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Original Citation:

Availability: This version is available at: 11577/3244021 since: 2019-04-09T15:59:50Z

Publisher:

Published version: DOI: 10.1016/j.fitote.2017.11.002

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Accepted Manuscript

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PII: DOI: Reference:	S0367-326X(17)31413-2 doi:10.1016/j.fitote.2017.11.002 FITOTE 3733
To appear in:	Fitoterapia
Received date: Revised date: Accepted date:	25 September 201726 October 20171 November 2017

Please cite this article as: Stefania Sut, Stefano Dall'Acqua, Valeria Baldan, Stephane L. Ngahang Kamte, Farahnaz Ranjbarian, Prosper C. Biapa Nya, Sauro Vittori, Giovanni Benelli, Filippo Maggi, Loredana Cappellacci, Anders Hofer, Riccardo Petrelli, Identification of tagitinin C from Tithonia diversifolia as antitrypanosomal compound using bioactivity-guided fractionation. The address for the corresponding author was captured as affiliation for all authors. Please check if appropriate. Fitote(2017), doi:10.1016/j.fitote.2017.11.002

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Identification of Tagitinin C from *Tithonia diversifolia* as antitrypanosomal compound using bioactivity-guided fractionation

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Tithonia diversifolia (Asteraceae), is used as traditional medicine in tropical countries for the treatment of various diseases, including malaria. Although numerous studies have assessed the antimalarial properties, nothing is known about the effect of T. diversifolia extracts on trypanosomiasis. In this study extracts of T. diversifolia aerial parts were evaluated for their bioactivity against Trypanosoma brucei. The activity was studied against bloodstream forms of T. brucei (TC221), as well as against mammalian cells (BALB/3T3 mouse fibroblasts), as a counter-screen for toxicity. Both methanolic and aqueous extracts showed significant effects with IC₅₀ values of 1.1 and 2.2 µg/mL against T. brucei (TC221) and 5.2 and 3.7 µg/mL against BALB/3T3 cells, respectively. A bioassay-guided fractionation on the methanolic extract yielded in identification of active fractions (F8 and F9) with IC₅₀ values of 0.41 and 0.43 μ g/mL, respectively, against T. brucei (TC221) and 1.4 and 1.5 µg/mL, respectively, against BALB/3T3 cells,. The phytochemical composition of the extracts and the purified fractions were investigated using HPLC-ESI-MS/MS and 1D and 2D NMR spectra showing the presence of sesquiterpene lactones that in turn were subjected to the isolation procedure. Tagitinin A and C were rather active but the latter presented a very strong inhibition on T. brucei (TC221) with an IC₅₀ value of 0.0042 µg/mL. This activity was 4.5 times better than that of the reference drug suramin. The results of this study shed light on the antitrypanosomal effects of T. diversifolia extracts and highlighted tagitinin C as one of the possible responsible for this effect. Further structure activity relationships studies on tagitinins are needed to consider this sesquiterpenes as lead compounds for the development of new antitrypanosomal drugs.

Keywords: BALB/3T3 mouse fibroblasts; Mexican sunflower; HPLC-ESI-MS/MS;

NMR; Tagitinins A and C; Sesquiterpene lactones

Abbreviations:

1D and 2D NMR: One and two-dimensional nuclear magnetic resonance

BALB//3T3 = Mouse fibroblasts

CH₂Cl₂/MeOH: Methylene chloride/Methanol

DMSO: Dimethylsulfoxide

HAT: Human African Trypanosomias is

NTDs: Neglected tropical diseases

SAR: Structure-activity relationship

SI: Selectivity index

STLs : Sesquiterpene lactones

T. brucei: Trypanosoma brucei

TC221: T. brucei bloodstream-form parasites

TLC: Thin layer chromatography

WHO: World Health Organization

Introduction

Neglected tropical diseases (NTDs) are classified as a subgroup of infectious illnesses affecting more than a billion people worldwide and causing significant public health problems, particularly in marginalized populations of rural developing countries (Benelli and Mehlhorn, 2016). People suffering from NTDs live in close contact with infectious vectors and domestic livestock and represent an unattractive market to pharmaceutical industry investments. For this reason, in 2010 the World Health Organization (WHO) released its first report on the global treat posed by NTDs. In this report WHO prioritized 17 NTDs. Of these 17, the three major protozoan diseases, namely human African trypanosomiasis (HAT), Chagas disease and leishmaniasis, have been accounted as life-threatening infections (Mehlhorn 2015).

