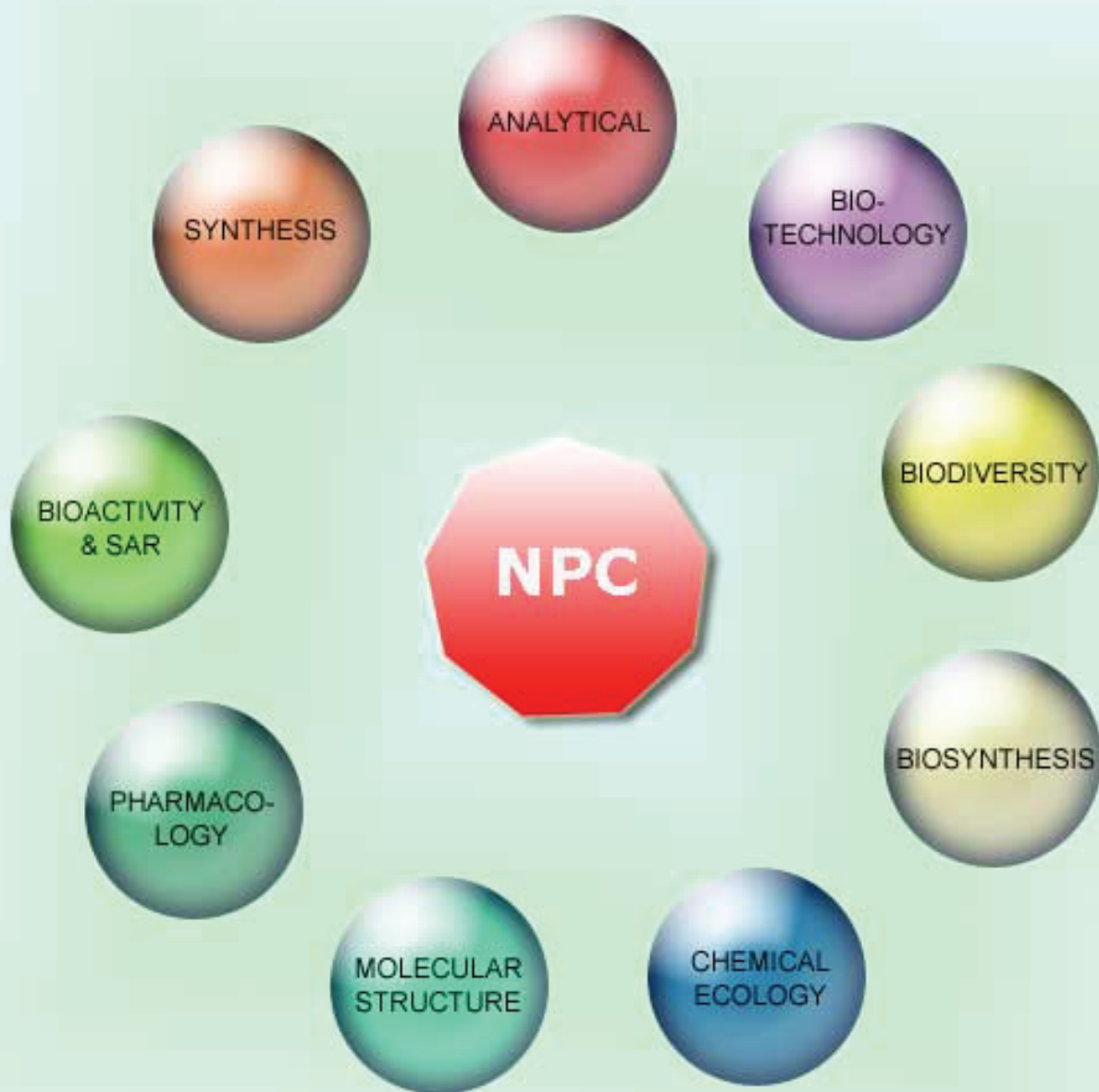


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Synergism between Terbinafine and a *Neo*-clerodane Dimer or a Monomer Isolated from *Baccharis flabellata* against *Trichophyton rubrum*

