

## ARTICLE / INVESTIGACIÓN

## Computational discovery of novel anthelmintic natural compounds from *Agave Brittoniana* trel. Spp. *Brachypus*

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**Abstract:** Helminth infections are a medical problem in the world nowadays. This report used bond-based 2D quadratic indices, a bond-level QuBiLS-MAS molecular descriptor family, and Linear Discriminant Analysis (LDA) to obtain a quantitative linear model that discriminates between anthelmintic and non-anthelmintic drug-like organic-compounds. The model obtained correctly classified 87.46% and 81.82% of the training and external data sets, respectively. The developed model was used in a virtual screening to predict the biological activity of all chemicals (19) previously obtained and chemically characterized by some authors of this report from *Agave brittoniana* Trel. spp. *Brachypus*. The model identified several metabolites (12) as possible anthelmintics, and a group of 5 novel natural products was tested in an in vitro assay against *Fasciola hepatica* (100% effectivity at 500 µg/mL). Finally, the two best hits were evaluated *in vivo* in bald/c mice and the same helminth parasite using a 25 mg/kg dose. Compound 8 (Karatavinoside A) showed an efficacy of 92.2% *in vivo*. It is important to remark that this natural compound exhibits similar-to-superior activity as triclabendazole, the best human fasciolicide available in the market against *Fasciola hepatica*, resulting in a novel lead scaffold with anti-helminthic activity.

**Key words:** TOMOCOMD-CARDD Software, QuBiLS-MAS, nonstochastic and stochastic bond-based quadratic indices, LDA-based QSAR model, Computational Screening, Anthelmintic Agent, *Agave brittoniana* Trel. spp. *Brachypus*, *Fasciola hepatica*.

### Introduction

Helminths remain among the most common chronic infections, with more than one-third of the world's population infected at any time<sup>1</sup>. Currently, the high cost and toxicity of anthelmintics as well as the emergence of resistant strains of pathogenic helminths, have stimulated the desire to search for additional chemotherapeutic agents allowing a more efficient control of these parasites<sup>2-4</sup>. A practical solution to this problem is to develop effective drugs from less expensive and more available raw materials<sup>5</sup>. Natural products (NP) can be one of these materials for various reasons: 1) They inspired most of the active ingredients in medicines, 2) NP exhibit enormous structural diversity, 3) NP are the result of centuries of evolutionary pressure to create biologically active molecules, 4) the structural similarity of protein targets across many species, and so on. 5) It is extensively known that NP share more similar than synthetic compounds to the 'chemical space' of drug molecules<sup>6-18</sup>. Unfortunately, only a small proportion of that diversity has

been extensively explored for its pharmacological potential so far<sup>19-21</sup>.

Until now, the search for new anti-helminthic compounds from natural origin has generally been based on traditional trial-and-error methods<sup>5,22</sup>. Unfortunately, these methods are highly inefficient and expensive<sup>9,23</sup>. For this reason, new technologies have emerged to replace these old "hand-crafted" approaches for synthesis and testing new chemical entities<sup>12,24-26</sup>. Virtual screening is an example of these modern approaches. Specifically, Quantitative Structure-Activity Relationships (QSAR) predictive models have been extensively used to filter large databases of compounds to identify new bioactive chemicals<sup>27-34</sup>. Compared to other areas of pharmaceutical research; however, the screening of NPs has suffered from a lack of data in an appropriate format. Such information can significantly impact virtual screening, where new natural agents would be identified as potential therapeutic anthelmintics.

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On the other hand, some authors of this report used an in-house computational approach to discover new anthelmintic synthetic compounds with rather good results<sup>35-37</sup>. A similar approach has been used to find new tyrosinase inhibitors from natural origin<sup>38,39</sup>. However, no scientific report about discovering NPs with an anti-helminthic activity using an analogous computational strategy has been published.

This report presents the creation/validation of the QSAR model able to identify potential anthelmintic compounds. Next, we used this model in the virtual screening of NPs previously obtained and chemically characterized from *Agave brittoniana* Trel. spp. *Brachypus*. Finally, the identification/selection of the most promising anti-helminthic NPs for *in vitro* and *in vivo* experimental evaluation and the results of these evaluations are presented.

## Materials and methods

### Experimental Section

#### Materials

Compounds 1-5 were derived from previous studies made with *Agave brittoniana* Trel. spp. *Brachypus*<sup>63</sup>. The rest of the chemicals were obtained using a similar approach described by the same research team<sup>63</sup>. The extraction and purification of all compounds with a purity higher than 99% were carried out employing previously described methods.<sup>63</sup> To obtain the initial dissolutions, each product dissolved in water at a concentration of 10 mg/mL (1%). The insoluble products were first dissolved in dimethylsulphoxide (DMSO) so that the concentrations of this product in the final solution did not exceed 1%. The necessary dilutions of each product to make possible the biological evaluation was obtained starting from the initial solutions. In addition, a solution of TCB was utilized as reference drug.

#### Animals

Healthy balb/c mice of both sexes (body weight: 0.018±0.001 Kg) and food were purchased from the National Center for Laboratory Animal Production (CENPALAB, Havana, Cuba). Quarantine, labeling, climatization and good maintenance conditions of animals were strictly obeyed.

#### General Experimental Procedures

To measure the chemical effectiveness against *F. hepatica*, an experimental technique reported in the literature was selected for biological material processing and *F. hepatica* egg extraction<sup>66</sup>. Mitterpak *et al.*'s technique for the host (*Lymnaea cubensis*) invasion was carried out<sup>67</sup>. Afterwards, we followed the steps reported by Olazábal *et al.*<sup>68</sup> to obtain the metacercariae. Metacercariae were conserved in the cold until the *in vivo* experiment<sup>66</sup>.

#### Biological Experiments

The anthelmintic activity of the compounds was evaluated, first, against *F. hepatica* in an *in vitro* assay using an earlier described procedure and second, against metacercariae of the same pathogen in an *in vivo* experiment, applying another well-established procedure.

Several treatment groups with ten mice per group were created. One group (infected control group) was treated with Miglyol 810N (administration vehicle). The second group

was neither infested nor treated. The remaining groups were treated with new chemicals. All mice received the new compounds through an oral route. Mouse invasion with metacercariae of *F. hepatica*, 2 weeks old, 14 days before drug administration, was carried out by Corba *et al.*'s method<sup>69</sup>. The effectiveness was evaluated based on the following:

1) determination of the E% index. This is a quantitative indicator of effectiveness introduced by Steward<sup>70</sup> and defined as  $E\% = [(XC-XT)/XC] \times 100$ <sup>71</sup>. Here, E% is the percentage of effectiveness, XC is the average amount of *Fasciola* in the control group, and XT is the average amount of *Fasciola* in the treated group. Effectiveness was measured based on the elimination or not of *F. hepatica*, in its juvenile stage, as shown by laboratory diagnostics, using the helminthological necropsy on day 7 after the inoculated treatment<sup>69</sup>.

2) Determination of the hepatic index<sup>72</sup>, by mean of the formula  $A = (B/C) \times 100$ . In this case, A = hepatic index, B = liver weight and C = body weight.

