previous studies of 3-aminoflavonoid substances have focused on the basis of semi-empirical calculation (Tat, 2009a). These studies have shown a way for designing new drugs efficiently with the assistance of computers. The QSAR model can be predicted the biological activity of new drugs from the atomic charges in the molecule. This method allows for the identification of an active-central location of molecule.

The set of flavones and isoflavones is known to have an important activity against cervical cancer cells (Chen, 2008; Liao et al., 2005; Liao, Chen, Qian, Shen, & Zheng, 2008). This group is currently of interest for researching in different directions such as the synthesis and metabolizing of natural products or extracting them from plants (Loi, 2006). The consideration of the quantitative relationship between the structure of flavones and isoflavones with activity against cancer is an important issue in searching for the flavone and isoflavone derivatives to be valid.

In this work, we report the use of semi-empirical quantum calculations and construction of quantitative structure-activity relationship (QSAR) models using 32 flavone and isoflavone derivatives (Chen, 2008; Liao et al., 2005, 2008). The flavones and isoflavones are constructed and optimized by means of molecular mechanics MM+. The atomic charge descriptors resulting from Parametric Model number 3 (PM3) method are used to build the multivariate QSAR models such as multiple linear regression (MLR), partial least squares regression (PLS), and artificial neural network (ANN). Anti-cancer activities  $GI_{50}/\mu\text{M}$  of flavones and isoflavones in the test group and the new flavonoids quercetin and cynaroside isolated from the leaves of *cynara scolymus* L and *artocarpus incisa* L are predicted from QSAR models and compared with those from experimental data.

## 2. Computational details

### 2.1. Materials and means

To ensure the accurate level of QSAR models, structural data and anticancer activities  $GI_{50}/\mu\text{M}$  for HeLa cells ( $GI_{50}$  is the concentration for 50% of maximal inhibition of cell proliferation) for flavones and isoflavones are taken from the data source of Wang and et al. (Chen, 2008; Liao et al., 2005, 2008) as pointed out in Figure 1 and Table 1. The anti-cancer activities are transformed into following value  $pGI_{50}$

$$pGI_{50} = -\log GI_{50} \quad (1)$$

The atomic charge parameters on molecules are calculated by means of program HyperChem v8.0 (HyperChem Release 8.03, 2008). The multiple linear regression ( $QSAR_{MLR}$ ) and the partial least squares regression  $QSAR_{PLS}$  models are built with program Origin 2015 (Tat, 2009b). The artificial neural network ( $QSAR_{ANN}$ ) models are constructed with Visual Gene Developer 1.7 (Jung & McDonald, 2011).

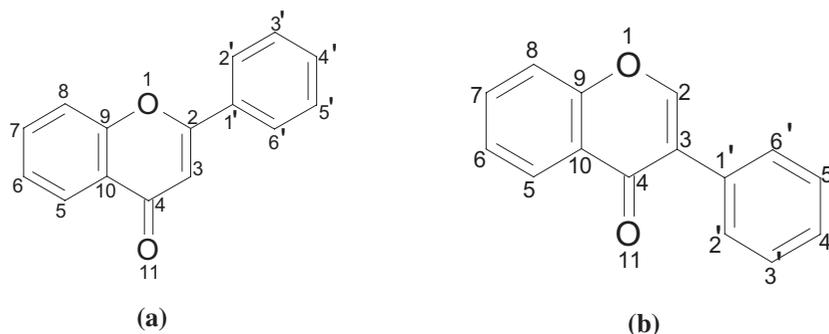
### 2.2. Isolated technology of quercetin and cynaroside

#### 2.2.1. Chemicals and equipment

In this work, we use the chemicals and the equipment for isolating and purifying two flavonoids quercetin and cynaroside before determining the substance structures by  $^1\text{H-NMR}$  and  $^{13}\text{C-NMR}$  spectrum:

- Silica gel with the particle size in range 0.04–0.06 mm was used for ordinary and Rp18 phase chromatography.
- Thin-layer chromatography was implemented by the thin plate DC-Alufolien F254 (Merck) for the ordinary phase and Rp18 F254s (Merck) for the reverse-phase chromatography.
- Solvents were used for the isolation processes: hexane, petroleum ether, chloroform, methanol, ethyl acetate, ethanol, acetone, distilled water.
- Reagent was used to trace out the compound coloration on plate: using  $\text{H}_2\text{SO}_4/\text{EtOH}$ ;  $\text{FeCl}_3/\text{EtOH}$ .
- UV handheld lamps, 254 and 365 nm UVITEC effect.
- Vacuum Evaporators Buchi-111.
- Water Bath cooker JULABO 461.
- Infrared heating equipment SCHOTT.
- Chromatography column with diameter range 2–5.5 cm.
- Analytical Balances AND HR 200.

Figure 1. Molecular skeleton: (a) flavone and (b) isoflavone.