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The new *neo*-clerodane dimer (DACD) likely formed by a [4+2] photo-cycloaddition between two molecules of ent-15,16-epoxy-19-hydroxy-1,3,13(16),14-clerodatetraen-18-oic acid (DAC), both isolated from *Baccharis flabellata* Hook et Arn. var. *flabellata* (Asteraceae), were tested in combination with terbinafine (Terb) against *Trichophyton rubrum*. The interactions were assessed with different methodologies such as 3D *CombiTool* software, Loewe Combination Index (CI) and isobolograms. Results showed that the monomer DAC as well as its dimer DACD act synergistically with Terb against *T. rubrum*. DAC/Terb and DACD/Terb showed eight and twelve synergistic mixtures respectively in the *CombiTool* graphics thus suggesting that DACD/Terb was more synergistic with Terb than DAC. These results were corroborated by the CI values and the isobolograms of the detected combinations. DACD/Terb showed much lower CIs (0.34-0.47) than DAC/Terb (0.57-1.15) at the different inhibition percentages tested. Concomitantly, DACD/Terb isobolograms were more concave than for DAC/Terb at the different inhibition percentages. This is the first report showing antifungal synergistic interactions of *neo*-clerodanes dimers and add new evidences that *neo*-clerodanes monomers interact synergistically with Terb against *T. rubrum*.

Keywords: Terbinafine, *Trichophyton rubrum*, *Neo*-clerodane monomer, *Neo*-clerodane dimer, Synergism.

Diterpenoid dimers are a rather uncommon subclass of diterpenoids which are composed of two 20-carbon diterpenoid units linked through either one or two C-C bond, an ester bond or a ring moiety and are naturally synthesized mainly by an enzyme-catalyzed Diels Alder cycloaddition [1]. Among the great structural diversity of diterpenoid dimers, natural *neo*-clerodanes diterpene dimers have been scarcely reported up to date. A new dimer (DACD, Figure 1) was recently isolated by us from *B. flabellata* [2]. It showed to be formed via a [4+2] cycloaddition of ent-15,16-epoxy-19-hydroxy-1,3,13(16),14-clerodatetraen-18-oic acid (DAC) (Figure 1), one of the major *B. flabellata* *neo*-clerodanes. The structure elucidation and absolute configuration of DAC and DACD have been recently reported [2] and can be found as Supplementary data.

In another previous paper, we have reported that the *neo*-clerodanes monomers bacchotricuneatin A and bacrispine, isolated from *Baccharis tricuneata* (L.f.) Pers [3] and *Baccharis crispa* Spreng. respectively among other spp. [4,5] showed moderate antifungal activity against *Trichophyton rubrum* [6]. However, when tested in combination with terbinafine (Terb), their activities were highly enhanced. Considering this previous finding, the antifungal activity of the clerodane dimer DACD as well as its monomer DAC were tested here alone and in combination with Terb against *T. rubrum* in order to have a look on the anti-*T. rubrum* enhancing behavior of a *neo*-clerodane dimer in comparison with the monomer.

T. rubrum is the major dermatophyte that cause human skin, nail and eye infections [7] and although superficial fungal infections are not life threatening, they usually are very difficult to eradicate and new antifungal compounds are highly needed.

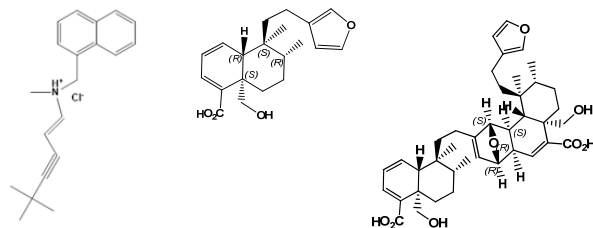


Figure 1: Structures of terbinafine (left), DAC (medium) and DACD (right)

Regarding the selected antifungal drug, it is well-known that Terb (Figure 1) is the preferred antifungal drug to treat superficial mycoses produced by dermatophytes [8] although it shows various side-effects that include liver problems and allergic reactions.

Considering that the combination of a single natural product with an antifungal chemosynthetic drug can improve the effectiveness of the antifungal drug at a lower dose [9], combination of Terb/DAC or Terb/DACD could take advantage of the high activity of Terb highly diminishing its side effects and also preventing or delaying the emergence of resistant populations of the pathogenic fungus.

