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Chemo-Informatic Comparison of Gibberellins and Anti-Gibberellins

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ABSTRACT

Molecular descriptors to differentiate four gibberellins and seven anti-gibberellins were studied. DRAGON software and Cambridge Soft ChemOffice were used to calculate 212 descriptors. Of them, 48 showed statistically significant differences between gibberellins and anti-gibberellins which can be summarized as follows. Gibberellins contain as average 14.3 times more 7-membered rings, 14.3 times more 9-membered rings, 7.4 times more ring tertiary carbon (sp3), 6.1 times more terminal tertiary carbon (sp³) and 6.0 times more terminal quaternary carbon (sp³) than anti-gibberellins. Also gibberellins usually have ring quaternary carbon (sp³) and aliphatic secondary carbon (sp²) while anti-gibberellins usually do not. On the other hand, anti-gibberellins generally have aromatic ratio, aromatic bonds, nitrogen atoms, chlorine atoms, halogen atoms, benzene-like rings, aromatic carbon (sp²), unsubstituted benzene carbon (sp²) and substituted benzene carbon (sp²) while gibberellins usually do not. A dendrogram was obtained after conducting a hierarchical cluster analysis with data of chemical molecular descriptors with statistical significant differences (48). The dendrogram correctly classified gibberellins and anti-gibberellins in two independent branches.

Keywords: Chemo-informatics, Molecular descriptors, Plant growth regulators

INTRODUCTION

Gibberellins are among the most important substances for regulating growth and morphogenesis in plant cell, tissue and organ culture [1,2]. They have been used, for instance, to control in vitro morphogenesis of sugarcane [3] pineapple [4], potato [5] bromeliads [6,7]. Modification of gibberellic acid levels are able to alter the biomass production, its allocation and may affect chemical resistance, but not tolerance [8]. The applications with gibberellic acid and abscisic acid to grapevines cv. Malbec may improve the transport of photo-assimilates from leaves to fruits by upregulation of sugar transporters at different phenological stages [9].

On the other hand, a wide range of synthetic substances, often called "anti-gibberellins", acts by blocking biosynthesis pathways. These were in general developed to achieve desirable agricultural outcomes, such as dwarfing. Anti-gibberellins are classified into four categories [10]. A number of quaternary ammonium, phosphonium and sulphonium salts act by inhibiting the cyclization process. An example of this type is chlormequat chloride (CCC), very used to regulate tomato growth and yield [11]. Certain heterocyclic nitrogen-containing compounds such as

ancymidol, paclobutrazol, uniconazole-P and tetcyclacis appear to act by inhibiting ent-kaurene oxidase [12] A further group of inhibitors are acyl cyclohexanedione derivate, for example prohexadione and diaminozide, which affect the later steps of gibberellin biosynthesis involving hydroxylases [13].

Agricultural effects of gibberellins and anti-gibberellins have been widely documented but their chemical contrasts need more studies. The present study compared the molecular descriptors of four gibberellins and seven antigibberellins (Figure 1).

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Figure 1. Gibberellins and anti-gibberellins compared.

MATERIALS AND METHODS

Our work was conducted with DRAGON software (version 5.5, 2007) and Cambridge Soft ChemOffice (version 12, 2010) with ChemDraw and Chem3D. We calculated 212 molecular descriptors. SPSS (Version 8.0 for Windows, SPSS Inc., New York, NY) was used to perform t-tests (p=0.05). The overall coefficients of variation (OCV) were calculated as described by Lorenzo et al. [14] (standard deviation/average) * 100). We considered the average values of the two growth regulators compared (gibberellins and anti-gibberellins) to calculate the standard deviation and average. The higher the difference between gibberellins and anti-gibberellins, the higher is the OCV. A dendrogram was obtained after conducting a hierarchical cluster analysis with data of chemical molecular descriptors. All variables were standardized (0-1) [15]