**Table 1. Molecular structure and anticancer activities  $GI_{50}$  ( $\mu\text{M}$ ) of flavones and isoflavones (Chen, 2008; Liao et al., 2005, 2008).**

Substance	Name	Substitutive site	Substitutes R	$GI_{50}$ ( $\mu\text{M}$ )	
<b>Training set for establishing QSAR models</b>					
Fla1	1a-1	Flavone	$C_3$	$-\text{OCH}_2\text{CCH}_3=\text{NOH}$	2,0
Fla3	2a-3	Flavone	$C_7$	$-\text{OCH}_2\text{CCH}_3=\text{NOH}$	2,0
Isofla4	3a-4	Isoflavone	$C_7$	$-\text{OCH}_2\text{CCH}_3=\text{NOH}$	9,8
Fla5	4a	Flavone	$C_3$	$-\text{OCH}_2\text{CCH}_3=\text{NOCH}_3$	2,0
Fla6	5a	Flavone	$C_3$	$-\text{OCH}_2\text{CCH}_3=\text{NOCH}_3$	0,9
Fla7	6a	Flavone	$C_7$	$-\text{OCH}_2\text{CCH}_3=\text{NOCH}_3$	2,2
Isofla8	7a	Isoflavone	$C_7$	$-\text{OCH}_2\text{CCH}_3=\text{NOCH}_3$	8,5
Fla10	8a	Flavone	$C_3$	$-\text{OCH}_2\text{C}(4-\text{F}-\text{C}_6\text{H}_4)=\text{NOH}$	2,1
Fla11	9a	Flavone	$C_3$	$-\text{OCH}_2\text{C}(4-\text{CH}_3\text{O}-\text{C}_6\text{H}_4)=\text{NOH}$	2,0
Fla13	10a	Flavone	$C_6$	$-\text{OCH}_2\text{C}(4-\text{F}-\text{C}_6\text{H}_4)=\text{NOH}$	1,6
Fla14	11a	Flavone	$C_3$	$-\text{OCH}_2\text{C}(4-\text{CH}_3\text{O}-\text{C}_6\text{H}_4)=\text{NOH}$	1,0
Fla17	12a	Flavone	$C_7$	$-\text{OCH}_2\text{C}(4-\text{CH}_3\text{O}-\text{C}_6\text{H}_4)=\text{NOH}$	2,0
Isofla18	13a	Isoflavone	$C_7$	$-\text{OCH}_2\text{C}(\text{C}_6\text{H}_5)=\text{NOH}$	9,0
Isofla19	14a	Isoflavone	$C_7$	$-\text{OCH}_2\text{C}(4-\text{F}-\text{C}_6\text{H}_4)=\text{NOH}$	7,8
Isofla20	15a	Isoflavone	$C_7$	$-\text{OCH}_2\text{C}(4-\text{CH}_3\text{O}-\text{C}_6\text{H}_4)=\text{NOH}$	7,6
Fla21	16a	Flavone	$C_3$	$-\text{OCH}_2\text{C}(\text{C}_6\text{H}_5)=\text{NOCH}_3$	1,6
Fla22	17a	Flavone	$C_3$	$-\text{OCH}_2\text{C}(4-\text{F}-\text{C}_6\text{H}_4)=\text{NOCH}_3$	2,0
Fla23	18a	Flavone	$C_3$	$-\text{OCH}_2\text{C}(4-\text{CH}_3\text{O}-\text{C}_6\text{H}_4)=\text{NOCH}_3$	2,0
Fla24	19a	Flavone	$C_6$	$-\text{OCH}_2\text{C}(\text{C}_6\text{H}_5)=\text{NOCH}_3$	2,4
Fla25	20a	Flavone	$C_6$	$-\text{OCH}_2\text{C}(4-\text{F}-\text{C}_6\text{H}_4)=\text{NOCH}_3$	2,3
Fla26	21a	Flavone	$C_6$	$-\text{OCH}_2\text{C}(4-\text{CH}_3\text{O}-\text{C}_6\text{H}_4)=\text{NOCH}_3$	2,0
Fla27	22a	Flavone	$C_7$	$-\text{OCH}_2\text{C}(\text{C}_6\text{H}_5)=\text{NOCH}_3$	6,6
Fla28	23a	Flavone	$C_7$	$-\text{OCH}_2\text{C}(4-\text{F}-\text{C}_6\text{H}_4)=\text{NOCH}_3$	2,7
Fla29	24a	Flavone	$C_7$	$-\text{OCH}_2\text{C}(4-\text{CH}_3\text{O}-\text{C}_6\text{H}_4)=\text{NOCH}_3$	2,5
Isofla30	25a	Isoflavone	$C_7$	$-\text{OCH}_2\text{C}(\text{C}_6\text{H}_5)=\text{NOCH}_3$	8,2
Isofla31	26a	Isoflavone	$C_7$	$-\text{OCH}_2\text{C}(4-\text{F}-\text{C}_6\text{H}_4)=\text{NOCH}_3$	6,4
<b>Test set for validating QSAR models</b>					
Fla2	1b	Flavone	$C_6$	$-\text{OCH}_2\text{CCH}_3=\text{NOH}$	1,2
Fla9	2b	Flavone	$C_3$	$-\text{OCH}_2\text{C}(\text{C}_6\text{H}_5)=\text{NOH}$	1,8
Fla12	3b	Flavone	$C_6$	$-\text{OCH}_2\text{C}(\text{C}_6\text{H}_5)=\text{NOH}$	0,8
Fla15	4b	Flavone	$C_7$	$-\text{OCH}_2\text{C}(\text{C}_6\text{H}_5)=\text{NOH}$	2,0
Fla16	5b	Flavone	$C_7$	$-\text{OCH}_2\text{C}(4-\text{F}-\text{C}_6\text{H}_4)=\text{NOH}$	2,0
Isofla32	6b	Isoflavone	$C_7$	$-\text{OCH}_2\text{C}(4-\text{CH}_3\text{O}-\text{C}_6\text{H}_4)=\text{NOCH}_3$	7,3