3) Degrees of lesions of the liver<sup>69</sup>.

4) Spleen relative weight<sup>73</sup>.

5) Intensity of invasion making use of the formula  $I = A/B$ , where A = total amount of parasites, B = total amount of positives.

6) Extension of invasion by use of the formula  $\%E.I = [T(t)/T(a)] \times 100$ , where %E.I is the percent of invasion extensity, T(t) = number of total positives, and T(a) = total of infected animals<sup>74</sup>.

7) Gain of weight (final weight) (initial weight).

From these different effectiveness indexes<sup>72-74</sup>, the E% index was selected.

#### Computational method

In the present report, we used a defined mathematical algorithm, which is characterized in this case by bond-based QuBiLS-MAS (acronym for Quadratic, Bilinear and N-Linear mapS based on graph-theoretic electronic-density Matrices and Atomic weightingS) MDs family (bond-level nonstochastic quadratic indices) to encode the chemical information in numbers<sup>50-52</sup>. The CARDD extension of the TOMOCOMD approach has been previously successfully used to discover new bioactive molecular entities<sup>35,36,38,44-49</sup>. The general principles of these indices and the main steps for the application of the QuBiLS-MAS<sup>50</sup> software (<http://tomocomd.com/software/qubils-mas>) in QSAR/QSPR for drug design have been described in detail elsewhere<sup>35,36,38,44-49</sup>.

To find the classification function that discriminates between active and inactive compounds, we select the LDA because it is one of the most broadly used and straightforward techniques to obtain QSAR equations<sup>35,36,48,49,75-85</sup>. It was carried out with the STATISTICA software<sup>53</sup>. Forward-stepwise and best subset search procedures were fixed as the strategy for variable selection. The best model was selected considering the principle of parsimony (Occam's razor). The considered tolerance parameter was the default value for minimum acceptable tolerance, which is 0.01. The quality of the model was determined by examining Wilks'  $\lambda$  parameter (*U* statistic), the square Mahalanobis distance ( $D^2$ ), the Fisher ratio (*F*), and the corresponding *p* level [*p*(*F*)] as well as the percentage of good classification (accuracy) in the training and test sets (see Schemes 1 and 2). The classification of cases was performed by means of the posterior classification probabilities where one compound can then be classified as active if  $\Delta P\% > 0$ , being  $\Delta P\% = [P(\text{Active}) - P(\text{Inactive})] > 100$ , or as inactive otherwise.  $P(\text{Active})$

and  $P(\text{Inactive})$  are the probabilities with which the equation classifies a compound as active or inactive, respectively. On the other hand, the probability density approach implemented in the Ambit Disclosure software was used to evaluate the applicability domain of the model developed<sup>60</sup>.

## Results and discussion

### *In silico* study and virtual screening

#### Developing and validating linear QSAR models

To obtain a mathematical relationship between chemical structures and biological activity, the chemical information contained in many compounds must be statistically processed. Therefore, we build a data set containing 212<sup>40-43</sup> and 305<sup>40,41</sup> inactive compounds from the literature. It was build including 517 (active + inactive) compounds and was randomly divided into two subgroups: a set of 352 compounds (138 active and 214 inactive) that was used as the training set for developing the classification model and a second set of 165 compounds (74 active and 91 inactive) that was used as a test set for testing the predictive power of the model developed (see figure 1).

Each structure was parameterized by using one TO-MOCOMD-CARDD<sup>35,36,38,44-49</sup> molecular descriptor (MDs) family, named bond-based nonstochastic 2D quadratic indices (QuBiLs-MAS Software)<sup>50-52</sup> (see the experimental section for more details). Linear discriminant analysis (LDA), implemented on the STATISTICA software, was used as the statistical technique for model building<sup>53</sup>. The best classification model obtained is given below, together with the LDA-statistical parameters:

$$\begin{aligned} \text{Class} = & -2.2558 + 0.00001^M q_{5L}^H(\bar{x}) - 0.00024^M q_3^H(\bar{x}) + 0.06528^P q_1(\bar{x}) \\ & - 0.00280^M q_{1L}(\bar{x}) - 0.00360^P q_3^H(\bar{x}) + 0.00447^M q_3^H(\bar{x}) \end{aligned} \quad (1)$$

$$N=352 \quad \lambda=0.443 \quad D^2=5.261 \quad F(6,344) = 72.196 \quad p < 0.0001$$

where,  $N$  is the number of compounds,  $\lambda$  is the Wilks' statistic,  $D^2$  is the squared Mahalanobis distance and  $F$  is the Fisher ratio.

The Wilks' parameter is equal to the proportion of the total variance in the discriminant scores not explained by differences among the groups. Smaller values of Wilks' lambda indicate the greater discriminatory ability of the function.

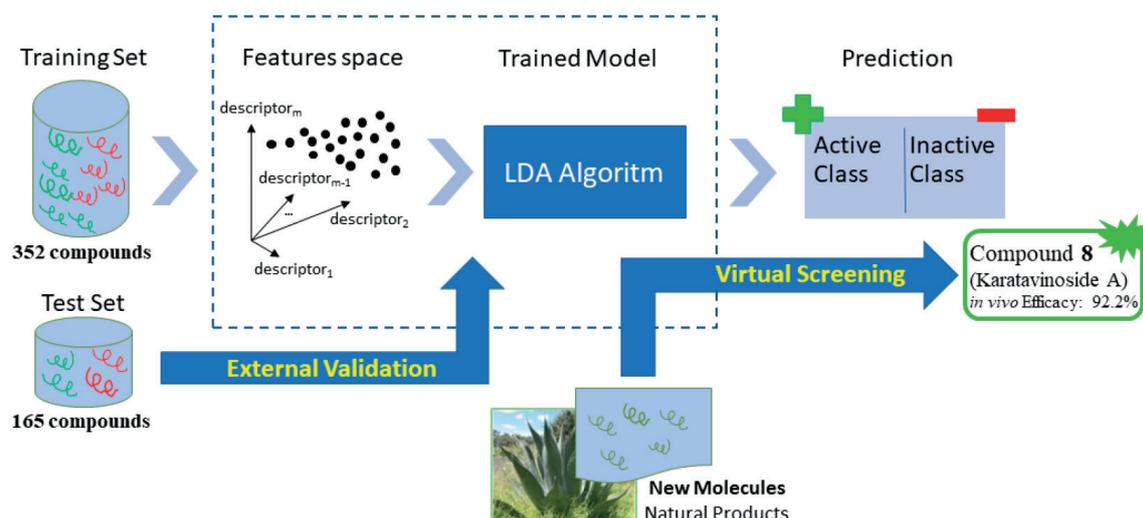
Its statistic parameter can take values in the range of 0 (perfect discrimination) to 1 (no discrimination)<sup>54</sup>. That is, Wilks' lambda is a direct measure of the proportion of variance in the combination of dependent variables unaccounted for by the independent variable (the grouping variable). Suppose a large proportion of the variance is accounted for by the independent variable. In that case, it suggests an effect from the grouping variable and that the groups (active and inactive) have different mean values. The Mahalanobis distance is a statistical technique that can be used to measure how distant a point is from the centre of a multivariate normal distribution, and its parameter indicates the separation between the respective groups<sup>55</sup>. It shows whether the model has an appropriate discriminatory power for differentiating between the two respective groups. The classification of cases was carried out by means of the posterior classification probabilities. Using the Mahalanobis distances to do the classification, we can now derive probabilities. The probability that a case belongs to a particular class is basically proportional to the Mahalanobis distance from that group centroid. In summary, the posterior probability is the probability, based on our knowledge of the values of other variables, that the respective case belongs to a particular group.

