Supplementary Information (SI) for New Journal of Chemistry. This journal is © The Royal Society of Chemistry and the Centre National de la Recherche Scientifique 2024

Supplementary Material

Development of α -acyloxycarboxamides targeting *Leishmania amazonensis* parasite

Saraliny B. França ^a, Jamilly E. da Silva ^a, Leandro R. Silva ^a, Emanuelly K. A. Padilha ^a, Fernando Almeida-Souza ^{b,c}, Lucas S. Barbosa ^b, Katia S. Calabrese ^b, Dimas J. P. Lima ^{a*} and Edeildo F. da Silva-Júnior ^{a*}

^a Institute of Chemistry and Biotechnology, Federal University of Alagoas, Maceió, Alagoas, 57072-970, Brazil

^b Laboratory of Protozoology – Oswaldo Cruz Institute, Oswaldo Cruz Foundation, Rio de Janeiro, 21041-250, Brazil

^c Post-graduate in Animal Science, State University of Maranhão, São Luís, 65055-310, Brazil

* Corresponding authors: edeildo.junior@iqb.ufal.br (E.F, Silva-Júnior), dimas.lima@iqb.ufal.br (D.J.P. Lima)

Spectroscopic analysis of cinnamic acid derivatives

Figure 1. Infrared (ATR - FTIR) spectrum of ethyl cinnamate (3a).





Figure 2. ¹H NMR spectrum of ethyl cinnamate (400 MHz, CDCl₃) (3a)







Figure 6. Infrared (ATR - FTIR) spectrum of ethyl *p*-chlorocinnamate (3d).



Figure 7. ¹H NMR spectrum of ethyl *p*-chlorocinnamate (400 MHz, CDCl₃) (3d).





Figure 9. ¹H NMR spectrum of ethyl *p*-trifluoromethoxycinnamate (400 MHz, CDCl₃) (3e).





Figure 11. Infrared (ATR - FTIR) spectrum of Cinnamic acid (4a).









Figure 15. ¹³C NMR spectrum of *o*-phenylcinnamate acid (4b) (150 MHz; CDCl₃/CD₃OD).

170	160	150	140	130	120	110	100	90	80	70	60	50	40	30	20	10	0	ppm
												Ţ						
											(\sim		Н			
		×143.94	/ 142.11 / 139.16	127.63	126.84	~117.05												





Figure 17. ¹H NMR spectrum of *p*-chlorocinnamic acid (4d) (600 MHz; DMSO-*d6*).







Figure 19. ¹H NMR spectrum of *p*-trifluoromethoxycinnamic acid (4e) (600 MHz; DMSO-



Figure 20. ¹³C NMR spectrum of *p*-trifluoromethoxycinnamic acid (**4e**) (150 MHz; CDCl₃/ DMSO-*d6*).



Figure 21. Infrared (ATR - FTIR) spectrum of 2 - (tert-Butylamino) - 2 - oxo - 1 - phenylethyl cinnamate (7a₁).





Figure 23. ¹³C NMR spectrum of 2 - (tert-Butylamino) - 2 - oxo - 1 - phenylethyl cinnamate (7a₁) (150 MHz; CDCl₃).

5 1	00140100			
40	00022002000	\sim	00	
• •			с С	
5 7	0040000000000000000000000000000000000	•	•	•
99	4 0 0 0 0 0 0 0 0 0 0	L)	\leftarrow	00
- $-$		2	ک	\sim
		1	1	1
\ /				
	· · · · · · · · · · · · · · · · · · ·	1	1	



Figure 24. ESI-TOF (m/z) - $[M+Na]^+$ spectrum of 2 - (tert-Butylamino) - 2 - oxo - 1 - phenylethyl cinnamate (7a₁).



Figure 25. Infrared (ATR - FTIR) spectrum of 2-(*tert*-butylamino)-1-(4-chlorophenyl)-2-oxoethyl cinnamate (**7a2**).