African trypanosomiasis, which includes African sleeping sickness in humans (HAT) and "Nagana" in cattle, are fatal vector-borne diseases caused by the kinetoplastid protozoan pathogen *Trypanosoma brucei* (*T. brucei*). HAT is endemic in 36 sub-Saharan African countries and is transmitted to the host by the bites of tsetse flies (*Glossina* spp., Glossinidae). The disease progresses through two stages, *i.e.* from a haemolymphatic first stage to a meningoencephalitic second stage, in which the parasites enter the central nervous system. HAT is caused by two subspecies of the parasite *T. brucei: T. b. gambiense* (West Africa) and *T. b. rhodesiense* (East Africa). *T. b. rhodesiense* as well as other African trypanosomes such as *T. congolense*, *T. vivax* and *T. brucei brucei* infect wild and domestic animals causing "Nagana", which has a significant impact on socioeconomic development in many parts of rural Africa. In

about eleven million square kilometres of farming land infested by the tsetse flies, cattle represent an important zoonotic reservoir for human infections.

There is a pressing need to develop novel and cost-effective drugs to treat HAT, especially at later stages when the parasites infect the brain. The lack of potential vaccine candidates together with the uncertainty for a successful chemotherapy against HAT add impetus to explore naturally occurring compounds as inspiring leads for designing future trypanocidal drugs (Scotti et al., 2016).

Among several medicinal plants traditionally used in Africa to treat different diseases and having potential as a source of trypanocidal compounds, our attention has been attracted by Tithonia diversifolia (Hemsl.) A. Gray. This plant, belonging to the Asteraceae family, is also known as Mexican sunflower, being native to central America. It is distributed in tropical areas of South America, Asia and Africa where it can be considered as a weed of cultivated and non-cultivated lands (Chagas-Paula et al., 2012; Ayeni et al., 1997). Given its abundance, T. diversifolia enjoys a long use as a traditional remedy in tropical regions (Heinrich, 2000). In particular, the plant leaves are used in Africa to treat malaria (Heinrich et al., 1998; Njoroge and Bussman, 2006). Noteworthy, this use was then confirmed by scientific studies (Goffin et al., 2002; Elufioye and Agbedahunsi, 2004; Oyewole et al., 2008) where the bioactive compounds were identified as germacranolide-type sesquiterpene lactones called tagitinins, with tagitinin C as the lead compound (Goffin et al., 2002; Ferreira et al., 2005). These sesquiterpene lactones (STLs) were also proven to exhibit anti-inflammatory/analgesic (Owoyele et al., 2004), antiproliferative (Liao et al., 2011), antifeedant (Ambrósio et al., 2008) and insecticidal activities (Castaño-Quintana et al., 2013). Other secondary

metabolites characteristics of *T. diversifolia* are diterpenoids, flavonoids, chlorogenic acid derivatives and essential oils (Orsomando et al., 2016).

Given the potential of *T. diversifolia* secondary metabolites to interact with protozoan parasites (Goffin et al., 2002), here we have investigated the effect of a *T. diversifolia* methanolic leaf extract against *T. brucei* using a bioassay-guided fractionation approach, leading to the identification of tagitinin A and C as main active constituents. To our knowledge, nothing is known about either the anti-trypanosomal activity of *T. diversifolia* extracts or the components responsible for this effect.

Different techniques were applied for the identification of phytoconstituents. Methanolic extract was initially studied by ${}^{1}HNMR$ and 2D NMR to obtain information about the main compounds. The total extract was subsequently fractionated by flash chromatography in 19 fractions of increasing polarity which were tested against *T*. *brucei in vitro*. A few fractions showed antitrypanosomal activity. Active fractions were analysed by HPLC MS-MS and by NMR to find the phytoconstituents responsible for activity towards the protozoan parasite. The two most abundant sesquiterpene lactones in the active fractions were finally purified and assessed for their antitrypanosomial activity.

Material and Methods

Chemicals and reagents

LC–MS grade acetonitrile (ACN) and methanol (MeOH) were obtained from J. T. Baker (Phillipsburg, USA). HPLC-grade formic acid was purchased from Dikma

Tech. Inc. (Beijing, China). Cycloexane was obtained from Scharlau. Deuterated chloroform was purchased from Sigma-Aldrich. Water (H₂O) was purified by a Milli-Q system (Millipore, Billerica, MA, USA) in our laboratory. Other reagents were of analytical grade.

