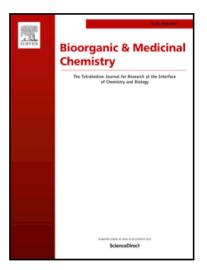
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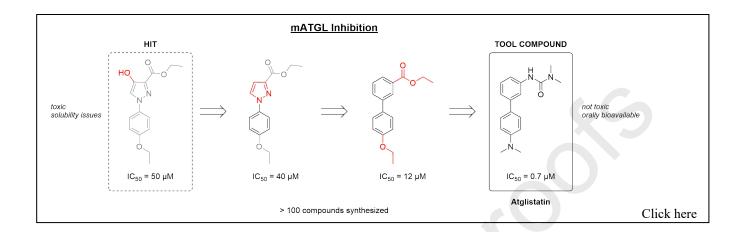


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Structure-activity relationship studies for the development of inhibitors of murine adipose triglyceride lipase (ATGL)

Nicole Mayer^{a,#}, Martina Schweiger^{b,c,#}, Elisabeth Fuchs^a, Anna K. Migglautsch^a, Carina Doler^a, Gernot F. Grabner^b, Matthias Romauch^b, Michaela-Christina Melcher^a, Rudolf Zechner^{b,c}, Robert Zimmermann^{b,c,*}, Rolf Breinbauer^{a,c,*}

^aInstitute of Organic Chemistry, Graz University of Technology, Stremayrgasse 9, A-8010 Graz, Austria ^bInstitute of Molecular Biosciences, University of Graz, Heinrichstraße 31, A-8010 Graz, Austria ^cBIOTECHMED Graz, A-8010, Austria

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ABSTRACT

High serum fatty acid (FA) levels are causally linked to the development of insulin resistance, which eventually progresses to type 2 diabetes and non-alcoholic fatty liver disease (NAFLD) generalized in the term metabolic syndrome. Adipose triglyceride lipase (ATGL) is the initial enzyme in the hydrolysis of intracellular triacylglycerol (TG) stores, liberating fatty acids that are released from adipocytes into the circulation. Hence, ATGL-specific inhibitors have the potential to lower circulating FA concentrations, and counteract the development of insulin resistance and NAFLD. In this article, we report about structure-activity relationship (SAR) studies of small molecule inhibitors of murine ATGL which has proven useful for the validation of ATGL as a potential drug target.

lipolysis NAFLD Atglistatin small molecule inhibitor

^{*} Corresponding authors.

E-Mail addresses: robert.zimmermann@uni-graz.at (R. Zimmermann), breinbauer@tugraz.at (R. Breinbauer).

[#] Co-first author, contributed equally.

Adipose tissue (AT) constitutes the body's largest energy store. The prevailing cell type of AT is the adipocyte which major function is to store and to liberate energy substrates in form of free fatty acids (FA) on demand. To fulfill their functions, adipocytes comprise a set of lipogenic and lipolytic enzymes and regulatory proteins, which are tightly regulated by feeding/fasting cycle of the organism. In the postprandial state a surplus of energy is stored as triacylglycerol within cytoplasmic lipid droplets. Upon fasting, adipose triglyceride lipase (ATGL) initiates the breakdown of stored TG to diacylglycerol (DG) and FA.¹ Subsequently the activities of hormone-sensitive lipase (HSL) and monoacylglycerol lipase (MGL) generate glycerol and two FA.² Adipocytes release the lipolytic products into the circulation to provide energy substrates for oxidative tissues. A constantly positive energy balance causes an increase in AT mass due to an increase in adipocyte number and adipocyte size, and obesity. However, the expandability of AT is not infinite. If the supply of energy substrates exceeds the storage capacity of AT, FA concentrations in the circulation increase and lipids are deposited in ectopic tissues, like muscle and liver. These ectopically stored lipids are responsible for the development of obesity associated insulin resistance and non-alcoholic fatty liver disease (NAFLD). finally leading to tissue dysfunction and early death.^{3,4} It is suggested that an increased release of FA from hypertrophic adipocytes contributes to the elevation of plasma and ectopic lipid concentrations.⁵ As ATGL is the initiating enzyme in the liberation of FA from TG stores of adipocytes, reducing FA in the circulation by inhibiting ATGL represents an attractive strategy to counteract the metabolic complications of obesity. Indeed, as mouse models of global as well as adipose tissue specific genetic ATGL deficiency showed, reduced plasma FA concentrations accompanied by a resistance to diet induced obesity and insulin resistance.6,7