This equation can correctly classify 87.46% (307/352) of the compounds in the training set and showed values of the Matthews correlation coefficients of 0.74 on it. More important, the model achieves a balanced classification accuracy in each group.

The results of the most relevant statistical parameters for this model are presented in Table 1, and the classification of compounds in the training set using Eq. 1 is presented in Table 2.

Once a model is trained, its validation is another crucial aspect in this kind of analysis which can be performed

by internal and external validation techniques (see Scheme 2)<sup>56,57</sup>. Here, a leave-many-out (LMO) cross-validation technique was carried out where groups of 176, 117, 70, 35, and 17 compounds of the training data (352 chemicals) were taken like cancellation groups and at each step. Then, the newly trained model was used to predict the left-out compounds. The results of this analysis are shown in Table 3,



**Figure 1.** Schematic representation of the process used to design training and test sets.

Matthews Corr. Coefficient	Accuracy 'Q <sub>Total</sub> '	Sensitivity 'True Positive rate' (%)	Specificity 'True Negative rate' (%)	False positive rate 'false alarm rate' (%)
<i>Training Set</i>				
0.74	87.46	87.59	87.38	13
<i>Test Set</i>				
0.64	81.82	83.78	80.22	20

**Table 1.** Prediction Performances and Statistical Parameters for QSAR Models in the Training and Test Sets.

and the model's parameters and predictions are rather stable when a perturbation is applied to the training set. This proves that our model is robust.

In addition, to check the possibility of random correlations, the Y-randomization test (Y-scrambling) was performed by calculating the quality of the model randomly modifying the sequence of the response vector *y* (binary response: active or inactive) of the 5%, 10%, 20%, 30% y 40% of the compounds in the training set and recalculating the statistical parameters of the obtained models<sup>57</sup>. The final conclusions of this test are present in Figure 2, indicating that the achieved level of random correlation is significantly lower than the original regression, leading to the conclusion that the models are not random.

A more strict performance evaluation of a model is provided by an external validation where the model predictively is a challenge by compounds (external test set) that were not used in the model training (see Figure 3)<sup>57</sup>. Therefore, the equation obtained was evaluated in the test set (external prediction), showing accuracies of 81.82 % (135/165) and values of the Matthews correlation coefficients of 0.64. In addition to the external validation, the results of the statistical parameters described in Table 1 show that our model is not only robust but also predictive; therefore, it can be used in ligand-based virtual screening. The classification of both compounds in the external prediction set are depicted in Table 2.

Finally, to define the applicability domain<sup>57</sup> of Eq 1, a city-block distance-based approach<sup>58,59</sup> implemented in the Ambit program<sup>60</sup> was used. The model's applicability domain was defined from the training set, and all compounds belonging to the external test series were inside it.

#### In silico identification of active compounds from natural products

Taking into consideration that NPs have inspired most of the active ingredients in medicines<sup>10</sup>, in the last years a number of recent investigation was carried out to discover new active compounds from the natural origin using computational strategy<sup>61</sup>. In our research, the developed model (Eq. 1) was used to filter an extensive database of NPs. All details of this database and other active (anthelmintics) NPs discovered by using our approach will be shown in the following reports.

Here, we only present the discovery of novel anthelmintic compounds from *Agave brittoniana* Trel. spp. *Brachypus*: a plant that grows like one of two endemic subspecies (ssp. *Brachypus* and ssp. *Spirituana*) of *Agave brittoniana* Trel. in the central region of Cuba<sup>62</sup>. A group of nineteen compounds composed by 12 steroidal saponins, 6 steroidal saponinins and 1 phytosterol (see Figure 4) that have been previously obtained and chemically characterized from this subspecies of *Agave* was evaluated *in silico* using the Eq. 1. These compounds were: agabrittonosides A–D<sup>63</sup>, agabrittonosides E–K, karatavioside A<sup>64</sup>, Diosgenin, Chlorogenin,

Hecogenin, Tigogenin, Rockogenin, and β-Sitosterol.

As result of this virtual screening, twelve compounds were identified by the model as potential anti-helminthic hits (see Table 4).

However, it is generally acknowledged that QSARs are valid only within the same domain for which they were developed. Even if the models are developed on the same chemicals, the DA for new chemicals can differ from model to model, depending on the specific MDs. One of the present reports aims is to develop a model for predicting the anthelmintic activity of NP at the early stages of the drug discovery and development pipelines. Therefore, the chemicals selected in this study were only evaluated *in vitro* after plotting them into the model's previously obtained AD. In this analysis, all compounds were inside the DA of the model, ensuring excellent reliability for the prediction of this kind of lead used in the virtual screening. Moreover, all new leaders fall within the model's DA, so the predictions are reliable.

#### Experimental corroboration

##### In vitro assay

Compounds were limited in availability; therefore, not all compounds were experimentally tested. Only three of the compounds detected *in silico* as potential anti-helminthic hits (Karataviosido A, Agabrittonósido A, Agabrittonósido B) and a mixture of Agabrittonósidos D and Agabrittonósidos E could be tested *in vitro* against *F. hepatica* at  $5 \times 10^{-1}$ ,  $5 \times 10^{-2}$ ,  $5 \times 10^{-3}$ ,  $5 \times 10^{-4}$ ,  $5 \times 10^{-5}$  and  $5 \times 10^{-6}$  mg/mL. Triclabendazole (TCB) was included in this experiment as a reference drug because it is the one of choice in treating human fascioliasis<sup>65</sup>. Besides, Yucagenin (predicted as inactive) was also included in determining the influence of the glycoside moiety in the anti-helminthic activity. The biological *in vitro* evaluation results can also be seen in Table 4.