Plant material

Leaves of *T. diversifolia* (Asteraceae) were collected in the city of Dschang, western region of Cameroon (N 05°26'18", E 10°04'07", 1450 m a.s.l) by Saague T. Maximiliene and Prosper C. Biapa Nya in January 2016 during the dry season. Botanical authentication was performed by plant taxonomist Mr. Nana of the National Herbarium, Yaoundé, Cameroon, where a voucher specimen coded 10196/HNC was deposited.

Preparation of plant extracts

T. diversifolia leaves were air-dried in the shade at room temperature ($\approx 25^{\circ}$ C) for 3 days and conserved in wrapping papers before extraction. Fifty g of dry leaves were reduced into powder using a blender MFC DCFH 48 IKA-WERK (D-Staufen) equipped with sieves of 2-mm size in diameter. The powder was subsequently macerated in 500 ml of methanol (MeOH) for 24 h and filtered. The filtrate was concentrated under reduced pressure at 30°C with a rotary evaporator and freeze-dried to obtain a crude MeOH extract (2.64 g, 5.3% yield, Tith-MeOH). Using the same protocol, an aqueous extract was obtained by maceration of 5 g of leaves in 50 mL of

deionized water to get 0.42 g (8.4%, Tith- H_2O) of extract. The extracts were kept in glass vials protected from light at -20°C before chemical analysis and anti-trypanosomal experiments.

Apparatus and chromatographic conditions

Detection and quantification of the compounds were performed on an LC-MS system. The LC-MS equipment (Varian Inc.) contained a chromatographic system (Varian LC-212) coupled with a Varian 500-MS (ion trap) mass spectrometer fitted with an ESI source (Varian). MS conditions were the following: needle potential -5.0kV, shield 600 V, spray chamber temperature 50°C, drying gas pressure 10 psi, drying gas temperature 350°C, capillary voltage 80 V, RF loading 100, and MS range 150-2000 Da. MSⁿ spectra were obtained during the chromatography run by using the turbodds (tdds) utility of the instrument. HPLC-DAD analysis was carried out by an Agilent 1100 series liquid chromatograph equipped with an Agilent 1100 Diode Array (DAD). An Eclipse XDB-C8, 5 µm, 4.6 x 150 mm column (Agilent Technologies) was used. The mobile phase consisted of (A) aqueous formic acid (0.1 %) and (B) acetonitrile. Gradient conditions were: 0-30 min, linear gradient from 10 % to 100 % of B; 30-35 min, isocratic conditions at 100 % of B; 35-36 min, linear gradient from 100% to 10% of B; 36-40 min, isocratic conditions at 10% of B. Flow rate: 1 mL/min. Calibration curves were obtained by standard solutions of rutin for flavonoid derivatives (UV detection at 350 nm), chlorogenic acid for caffeoylquinic derivatives (UV detection at 330 nm) and gallic acid for small phenols (UV detection at 280 nm). The concentration ranges were 11.7-117, 13.2-132, and 14.6-146 µg/mL for chlorogenic acid, rutin and

gallic acid, respectively. The limits of detection (LOD) and quantification (LOQ) were 1.5 and $4.0 \mu \text{g/ml}$ and 0.5 and $1.5 \mu \text{g/ml}$ for chlorogenic acid and rutin, respectively.

Fractionation of the extract, isolation of Tagitinin A and C

The crude methanolic extract of *T. diversifolia* leaves (1 g) was dissolved in methanol and adsorbed on silica gel (40 mesh, 2 g). Subsequently, the solvent was removed under vacuum to obtain a dried powder. This powder was packed in a precolumn, which was fixed on a Silica column (Buchi Sepacore[®] silica 12 g). Separation was performed on a Varian Intelliflash Flash chromatograph. The eluent was initially cyclohexane followed by an increasing percentage of methanol up to 30 % in 120 min. The flow rate was 1 mL/min and the chromatogram was monitored at 220 and 340 nm. Eluted fractions were checked by TLC and collected in 19 groups. The solvent was removed under vacuum and the weight of each fraction obtained was as follows: 1 (25.8 mg), 2 (15.0 mg), 3 (17.1 mg), 4 (18.5 mg), 5 (23.1 mg), 6 (26.7 mg), 7 (13.9 mg), 8 (16.7 mg), 9 (19.4 mg), 10 (18.0 mg), 11 (8.9 mg), 12 (5.3 mg), 13 (7.0 mg), 14 (4 mg), 15 (104.6 mg), 16 (128.4 mg) 17 (78.7 mg), 18 (59.7 mg), 19 (38.2 mg).