These findings prompted us to develop and characterize small molecule inhibitors of ATGL. The versatile synthetic strategies we applied resulted in the development of Atglistatin. Atglistatin selectively, competitively, and transiently inhibits murine ATGL (mATGL) activity *in vitro*, in cultured cells, and *in vivo*.⁸ In contrast to complete ATGL deficiency in mice and humans, long term Atglistatin treatment does not cause cardiac steatosis, cardiac failure or a premature death. Importantly, Atglistatin treatment reduced circulating plasma lipids and protected from diet induced obesity, insulin resistance and NAFLD in mice.⁹ At present no 3D-structure of ATGL is available and any effort to identify inhibitors of ATGL has to rely on the traditional approach of synthesizing and testing compounds.¹⁰ In this manuscript we describe the structure-activity relationship (SAR) studies which allowed us in

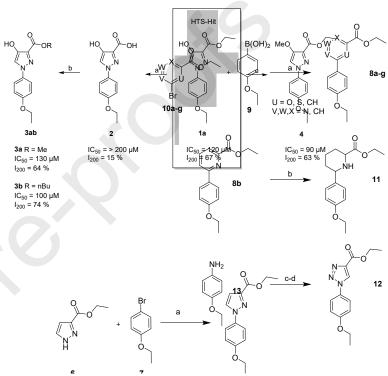
Scheme 1. HTS-hit and explorative modifications. Reagents and conditions: (a) 5 M NaOH, EtOH, 60 °C, 30 min, 41%; (b) R-OH, cat. conc. H₂SO₄, reflux; (c) 1.5 equiv K₂CO₃, DMF, overnight, 93%.

iterative rounds of optimization to develop biaryl hit compound 1 (IC₅₀ (half maximal inhibitory concentration) = 120 μ M, resulting from a high-throughput screening (HTS) campaign originally aimed to identify HSL inhibitors) to Atglistatin, which has proven to serve as a valuable tool compound to validate ATGL as a drug target. The presented SAR-data give valuable insight about the nature of the ATGL binding pocket and might provide opportunities for applying *in silico* methods to develop ATGL inhibitors.

2. Results and discussion

2.1. Preliminary experiments

H1S hit compound 1a to modify the substituents of the pyrazole moiety (Scheme 1). Saponification of the ethyl ester delivered carboxylic acid 2, which did not show any substantial mATGL inhibitory activity anymore. The importance of the ester moiety was further corroborated by the decent inhibitory activities of the methyl and butyl esters 3a-b. Alkylation of the parent HTS-hit compound 1a with methyliodide produced methoxypyrazole 4, which showed a significantly improved IC₅₀ value. In addition to IC₅₀ values, activity of new compounds was further determined as % of inhibition at a single concentration of 200 μ M (I₂₀₀). Variation of the substitution pattern (1b-p) of the bottom ring of the parent hit structure 1a led to activity loss (see SI). Further we tested if a substituent is necessary at all at the 4-position. The



Scheme 3. Introducing variations into the top ring fragments. Reagents and conditions: (a) 0.05 equiv PdCl₂(dppf)*DCM, 2.1 equiv CsF, DME, 80 °C; (b) 1.6 equiv PtO₂, 1 bar H₂, EtOH/DCM, RT, 27 h, 59%; (c) 1.5 equiv NaNO₂, H₂O, conc. HCl, 0 °C, 1 h, then 2.0 equiv NaN₃, H₂O, 0 °C to RT, 1 h, 84%; (d) 1.0 equiv ethyl propiolate, 0.2 equiv sodium ascorbate, 0.07 equiv CuSO₄*5H₂O, H₂O/ACN, RT, 15 h

compound **5** was prepared via CuI-catalyzed *N*-arylation of ethyl 3-pyrazolecarboxylate (**6**) and *p*-bromophenetole (**7**) (Scheme 2).¹¹ **5** exhibited the best inhibitory activity so far. Unfortunately, we also observed a considerable inhibitory activity of **5** against MGL (see SI, Figure S1).