Regarding the several methods of choice to test compounds in combination [10,11], we first selected a three-dimensional (3D) study with the *CombiTool* software [12]. The 3D-model is considered the most complete way to identify the regions of significant synergy and antagonism, and presents a complete map of drug interactions in a way that can be easily interpreted.

CombiTool software first needs the construction of single-agent dose-response curves for the calculation of zero interaction response surfaces [12]. The parameters 'a' and 'm' obtained from these curves are used by the software to generate a 3D-zero interaction surface (represented in green in Figure 2), according to the Loewe additivity criteria.

Then, the experimental effects of DAC/Terb (A) and DACD/Terb (B) combinations (formed with different ratios of each partner) were introduced in *CombiTool* software. Any statistically significant deviation of these combinations from the zero interaction surface is indicated by the position of red balls, each representing one combination (Figure 2). Balls over the zero interaction surface are indicative of synergism; within the surface, are indicative of additivism and below the surface, of antagonism. The red balls in Figure 2 for DAC/Terb and DACD/Terb show the interactions at all effect levels (from 1 to 100 % inhibition). However, for the sake of clarity, only the synergistic combinations with > 50 % inhibition level are represented in Figure 2.

Figure 2 shows that eight combinations of DAC/Terb and twelve combinations of DACD/Terb showed synergism, thus suggesting that not only the monomer DAC but its dimer DACD interact synergistically with Terb, although DACD shows more synergistic combinations with an inhibition effect ≥ 50 %. In Table 1, the composition of the combinations and the percentages of inhibition at which they showed synergism are detailed.

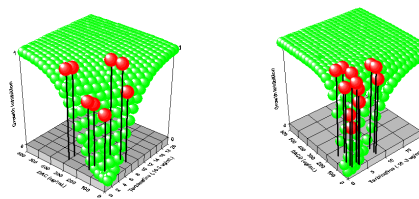


Figure 2: Surfaces of experimental effects for DAC/Terb (left), DACD/Terb (right) obtained with *CombiTool* software. Red spheres over the surface correspond to combinations that show synergism with an inhibition effect ≥ 50 %.

In a second step, the CI of the synergistic combinations detected with the *CombiTool* software were calculated as follows: an inoculum of *T. rubrum* was incubated with serial dilutions of Terb (7.8, 3.9 and 1.56×10^{-3} $\mu\text{g/mL}$), added with fixed sub-inhibitory concentrations of DAC (300, 150, 75, 37.5, 18.75 or 9.38 $\mu\text{g/mL}$) or DACD (150, 75, 37.5, 18.75 or 9.38 $\mu\text{g/mL}$). The % of growth for Terb and each combination were represented in dose-response curves (Figure 3). In the two-drug combinations curves of Terb added with different fixed concentrations of DAC (Figure 3A) or DACD (Figure 3B), it is clear that both *neo*-clerodanes decreased the MIC₅₀, MIC₇₀, MIC₉₀, and MIC₁₀₀ (dotted lines a-d) of Terb, thus enhancing its antifungal capacity with lower doses. The MIC₅₀, MIC₇₀, MIC₉₀, and MIC₁₀₀ values of Terb, DAC and DADC alone and in dual combinations DAC/Terb and DACD/terb along with the Dose Reduction Index (DRI) and the CI are shown in Table 2.

Table 1: Synergistic combinations of DAC/Terb and DACD/Terb detected with the *CombiTool* software. The number between each bracket indicates the percentage of inhibition at which each combination showed synergism (S).

Terb	Conc. $\mu\text{g/mL}$											
	300		150		75		37.5		18.75		9.38	
	DAC	DACD	DAC	DACD	DAC	DACD	DAC	DACD	DAC	DACD	DAC	DACD
7.80×10^{-3}	-	-	S (100)	-	S (100)	S (100)	S (70)	S (100)	-	S (90)	-	-
3.12×10^{-3}	S (90)	-	S (70)	S (100)	S (70)	S (100)	-	S (90)	-	S (90)	-	S (70)
1.56×10^{-3}	S (90)	-	S (70)	S (100)	-	S (90)	-	S (50)	-	S (50)	-	-

Table 2: Antifungal activity of terbinafine (Terb) and the *neo*-clerodane monomer (DAC) and dimer (DACD) at different inhibition percentages (MIC₁₀₀, MIC₉₀, MIC₇₀ and MIC₅₀) either alone or in the two-drug combinations DAC/Terb and DACD/Terb. The Combination Index (CI) and the Dose-Reduction Index (DRI) in combination were calculated for *Trichophyton rubrum* fungal sp.