RESULTS AND DISCUSSION

In spite of 48 out of 212 molecular descriptors showed statistically significant differences between gibberellins and anti-gibberellins (Table 1), only 16 showed "High" OCVs (101.72 to 141.42%). Gibberellins have 14.3 times (2.00/0.14) more 7-membered rings, 14.3 times (2.00/0.14) more 9-membered rings, 7.4 times (5.25/0.71) more ring tertiary carbon (sp³), 6.1 times (5.25/0.86) more terminal tertiary carbon (sp^3) and 6.0 times (1.75/0.29) more terminal quaternary carbon (sp³) than anti-gibberellins. Also gibberellins usually have ring quaternary carbon (sp³) and aliphatic secondary carbon (sp²) while anti-gibberellins usually do not. On the other hand, anti-gibberellins generally have aromatic ratio, aromatic bonds, nitrogen atoms, chlorine atoms, halogen atoms, benzene-like rings, aromatic carbon (sp^2) , unsubstituted benzene carbon (sp^2) and substituted benzene carbon (sp^2) while gibberellins usually do not.

With "Medium" OCVs (64.82 to 98.08%), some molecular descriptors also showed statistically significant differences between gibberellins and anti-gibberellins. Gibberellins averaged 5.54 times (15.00/2.71) more circuits, 4.26 times (5.50/1.29) more ring secondary carbon (sp^3) , 4.23 times (3.00/0.71) more 5-membered rings, 3.45 times (1.00/0.29) more 8-membered rings, 3.45 times (1.00/0.29) more carboxylic acid (aliphatic), 3.45 times (1.00/0.29) more secondary alcohols, 3.52 times (2.50/0.71) more hydroxyl groups, 3.22 times (5.50/1.71) more oxygen atoms, 2.91 times (3.75/1.29) more double bonds, 2.91 times (2.50/0.86) more donor atoms for H-bonds (N and O), 2.80 times (1.29/0.46) more hydrophilic factor, 2.69 times (5.00/1.86) more rings, 5.00 times (0.10/0.02) less rotatable bonds fraction and 3.14 times (3.14/1.00) less rotatable bonds than anti-gibberellins (Table 1).

Data of Table 1, used in the hierarchical cluster analysis, generated the dendrogram shown in Figure 2. The two groups of regulators were appropriately congregated in two independent branches. Molecular descriptors have been applied to describe biological activities in many studies [16,17] showing their applicability as an attractive tool for efficient (e.g.) drug design process. It has been studied the potential of innovative computational tools in processing of structurally complex natural products to predict their macromolecular targets and attempt to forecast their role in drug discovery. Rodrigues et al. [18] and Faulon et al. [19] proposed a unified method for predicting protein-chemical interactions based on the representation of a protein using its atomistic structure. There are models that are useful in identifying compounds with potential risk of inhibiting the CYP3A4 enzyme. Arimoto et al. [20] and Kombo et al. [21] showed that shape-based descriptors ROG and SXL correlate with off-target activity and solubility, which in turn influence clinical success. They searched a potential

correlation between these shape-based descriptors and clinical success. With the rapid growth of public biological databases and biology-related web resources, abundant bioactivity data of small molecules and their targets are now available to the entire research community [22].



Figure 2. Hierarchical cluster analysis using the molecular descriptors for gibberellins and anti-gibberellins. Only those descriptors with statistical significant differences between gibberellins and anti-gibberellins were included (**Table 1**). The dendogram was built using average linkage (between groups). Variables were standardized to vary from 0 to 1 according to Kantardzic [15].

From the biochemical point of view, it is important to note that it is not possible to justify dissimilarities between different compounds that act completely different. Gibberellins are hormones and these molecules are only active in plants if they link to its receptor so its role is very specific and this specificity is due to the structure of this molecule [16]. On the other hand, the anti-gibberellins studied here are compounds that act directly in the action of some enzymes involved in gibberellins biosynthesis. So, in this case, it does not matter their structure as there is no connection with gibberellin receptors and whatever the structure of these compounds, they act blocking enzymes from gibberellin biosynthesis [10,13].