Figure 27. ¹³C NMR spectrum of 2-(*tert*-butylamino)-1-(4-chlorophenyl)-2-oxoethyl cinnamate (7a2) (150 MHz; CDCl₃).



Figure 28. ESI-TOF (m/z) - $[M+Na]^+$ spectrum of 2 - (tert-Butylamino) - 2 - oxo - 1 - phenylethyl cinnamate (7a₂).



Figure 29. Infrared (ATR - FTIR) spectrum of (*E*)-2-(*tert*-Butylamino)-1-(3,4-dichlorophenyl)-2-oxoethyl 3-([1,1'-biphenyl]-2-yl)acrylate (**7b**₁).



Figure 30. ¹H NMR spectrum of *E*)-2-(*tert*-Butylamino)-1-(3,4-dichlorophenyl)-2-oxoethyl 3-([1,1]-biphenyl]-2-yl)acrylate (**7b**₁) (600 MHz; CDCl₃).



Figure 31. ¹³C NMR spectrum of *E*)-2-(*tert*-Butylamino)-1-(3,4-dichlorophenyl)-2-oxoethyl 3-([1,1]-biphenyl]-2-yl)acrylate (7b₁) (150 MHz; CDCl₃).



Figure 32. ESI-TOF (m/z) - $[M+Na]^+$ spectrum of 2 - (tert-Butylamino) - 2 - oxo - 1 - phenylethyl cinnamate (7b₁).



Figure 33. Infrared (ATR - FTIR) spectrum of (*E*)-2-(*tert*-Butylamino)-1-(3,4-dimethoxyphenyl)-2-oxoethyl 3-([1,1'-biphenyl]-2- yl)acrylate (**7b**₂).



Figure 34. ¹H NMR spectrum of (*E*)-2-(*tert*-Butylamino)-1-(3,4- dimethoxyphenyl)-2- oxoethyl 3-([1,1'-biphenyl]-2- yl)acrylate (**7b**₂) (600 MHz; CDCl₃).



Figure 35. ¹³C NMR spectrum of (E)-2-(*tert*-Butylamino)-1-(3,4- dimethoxyphenyl)-2- oxoethyl 3-([1,1'-biphenyl]-2- yl)acrylate (7b₂) (150 MHz; CDCl₃).



Figure 36. Infrared (ATR - FTIR) spectrum of (*E*)-2-(*tert*-Butylamino)-1-(3,4-dichlorophenyl)-2-oxoethyl 3-(naphthalen-2-yl)acrylate (**7cb1**).



Figure 37. ¹H NMR spectrum of (*E*)-2-(*tert*-Butylamino)-1-(3,4-dichlorophenyl)-2-oxoethyl 3-(naphthalen-2-yl)acrylate (**7cb1**) (600 MHz; CDCl₃).



Figure 38. ¹³C NMR spectrum of (*E*)-2-(*tert*-Butylamino)-1-(3,4-dichlorophenyl)-2-oxoethyl 3-(naphthalen-2-yl)acrylate (**7cb1**) (150 MHz; CDCl₃).



Figure 39. ESI-TOF (m/z) - $[M+Na]^+$ spectrum of (E)-2-(tert-Butylamino)-1-(3,4- dichlorophenyl)-2-oxoethyl 3-(naphthalen-2-yl)acrylate (7cb1).



Figure 40. Infrared (ATR - FTIR) spectrum of (*E*)-2-(*tert*-Butylamino)-1- (3,4dichlorophenyl)-2 oxoethyl 3-(4-chlorophenyl)acrylate (**7db**₁).



Figure 41. ¹H NMR spectrum of *(E)-2-(tert*-Butylamino)-1-(3,4dichlorophenyl)-2 oxoethyl 3-(4-chlorophenyl)acrylate (7**db**₁). (600 MHz; CDCl₃).