For tagitinin isolation 250 g of dried plant material was extracted using dichloromethane (550 mL x 3 times) using ultrasound bath. Liquids were collected, filtered and pooled and solvent was evaporated under vacuum. The crude extract (9.5 g) was used for chromatographic separation. A silica gel column (10 x 100 cm) was packed using hexane as solvent. Extract was charged in the column and hexane was used for elution, then hexane/ethyl acetate with an increasing ratio, starting from 6/1 to 1/1, was used as eluent system. Fifty-five fractions were pooled on the basis of their

chromatographic behaviour. Further separations were achieved on Sephadex LH20 column (5 x 40 cm) eluting with dichloromethane. Tagitinin A and tagitinin C were isolated as pale oil (600 and 250 mg, respectively). Their spectral data were compared with those of literature (Baruah et al., 1979).

NMR analysis was obtained on a Bruker AVANCE III spectrometer operating at 400.13 MHz for ¹H-NMR. 2D spectra, HSQC-DEPT, HMBC, COSY and TOCSY were used for identification of compounds occurring in mixtures. Samples were dissolved in deuterated chloroform and used for analysis. For quantitative purposes, previously published conditions were used (Comai et al., 2010).

T. brucei and mammalian cell culture

The cell culture conditions and the growth inhibition assay on T. brucei and Balb/3T3 cells were performed as described before (Petrelli et al., 2016). *T. brucei* bloodstream forms (TC221) were cultured at 37 °C with 5 % CO₂ in HMI-9 medium supplemented with 10 % (v/v) heat-inactivated fetal bovine serum (Gibco). Mouse BALB/3T3 fibroblast (ATCC no CCL-163) were cultivated at 37 °C and 5 % CO₂ in Dulbecco's modified Eagle's medium (Sigma), supplemented with 10 % (v/v) heatinactivated fetal bovine serum, glutamine (0.584 g/L) and 10 mL/L of 100 × penicillinstreptomycin (Gibco BRL) (Hirumi and Hirumi, 1989).

Growth inhibition assay on T. brucei and BALB/3T3 cells

The methanolic extract and its fractions were dissolved in dimethyl sulfoxide (DMSO). They were serially diluted with growth medium to concentrations ranging

from 2×10^{-5} to 200 µg/mL in 96-wells microtiter plates (100 µL/well). Subsequently, 100 µL of *T. brucei* or mammalian cell culture was added to each well (20,000 cells/well). After 48 h incubation, the plates were treated for 24 h with 20 µL of 0.5 mM resazurine (Sigma-Aldrich). They were subsequently quantified by fluorescence (540 nm excitation and 590 nm emission) using an Infinite M200 microplate reader (Tecan Group, Ltd.). The IC₅₀ values were calculated by fitting the data to a log inhibitor vs. response curve (variable slope, four parameters) using the GraphPad Prism 5.04 software.

Results and Discussion

Characterization of T. diversifolia extracts by NMR and HPLC-MSⁿ

In order to establish the phytochemical composition of the Tith-MeOH extract, we decided to use a multi-technique approach. As a first step the Tith-MeOH extract was dissolved in deuterated chloroform (final concentration 20 mg/mL) and different spectra were acquired namely ¹H-NMR, HSQC-DEPT, HMBC and COSY experiments. Diagnostic signals of sesquiterpene sp² protons were detected, and assigned on the basis of the ¹H, HSQC, HMBC and COSY data as compared with the literature (Baruah et al., 1979). These results are reported in Table 1. A representative portion of a HSQC-DEPT spectrum is shown in Figure 1 indicating the assigned positions.