The experimental results agreed with the virtual screening predictions. As predicted, Yucagenin is not active at any test concentrations. However, its glycoside derivative (8, 9) had a bioactivity profile as TCB. This first saponin (8) has a glycoside rest joined to the C-3 atom identical to compound 9, its structural difference in the opening of the ring F and the glycosidation in the C-26 atom. The responsible for the little activity of 10, can be this structural modification or the increase of polarity of this zone. The mixture of compounds 12 and 13 presented *in vitro* activity higher than that observed for TCB. Compounds 12 and 13 are very similar structurally; both have the diosgenin-like central scaffold, but in compound 13 one xylose unit in 12 is substituted by a rhamnose group. In addition, 12 have a hydroxyl moiety in C-2, which is the only difference from 9. The combination of these subtle changes notably increases the activity of 12 and 13 concerning 9.

name	$\Delta P\%^a$	name	$\Delta P\%^a$
<i>Training active group</i>			
Tetrachloromethane	99.96	Benacyl	64.94
Hexachloroethane	99.99	Imcarbofos	-11.47
Antimony sodium thioglycollate	-12.40	Etibendazole	58.88
Dichlorvos	95.20	Fluranteel	99.92
1-Chlorobutane	-84.87	Pararosaniline embonate	88.20
Dimetrizadole	-52.17	Amphotolide	81.43
Disoferol	-57.14	Praziquantel	67.48
Lindane	19.06	Diuredosan	58.21
Mindazole	8.40	153C51	74.88
Fospirate	87.49	Trichlorophen	99.75
Oltipraz	96.40	Febantel	61.19
Certuna	-37.26	Lucanthone	81.00
Butonate	99.95	Miracil A	56.87
Antienite	73.81	Diamphenetide	48.22
Sodium antimony dimethylcysteine tartrate	53.94	Alazanine triclofenate	98.23
Wormin	35.99	Antelmycin	-42.25
Nitrodam	83.60	Diospyrol	96.63
Lobendazole	-58.33	Becantone	91.94
Bromothymol	81.12	Desaspidin	97.44
Iodothymol	70.57	Methylrosanilidium chloride	-5.71
Famophos	12.06	Bunamide	84.38
Thiacetalsamide	4.78	Bidimazium oidide	94.02
Antazonite	76.66	Pyruinium chloride	74.74
Vincofos	98.97	Desoine	34.28
Niclofolan	98.21	Thymoloverm	97.10
Bromofenofos	80.53	Teroxalene	96.95
Sinid	99.82	Pretamazium iodide	97.82
Phenotiazine	79.08	Hedaquinium chloride	97.71
Nitazoxanide	75.50	Dryocrassin	99.97
Bendamizole	38.85	Netobimin	4.02
Tioxidazole	56.19	Phenithionate	91.48
Phoxim	70.22	Artesunate	73.02
Albendazole	20.75	Abamectin B	99.45
Anthiolimine	67.17	Alantolactone	48.31
Carbantel	40.67	Antimony potasium tartrate	90.38
Cetovex	-33.89	Arecoline	-70.97
4-Hexylresorcinol	25.63	Aspidin	96.52
Crufomate	86.81	Benomyl	-6.42
Butynorate	-13.36	1,4-Bistrichloromethylbenzene	100.00
Hexachlorophenimonophosphate	97.18	Carbacol	36.93
Niclosamide	96.88	Triclabendazole sulphoxide	95.99
Nitroscanate	86.36	Epritantel	53.65
G-572	76.60	Acid filix (c)	99.91
Resurantel	87.70	Oentian violet	-32.44
Phenzidole	45.11	Hectolin	98.35
Ontianil potasium	77.13	Hidroxyquinoline	7.47
Atractyl	57.00	Urea stibamine	-81.83
Amidantel	1.49	alfa-Kosin	95.91
Mirasan	-4.31	Mandelic acid isoamyl ester	72.81
Triclabendazole	96.56	Mibbemycin Oxime A3	96.35
Nocodazole	19.80	Moxidectin	98.94
Styrylpyridiniumchloride	69.52	Naftalofos	85.30
Haloxon	87.78	Naphthalene	13.12
Coumafos	95.29	Nicotine	-61.44
Coralop	93.40	Nitroxymil	53.93

**Table 2.** Results of the Classification of Compounds in the Training and Test Set using QSAR Models.

name	$\Delta P\%^a$	name	$\Delta P\%^a$
Evultin	74.23	Paraherquanude	51.83
Oxamniquine	28.86	Thymol	35.82
Brotianide	98.98	alfa-Santonin	42.82
Afesal	98.55	Arsamilate	-88.07
Furodazole	78.42	Quinacrine	79.37
Oxfendazole	57.98	Kaimic acid	-63.64
Dribendazole	53.95	Hexachlorophene	99.65
Butamisol	80.95	Tetrachloro-difluoroethane	99.70
Spirazine	70.66	Diamphenethide Deacetylated (metabolite)	43.69
Lubisan	17.98	Doramectin	99.57
S72014	86.50	Dymanthine	-24.86
Flubendazole	75.13	Bromoxanide	99.91
Mebendazole	54.79	Tetrachloroethane	-17.65
Egresin	58.67	Paramomycin	-12.13
<i>Training inactive group</i>			
3-Episiostatin B	-66.80	2-etoxybenzamide	-95.05
Thiacetazone	-63.38	2-isopropil-4-pentenoyl urea	-93.63
ZDPFA	-3.61	Carbetimer	-65.19
Foscarnet	-98.33	Etimidin	-82.76
Ethoxene	-82.33	Leucenol	-89.40
Arildone	87.74	Azatepa	-54.26
Fanciclovir	-27.92	Asperlin	-53.90
Valerium Paul Thibault	-63.43	Pipobroman	-99.98
Ferriscorbone magnesienne	-53.98	Glucin	-98.56
Aponal	-51.73	Isomalt	-86.18
Carbavin	-51.30	S-2346	-96.34
Zonisamide	-35.28	DL 204 IT	-97.47
Atrolactamide	-38.70	Equilenin	-96.11
Beclamide	-53.84	Ethyllysergamide	-99.39
Tetrantoin	-19.04	Trenbdone	-96.14
Tiletamine	13.30	Methallenestril	-94.65
Ferro-Drops	-79.75	Estradiol	-97.10
Cobalti glutamas	-72.22	Metexaminum	-82.55
Fructosum Ferricum	-80.55	Solution A40	-94.81
Ferromaltose	-74.93	Proxamine	-92.92
Erythrophyll	99.96	Clonazoline	-94.53
Butanolium	-74.30	Tymazoline	-91.30
A-Peest	-71.84	Isopropylmethoxamine	-96.06
Iprazochrome	-35.89	Angiotensinamide	-99.79
Esculamine	35.44	Terlipressin	-97.99
Morfafen	95.81	Isomylamine	-88.00
Acetazolamide	-57.16	Dimebamate	-78.19
Trometamol	-82.01	DEP	-85.12
Butazolamidum	-32.90	Silamprobamate	-89.54
Merbiurelidin	-95.50	Chlorphenesin carbamate	-81.89
Chlorothiazide	-3.72	Guaifenesin	-93.80
Disulfamide	-40.20	Murexine	-99.52
Trichlormethiazide	83.08	Nafomine	-97.56
WR-2823	-97.28	AHR-2666	-82.70
Peucedanin	87.50	Pifexole	-83.22
Batilol	21.98	Tybamate	-85.86
Glipentide	61.98	Carmecolina cloruro	-99.27
Glisindamide	76.76	Eseridine	-97.46
Glimepiride	79.06	Distigmine bromide	-99.46
PIDH	-11.49	D-935	-95.24
Carbutamide	-23.95	Ergocristine	-96.60
Ag-307	-65.39	Etilefrine pivalate	-74.11