2.2. Exchange of the top ring system

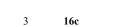
In order to improve lipase selectivity, the pyrazole ring was replaced by several different ring systems, while keeping the 1,3-arrangement of the 4-ethoxyphenyl and ethyl ester substituents constant (**8a-g**). 3-Bromoaryl esters **10a-g** were coupled with 4-ethoxyphenylboronic acid (**9**) in a Suzuki coupling reaction with PdCl₂(dppf)*DCM as catalyst and CsF as base in DME (Scheme 3).^{12,13} In order to test if a non-planar compound could also fit into the binding pocket, piperidine derivative **11** was produced from pyridine ester **8b** via catalytic hydrogenation.¹⁴ Triazole ester **12** was produced by converting aniline **13** via diazonium chemistry into the corresponding azide,¹⁵ which was reacted with ethyl propiolate in a Cu(I)-catalyzed azide-alkyne 1,3-dipolar

acetonitrile and water.10

In the biological screening data (Table 1) we noticed a strong influence of the top ring. Replacing the pyrazole ring with a phenyl ring (8e) resulted in a similar IC₅₀ value (50 μ M) and inhibition

Entry	No.	R	IC ₅₀ [µM]	I ₂₀₀ [%]
1	8a		105	72
2	8b		60	79
3	8c		80	87
4	8d	N C C C C C C C C C C C C C C C C C C C	130	70
5	8e		50	87
6	8f	s s	70	81
7	8g		50	93
10	8h	NH	> 200	44
8	11		> 200	48
9	12*	N N N N	> 200	43

effect (87%). Also, the pyridine ring with the nitrogen in 2position (8b) and the furan ring (8g) showed comparably good results. From these compounds we chose the presumably more drug-like structure 8e for further studies. 8e showed no cellular toxicity and very good selectivity in an ex vivo assay by taking tissue pieces of gonadal fat and incubating them with the inhibitor compound (see SI, Figure S2 and Figure S3). Gonadal fat is the tissue of choice to determine the release of lipolytic products as a measure of tissue lipolysis as it is a homogenous tissue consisting mostly of white adipocytes in contrast to other adipose tissue depots, like subcutaneous adipose tissue.17,18



Pd(OAc)₂, toluene, 100 °C

No.

Entry

			50 [1]	
1	16a	но−€	100	77
2	16b	`o-√}_ŧ	80	77
3	16c		200	52
4	16d	N-{-}	40	87
5	16e		130	68
6	16f	O ₂ N	95	63
7	16g	N	> 200	40
8	16h	s	75	67
9	16i		90	55
10	16 j		140	63
11	16k*	Br→	> 200	34
12	17a		> 200	33
13	17b		75	68
14	17c		150	67
15	17d		> 200	26
16	17d*		> 200	32
17	17e		100	68
18	17e*		200	54
19	18	HN	60	80
20	19	HN	120	66
21	20	N-	75	84
22	21		200	44

*reaction conditions: 2.0 equiv. K₃PO₄, 0.02 equiv. SPhos, 0.01 equiv.

IC50 [µM]

R

Scheme 2. Synthesis of unsubstituted derivative 5. Reagents and conditions: (a) 0.2 equiv CuI, 2 equiv Cs₂CO₃, MeCN/DMF, 120 °C, 65 h, 8%

2.3. Variations of bottom ring substituents



I200 [%]

investigate the SAR of this morety by varying the bottom ring fragment. Again Pd-catalyzed cross-coupling provided efficient synthetic access to many of the test compounds (Scheme 4). Some others had to be synthesized by subsequent modification via Buchwald-Hartwig-amination, reductive amination, or alkylation of amines (see SI).