However, the procedure described here is effective to differentiate (chemically) gibberellins and anti-gibberellins. New potential growth regulators can be identified, although this preliminary result should be later tested experimentally. A similar chemo-informatic procedure was previously used by our group to compare auxins, cytokinins and gibberellins; although different molecular descriptors were analyzed [23-25].

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Table 1. Comparison of molecular descriptors for gibberellins and anti-gibberellins.															
	Gibberellins				Anti-gibberellins					. £,1	1S 2,3		f		
	GA1	GA3	GA4	GA7	CCC	Ancymidol	Paclobutrazol	Uniconazole-P	Tetcyclasis	Prohexadione	Daminozide	Gibberellins (average ± SE)	Anti-gibberellir (average ± SE)	$0CV^4$	Classification o OCV ⁵
Aromatic ratio	0	0	0	0	0	0.571	0.524	0.524	0.261	0	0	$\begin{array}{c} 0.00 \pm \\ 0.00 \end{array}$ b	0.27 ± 0.10 a	141.42	High
Number of aromatic bonds	0	0	0	0	0	12	11	11	9	0	0	0.00 ± 0.00 b	5.71 ± 2.15 a	141.42	High
Number of ring quaternary C (sp ³)	2	1	2	7	0	0	0	0	0	0	0	1.75 ± 0.25 a	0.00 ± 0.00 b	141.42	High
Number of nitrogen atoms	0	0	0	0	1	2	ω	ε	5	0	5	$0.00 \pm 0.00 b$	2.29 ± 0.61 a	141.42	High
Number of chlorine atoms	0	0	0	0	1	0	1	1	1	0	0	0.00 ± 0.00 b	0.57 ± 0.20 a	141.42	High
Number of halogen atoms	0	0	0	0	1	0	1	1	1	0	0	$\begin{array}{c} 0.00 \pm \\ 0.00 \end{array}$ b	0.57 ± 0.20 a	141.42	High
Number of benzene- like rings	0	0	0	0	0	1	1	1	1	0	0	$\begin{array}{c} 0.00 \pm \\ 0.00 \end{array}$ b	0.57 ± 0.20 a	141.42	High
Number of aromatic C (sp ²)	0	0	0	0	0	10	×	∞	6	0	0	0.00 ± 0.00 b	4.57 ± 1.67 a	141.42	High
Number of unsubstituted benzene C (sp ²)	0	0	0	0	0	4	4	4	4	0	0	$0.00\pm0.00~b$	2.29 ± 0.81 a	141.42	High

Number of substituted benzene C (sp ²)	0	0	0	0	0	7	7	7	7	0	0	0.00 ± 0.00 b	1.14 ± 0.40 a	141.42	High
Number of aliphatic secondary C (sp ²)	2	2	0	2	0	0	0	0	0	0	0	1.50 ± 0.50 a	$\begin{array}{c} 0.00 \pm \\ 0.00 \end{array}$ b	141.42	High
Number of 7- membered rings	2	2	2	2	0	0	0	0	1	0	0	2.00 ± 0.00 a	$\begin{array}{c} 0.14 \pm \\ 0.14 \ b \end{array}$	122.57	High
Number of 9- membered rings	2	2	2	2	0	0	0	0	1	0	0	2.00 ± 0.00 a	$\begin{array}{c} 0.14 \pm \\ 0.14 \ b \end{array}$	122.57	High
Number of ring tertiary C (sp ³)	5	6	5	5	0	1	0	0	2	7	0	5.25 ± 0.25 a	$0.71 \pm 0.36 b$	107.55	High
Number of terminal tertiary C (sp ³)	5	6	5	5	0	5	0	0	2	7	0	5.25 ± 0.25 a	$\begin{array}{c} 0.86 \pm \\ 0.40 \ b \end{array}$	101.72	High
Number of terminal quaternary C (sp ³)	2	1	2	2	0	0	1	1	0	0	0	1.75 ± 0.25 a	$\begin{array}{c} 0.29 \pm \\ 0.18 \end{array} \\ \end{array}$	101.72	High
Number of circuits	15	15	15	15	0	3	3	7	11	1	0	15.00 ± 0.00 a	2.71 ± 1.44 b	98.08	Medium
Rotatable bond fraction	0.020	0.021	0.019	0.020	0.053	0.108	0.122	0.103	0.029	0.111	0.182	0.02 ± 0.00 b	0.10 ± 0.02 a	94.73	Medium
Number of ring secondary C (sp ³)	5	5	7	5	0	7	0	0	5	7	0	5.50 ± 0.50 a	$\begin{array}{c} 1.29 \pm \\ 0.71 \end{array} b$	87.83	Medium
Number of 5- membered rings	3	3	3	3	0	0			ю	0	0	3.00 ± 0.00 a	$\begin{array}{c} 0.71 \pm \\ 0.42 \ b \end{array}$	87.03	Medium