		MICs in $\mu\text{g/mL}$				CI	DRI Terb
		MICs at different inhibition %	MIC _{DAC} alone	MIC _{Terb} alone $\times 10^{-3}$	MIC _{DAC} in comb		
DAC/ Terb	MIC ₁₀₀		558.0 ± 10.2	15.6 ± 1.1	75.0 ± 1.1	7.8 ± 0.6	2.0
	MIC ₉₀		409.0 ± 7.9	14.4 ± 0.8	300.0 ± 0.1	3.1 ± 0.0	4.6
	MIC ₇₀		312.0 ± 5.3	9.4 ± 0.2	75.0 ± 0.9	3.1 ± 0.1	3.0
	MIC ₅₀		258.0 ± 6.3	7.2 ± 0.0	18.5 ± 0.8	7.8 ± 0.1	0.9
DACD/ Terb			MIC _{DACD} alone		MIC _{DACD} in comb		
	MIC ₁₀₀		274.0 ± 3.5	15.6 ± 1.4	75.0 ± 3.8	3.1 ± 0.5	5.0
	MIC ₉₀		145.0 ± 2.4	14.4 ± 0.8	18.8 ± 1.5	3.1 ± 0.1	4.6
	MIC ₇₀		99.0 ± 4.2	9.4 ± 0.8	9.4 ± 2.3	3.1 ± 0.1	3.0
	MIC ₅₀		77.0 ± 3.1	7.2 ± 0.0	18.8 ± 1.1	1.6 ± 0.0	4.6

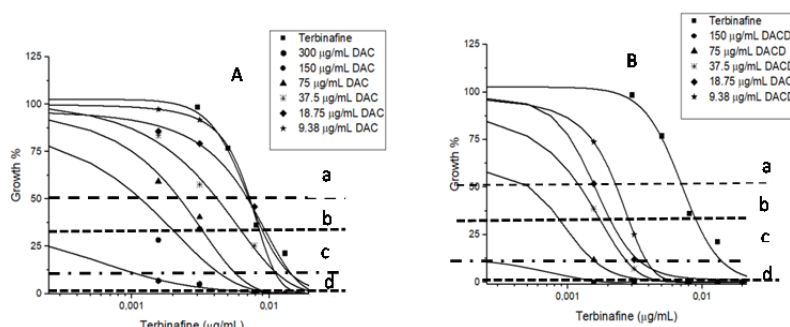


Figure 3: Dose-response curves of terbinafine with fixed concentrations of the monomer DAC (A) and the dimer DACD (B) against *Trichophyton rubrum*. The response is presented as percentage of growth. The horizontal dotted lines at 50 % inhibition (50% growth) (a); 70 % inhibition (30 % growth) (b); 90 % inhibition (10 % growth) (c) and 100 % inhibition (0 % growth) (d) were drawn for an easier comparison of DAC's and DACD's effects against *T. rubrum*.

From Table 2 it is clear that DACD/Terb is more synergistic than DAC/Terb since DACD/Terb showed much lower CIs (0.3-0.5) than DAC/Terb (0.6-1.2), being the ratio DACD (18.75 $\mu\text{g/mL}$)/Terb ($3.12 \times 10^{-3} \mu\text{g/mL}$) the one that showed the most synergistic interaction for displaying a percentage of inhibition of 90 %. For achieving a 100 % inhibition, the ratio should be 75 $\mu\text{g/mL}$ DACD/ $3.12 \times 10^{-3} \mu\text{g/mL}$ Terb. In this combination the MIC_{100} of Terb decreased 5-fold (from 15.60 to $3.12 \times 10^{-3} \mu\text{g/mL}$, DRI = 5) and the MIC_{90} , from 14.37 to $3.12 \times 10^{-3} \mu\text{g/mL}$ with DRI = 4.6. In contrast, the MIC_{100} and MIC_{90} of Terb decreased 2- and 4.6-fold (DRI = 2.0 and 4.6) when combined with DAC showing lower synergistic interaction with CIs = 0.6 and 0.8 respectively.