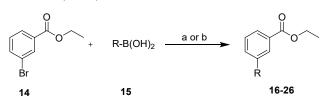


 Table 2. Screening results of 3-arylphenyl esters (obtained in assays with Cos-7 cells).



*values measured with E.coli lysate

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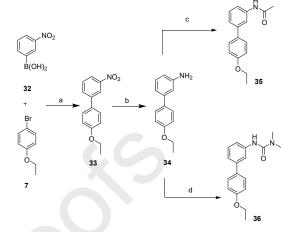
Table 2. Screening results of 3-arylphenyl esters (obtained in assays with Cos-7 cells), continued.



Entry	No.	R	IC50 [µM]	I ₂₀₀ [%]
23	22		140	59
24	23a		190	53
25	23b	N N S	120	67
		\$ ⁰		
26	23c	O'C'N N N O F	100	66
27	24	N-	200	20
28	24*	N	200	4
29	25	° → − Į	200	36
30	25*		200	17
31	26a		200	50
32	26a*		200	9
33	26b		90	69

Scheme 4. Synthesis of 3-arylphenyl esters. Reagents and conditions: (a) 0.05 equiv PdCl₂(dppf)*DCM, 2.1 equiv CsF, DME, 80 °C, (b) 2.0 equiv. K₃PO₄, 0.02 equiv. SPhos, 0.01 equiv. Pd(OAc)₂, toluene, 100 °C.

In order to increase reproducibility, to improve the yield of recombinant protein and to reduce background activity, we switched the ATGL-overexpression system from eukaryotic



Scheme 7. Synthesis of 4'-ethoxybiphenyl acetamido and urea derivatives. Reagents and conditions: (a) 2 equiv K_3PO_4 , 0.01 equiv Pd(OAc)₂, 0.02 equiv S-Phos, toluene, 100 °C, overnight, 68 %; (b) 10% Pd/C, 1 bar H₂, EtOAc, RT, overnight, 85 %; (c) 1.2 equiv. Ac₂O, DCM, RT, 2 h 74 %; (d) 1.2 equiv. Me₂NCOCl, 1 equiv. Et₃N, EtOAC, 0 °C to reflux, overnight, 27 %.

Cos-7 cells to bacteria (*E.coli*). In the following screenings (Tables 2-4) SAR studies marked with * have been performed using *E.coli* lysates containing recombinant mATGL. The screening results (Table 2) showed a preference for electron-rich biarylic systems with electron donating substituents, such as dimethylamine **16d**, and indoles (compounds **18** and **19**). By introducing substituents bridged with a CH₂-group to the biaryl (Table 2, **17a-e** and **23a-c**) or keto- (Table 2, **24-25**) and methine-moieties (Table 2, **26a-b**) the biological activity was decreased suggesting a negative influence of either increased flexibility or steric bulk. The best results were achieved for **16d** and **18** with IC₅₀ values from 40-60 μ M. Compound **16d** was further characterized for its effect on ATGL and HSL activity *in vitro*, on the release of FA from fat explants from WT and ATGL-ko mice, and cultured 3T3-L1 adipocytes (see SI, Figure S3, S4 and S5) in which **16d** showed

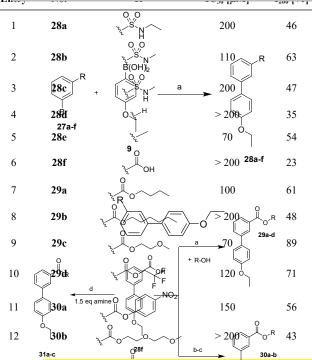
Scheme 6. Synthesis of 4'-ethoxybiphenyl esters and amides. Reagents and conditions: (a) 14 equiv ROH, 3 equiv H_2SO_4 , reflux, 3-5 h; (b) 15 equiv SOCl₂, 2 drops DMF, reflux, 23 h, quant.; (c) 1 equiv ROH, 2 equiv pyridine, DCM, reflux; (d) 1.5 equiv amine, 1.2 eq. EDC, THF, 0 °C to RT, 24 h.

improved selectivity compared to 8e.