Figure 42. ¹³C NMR spectrum of (*E*)-2-(*tert*-Butylamino)-1-(3,4dichlorophenyl)-2 oxoethyl 3-(4-chlorophenyl)acrylate ($7db_1$) (150 MHz; CDCl₃).



Figure 43. ESI-TOF (m/z) - $[M+Na]^+$ spectrum of (E)-2-(*tert*-Butylamino)-1- (3,4dichlorophenyl)-2 oxoethyl 3-(4-chlorophenyl)acrylate (7db₁).



Figure 44. Infrared (ATR - FTIR) spectrum of (*E*)-2-(*tert*-Butylamino)-1-(4-chlorophenyl)-2 oxoethyl 3-(4-(trifluoromethoxy) phenyl)acrylate (**7ea2**).



Figure 45. ¹H NMR spectrum of (*E*)-2-(*tert*-Butylamino)-1-(4-chlorophenyl)-2 oxoethyl 3-(4-(trifluoromethoxy) phenyl)acrylate (**7ea2**) (600 MHz; CDCl₃).



Figure 46. ¹³C NMR spectrum of (*E*)-2-(*tert*-Butylamino)-1-(4-chlorophenyl)-2 oxoethyl 3-(4-(trifluoromethoxy) phenyl)acrylate (**7ea2**) (150 MHz; CDCl₃).



Figure 47. ESI-TOF (m/z) - $[M+Na]^+$ spectrum of (E)-2-(*tert*-Butylamino)-1-(4-chlorophenyl)-2 oxoethyl 3-(4-(trifluoromethoxy) phenyl)acrylate (**7ea2**).



Key targets	Target name (gene)	UniProt code	Description	Free energy of binding (kcal/mol)
O93874_COCLU	17HSDcl	93874	17beta-hydroxysteroid dehydrogenase	-7.361
Q8ILP4_PLAF7		Q8ILP4	Pantothenate quinase, putative	-7.873
ACER2_HUMAN	ACER2	Q5QJU3	Alkaline ceramidase 2	-8.099
STXA_SHIDY	stxA	Q9FBI2	Subunit A of Shiga toxin	-8.597
PPAP_HUMAN	ACP3	P15309	Prostatic acid phosphatase	-7.480
AMD_HUMAN	PAM	P19021	Peptidyl-glycine alpha-amidant monooxygenase	-7.667
CA2D1_RAT	Cacna2d1	P54290	Alpha-2/delta-1 subunit of the voltage-dependent calcium channel	-7.200
AK1BA_HUMAN	AKR1B10	60218	Aldo-keto reductase family 1 member B10	-7.622
MALX3_YEAST	IMA1	P53051	Oligo-1,6-glucosidase IMA1	-8.044
KCNQ2_MOUSE	Kenq2	Q9Z351	Subfamily of voltage-gated potassium channels, KQT member 2	-7.923
CACB1_RAT	Cacnb1	P54283	Beta-1 subunit of the voltage-dependent L-type calcium channel	-7.412
VMAT2_HUMAN	SLC18A2	Q05940	Synaptic vesicular amine transporter	-8.525
RBBP9_HUMAN	RBBP9	75884	Serine hydrolase RBBP9	-8.246
A0A060INS8_CAPHI	CTSH	A0A060INS8	Cathepsin H	-8.179
CP24A_HUMAN	CYP24A1	Q07973	1,25-dihydroxyvitamin D(3) 24-hydroxylase, mitochondrial	-7.758
MMP1_HUMAN	MMP1	P03956	Interstitial collagenase	-7.697
MMP9_HUMAN	MMP9	P14780	Matrix metalloproteinase-9	-7.399
CP27B_MOUSE	Cyp27b1	35084	25-hydroxyvitamin D-1 alpha-hydroxylase, mitochondrial	-8.381
MMP2_HUMAN	MMP2	P08253	72 kDa type IV collagenase	-8.155
C3TDZ2_ECOLX	fabH	C3TDZ2	3-oxoacyl-[acyl-transporter protein] synthase 3	-8.287
AMPL_PIG	LAP3	P28839	Cytosol aminopeptidase	-8.608

Table 1. Results of the virtual screening of biological targets present in *Leishmania* of compound 7a1 based on reverse docking.