Many other signals were detected and may be linked to other phytoconstituents. For this reason, HPLC-MSⁿ analysis was performed in order to identify other secondary metabolites occurring in the crude extract. The fragmentation pathways for the main

sesquiterpenes of *T. diversifolia* were also studied and are reported in Figure 2. Six different sesquiterpenes were detected and their structures were determined on the basis of HPLC-MSⁿ analysis. Limited information is available related to ESI/MS behaviour of these sesquiterpenes. Abea et al. (2015) reported proposed fragmentation pathways of tagitinin F. In our analysis, we observed that the tirotundin derivatives (compounds **4** and **5**, Figure 3) are characterized in MS² by three main fragments ascribable to the loss of an ester side chain, a water molecule and an ether bridge or lactone ring with rearrangement at m/z 265, 247 and 229, respectively (Figure 2). The same fragments are detectable in the MS² spectra of compound **6** at m/z 279, 261 and 243 (Figure 2) presenting additional 14 Da due to the presence of a methyl group. Polyphenol derivatives, mainly caffeic acid derivatives and flavonoids, were also identified. Table 2 reports the different constituents identified. Phytochemical analysis showed that the *T. diversifolia* extract is complex and contains sesquiterpenes, phenylpropanoids and flavonoid derivatives as main constituents.

In vitro evaluation of antiprotozoal activity of bioactive fractions, tagitinin A and tagitinin C

Two extracts (Tith-MeOH and Tith-H₂O) were prepared from the aerial parts of *T. diversifolia* to assess their *in vitro* antitrypanosomal activity (Table 3). The methanolic extract was two times more active against *T. brucei* than the aqueous extract ($IC_{50} = 1.08$ and 2.18 µg/mL, respectively) and was therefore chosen for further fractionation. The methanolic extract was subjected to chromatographic separations on silica gel and the fractions with increasing polarity collected after each purification step (F1-F19) were tested against the TC221 cells with the purpose of following the

antitrypanosomal activity. The bioassay-guided purification of the Tith-MeOH extract gave two fractions (F8 and F9, Table 3) showing significantly higher antitrypanosomal effects in our cell-based assay ($IC_{50} = 0.41$ and $0.43 \mu g/mL$, respectively). Bioassays performed on the fractions from Silica-based chromatography of Tith-MeOH extracts showed the effect on *T. brucei vs.* mammalian cell proliferation (Table 3).

¹H-NMR of the 16 assayed fractions (Figure 4) clearly shows the presence of signals ascribable to sesquiterpene derivatives as well as other aromatic compounds in the most active mixtures, namely F7 to F11. In all the fractions, the signals ascribable to phenolic and phenylpropanoid derivatives were not detectable or poorly detectable. We decided to consider the most active fractions, thus the ones showing $IC_{50} < 1.6 \mu g/mL$, i.e. fractions 8-11, which were analysed for the contents of tagitinins A and C. Due to the non-commercially availability of tagitinins as reference compounds, the quantification of the main derivatives, namely tagitinins A and C, was performed using an ¹H-NMR approach using caffeine as internal standard (Comai et al., 2010) and the amounts of these STLs in the active fractions were reported in Table 4. Results showed that bioactivity of the obtained fractions can be in part ascribed to STLs constituents. Isolated tagitinin A and C were also tested. As reported in Table 5, tagitinin C resulted by far the most active compound compared with active fractions. Notably, tagitinin C was 4.5 times more active than the reference compound suramin. Furthermore, selective index for tagitinine C was 3 showing preferential effect on target cells compared with tagitinin A (1.3). The large difference in bioactivity observed for the tagitinin C and A indicates the need for the evaluation of other tagitinine derivatives in order to explore possible structure activity relationships.

Potent and selective antitrypanosomal activity of different plant sequiterpenes such as α -eudesmol, hinesol, nardosinone and 4-peroxy-1,2,4,5-tetrahydro- α -santonin (Otoguro et al. 2011) has been previously reported, as well as the possible usefulness of sesquiterpenes as lead compounds for the development of new antitrypanosomal drugs (Saeidnia et al. 2013).

As regards the mode of action of T. diversifolia STLs on trypanosome cell, some considerations can be made. As reported by Schimdt et al. (1999), STLs possessing an α,β -unsaturated moiety in the form of α -methylene- γ -lactone can act as a Michael acceptor, reacting with nucleophiles such as free thiol groups (R-SH) in proteins leading to macromolecular dysfunction, oxidative stress and genetic mutations as a result of the oxidative stress. The presence of one potentially reactive α_{β} -unsaturated carbonyl group as a pharmacophore is often associated with significant antiprotozoal and cytotoxic activities. Tagitinin A has α,β -unsaturated carbonyl group in the γ -lactone ring, whereas tagitinin C has a carbonyl group conjugated with two double bonds in the germacrene ring and an α,β -unsaturated carbonyl group in the γ -lactone ring. The measured activity of the two compounds showed the importance of the carbonyl group conjugated with two double bonds because of the greater antitripanosomal activity of tagitinin C. The mechanism of these compounds against trypanosomes may be related to the capability to form thiol adducts with components found in the cellular environment such as trypanothione, glutathione and SH-groups in proteins. The lack of reduced trypanothione and other SH-group containing substances may in turn lead to that the parasite cells become more vulnerable to oxidative stress (Wink, 2008). The same

mechanism of action may occur in the mammalian fibroblasts (BALB/ 3T3), where glutathione depletion renders the cells prone to oxidative stress.