In orienting experiments, the inhibitor **16d** was injected intraperitoneally into mice (with a concentration of 200 μ M) and plasma FA concentration was determined after 2, 4 and 8 h to measure FA release from adipose tissue. We found, that FA release was decreased by 40% 8 h after inhibitor injection (see SI; Figure S6). The inhibition of lipolysis by the use of **16d** could be increased to 40% *in vivo*, which was an encouraging result considering the maximum inhibition of lipolysis by ~60% in ATGL-ko-mice.¹⁹

2.4. Variations of top ring side chain

In a next step to further improve the activity and selectivity of the inhibitors, the 4'-ethoxybiphenyl moiety was decorated with sulfonamides or alkyl groups (Scheme 5), different ester and amide groups (Scheme 6), and acetamido and urea derivatives (Scheme 7).



Scheme 5. Synthesis of 4'-ethoxyphenyl derivatives with sulfonamide and alkyl side chains. Reagents and conditions: (a) 0.05 equiv PdCl₂(dppf)*DCM, 2.1 equiv CsF_DME_80 °C.

2.1 equiv	CsF, DME,	80 °C.	0		
14	310	N H	200 × 200	40	
15	31c	O V V N H	90	62	
16	33	NO2	> 200	35	
17	34	NH2	100	65	
18	35	HN N O	200	52	
19	36	N N N	70	68	
20	36*	[−] ² ² N N N	9	83	

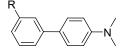
While the amides (Table 3, **31a-c**) and sulfonamides (Table 3, **28a-c**) resulted in inferior activity, various ester derivatives (e.g. **29c**) performed quite well. To our delight, the urea compound **36** (Table 3) showed quite promising results.

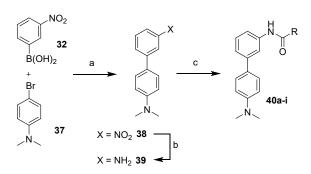
Table 3. Screening results 3-(4'-ethoxyphenyl)-substituted phenyl derivatives

2.5. Combination of best results

Hoping for an additive effect we decided to combine the best features of the optimization of the top ring, bottom ring substituent side chain and the top ring proposing 3-(4)dimethylamino)biphenyl-3-yl)-1,1-dimethylurea (Table 4, 40a) as a target structure (Scheme 8). The synthesis of this final compound starts with the known Suzuki coupling reaction between 3nitrophenylboronic acid (32) and 4-bromo-N,N-dimethylaniline (37) using PdCl₂(dppf)*DCM catalyst and CsF in DME. After reaction overnight biphenyl 38 could be isolated in 90% yield after purification via column chromatography. The nitro group was reduced by hydrogenation with 10% Pd/C and hydrogen stream²⁰ overnight furnishing 95% 4'-(1,1-dimethylamino)biphenyl-3amine (39). Amine 39 was directly converted to the urea without further purification. It was reacted with 1.5 eq dimethylcarbamoyl chloride and 1.0 equiv. triethylamine in DCM²¹ at 50 °C for 5 d. After reaching full conversion the final inhibitor compound could be isolated and purified via column chromatography yielding urea 40a in 54% yield. In addition to the previously tested dimethylurea, also other related side chains were incorporated (Table 4, 40b-i, 42, 43). However, 40a proved to be the most potent tested compound and was named Atglistatin®.