**Figure 2. Separate equipment for two flavonoids quercetin and cynaroside.**



a) Vacuum Evaporators

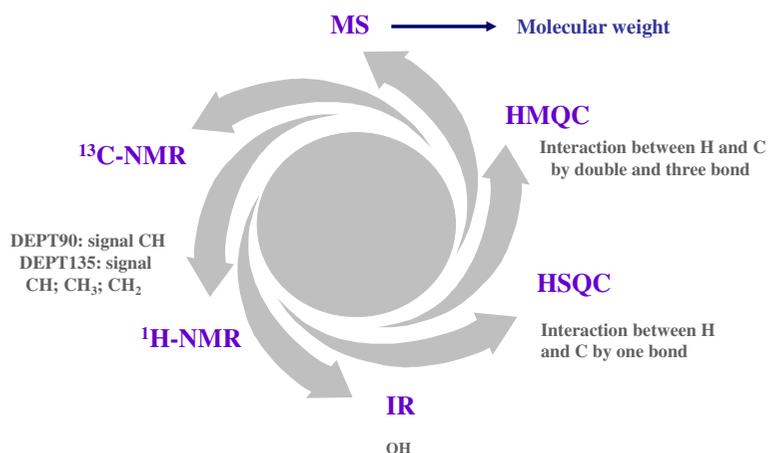


b) Column chromatography at atmospheric and high pressure



c) Thin-layer chromatography

**Figure 3. General diagram of structure identification for substances quercetin and cynaroside.**



### 2.2.2. Isolation and identification of flavonoids

To isolate and purify the flavonoid compounds, we used the techniques of thin-layer and column chromatography, as exhibited in Figure 2. After isolating the compounds, they were identified with the structure using different spectrums:

- Melting temperature carried out on Electrothermal IA 9000 series, using unadjusted capillary.
- Column chromatography with silica gel for ordinary-phase, reverse-phase chromatography Rp 18 and Sephadex techniques combined with thin-layer chromatography.
- Substances were detected by ultraviolet light at wavelengths 254 and 365 nm or reagent used is liquid H<sub>2</sub>SO<sub>4</sub>/EtOH or FeCl<sub>3</sub>/EtOH.
- Nuclear magnetic resonance spectrum (NMR) <sup>1</sup>H-NMR (500 MHz) and <sup>13</sup>C-NMR (125 MHz) implemented on Bruker AM500 FT-NMR Spectrometer (Figure 3).

### 2.3. Constructing QSAR models

The fitness models were proved using different validations. To validate QSAR models, the method was carried out: (1) leave-one-out cross-validation technique, (2) validation was implemented by dividing randomly the 32 compounds into training and test group. The developed model should be capable enough making accurate and reliable predictions of anticancer activities of new substances. So, the QSAR models that are constructed from a training set should be validated using the divided compounds in test group for testing the predictability of the developed models. The validation methods depicted the reliability of the developed models for their applicability on a group of new compounds, and confident level of predictability can thus be resolved (Tat, 2009). For all the new-constructed models, the multiple-determined coefficient ( $R^2$ ) and leave-one-out cross-validation value ( $R_{pred}^2$ ) for the training set were evaluated by (OriginLab Corporation, 2015; Tat, 2009). However, additionally, test set was used for calculation of ( $R_{pred}^2$ ) values. The acceptability and the predictability of the model are determined based on statistical parameters such multiple-correlation coefficient ( $R^2$ ) of training group and test group and cross-validation coefficient ( $R_{pred}^2$ ), Fisher value ( $F$ ), and standard deviation (SD). Multiple linear models were set (Tat, 2009).

$$Y = \sum_{i=1}^k a_i x_i + b \quad (2)$$

where  $Y$  is the biological activity  $pGI_{50}$  (the dependent variable),  $x_i$  is the atomic charge parameters (independent variable). Check the results from the model with experimental data based on single-factor analysis of variance (ANOVA). The correlation coefficient  $R^2$ , absolute relative error (ARE, %) and mean of absolute relative error (MARE,%) were calculated according to the Equation (3-5) (Tat, 2009).

$$R^2 = 1 - \frac{\sum_{i=1}^N (Y_i - \hat{Y}_i)^2}{\sum_{i=1}^N (Y_i - \bar{Y})^2} \quad (3)$$