**Table 2.** Results of the Classification of Compounds in the Training and Test Set using QSAR Models.

name	$\Delta P\%^a$	name	$\Delta P\%^a$
<b>Diazoxide</b>	48.58	Pivenfrine	-72.63
<b>Alarmino</b>	-67.86	Etafedrine	-98.00
<b>MK-534</b>	86.29	Dimetofrine	-97.44
<b>Oxonazine</b>	-78.29	Methoxyphenamine	-96.85
<b>Guanisoquine sulfate</b>	18.67	Clorprenaline	-85.85
<b>Stilonium iodide</b>	29.75	Phenylethanolamine	-94.85
<b>Pancuronium bromide</b>	-70.89	Norfefrine	-93.92
<b>Dimecolonium iodide</b>	-97.46	Oxidopamine	-90.52
<b>Chlorisondamine chloride</b>	-62.91	Pacamine	-90.78
<b>Mecamylamine</b>	-77.27	Metaraminol	-94.36
<b>Methylene chloride</b>	-91.59	Synephrine	-96.92
<b>Vinyl ether</b>	-72.01	Corbadrine	-93.16
<b>Neothyl</b>	-83.80	Phenamazoline	-97.60
<b>Novasil</b>	-39.87	Pholedrine sulfate	-95.71
<b>Acidum isobutiaticicum</b>	-14.08	Etilefrine	-97.13
<b>Iodothiouracil</b>	-39.43	Para-Aminoephedrine	-97.11
<b>Basthioryl</b>	-94.03	AMT	-96.63
<b>Pyrglutargine</b>	-68.78	Aphidicolin	-93.27
<b>Acustasin</b>	-76.09	Pipratecol	-98.77
<b>Gluronsan</b>	-59.23	Ftorin	-78.82
<b>Dicumarol</b>	96.78	Alprostadil	-92.58
<b>Chloracyzine</b>	81.07	Prostaglandin F1alpha	-95.10
<b>Aminoethylnitrate</b>	-58.55	Azaclorzine	-71.68
<b>Nitrodimehtylin</b>	-69.40	Aceperone	-97.98
<b>Propatyl nitrate</b>	72.94	Prenylamine	-99.48
<b>Carpronium chloride</b>	-96.05	Ericolol	-50.93
<b>6,9-Didesmethylartemisinin</b>	28.85	Ancarolol	-86.07
<b>Strychnobrasiline</b>	-51.99	Cicloprolol	-94.58
<b>Cilional</b>	-42.36	Pafenolol	-97.54
<b>7,7 Difluoro-<math>\beta</math>-arteether</b>	59.45	Nafetolol	-87.75
<b>Mezepine</b>	25.77	Sulfinalol	-98.59
<b>Tandamine</b>	17.54	ROM-203	-99.01
<b>Perafensine</b>	50.44	Spirendolol	-84.25
<b>Doxepin</b>	16.96	Flusoxolol	-95.37
<b>Ketimipramine</b>	10.33	Dobutamine	-96.87
<b>Tisocromide</b>	93.66	Carbazeran	-97.71
<b>Cotriptyline</b>	42.34	Bufalin	-97.30
<b>Diminazone</b>	38.39	Niludipine	-81.15
<b>Lauroguadine</b>	-31.96	Locundieside	-95.58
<b>Glycobiazol</b>	74.00	Covallatoxal	-96.67
<b>Clioquinol</b>	61.50	Peruvoside	-96.94
<b>Noscapine</b>	-97.31	Olitoriside	-94.64
<b>Tilidine</b>	-99.19	Deltamethrin	100.00
<b>Bextrometorfano</b>	-98.11	Phenothrin	-88.12
<b>Levallorphan</b>	-98.61	KC-8973	-90.43
<b>Fenytiltoxamine</b>	-99.27	Lipothiamine pyrophosphate	-99.45
<b>Clorfenoxamine</b>	-98.03	Xanthine	-94.25
<b>Medrylamine</b>	-99.58	Methioninol	-97.15
<b>Homoclorciclicine</b>	-97.98	Silidianin	-95.38
<b>Pimetixene</b>	-96.08	Bietamiverine	-98.35
<b>Borimamide</b>	-98.97	Cimetropium Bromide	-95.13
<b>Nigrifactin</b>	-94.21	Diponium Bromide	-97.74
<b>Octastine</b>	-94.55	Feclomine	-99.66
<b>Acidum etidronicum</b>	-99.57	Flavoxate	-95.16
<b>Nibet</b>	-99.04	Flopropione	-86.14
<b>Refortan</b>	-26.69	Pipoxolan	-97.17
<b>Clorotepine</b>	-88.43	Fludalanine	-87.26

**Table 2.** Results of the Classification of Compounds in the Training and Test Set using QSAR Models.

name	$\Delta P\%^a$	name	$\Delta P\%^a$
<b>Docloxytepin succinate</b>	-87.52	Cryptargol	-93.20
<b>Diclorpromazine</b>	-79.47	Mepartricin A	-99.86
<b>5-fluorocytosine</b>	-83.27	Protoanemonin	-76.14
<b>5-nitro-2-firfurilmetil eter</b>	-84.77	Contramine B	-97.95
<b>4(2-aminoetil)imidazole</b>	-94.04	F-8	-95.26
<b>2 hidroxibenzamine</b>	-93.21	Nitrofurylather	-84.77
<b>3-hidroxyacetanilide</b>	-93.99	Protoxyl	-99.98
<b>O-acetilsalicylamide</b>	-90.46	Bismuth Cevitamate	-73.39
<i>Test active group</i>			
<b>Tetrachloroethylene</b>	98.42	Artemethei	9.05
<b>Metrifonate</b>	99.96	Santoperonin	99.88
<b>Stibomen</b>	46.92	Agrimonolide	80.21
<b>26354R-P</b>	13.02	Dibutyltin dilarate	100.00
<b>VUFB-7904</b>	-43.77	Uredofos	58.90
<b>Bitoscanate</b>	33.74	Salantel	97.29
<b>SQ18506</b>	-64.91	Miracil	85.40
<b>Carbendazim</b>	-44.87	Hycanthone	72.64
<b>Tiabendazole</b>	43.77	Miracil C	6.53
<b>F30066</b>	25.19	Dithiazanine iodide	98.37
<b>Ascaridole</b>	14.50	Dicroden	70.73
<b>Eucalyptol</b>	-16.31	RO2-9009	-49.19
<b>R8231</b>	85.91	Zilantel	99.77
<b>Nitramisole</b>	69.53	Bisbendazole	99.80
<b>Levamisole</b>	22.03	Chuanliansu	-86.51
<b>Pyrantel</b>	24.06	Stilbaziumiodide	90.80
<b>Clorsulon</b>	85.46	Coib	65.81
<b>Feniodiumchloride</b>	100.00	Luxabendazole	80.84
<b>Stibofen</b>	3.76	Abamectin A	99.31
<b>Sodium stibocaptate</b>	93.88	Amorcazine	91.74
<b>Oxibendazole</b>	-16.25	Aspidinol	37.83
<b>Morantel</b>	41.11	Bethionol	99.64
<b>Oxyclozanide</b>	99.38	Embelin	77.42
<b>Nitroclofene</b>	98.39	Acid filix (a)	99.94
<b>Amoscanate</b>	76.60	Acid filix (b)	99.93
<b>Dichlorophen</b>	94.80	Glycarsamide	-93.83
<b>Ciclobendazole</b>	-2.50	Dichlorophenersine	66.43
<b>Parbendazole</b>	-4.81	beta-Kosin	95.95
<b>SRC-4402</b>	94.38	Mibbemycin Oxime A4	96.09
<b>Cambendazole</b>	79.05	Moxidentin	99.47
<b>Thiophanate</b>	-57.67	2-Naphthol	18.87
<b>Cloixanide</b>	69.61	Albendazole sulphoxide	-2.36
<b>Fenbendazole</b>	81.78	Rafoxanide	98.65
<b>Santolactone</b>	42.82	Tenium Closylate	52.10
<b>Ticarbodine</b>	91.51	Dibromomsalan	97.71
<b>Meclorazepan</b>	92.18	Milbemycin D	98.09
<b>RP12869</b>	32.95	Tribromsalam	97.87
<b>2-Azamizoribine</b>	-47.22	Buparvaquone	94.66
<b>2-fluoroNpcA</b>	-14.23	Quinapyramine	-13.66
<b>Futhan</b>	72.13	Metadone	-99.57
<b>Ribavirin</b>	-64.08	Dextromoramide	-99.56
<i>Test inactive group</i>			
<b>Aciclovir</b>	-81.10	Antazoline	-98.95
<b>RS-21592</b>	-74.34	Histamithizine	-98.22
<b>Zalcitabine</b>	-46.89	Tironamine	-94.24
<b>Triclofos</b>	97.57	Clofibrique Acid	-34.00
<b>Chlorobutanol</b>	99.44	Nicofibrate	-47.60