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Number of 8- membered rings	-	1	1	1	0	0	0	0	2	0	0	1.00 ± 0.00 a	$\begin{array}{c} 0.29 \pm \\ 0.29 \end{array}$	78.57	Medium
Number of carboxylic acid (aliphatic)	-	1	1	1	0	0	0	0	0	-1	-	1.00 ± 0.00 a	$\begin{array}{c} 0.29 \pm \\ 0.18 \end{array}$	78.57	Medium
Number of secondary alcohols	1	1	1	1	0	0	1	1	0	0	0	1.00 ± 0.00 a	$0.29 \pm 0.18 b$	78.57	Medium
Number of hydroxyl groups	3	3	7	7	0	1	1	1	0	1	1	2.50 ± 0.29 a	$0.71 \pm 0.18 b b$	78.57	Medium
Number of oxygen atoms	6	6	5	5	0	2	1	1	0	S	ε	5.50 ± 0.29 a	$\begin{array}{c} 1.71 \pm \\ 0.68 \end{array} b$	74.21	Medium
Number of rotatable bonds	1	1	1	1	1	4	5	4	-1	ω	4	1.00 ± 0.00 b	3.14 ± 0.59 a	73.15	Medium
Number of double bonds	4	4	3	4	0	0	0	1	2	4	7	3.75 ± 0.25 a	$\begin{array}{c} 1.29 \pm \\ 0.57 \ b \end{array}$	69.21	Medium
Number of donor atoms for H-bonds	3	3	7	7	0	-	1	1	0	1	2	2.50 ± 0.29 a	$\begin{array}{c} 0.86 \pm \\ 0.26 \end{array} \\ \end{array}$	69.21	Medium
Hydrophili c factor	1.638	1.670	0.927	0.927	0.004	0.371	0.401	0.401	0.000	0.538	1.508	1.29 ± 0.20 a	$\begin{array}{c} 0.46 \pm \\ 0.19 \end{array}$	67.04	Medium
Number of rings	5	5	5	5	0	ю	3	3	5	1	0	5.00 ± 0.00 a	$\begin{array}{c} 1.86 \pm \\ 0.67 \ b \end{array}$	64.82	Medium
Number of terminal secondary C (sp ³)	5	5	7	5	0	2	ю	1	5	ю	7	5.50 ± 0.50 a	2.29 ± 0.61 b	58.38	Low