Here  $Y$  is  $\text{pGI}_{50}$  experimental value and  $\hat{Y}$  is  $\text{pGI}_{50}$  predicted value;  $\bar{Y}$ :  $\text{pGI}_{50}$  is mean value; The statistical error values ARE,% are determined by

$$\text{ARE, \%} = 100 |(\text{pGI}_{50,\text{exp}} - \text{pGI}_{50,\text{pred}}) / \text{pGI}_{50,\text{exp}}| \quad (4)$$

where  $\text{pGI}_{50,\text{exp}}$  and  $\text{pGI}_{50,\text{pred}}$  are experimental and prediction activities. The average value of absolute relative error MARE,% is used to assess the global uncertainty of QSAR model using the formula:

$$\text{MARE, \%} = \frac{100}{N} \left| \frac{(\text{pGI}_{50,\text{exp}} - \text{pGI}_{50,\text{pred}})}{\text{pGI}_{50,\text{exp}}} \right| \quad (5)$$

with  $N$  as number of activity values.

### 3. Results and discussion

#### 3.1. Calculation of charge parameters

The molecular structures were built and optimized by means of MM+ molecular mechanics. The atomic charge parameters on molecules were calculated by semi-empirical quantum chemistry method SCF PM3 using the optimized molecules. The program HyperChem 8.05 (HyperChem Release 8.03, 2008) was used for all these calculations. The atomic charge parameters were used to build the multiple linear regression (QSAR<sub>MLR</sub>), partial least squares regression (QSAR<sub>PLS</sub>), and artificial neural network (QSAR<sub>ANN</sub>) model.

#### 3.2. Constructing QSAR<sub>MLR</sub> and QSAR<sub>PLS</sub> model

Before conducting the QSAR<sub>MLR</sub> and QSAR<sub>PLS</sub> modeling, the activity values  $\text{GI}_{50}$  ( $\mu\text{M}$ ) are transformed into the values  $\text{pGI}_{50}$  to adapt the statistical properties. The activity values  $\text{pGI}_{50}$  ( $\mu\text{M}$ ) are the most appropriate value. The QSAR<sub>MLR</sub> models were established using the relationship of the atomic charge predictors and biological activities  $\text{pGI}_{50}$  (Tat, 2009). The change of values  $R^2$ ,  $R^2_{\text{pred}}$  and SE (standard error) in the QSAR<sub>MLR</sub> models with the atomic charge predictors, respectively, are pointed out in Table 2.

To have those QSAR<sub>MLR</sub> models, the atomic charge descriptors were selected using stepwise regression algorithm. The selection process for atomic charge descriptors is based on the change of

**Table 2. The QSAR<sub>MLR</sub> models (k from 1 to 10) with change of values  $R^2$ ,  $R^2_{\text{pred}}$  and SE**

$k$	The atomic charge predictors in models	$R^2$	SE	$R^2_{\text{pred}}$
2	$O_1, C_7$	0.816	0.139	0.765
3	$O_1, C_4, C_6$	0.860	0.124	0.800
4	$O_1, C_2, C_4, C_5$	0.901	0.107	0.829
5	$O_1, C_2, C_4, C_5, C_3$	0.924	0.096	0.873
6	$O_1, O_{11}, C_3, C_4, C_6, C_7$	0.938	0.089	0.903
7	$O_{11}, C_4, C_6, C_8, C_9, C_2, C_6$	0.959	0.074	0.879
8	$O_{11}, C_3, C_6, C_7, C_8, C_9, C_{10}, C_3$	0.970	0.065	0.696
9	$O_1, O_{11}, C_3, C_4, C_6, C_7, C_8, C_9, C_{10}$	0.978	0.057	0.563
10	$O_1, O_{11}, C_3, C_6, C_7, C_8, C_9, C_2, C_3, C_6$	0.978	0.059	0.358

the statistical values  $R^2$ , SE,  $R_{pred}^2$  and  $F$ -stat. The QSAR<sub>MLR</sub> models were cross-validated using leave-one-out (LOO) technique to determine  $R_{pred}^2$ . The 10 fitness models are shown in Table 2. The QSAR<sub>MLR</sub> models (with  $k$  from 2 to 10) that are arranged in an orderly change of values  $R^2$  and  $R_{pred}^2$ . From the models in Table 2, the QSAR<sub>MLR</sub> models (with  $k$  from 5 to 7) are shown the greater values of  $R_{pred}^2$  than others.

In particular, the QSAR<sub>MLR</sub> model with  $k = 6$  with value  $R^2$  of 0.938 gave the highest value  $R_{pred}^2$  of 0.903. So we selected the best models (with  $k$  of 5, 6, and 7) to determine the contribution percentage of atomic charges. The valuable contribution percentages  $MP_{m,x_k}$ %,  $GMP_{m,x_k}$ % and the statistical values of these models (with  $k$  of 5, 6, and 7), respectively, are exhibited in Table 3.