Key targets	Target name (gene)	Code in UniProt	Description	Free energy of binding (kcal/mol)
NR1H4_HUMAN	NR1H4	Q96RI1	Bile acid receptor	-7.217
Q94F81_MAIZE	hda106	Q94F81	Histone deacetylase type HD2 HDA106	-7.971
HDAC7_HUMAN	HDAC7	Q8WUI4	Histone deacetylase 7	-7.538
CAC1B_RAT	Cacna1b	Q02294	Alpha-1B subunit of the voltage-dependent N-type calcium channel	-7.312
H9B8T9_CAPHI	CTSB	H9B8T9	Cathepsin B	-7.014
Q72874_9HIV1	pol	-	Pol polyprotein	-8.120
Q7ZJM1_9HIV1	pol	-	Integrase	-7.560
LPXC_PSEAE	lpxC	P47205	UDP-3-O-acyl-N-acetylglucosamine deacetylase	-7.380
DHB3_RAT	Hsd17b3	54939	Testosterone 17-beta-desidrogenase 3	-7.298
TLR4_MOUSE	Tlr4	Q9QUK6	Receiver type Toll 4	-7.661
PSA_HUMAN	NPEPPS	P55786	Puromycin-sensitive aminopeptidase	-8.284
AK1C4_HUMAN	AKR1C4	P17516	Aldo-keto reductase family 1 member C4	-7.959
HDAC5_HUMAN	HDAC5	Q9UQL6	Histone deacetylase 5	-7.100
S12A2_HUMAN	SLC12A2	P55011	Solute carrier family 12 member 2	-7.700
TTHY_HUMAN	TTR	P02766	Transthyretin	-7.617
FOS_HUMAN	FOS	P01100	Proto-oncogene c-Fos	-7.153
KCNQ3_HUMAN	KCNQ3	43525	Subfamily of voltage-gated potassium channels, KQT member 3	-7.800
GABT_RAT	Abat	P50554	mitochondrial 4-aminobutyrate aminotransferase	-7.134
ALDR_HUMAN	AKR1B1	P15121	Aldo-keto reductase family 1 member B1	-7.059
AGP1_YEAST	AGP1	P25376	General amino acid permease AGP1	-7.937
HDAC1_MOUSE	Hdac1	9106	Histone deacetylase 1	-7.104
Р96830_МҮСТО		P96830	Protein containing TYR_PHOSPHATASE_2 domain	-8.293
CAH13_MOUSE	Ca13	Q9D6N1	Carbonic anhydrase 13	-8.732

 Table 1. Continued.

Key targets	Target name (gene)	Code in UniProt	Description	Free energy of binding (kcal/mol)
HDAC9_HUMAN	HDAC9	Q9UKV0	Histone deacetylase 9	-7.790
	PDB: 2JK6		Trypanothione reductase	-8.092
	-		Pantothenate quinase, putative	-7.200
	3UIB		Protein Kinase A	-7.865
	PDB: 4JZX		Ceramide sintase	-8.180
	PDB: 4C7H		N-myristoyltransferase	-8.980
	PDB: 50EY		Frutose-1,6-bisphosphatase	-7.510
	-		Secreted acid phosphatase	-8.208
	-		Serine hydrolase	-7.784
	Homology		Aspartyl aminopeptidase	-7.989
	Homology		Aldehyde dehydrogenase	-8.281
	Homology		Mitochondrial	-7.708
			Kinetoplastids (KMP-11)	-7.999
			Glycoprotein – gp46/M-2	-7.767
	Leishmanolysin (GP63)		Glycoproteins - gp63	-7.681