Conclusions

Overall our results, although preliminary, confirmed the usefulness of *T*. *diversifolia* as an effective herbal remedy in the African traditional medicine and highlighted the opportunity to use the tagitinins as inspiring leads for designing future trypanocidal drugs. Further studies are needed in particular to increase the knowledge about the mode of action and the possible molecular targets related to the antitrypanosomal activity of tagitinins. The opportunity to study different derivatives as well as strictly related sesquiterpene can be a good opportunity to explore structure activity relationships. Also synthetic or semisyntetic derivatives can be produced in order to follow a rational approach for the development of more active compounds. Finally further investigation will be needed in order to assess the real efficacy *vs*. toxicity through *in vivo* assays.

Conflict of interest

The authors declare no competing interests.

Funding

The authors would like to thank the University of Camerino (FAR 2014/15, Fondo di Ateneo per la Ricerca, FPI000044) for financial support. Anders Hofer also thanks the Swedish Research Council (2012-1932) and the Kempe Foundation. Giovanni Benelli is sponsored by PROAPI (PRAF 2015) and University of Pisa, Department of Agriculture,

Food and Environment (Grant ID: COFIN2015_22). Funders had no role in the study design, data collection and analyses, decision to publish, or preparation of the manuscript.

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 Table 1. Diagnostic assignments of tagitinins A and C from NMR spectra of Tith

 MeOH extract. Assignments were obtained comparing HSQC-DEPT, HMBC and

 COSY data with the literature (Baruah et al., 1979).

Tagitinin C				Tagitinin A	
Position	δΗ	δC	Position	δН	δC
1	6.93 d	159.6	1	4.23	78.4
	<i>J</i> =13.5				
2	6.21	129.5	6	4.55	81.8
13	6.36-5.81	124.0	8	5.59	69.9
5	5.88	137.6	13	6.25-5.53	121.7

6.3b-. 5.88 137.0

Ret.	Comment	Delesite	[M-H] ⁻ or	European ander	
time	Compound	Polarity	$[M+H]^+$	Fragments	
2.53	Quinic acid ^a	Negative	191	85	
12.0	Chlorogenic acid ^a	Negative	353	191-179	
13.0	Dutin ^a	Negative	600	301-271-255-	
13.7	i Xuuni	regative		179-151	
14.5	Quercetin-glucuronide	Negative	477	301-179-151	
14.6	1.5-Dicaffeovl quinic acid ^a	Negative	515	353-191-173-	
14.0	1,5 Deaneoyr quine acki	reguive	515	127	
14.8	3.4-Dicaffeovl quinic acid	Negative	515	353-191-179-	
1110	e, i Dicance yr quane ach	Trogentro	515	173	
15.7	3.5-Dicaffeovl quinic acid ^a	Negative	515	353-191-173-	
1017	e,e Dicanceyr quane uch	rieguire	010	127-111-85	
18.8	Isorhamnetin	Negative	315	300-272-228	
19.5	Tagitinin A	Negative	367	279-261-235	
		Positive	369	281-263-245	
19.7	Tirotundin 3- <i>O</i> -methylether	Positive	367	279-261-243-	
			501	233-215	
	~	Negative	365	277-233-215-	
		8		191-176	
20.8) 8 Hispidulin Nagativa 200		299	284-256-227-	
20.0		1,054410		212	
24.6	Tagitinin C	Positive	349	261-243-215-	

Table 2. HPLC-MS constituents identified in T. diversifolia methanolic extra	acts.
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				173
27 /	$4 \qquad 1 \theta Mathematikarsifalin \qquad Desitive$	305	293-275-261-	
27.4		1 OSLIVE	373	199-181
	8- β -O-(2-methylbutyroyl)-			
26.8	Tirotundin or $8-\beta$ -O-	Positive	367	265-247-229
	(isovaleroyl)-Tirotundin		Ó	
acc	onfirmed by the injection of a refer	ence compo	und	
			6	
			5	
		2		
	R			
	\mathbf{G}			
	A A			

Table 3. Activity of *T. diversifolia* leaves extracts and fractions 1-19 against *T. brucei* TC221 andBALB/3T3.