**Table 2.** Results of the Classification of Compounds in the Training and Test Set using QSAR Models.

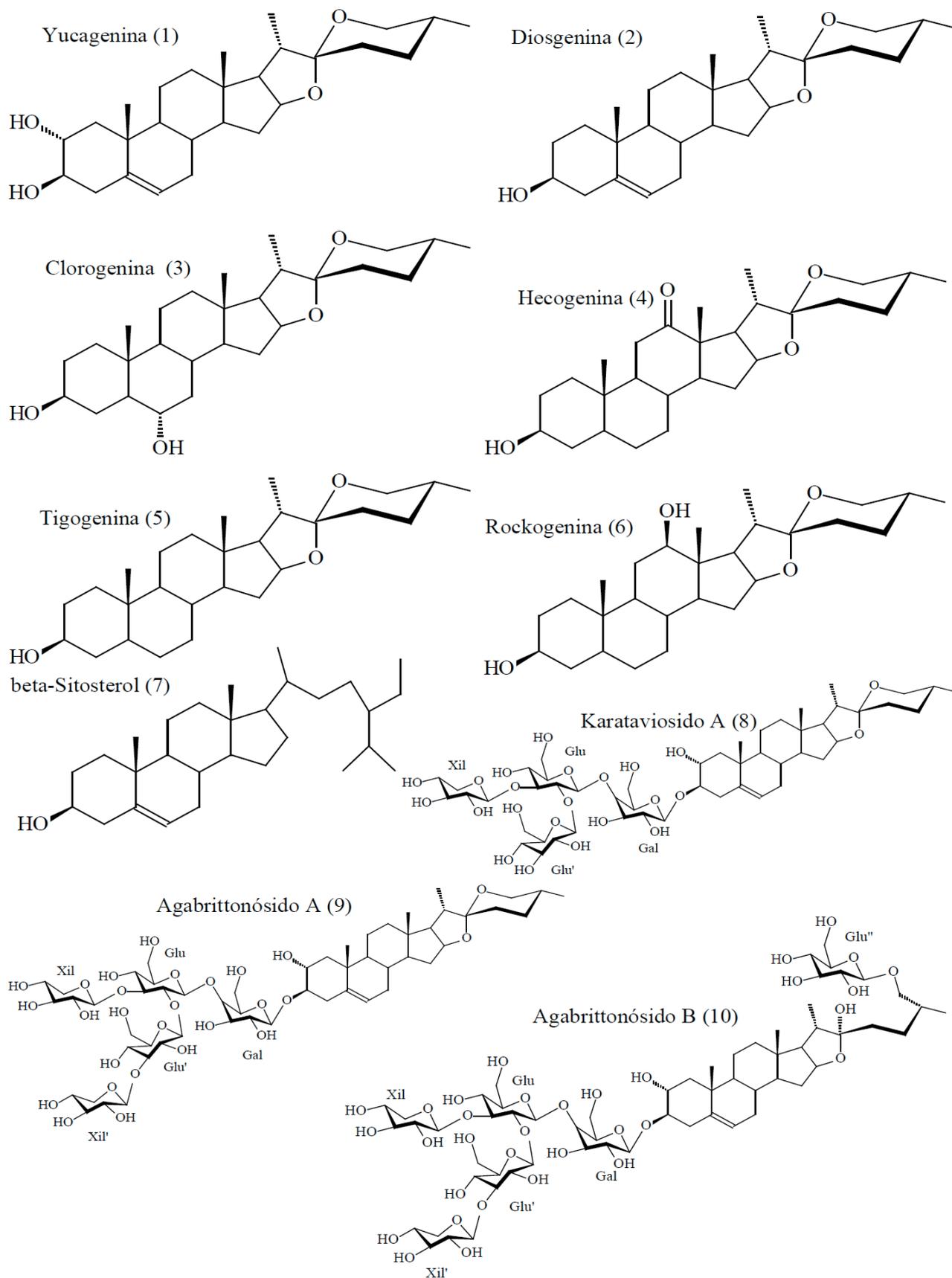
name	$\Delta P\%^a$	name	$\Delta P\%^a$
Bromisoval	-94.72	Flutizenal	8.37
Baldrianol	-66.64	Trimetilsulfonium hidroxide	-98.91
Ethchlorvynol	-51.93	l-fenilsemicarbazide	-98.35
Trimethadione	-19.96	p-Etoxyanilinometane sulfonic acid	-99.20
Phenacemide	-54.72	Pyrimazid	-95.70
UK-17022	71.11	Improsulphan	-99.42
Cinromide	51.40	Phenetylurea	-97.84
Zebromal	-99.43	Diethylstilbestrol	-97.20
Phenythilone	17.72	Norclostebol	-85.07
Iron aspartate	-75.26	Mephentermine	-92.42
Orotonsan Fe	-65.18	Mtrafazoline	-97.26
Ferrogluconat	-83.98	Chlorzoxazone	-61.01
Ferrotrenine	-77.68	Luvatren	-86.42
Cupriaseptol	-60.00	Furtrethonium iodide	-99.02
Besunide	53.38	Benzyllephedrine	-98.80
Pytamine	-8.01	Isoprenaline	-93.75
Lemidosul	48.68	Dopamine	-91.79
Propazolamide	-42.63	Octodrine	-74.53
Sulclamide	-17.71	Isometheptene	-88.25
Propamin soviet	-80.83	Metanephrene	-97.20
Calcii mesoxalas	-58.45	Bufeniode	-99.29
Mebetide	-18.70	Buphenine	-97.90
Olmidine	-37.91	Nimedipine	-95.34
Gangliefene	66.58	Odiphalin	-99.01
Gaplegin	-98.87	Acebutolol	-93.61
Trepirium	-98.48	Indopanolol	-95.92
Enflurane	23.59	Pirepolol	-98.34
Subcutin	-9.76	Colforsin	-63.68
Thiamazole	-88.68	Frutisin	-100.00
Orotic	-64.99	PR-H 286 BS	-97.99
Esorben	-78.28	Nicoxamat	-94.78
Efloxate	91.34	Aminopromazine	-99.13
Nitronal	0.11	Emepronium	-99.77
Berberine	67.88	Hymecromone	-82.19
Aecachinium	46.06	Taurultam	-97.54
Elanzepine	66.48	Fluoramphenicole	-85.13
Almoxatone mesilate	85.13		