For combinations that produced 70 or 50 % inhibition, also DACD showed better interaction effects (CIs = 0.4-0.5, with DRIs = 3.0 and 4.6) than DAC (CI = 1.2-0.6, with DRIs = 3.0-0.9). The corresponding isobolograms show the behavior of the combinations at the different effect levels [13]. The isobolograms at 100 % inhibition are presented in Figure 4. They are representative of the isobolograms at other effect levels such as 50, 70 and 90 % inhibition percentages.

The comparison of the isobologram of DACD/Terb (left) with that of DAC/Terb (right) at the same effect level shows that the isobologram of DACD/Terb have higher concavity than that of DAC/Terb, corroborating that DACD showed a greater synergism with Terb than DAC.

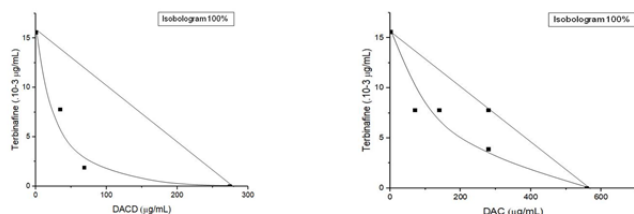


Figure 4: Isobolograms of the combinations DACD/Terb (left) and DAC/Terb (right) at 100 % effect level. MIC_{100} of Terb is plotted in the y-axis and of DACD or DAC on the x-axis. The lines connecting the MIC in each isobologram of each partner is the line of additivity. Points located below the line indicate synergy, above the line antagonism and on the line additivity.

As conclusions, we could demonstrate that the *neo*-clerodane monomer DAC as well its dimer DACD behave synergistically with Terb against *T. rubrum*. However, the dimer DACD showed a higher synergism than DAC clearly demonstrated by different methodologies such as 3D, CI and isobolograms.

Regarding the monomer DAC, the results of this work add new evidences that the monomers of *neo*-clerodanes interacts synergistically with Terb against *T. rubrum*. As for the dimer DACD, this is the first report on the synergism of a *neo*-clerodane dimer with Terb against *T. rubrum* and also the first demonstration that it is a better enhancer of the activity of the antifungal drug Terb. Both compounds, mainly the dimer, enhance the activity of Terb at lower doses than the required for acting alone. Since Terb produces unwanted side effects, the fact that it can be used in combination at lower amounts with the same effectivity, is an important finding for a future pharmaceutical development of these combinations.

Experimental

Chemicals: Terb hydrochloride ($\geq 98\%$) was purchased from Sigma-Aldrich (St Louis, MO, USA). DAC and DACD ($\geq 98\%$ each) were isolated from *B. flabellata* as reported in [2]. Their HPLC profiles are shown in Supplementary Material Figures 1S and

2S. One specimen of *B. flabellata* was deposited in the Herbarium of the National University of San Luis (UNSL voucher number L.A. Del Vitto & E.M. Petenatti # 9436).

Fungal strain and inoculum preparation: *T. rubrum* CCC 110 (CEREMIC, Centro de Referencia en Micología, Facultad de Ciencias Bioquímicas y Farmacéuticas, Rosario, Argentina) was used for the antifungal evaluation. This is the same strain used in the previous paper [6]. The growth of the strain was performed on Sabouraud-chloramphenicol agar during 7 d at 30°C , maintained on slopes of Sabouraud-dextrose agar (SDA) (Oxoid, Ontario, Canada). Inocula were adjusted to $1-5 \times 10^3$ spores with colony forming units (CFU)/mL [14].

Determination of MIC: Minimum Inhibitory Concentration (MIC) was determined according to the guidelines of the CLSI for filamentous fungi (M38 A2) [14]. Dilutions of stock Terb or pure compounds in DMSO (final concentration $\leq 1\%$) in RPMI-1640 (Sigma-Aldrich) (200 μL) were poured into the first well of a 96-well microplate, and then, 100 μL were transferred to the next well containing 100 μL of RPMI-1640 buffered to pH 7.0 with 4-morpholinepropanesulfonic acid (MOPS, Sigma-Aldrich). A volume of 100 μL of inoculum suspension was added to each well (for sterility control, sterile water was added to the well instead) rendering concentrations from $1-30 \times 10^{-3} \mu\text{g/mL}$ for Terb and from 800 to $31.25 \mu\text{g/mL}$ for pure compounds. Plates were incubated for 7 days at $28-30^\circ\text{C}$ in a moist, dark chamber. MIC_{100} was defined as the lowest concentration of Terb or pure compound resulting in total inhibition of visual growth compared to the growth in the control wells containing no antifungal. Tests were carried out in duplicate.