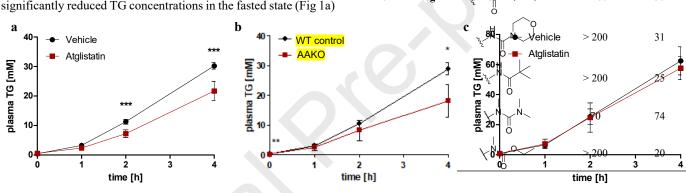
 Table 4. Screening results of 3-(4'-dimethylaminophenyl)-substituted phenyl derivatives





Scheme 8. Synthesis of Atglistatin and related derivatives. Reagents and conditions: (a) 0.05 equiv. PdCl₂(dppf)*DCM, 2.1 equiv. CsF, DME, 80 °C, overnight, 98%; (b) 10% Pd/C, 1 atm H₂, EtOH, EtOAc, RT, overnight, 95%; (c) 1.5 equiv. RCOCl, Et₃N, DCM or THF.

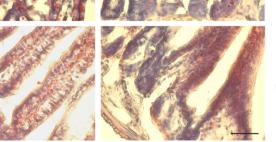
Journal Pr	e-proof	fs			
*values measured with <i>E.coli</i> lysate	Entry	No.	R	IC ₅₀ [µM]	I ₂₀₀ [%]
2.6. The Biological Data	1	38	NO2	> 200	21
The results of the <i>in vitro</i> screening of Atglistatin were comparable in both Cos-7 and <i>E.coli</i> lysates (Table 4). With an	2	39	NH₂	100	70
IC ₅₀ value below 5 μ M we could achieve up to 98% inhibition of the enzyme by applying an inhibitor concentration of 100 μ M	3	40a		< 5	98
instead of 200 μ M as for all compounds before. To further explore its <i>in vivo</i> activity, we administered Atglistatin at a final concentration of 200 μ M via oral gavage to <i>ad libitum</i> fed or 12 h	4	40a*	N N N	0.7	99
fasted mice and measured circulating TG, FA and glycerol 8 h later (Fig. 1d). In <i>ad libitum</i> fed mice, FA and glycerol concentrations	5	40b	N O O	5	95
were reduced by 27% and 23% (not significant) after Atglistatin gavage, respectively. In fasted mice Atglistatin gavage reduced FA and glycorel concentrations by 27%, and 26% recreatively, again	6	40c	N N N N N N N N N N N N N N N N N N N	3	93
and glycerol concentrations by 27%, and 26%, respectively - again validating Atglistatin as a tool compound to reduce plasma lipids. Interestingly, plasma TG content was 2-fold increased in the fed	7	40d	N O	82	69
and 37% decreased in the fasted animals in response to Atglistatin gavage. To investigate whether reduced TG plasma content is due	8	40e	zz-Nyo,↓	22	90
to reduced synthesis and/or secretion of lipoproteins from the liver or intestine we treated mice with Atglistatin and poloxamer to	9	40f	N O	73	75
inhibit lipoprotein lipase (LPL) activity, and determined plasma TG concentrations. Mice injected with Atglistatin showed significantly reduced TG concentrations in the fasted state (Fig 1a)	10	40g	N O	> 200	50



e

+ Atglistatin

Duodenum



control



Jejunum

lleum

40a

Same results were obtained in mice specifically lacking A1GL in adipose tissues (AAKO) (Fig. 1b), indicating that reduced VLDL synthesis is due to a reduced flux of FA from adipose tissue to the liver. No difference in plasma TG concentration was observed after an oral lipid load (200 μ L olive oil) indicating that chylomicron synthesis is not affected by Atglistatin (Fig 1c). However, but in line with a recent report on intestine specific ATGL/CGI-58 deficient animals, gavage of Atglistatin in olive oil led to an accumulation of intestinal lipids, primarily in the duodenum and jejunum indicating that ATGL is rather responsible for the degradation of lipid stores that are not directed to lipoprotein synthesis (Fig 1e).²²