Number of 6- membered rings	2	2	2	2	0	7	1	1	5	-	0	2.00 ± 0.00 a	$\begin{array}{c} 1.00 \pm \\ 0.31 \ b \end{array}$	47.14	Low
Topologic al polar surface area using	104.06	104.06	83.83	83.83	0	55.24	50.94	50.94	52.68	88.51	69.64	93.95 ± 5.84 a	52.56 ± 10.19 b	39.94	Low
Topologic al polar surface area using	104.06	104.06	83.83	83.83	0	55.24	50.94	50.94	52.68	88.51	69.64	93.95 ± 5.84 a	52.56 ± 10.19 b	39.94	Low
Number of non-H bonds	29	28	28	28	6	21	21	21	23	15	10	28.25 ± 0.25 a	16.71 ± 2.48 b	36.28	Low
Number of carbon atoms	19	18	19	19	5	15	15	15	13	10	9	18.75 ± 0.25 a	11.29 ± 1.64 b	35.15	Low
Sum of Kier-Hall electrotopological states	65.83	63.92	58.92	59.92	14.61	42.80	44.86	45.69	38.94	49.83	34.83	62.15 ± 1.64 a	38.79 ± 4.43 b	32.72	Low
Number of bonds	51	48	52	50	19	37	41	39	35	27	22	50.25 ± 0.85 a	31.43 ± 3.29 b	32.59	Low
Number of acceptor atoms for H-bonds (N, O, F)	6	6	5	5	0	4	3	3	S	S	5	$5.50 \pm 0.29 a$	$3.57 \pm 0.69b$	30.07	Low
Sum of conventional bond orders (H-depleted)	33	32	31	32	6	27	26.5	27.5	28	19	12	32.00 ± 0.41 a	20.86 ± 3.34 b	29.81	Low
Numb of non-N atoms	25	24	24	24	7	19	20	20	19	15	11	24.25 ± 0.25 a	15.86 ± 1.93 b	29.59	Low

Electronegativities (scaled on carbon	47.68	44.80	48.24	46.36	19.67	35.04	39.91	38.03	31.37	27.94	23.60	46.77 ± 0.77 a	$30.79 \pm 2.83 b$	29.13	Low
Number of atoms	47	44	48	46	20	35	40	38	31	27	23	46.25 ± 0.85 a	30.57 ± 2.87 b	28.86	Low
Sum of atomic van der Waals volumes (scaled on carbon atom)	28.65	27.05	28.73	28.13	10.58	22.20	24.57	23.98	21.06	16.15	12.51	$28.14 \pm 0.39 a$	$18.72 \pm 2.13 b$	28.42	Low
Sum of atomic polarizabilities (scaled on carbon atom)	30.10	28.34	30.41	29.65	11.81	23.25	26.18	25.42	21.93	16.84	13.18	29.63 ± 0.46 a	19.80 ± 2.21 b	28.11	Low
Number of hydrogen atoms	22	20	24	22	13	16	20	18	12	12	12	22.00 ± 0.82 a	14.71 ± 1.25 b	28.06	Low
Molecular weight	346.41	332.38	332.43	330.41	122.64	256.33	293.83	291.81	273.75	212.22	160.20	335.41 ± 3.70 a	230.11 ± 25.46 b	26.33	Low
Ghose-Crippen molar refractivity	86.414	81.913	84.078	84.887	32.130	72.148	81.795	83.946	77.388	49.650	38.851	$84.32 \pm 0.94 a$	$62.27 \pm 8.16 \text{ b}$	21.27	Low

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¹ Average information of GA1, GA3, GA4 and GA7

² Average information of 2-chloro-N,N,N-trimethylethanaminium chloride; ancymidol; paclobutrazol; uniconazole-P;

tetcyclasis; prohexadione and daminozide

³ Results with the same letter are not statistically different (t-test, p>0.05)

⁴ Overall coefficient of variation = (Standard deviation/Average)*100. To calculate this coefficient. Average values of gibberellins and anti-gibberellins were considered The higher the difference between these two averages, the higher the overall coefficient of variation

⁵ Classification of OCVs: "Low" from 21.27 to 58.38%; "Medium" from 64.82 to 98.08% and "High" from 101.72 to 141.42%

*GA1: Gibberellin 1; GA3: Gibberellic acid; GA4: Gibberellin 4; GA7: Gibberellin 7; CCC: 2-chloro-N,N,N*trimethylethanaminium chloride; ancymidol; paclobutrazol; uniconazole-P; tetcyclasis; prohexadione and daminozide

AUTHOR CONTRIBUTION

I.A., D.G., L.P. and J.C.L. designed the research, analyzed the data and wrote the paper. J.C.L. had primary responsibility for the final content. All authors have read and approved the final manuscript.

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COMPLIANCE WITH ETHICAL STANDARDS

Conflict of interest

Authors do not have any conflict of interests.

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