Determination of growth inhibition percentages of *T. rubrum* and MIC_{50} , MIC_{70} , MIC_{90} of DAC and DACD: *T. rubrum* growth inhibition was obtained by image analysis [6,15]. After incubation, the plates with the assays were placed on an Epichemi3 darkroom (UVP, Upland, CA, USA). Photographs of each plate were recorded with a high resolution cooled digital monochrome CCD camera (Hamamatsu C8484-51-03G). Pictures were analyzed using LabWorks version 4.6 (UVP) MacBiophotonics ImageJ and according to the steps previously described [6].

Analysis with CombiTool software: Eighteen combinations of Terb and each pure compound (DAC or DACD) were tested. Sub-inhibitory concentrations for DAC ($\text{MIC}_{100} = 558.95 \mu\text{g/mL}$) and DACD ($\text{MIC}_{100} = 274.84 \mu\text{g/mL}$) and 1/2, 1/5 and 1/10 MIC for Terb ($\text{MIC}_{100} = 15.6 \times 10^{-3} \mu\text{g/mL}$) were combined in a microtiter tray by duplicate as described for MIC_{100} determinations above. The inhibition percentage of each combination was determined after 7 days incubation time by image analysis. The program used to obtain graphs was the *CombiTool* kindly provided by Dr. Jürgen Sühnel (Jena, Germany) [12]. Dose-responses curves for DAC, DACD and Terb were performed and parameters “a” and “m” were determined by the median-effect equation [16] with the nonlinear regression module of the Origin software package (www.originlab.de). Loewe Additivity response surface (LA) was calculated for DAC, DACD and Terb with a and m parameters. The comparison of the surface obtained with experimental data vs the zero response surface allows the direct visualization of spheres which, if they fall into the surface are indicative of additivism, over the surface indicate synergism and below the surface show antagonism.

Analysis of the combination effect: The nature of the interaction (synergy, additivity, or antagonism) between an individual DAC or DACD and Terb as a function of their concentrations and fungal

growth inhibition (MIC₅₀, MIC₇₀, MIC₉₀, MIC₁₀₀) was assessed by the combination index (CI) [16] as follows:

$$CI = \frac{MIC_x \text{ Terb in comb}}{MIC_x \text{ Terb alone}} + \frac{MIC_x \text{ NC* in comb}}{MIC_x \text{ NC* alone}}$$

Where x refers to the inhibition percentage and can be 50, 70, 90 or 100 %; * NC represents one of the natural neo-clerodanes that can be DAC or DACD.

CI helps to identify synergistic (CI < 1), additive (CI = 1), and antagonistic interactions (CI > 1) [16].

The dose reduction index (DRI) is a measure of how many -fold the concentrations in the combination may be reduced as compared with the doses of each drug alone [16] and is calculated as follows:

$$DRI = \frac{MIC_x \text{ of Terb alone}}{MIC_x \text{ Terb in combination}}$$

DRI > 1 is beneficial, indicating synergism and is important in clinical situations, in which dose reduction leads to reduced toxicity while the therapeutic efficacy is retained [16].

Isobolograms: The isobologram illustrates the result of the checkerboard assay and the CI values. The axis of the isobologram represents the dose of DAC or DACD and the ordinate represents the dose of Terb. The straight line connecting the MIC points represents zero interaction [13]. Below and above this line the areas of synergistic and antagonistic interactions respectively are found. The concentrations of the two drugs used in combination to provide the same effect (50, 70, 90 or 100 % inhibition) were placed in the same plot.

Supplementary data: Median-effect equation parameters of pure compounds Terb, DAC and DACD and UV-Vis spectrum, HPLC-DAD of DAC and DACDs and isobolograms at all effect levels are available in electronic form on the publisher's website.

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