The valuable contribution percentages  $MP_{m,x_k}$ % of independent variables in each model QSAR<sub>MLR</sub> (with  $k$  of 5, 6, and 7) were determined from the contribution percentages  $P_{x_k}$ % of variables in each case, respectively (Tat, 2009). This value is determined by the total value of contribution  $C_{total}$  of variables in a substance (Chen, 2008). So the average contribution percentage  $MP_{m,x_k}$ % of each variable is defined by the formula (6) and the results are depicted in Table 3.

$$MP_{m,x_k}\% = \frac{1}{N} \sum_{j=1}^N \left( 100 \cdot \frac{|b_{m,i}x_{m,i}|}{C_{total}} \right) \text{ with } C_{total} = \sum_{j=1}^k |b_{m,k}x_{m,k}| \quad (6)$$

where  $N$  the total number of cases,  $m$  number of variables. The global average contribution percentage  $GMP_{m,x_k}$ % of each independent variable for three models is determined by the formula (7):

$$GMP_{m,x_k}\% = \frac{1}{n} \sum_{n=1}^3 MP_{m,x_k} \quad (7)$$

with  $n$  number of models.

**Table 3. Statistical values and valuable contribution percentages  $MP_{m,x_k}$ % and  $GMP_{m,x_k}$ % for atomic charges in the models QSAR<sub>MLR</sub> (with  $k$  of 5, 6, and 7)**

Variable	QSAR <sub>MLR</sub>			MP <sub>m,x<sub>k</sub></sub> %			GMP <sub>m,x<sub>k</sub></sub> %
	k = 5	k = 6	k = 7	k = 5	k = 6	k = 7	
$x_k$							
$R^2$	0.9243	0.9382	0.9589				
$R_{adj}^2$	0.9053	0.9186	0.9429				
SE	0.0957	0.0887	0.0743				
$R_{pred}^2$	0.873	0.903	0.879				
Constant	-0.9332	6.7116	4.714				
$O_1$	-101.2076	-42.3105	-	57.6024	24.6289	-	27.4104
$O_{11}$	-	-8.1592	-32.8026	-	18.6316	21.4621	13.3646
$C_2$	-15.4264	-	-	13.4176	-	-	4.4725
$C_3$	-	3.0139	-	-	4.2160	-	1.4053
$C_4$	-6.8735	-19.0370	-60.0703	15.1206	42.4467	38.3868	31.9847
$C_5$	-7.9686	-	-	2.0583	-	-	0.6861
$C_6$	-	6.6117	20.8772	-	6.5716	5.7785	4.1167
$C_7$	-	4.6038	-	-	3.5052	-	1.1684
$C_8$	-	-	16.9016	-	-	5.6960	1.8987
$C_9$	-	-	95.4205	-	-	22.5970	7.5323
$C_2$	-	-	-24.4720	-	-	2.4973	0.8324
$C_3$	-16.1166	-	-	11.8011	-	-	3.9337
$C_6$	-	-	-25.4219	-	-	3.5824	1.1941

The contribution percentages  $GMP_{m^k}$ , % in Table 3 display the important level of atomic charges in flavones and isoflavones. For 3 QSAR<sub>MLR</sub> models, the important level of atomic charges are that arranged by the values  $GMP_{m^k}$ , %:  $C_4 > O_1 > O_{11} > C_9 > C_2 > C_6 > C_3$ . The atom positions  $C_4$ ,  $O_1$ ,  $O_{11}$  are considered such as the most important positions in the molecules. Besides those atoms are in carbonyl group  $C_4 = O_{11}$  and atom  $O_1$  has free electron pair conjugating with  $\pi$  electronic bond  $C_2 = C_3$ , and  $C_4 = O_{11}$  to form a conjugate system. The carbonyl group  $C_4 = O_{11}$  has fully reactive natures of carbonyl substance. So, these important atoms are demonstrated quantitatively using the  $GMP_{m^k}$ , % values, % and this is also consistent with the verdicts from experimental evaluation (Lee et al., 2012; Liao et al., 2005, 2008). Also, the atomic position  $C_6$  is also an important position and is explored for attaching the new substitutes (Chen, 2008; Liao et al., 2005, 2008). The atomic positions  $C_9$  and  $C_3$  also represent the important impacts for biological activities  $GI_{50}$ , but the  $C_9$  atom is not vacant position so should not be selected for attaching the new substitutes. So the  $C_6$  is vacant position can be chosen to add the new substitutes to sample flavone in Table 1 or new flavonoid. Similarly, the position  $C_3$  is also empty and can be utilized to add the new substitutes. Those can hope to constitute a new compound with higher activity. From this orientation, a new flavonoid isolating from the leaf of *artocarpus incisa* L was selected such as sample substance to design new drugs with high activity. This is carried out in below discussion.

The QSAR<sub>PLS</sub> model is also built from the atomic charges, in which those were selected for the QSAR<sub>MLR</sub> model (Tat, 2009). The six variables  $O_1$ ,  $O_{11}$ ,  $C_3$ ,  $C_4$ ,  $C_6$ , and  $C_7$  are also used to build the QSAR<sub>PLS</sub> models. The present results of biological activities are depicted in  $R^2$  values in which those are consistent with experimental data. The partial least squares (QSAR<sub>PLS</sub>) model exhibited in the form:

$$Y = 5.168 - 20.643 \times O_1 - 0.358 \times C_3 - 7.892 \times C_4 + 0.425 \times C_6 - 0.583 \times C_7 - 3.465 \times O_{11} \quad (8)$$

with  $n = 26$ ;  $R^2 = 0.943$ ;  $SE = 0.360$ ;  $R^2_{pred} = 0.912$ .