**Table 2.** Results of the Classification of Compounds in the Training and Test Set using QSAR Models.

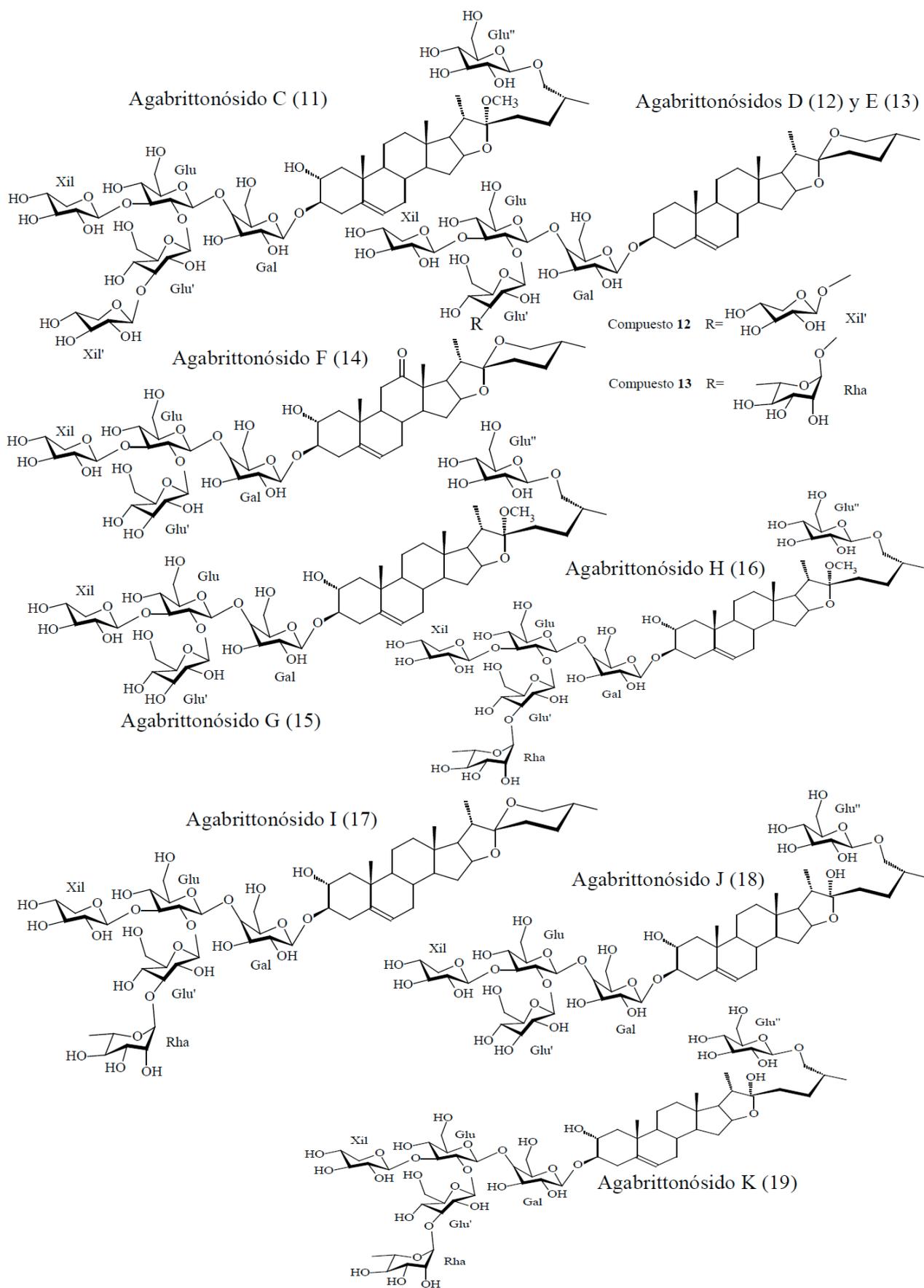
# compounds out in each step	$Q\%^a$	$\lambda$	$D^2$	$F$	$Q\%^b$
<i>LMO for the model obtained [Eq. (1)]</i>					
17	85.43	0.52	3.80	45.15	82.42
35	85.17	0.53	3.72	45.91	82.97
70	85.11	0.53	3.83	40.94	83.52
117	85.96	0.51	3.97	36.76	80.22
176	81.82	0.56	3.55	22.34	80.22
average	<b>84.70</b>	<b>0.53</b>	<b>3.77</b>	<b>38.22</b>	<b>81.87</b>
SD	<b>1.64</b>	<b>0.02</b>	<b>0.15</b>	<b>9.60</b>	<b>1.55</b>

<sup>a,b</sup> Global accuracy from both models in training and test sets (group out), respectively.

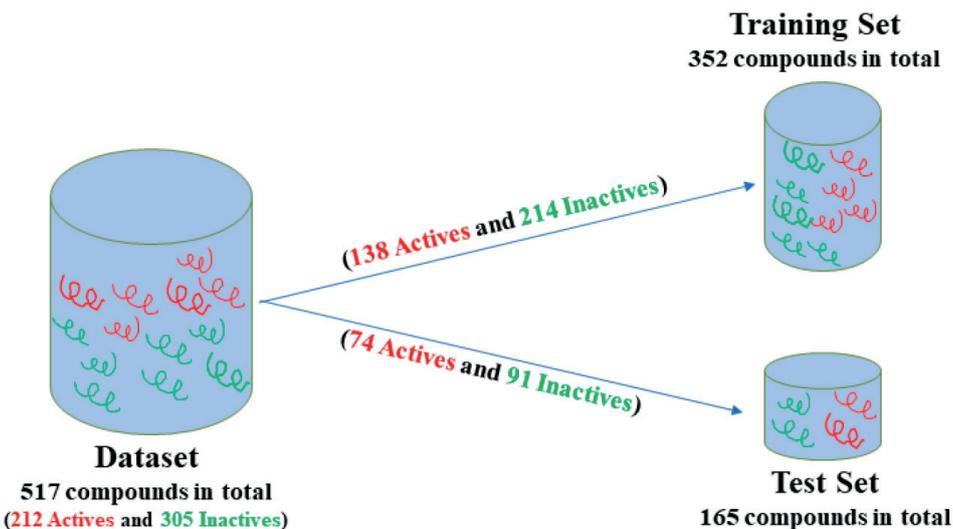
**Table 3.** Results of the Leave-Many-Out (LMO) Cross-Validation Analysis.