Samples	IC ₅₀ (µg/mL)	Selectivity	
Sumples	T. brucei (TC221)	BALB/3T3	Index (SI)
Crude methanol extract (Tith-MeOH)	1.05 ± 0.18	5.15 ± 0.82	4.9
Crude aqueous extract (Tith-H ₂ O)	2.18 ± 0.06	6.71 ± 0.48	3
Fractions from methanol extract (Tith-Me	OH)		
Fraction 1	62.05 ± 20.4	- 0	-
Fraction 2	13.69 ± 1.8	52.16 ± 3.1	3.8
Fraction 3	8.29 ± 0.79	17.52 ± 2.9	2.1
Fraction 4	8.28 ± 0.56	12.85 ± 1.1	1.5
Fraction 5	4.37 ± 1.98	7.65 ± 0.85	1.7
Fraction 6	1.66 ± 0.66	4.37 ± 0.61	2.6
Fraction 7	1.67 ± 0.51	2.78 ± 0.03	1.6
Fraction 8	0.41 ± 0.22	1.44 ± 0.04	3.3
Fraction 9	0.43 ± 0.09	1.47 ± 0.04	3.4
Fraction 10	1.09 ± 0.37	2.36 ± 0.17	2.1
Fraction 11	1.20 ± 0.21	2.66 ± 0.18	2.2
Fraction 12	2.34 ± 0.55	4.24 ± 0.49	1.8
Fraction 13	3.16 ± 0.89	6.03 ± 1.86	1.9
Fraction 14	2.02 ± 0.88	5.10 ± 0.64	2.5
Fraction 15	1.63 ± 0.61	3.17 ± 0.14	1.9
Fraction 16	7.65 ± 1.61	19.34 ± 2.76	2.5
Fraction 17	>100	-	-
Fraction 18	>100	-	-
Fraction 19	>100	-	-
Pure compounds	μg/mL (μM)	µg/mL (µM)	

Tagitinin A	$0.356 \pm 0.17 \ (0.97)$	0.467 ± 0.21 (1.27)	1.3
Tagitinin C	$\begin{array}{c} 0.0042 \pm 0.0005 \\ (0.0121) \end{array}$	$\begin{array}{c} 0.012 \pm 0.001 \\ (0.036) \end{array}$	3
Reference drug	$\mu g/mL \ (\mu M)$	$\mu g/mL~(\mu M)$	
Suramin	$\begin{array}{c} 0.0191 \pm 0.002 \\ (0.0147) \end{array}$	n.d.*	n.d.*

*n.d. not determined

Table 4. Amount of tagitinins in the bioactivity-guided selected fractions. The amount is shown as the percentage of the total volume, IC_{50} are expressed in µg/mL.

	Tagitinin	Tagitinin	T. brucei (TC221)
Samples	A %	С %	IC ₅₀ (µg/mL)
Crude methanol extract (Tith-	0.83 ± 0.01	1.10 ± 0.01	1.05 ± 0.18
MeOH)		X	
Active fractions from methanol e.	xtract (Tith-MeOH	[)	
Fraction 8	10.8 ± 0.01	9.39 ± 0.01	0.41 ± 0.22
Fraction 9	8.36 ± 0.01	8.18 ± 0.01	0.43 ± 0.09
Fraction 10	7.03 ± 0.01	7.09 ± 0.01	1.09 ± 0.37
Fraction 11	7.07 ± 0.01	$6.51{\pm}~0.01$	1.20 ± 0.21
	~		



Figure 1. HSQC-DEPT (MeOD) of a T. diversifolia extract. TA and TC indicate signals

of Tagitinin A and Tagitinin C, respectively.

A CERMIN





Figure 2. Mass spectra of the main sesquiterpenoids of *Tithonia diversifolia* and

proposed fragmentation pathways.

A CLANK



Figure 3. Structures of main sesquiterpenes identified in T. diversifolia extracts.



Figure 4. ¹H-NMR of *T. diversifolia* fractions 1-16, diagnostic signals indicating the presence of sesquiterpene and phenolics are highlighted with coloured rectangles.

Grafical abstract

Identification of Tagitinin C from *Tithonia diversifolia* as antitrypanosomal compound using bioactivity-guided fractionation