### 3.3. Building QSAR<sub>ANN</sub> model

The QSAR<sub>ANN</sub> model is built by the neuro-fuzzy technique with the genetic algorithms using program Visual Gene Developer v1.7 (Jung & McDonald, 2011). The artificial neural network architecture consists of three layers I(6)-HL(4)-O(1); the input layer I(6) includes six neurons as parameters  $O_1$ ,  $O_{11}$ ,  $C_3$ ,  $C_4$ ,  $C_6$ , and  $C_7$ ; the neuron on output layer O(1) is biological activity  $pGI_{50}$ ; the hidden layer HL(4) consists of four neurons. This multi-layer neural network employing backpropagation algorithm is used to train the network. The transfer function is sigmoid on each node of the network; the neural network parameters include the training rate of 0.7 and learning rate of 0.7; the goal monitoring error  $MSE = 0.000816$  with 10,000 iteration. After training the neural network,  $R^2$  value is 0.993 and  $R^2_{pred}$  of 0.971 while for QSAR<sub>MLR</sub> model, the value  $R^2$  is 0.938 and  $R^2_{pred}$  of 0.903.

### 3.4. Prediction of biological activity for new substance

The predictability of the models QSAR<sub>MLR</sub>, QSAR<sub>PLS</sub> and QSAR<sub>ANN</sub> are evaluated carefully using the leave-one-out (LOO) technique to determine the value  $R^2_{pred}$ ; the flavonoids were divided randomly from the data in Table 1 into the training group of 26 compounds and the test group of six compounds. The biological activities of six flavonoids in the test group in Tables 1 and 2 new flavonoids isolated from the leaves of *cynara scolymus* L and *artocarpus incisa* L (Loi, 2006) are predicted from models QSAR<sub>MLR</sub>, QSAR<sub>PLS</sub>, and QSAR<sub>ANN</sub>.

The predicted values of biological activities for those are compared with experimental values, as presented in Table 4. The substance cynaroside is isolated from the leaf of *cynara scolymus* L (Loi, 2006) and its structure is identified using the different spectra such as:  $^1H$ -NMR (500 MHz, DMSO,  $\delta$  ppm):  $\delta$  6.39 (s,  $H_3$ ); 6.44 (d,  $J = 2$  Hz,  $H_6$ ); 6.78 (d,  $J = 2$  Hz,  $H_8$ ); 5.08 (1H, d,  $J = 7.5$  Hz,  $H_{1'}$ ); 7.41 (d,  $J = 8.5$  Hz); 6.90 (d,  $J = 8.5$  Hz); 7.44 (dd,  $J = 8.5$  Hz,  $J = 2$ ); 3.489 (1H, m,  $H_{2'}$ ); 3.476 (1H, m,  $H_{4'}$ ); 3.466 (1H, m,  $H_{3'}$ ); 3.500 (1H, m,  $H_{5'}$ ); 3.725 (1H, dd,  $J = 12.3$ ; 2.5 Hz,  $H_{6'}$ ); 3.702 (1H, dd,  $J = 12.4$ ; 6.3 Hz,  $H_{6''}$ ); 12.8 ( $C_5OH$ ); 3.466–3.725 (7H, m, glucose protons). The associated spectrum was also used to

**Table 4. Activities pGI<sub>50</sub> of test group resulting from models QSAR<sub>MLR</sub>, QSAR<sub>PLS</sub> and QSAR<sub>ANN</sub>**

Substance	Reference	pGI <sub>50,exp</sub>	pGI <sub>50,pred</sub>			ARE (%)		
			QSAR <sub>MLR</sub>	QSAR <sub>PLS</sub>	QSAR <sub>ANN</sub>	QSAR <sub>MLR</sub>	QSAR <sub>PLS</sub>	QSAR <sub>ANN</sub>
Fla2	(Lee et al. 2012; Liao et al., 2005, 2008)	5.9208	6.0079	5.8012	5.8509	1.4719	2.0200	1.1802
Fla9	(Lee et al. 2012; Liao et al., 2005, 2008)	5.7447	5.6915	5.6082	5.7407	0.9252	2.3758	0.0698
Fla12	(Lee et al. 2012; Liao et al., 2005, 2008)	6.0969	5.7587	5.8416	5.8136	5.5478	4.1875	4.6463
Fla15	(Lee et al. 2012; Liao et al., 2005, 2008)	5.6990	5.6511	5.6518	5.7124	0.8402	0.8275	0.2357
Fla16	(Lee et al. 2012; Liao et al., 2005, 2008)	5.6990	5.6514	5.6549	5.7188	0.8350	0.7746	0.3466
Isofla32	(Lee et al. 2012; Liao et al., 2005, 2008)	5.1367	5.0917	5.0830	5.1115	0.8767	1.0457	0.4900
Cynaroside	This work	5.3260	5.1910	5.6317	5.3186	2.5350	5.7393	0.1388
Quercetin	This work	5.3790	4.5858	5.5355	5.3591	14.7455	2.9094	3.9388
					MARE (%)	3.4722	2.4850	1.3808