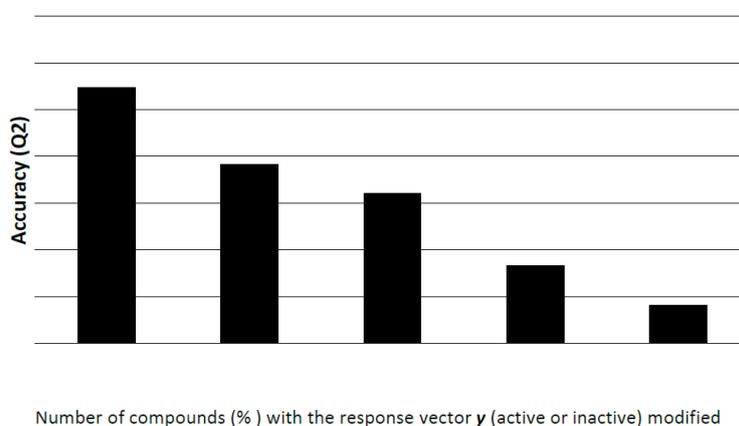
**Figure 2.** Chemical Structures of Compounds Evaluating in the in silico Experiment from *Agave brittoniana* Trel. spp. *Brachypus*.



**Figure 2.** Chemical Structures of Compounds Evaluating in the in silico Experiment from *Agave brittoniana* Trel. spp. *Brachypus*.



**Figure 3.** General overview of the computational procedure.



**Figure 4.** Behavior of the Percentage of Good Classification in the Y-scrambling Analysis.

Compounds	<i>In silico</i>	<i>In vitro</i> *						<i>In vivo</i> **
	$\Delta P$ % (Ec.1)	$5.10^{-1}$	$5.10^{-2}$	$5.10^{-3}$	$5.10^{-4}$	$5.10^{-5}$	$5.10^{-6}$	Efficacy (%)
Yucagenin (1)	-51.28	0	0	0	0	0	0	-
Diosgenin (2)	-40.23	-	-	-	-	-	-	-
Clorogenin (3)	-71.14	-	-	-	-	-	-	-
Hecogenin (4)	-50.95	-	-	-	-	-	-	-
Tigogenin (5)	-59.48	-	-	-	-	-	-	-
Rockogenin (6)	-81.05	-	-	-	-	-	-	-
$\beta$ -Sitosterol (7)	-42.83	-	-	-	-	-	-	-
Karatavinoside A (8)	56.69	100	100	100	100	0	0	92.2
Agabrittonoside A (9)	99.30	100	100	100	100	0	0	52.9
Agabrittonoside B (10)	61.41	100	0	0	0	0	0	-
Agabrittonoside C (11)	71.16	-	-	-	-	-	-	-
Agabrittonoside D (12)	77.75	100	100	100	100	100	100	-
Agabrittonoside E (13)	82.35	-	-	-	-	-	-	-
Agabrittonoside F (14)	64.46	-	-	-	-	-	-	-
Agabrittonoside G (15)	56.10	-	-	-	-	-	-	-
Agabrittonoside H (16)	76.95	-	-	-	-	-	-	-
Agabrittonoside I (17)	77.31	-	-	-	-	-	-	-
Agabrittonoside J (18)	42.95	-	-	-	-	-	-	-
Agabrittonoside K (19)	68.80	-	-	-	-	-	-	-
Triclabendazole (TCB)	96.56	100	100	100	100	0	0	92.2

\*Concentration used (in mg/mL). \*\*Compounds 8 and 9 as well as TCB were tested at 10 mg/Kg of weight of balb/c mice.

**Table 4.** Results of the in silico Classification and Percentages of Anthelmintic Activity of the Selected Compounds from *Agave brittoniana* Trel spp. *Brachypus* in vitro and in vivo Assayed.

### In vivo assay

An *in vivo* experiment using Bald/c mice-like biological models was conducted to obtain more profound conclusions about the pharmacological activity of *in vitro* hits. In this case, we only include in this experiment the two more active and pure substances (8 and 9) at doses of 3 mg/Kg. Table 4 shows the results of this study, where compound 8 was more active (92.16 % of efficacy) than 9 (52.94 %). The *in vivo* efficacy of compound 8 was identical to that of the control TCB. It is important to emphasize that this experiment was performed with a reduced dose (3 mg/kg). For instance, the TCB (the best human fasciolicide on the market<sup>65</sup>) is only wholly effective at 10 mg/kg. In addition, the few injuries in the livers and low inflammation of the spleens observed during the postmortem examination are qualitative criteria that positively appraise the effect of the tested compounds.

### Conclusions

Today virtual screening has become an essential tool in drug discovery protocols. Here, bond-level quadratic indices (QuBiLS-MAS software, <http://tomocomd.com/software/qubils-mas>) and LDA were used to obtain a QSAR model that discriminates anthelmintic from inactive ones. Virtual screening of several metabolites from *Agave brittoniana* Trel. spp. *Brachypus* was carried out to discover new lead scaffold anthelmintics, and experimental corroboration showed that Karatavinoside A (8) exhibits similar-to-superior activity as triclabendazole (fasciolicide reference drug), with 100% *in vitro* effectivity (at 500 µg/mL) against *Fasciola hepatica* and 92.2% *in vivo* efficacy (25 mg/kg). This natural compound has been identified as a promising starting point for the rational optimization/design of new chemical derivatives with more potent anthelmintic activity.

### Program availability

The QuBiLS-MAS software (portable standalone) and the respective user manual are freely available online at <http://tomocomd.com/software/qubils-mas50>

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One of the present authors (M-P. Y) thanks the program 'Estades Temporals per a Investigadors Convidats' for a fellowship to work at Valencia University (2020). Y.M.-P. and Noel Pérez acknowledge the support from Collaboration Grant 2019–2020 (Project ID16897) and Med Grant 2019–2020 (Project ID16911).

### Competing interests

The authors declare no conflict of interest.

### Author Contributions Statement

YG-C, FP-G, JOG, YE-D, NP and YM-P proposed the computational applications, QSAR modeling and performed the statistical analysis as well as prepared the manuscript. AMS, JOG, FAM, and CMN worked in the chemical methods and prepared the manuscript. EO and HS worked in the Parasitology tests. YG-C, FP-G, YE-D, NP and YMP worked in the QSAR modeling and performed the statistical analysis. All authors read and approved the final manuscript.

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