The data from this report and our previous studies9 indicate that Atglistatin exerts its main metabolic activity by the inhibition of ATGL specifically in adipose tissue. With regard to non-adipocyte cells it has been shown that ATGL inhibition leads to the accumulation of arachidonic acid thereby preventing an inflammatory response in immune cells.²³ Moreover, pharmacological inhibition of ATGL decreased the regeneration of corneal nerves after injury, presumably by reducing the release of DHA.24 Interestingly, while it is reported that Atglistatin does not inhibit human ATGL,9 a recent study showed that the compound reduces TG hydrolase activity as well as TG rich lipoprotein secretion from human hepatoma cells.25 Further studies are needed, to evaluate the potential off target effects of Atglistatin in non-adipocyte cells. In contrast to ATGL deficiency, Atglistatin treatment protected from cardiac and hepatic steatosis on a high fat diet (HFD).9 In this regard recent studies demonstrated that Atglistatin rescues mice from heart failure by the inhibition of ATGL in adipocytes.²⁶ Moreover, inhibition of ATGL by Atglistatin reduced immune cell abundance in adipose tissue and completely blocked the IL-6 mediated induction of lipolysis. Hence, the unfavorable immunometabolic phenotypes associated with the complete absence of ATGL activity are not observed by Atglistatin treatment. This is most likely due to a transient and not a permanent inhibition of ATGL, and the preference of the drug to

fasted-gavage-8h fasted d fed-gavage-8h fasted control control 1.0 2.0 40a 40a 0.8 FA (mmol/L) (mmol/L) 0.6 1.0 04 FA 0.5 0.2 0.0 0.0 control control 0.3 0.6 40a 40a Jlycerol (mmol/L) jlycerol (mmol/L) 0.2 0.4 0.2 0.1 0.0 0.0 contro control 40a 40a 0.6 1.0 0.8 TG (mmol/L) TG (mmol/L) 0.4 0.6 0.4 0.2 0.2

taken into consideration for future drug development.²⁷

3. Conclusion

In this manuscript we have described how a classic medicinal chemistry approach led us to the development of the first inhibitor of mATGL, Atglistatin (**40a**), which – despite efforts to modify its structure^{28,29} – remains the reference tool compound for studying ATGL function in adipocytes, hepatocytes, corneocytes, cardiomyocytes, cancer cells, *in vitro* and *in vivo*. The data gained with Atglistatin in mice demonstrate that the pharmacological intervention of ATGL function offers an opportunity to develop drugs against several diseases, such as diabetes type 2, non-alcoholic fatty liver disease, heart failure and infectious disease. As Atglistatin does not inhibit human ATGL,³⁰ it will be necessary to develop a new class of compounds which target the human enzyme. Efforts in this direction are currently actively pursued in our labs and will be reported in due course.

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Declaration of Competing Interest

The authors declare no conflict of interest.

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Supplementary Material

Supplementary data to this article can be found online at xy

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Figure 1. (a) Plasma TG of fasted mice treated with Atglistatin. (b) Plasma TG of fasted WT control and adipose tissue specific ATGL-k.o. (AAKO) mice. (c) Plasma TG of fed mice treated with Atglistatin. (d) Plasma TG, FA and glycerol after oral gavage of Atglistatin in mice. n = 4-5. Data represent mean \pm s.d. Statistical significance was determined by unpaired Student's *t*-test (*P ≤ 0.05 , **P ≤ 0.01); ***P ≤ 0.001). (e) Neutral lipid accumulation in small intestine of mice. Mice received Atglistatin by oral gavage and were sacrificed after 4 h. Cryo-sections of duodenum, jejunum, and ileum were stained for neutral lipids with OilRedO. Nuclei were counterstained with hematoxylin. Bar = 50µm.

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Declaration of interests

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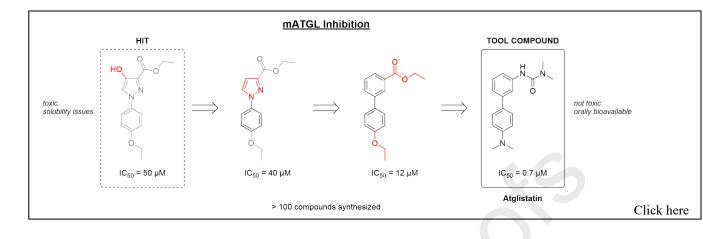
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relationships that could have appeared to influence the work reported in this paper.

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests:

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