have more structural information such as: <sup>13</sup>C-NMR (DMSO, δ ppm, 125 MHz), DEPT: δ 164.5(C<sub>2</sub>); 105.3(C<sub>3</sub>); 181.785 (C<sub>4</sub>); 161.2 (C<sub>5</sub>); 100.2 (C<sub>6</sub>); 162.9 (C<sub>7</sub>); 95 (C<sub>8</sub>); 156.9 (C<sub>9</sub>); 103.1 (C<sub>10</sub>); 121.4 (C<sub>1</sub>); 113.6 (C<sub>2</sub>); 145.7 (C<sub>3</sub>); 149.9 (C<sub>4</sub>); 115.9 (C<sub>5</sub>); 119.2 (C<sub>6</sub>); 100.0 (C<sub>1</sub>); 76.4 (C<sub>2</sub>); 77.2 (C<sub>3</sub>); 69.5 (C<sub>4</sub>); 78 (C<sub>5</sub>); 60.6 (C<sub>6</sub>). Interaction of C and H in spectrum HMBC and HSQC are also pointed out: H<sub>6</sub>-C<sub>5</sub>-C<sub>7</sub>-C<sub>8</sub>-C<sub>10</sub>; H<sub>8</sub>-C<sub>6</sub>-C<sub>7</sub>-C<sub>9</sub>-C<sub>10</sub>; H<sub>2</sub>-C<sub>2</sub>-C<sub>1</sub>-C<sub>3</sub>-C<sub>4</sub>-C<sub>6</sub>; H<sub>5</sub>-C<sub>1</sub>-C<sub>3</sub>-C<sub>4</sub>-C<sub>6</sub>; H<sub>6</sub>-C<sub>2</sub>-C<sub>1</sub>-C<sub>2</sub>-C<sub>4</sub>-C<sub>5</sub>.

The substance quercetin is isolated from the leaf of *artocarpus incisa* L (Loi, 2006) and its structure is also identified using the spectrum <sup>1</sup>H-NMR (DMSO-d<sub>6</sub>, 500 MHz, δ ppm) combining with spectrum HSQC, HMBC: δ 6.26 (1H, d, J = 1.5 Hz, H<sub>6</sub>); δ 6.52 (1H, s like t, H<sub>8</sub>); δ 7.82 (1H, d, J = 1.5 Hz, H<sub>2</sub>); δ 7.68 (1H, dd, J = 8.5 và 2 Hz, H<sub>5</sub>); δ 6.98 (1H, d, J = 8.5 Hz, H<sub>6</sub>); δ 12.16 (1H, s). Also, using spectrum <sup>13</sup>C-NMR (DMSO-d<sub>6</sub>, 125 Hz) combining with spectra DEPT, HSQC, HMBC: δ<sub>C</sub> 146.9 (C<sub>2</sub>); δ<sub>C</sub> 136.6 (C<sub>3</sub>); δ<sub>C</sub> 176.5, (C<sub>4</sub>); δ 162.2 (C<sub>5</sub>); δ 99.1 (C<sub>6</sub>); δ 164.9 (C<sub>7</sub>); δ 94.4 (C<sub>8</sub>); δ 157.7 (C<sub>9</sub>); δ 104.0 (C-10); δ<sub>C</sub> 121.4 (C<sub>1</sub>); δ 116.1 (C<sub>2</sub>); 145.7 (C<sub>3</sub>); δ 148.2 (C<sub>4</sub>); δ 115.6 (C<sub>5</sub>); δ<sub>C</sub> 123.7 (C<sub>6</sub>).

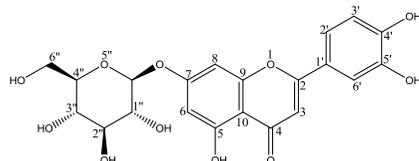
**Figure 4. The molecular structures of: (a) cynaroside and (b) quercetin.**



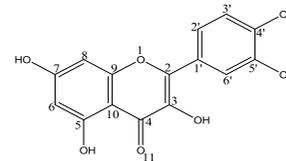
a) leaf of *cynara scolymus* L [1]



b) leaf of *artocarpus incisa* L [1]



Cynaroside with GI<sub>50,exp</sub> (μM) = 4.72 ± 0.280



Quercetin with GI<sub>50,exp</sub> (μM) = 4.18 ± 0.327

**Table 5. The anti-cancer activities  $GI_{50}$  ( $\mu\text{M}$ ) of five new flavonoids (n) designing from the vacant positions  $C_6$  và  $C_3$ , on quercetin, resulting from QSAR<sub>ANN</sub> model**

New substance	Substitutes at $C_6$	Substitutes at $C_3$	$GI_{50}$ ( $\mu\text{M}$ )	Method in this work
Quercetin	-H	-H	$4.18 \pm 0.327$	<i>in vitro</i> test on Hela
Fla-1(n)	$-\text{OCH}_2\text{CONHCH}_3$	-OH	0.1539	QSAR <sub>ANN</sub>
Fla-2(n)	$-\text{OCH}_2\text{CONHC}_6\text{H}_4\text{F}$	-H	0.1487	QSAR <sub>ANN</sub>
Fla-3(n)	-OH	$-\text{OCH}_2\text{CONHCH}_3$	0.1247	QSAR <sub>ANN</sub>
Fla-4(n)	$-\text{OCH}_2\text{CH}_3\text{C}=\text{NOH}$	-OH	0.1233	QSAR <sub>ANN</sub>
Fla-5(n)	$-\text{OCH}_2\text{CONHC}_6\text{H}_4\text{OCH}_3$	-H	0.1174	QSAR <sub>ANN</sub>

The molecular structures of substances cynaroside and quercetin are shown in Figure 4.

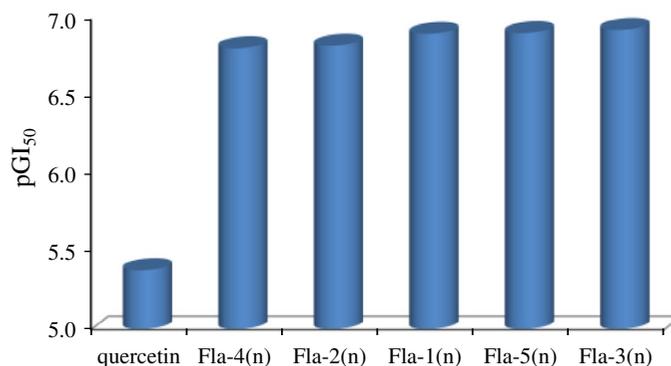
After isolation of two new flavonoids cynaroside and quercetin, their activities  $pGI_{50}$  were conducted to test *in vitro* toxicity on Hela cells in the laboratory of molecular biology, university of natural sciences. The activity values  $pGI_{50}$  of two these flavonoids were also predicted from the models QSAR<sub>MLR</sub>, QSAR<sub>PLS</sub>, and QSAR<sub>ANN</sub>, as shown in Table 4. Those were compared with experimental activities and with each other based on the average value of the absolute relative error MARE,%. The predictability of the model QSAR<sub>MLR</sub> is lower than models QSAR<sub>PLS</sub> and QSAR<sub>ANN</sub>, respectively, as given in Table 4.

The QSAR<sub>ANN</sub> model has the valuable error MARE,% of 1.3808. This is smaller than values MARE,% of both models QSAR<sub>MLR</sub> and QSAR<sub>PLS</sub>. So, the predictability of QSAR<sub>ANN</sub> model is better than models QSAR<sub>MLR</sub> and QSAR<sub>PLS</sub>. After using the models QSAR<sub>MLR</sub>, QSAR<sub>PLS</sub>, and QSAR<sub>ANN</sub> to predict the biological activities  $pGI_{50}$  of six compounds in test group and two new flavonoids quercetin and cynaroside, the accurate level of the predicted results is exhibited in the acceptable errors within the uncertainty of experimental measurements. Thus, the models QSAR<sub>MLR</sub>, QSAR<sub>PLS</sub> and QSAR<sub>ANN</sub> are good adaptable for predicting the biological activities of new substances.

In this work, we selected the new flavonoid quercetin with the vacant positions  $C_6$  and  $C_3$ , such as sample compound for designing five new compounds. The substitutes were attached to two vacant positions  $C_6$  and  $C_3$ , as shown in Table 5. The new designed compounds were also predicted with the biological activities  $pGI_{50}$  using the QSAR<sub>ANN</sub> model. Then, the predictive activities  $pGI_{50}$  were recovered in the original form  $GI_{50}$  ( $\mu\text{M}$ ), as given Table 5.

The predicted results  $pGI_5$  for new substances are transformed into values  $GI_{50}$  ( $\mu\text{M}$ ) and compared with experimental activity of sample quercetin, as depicted in Figure 5. Thus, the five new compounds designing from the  $C_6$  and  $C_3$  positions on quercetin displayed stronger activity  $GI_{50}$  ( $\mu\text{M}$ ) than sample quercetin. Herein, the new designed compounds will promise to forward a designing plan for the new pharmaceutical products from natural products.

**Figure 5. Comparison between values  $pGI_{50}$  of five new flavonoids with quercetin.**



#### 4. Conclusion

We used the quantum chemistry calculations, multivariable regression and artificial neural network to construct successfully the quantitative relationships between the partial atomic charges and anti-cancer activities  $GI_{50}$  ( $\mu\text{M}$ ) of flavonoids. The models QSAR<sub>MLR</sub> showed six important sites  $O_1$ ,  $O_{11}$ ,  $C_3$ ,  $C_4$ ,  $C_6$ , and  $C_7$  on flavonoids.

The QSAR<sub>MLR</sub> model found out the most important positions  $C_6$  and  $C_3$ , to add the new substitutes to create five new flavonoids with higher activity of quercetin isolating from the leaf of *artocarpus incisa* L. The QSAR<sub>ANN</sub> model with architecture I(6)-HL(4)-O(1) is better predictable for flavonoids.

#### Supplementary material

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