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Total number of authors:
20

Published in:
Angewandte Chemie - International Edition

Link to article, DOI:
[10.1002/anie.201909518](https://doi.org/10.1002/anie.201909518)

Publication date:
2019

Document Version
Publisher's PDF, also known as Version of record

[Link back to DTU Orbit](#)

Citation (APA):
Ceballos, J., Schwalfenberg, M., Karageorgis, G., Reckzeh, E. S., Sievers, S., Ostermann, C., Pahl, A., Sellstedt, M., Nowacki, J., Carnero Corrales, M. A., Wilke, J., Laraia, L., Tschapalda, K., Metz, M., Sehr, D. A., Brand, S., Winklhofer, K., Janning, P., Ziegler, S., & Waldmann, H. (2019). Synthesis of Indomorphan Pseudo-Natural Product Inhibitors of Glucose Transporters GLUT-1 and -3. *Angewandte Chemie - International Edition*, 58(47), 17016-17025. <https://doi.org/10.1002/anie.201909518>

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Antitumor Agents

International Edition: DOI: 10.1002/anie.201909518
German Edition: DOI: 10.1002/ange.201909518

Synthesis of Indomorphan Pseudo-Natural Product Inhibitors of Glucose Transporters GLUT-1 and -3

Javier Ceballos⁺, Melanie Schwalfenberg⁺, George Karageorgis, Elena S. Reckzeh, Sonja Sievers, Claude Ostermann, Axel Pahl, Magnus Sellstedt, Jessica Nowacki, Marjorie A. Carnero Corrales, Julian Wilke, Luca Laraia, Kirsten Tschapalda, Malte Metz, Dominik A. Sehr, Silke Brand, Konstanze Winklhofer, Petra Janning, Slava Ziegler, and Herbert Waldmann*

Abstract: Bioactive compound design based on natural product (NP) structure may be limited because of partial coverage of NP-like chemical space and biological target space. These limitations can be overcome by combining NP-centered strategies with fragment-based compound design through combination of NP-derived fragments to afford structurally unprecedented “pseudo-natural products” (pseudo-NPs). The design, synthesis, and biological evaluation of a collection of indomorphan pseudo-NPs that combine biosynthetically unrelated indole- and morphan-alkaloid fragments are described. Indomorphane derivative Glupin was identified as a potent inhibitor of glucose uptake by selectively targeting and upregulating glucose transporters GLUT-1 and GLUT-3. Glupin suppresses glycolysis, reduces the levels of glucose-derived metabolites, and attenuates the growth of various cancer cell lines. Our findings underscore the importance of dual GLUT-1 and GLUT-3 inhibition to efficiently suppress tumor cell growth and the cellular rescue mechanism, which counteracts glucose scarcity.

Introduction

Strategies for the design and discovery of novel chemical matter endowed with bioactivity can draw from previous insight about the biological relevance of compound classes. This reasoning underlies, for example, biology-oriented synthesis (BIOS)^[1] and ring-distortion and/or -modification approaches (“complexity to diversity”; CtD).^[2] Limitations of BIOS derive from restricted coverage of natural product like chemical space and biological target space. These limitations may be overcome by the combination of BIOS with fragment-based design^[3] to arrive at “pseudo-natural products”.^[4] Pseudo-natural products are obtained by the de novo combination of natural product fragments^[5] that generate unprecedented compound classes not accessible by known biosynthesis pathways, and, therefore, go beyond the chemical space explored by nature. They may have new biological targets and modes of action, such that their bioactivity will best be evaluated in unbiased target-agnostic cell-based assays covering a wide range of biological processes.^[6]

Here we describe the design, synthesis, and biological investigation of indomorphan pseudo-natural products that

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<https://doi.org/10.1002/anie.201909518>.

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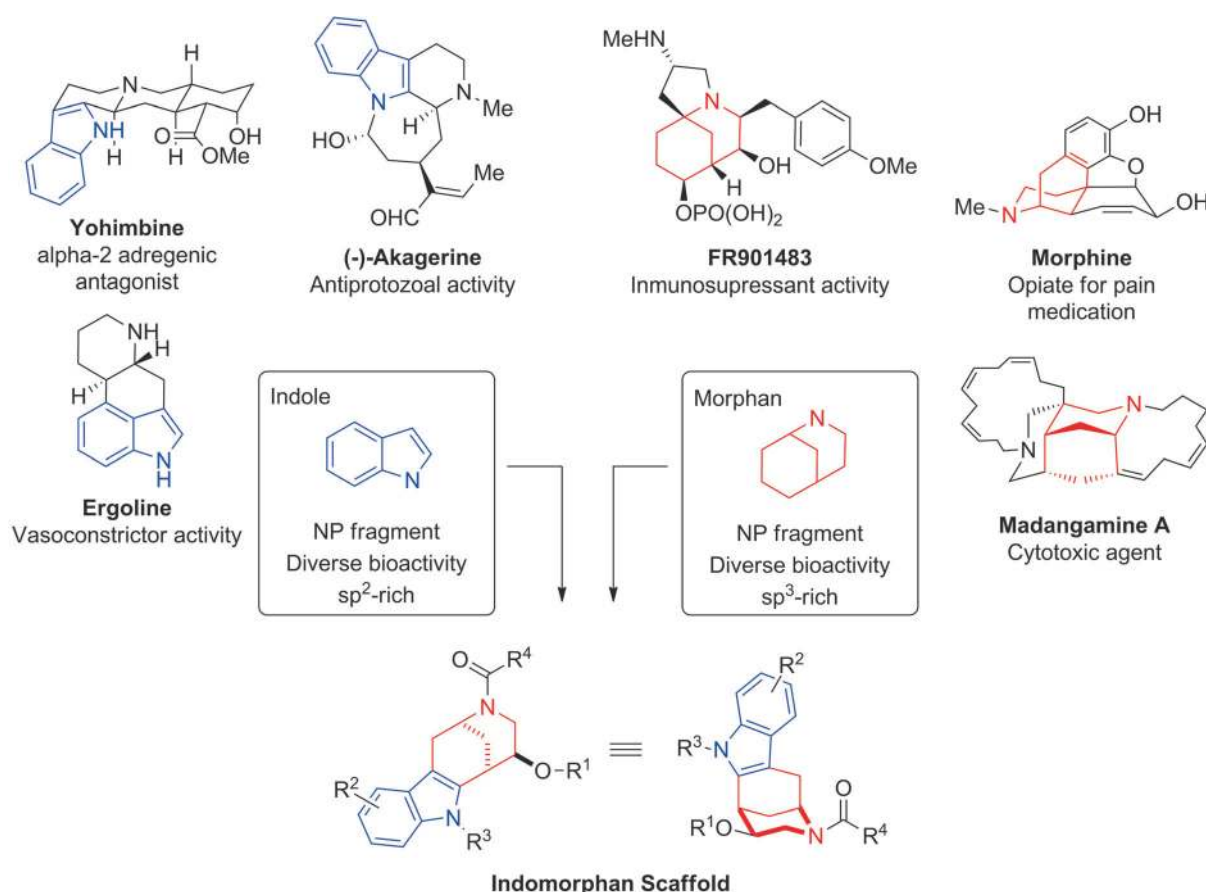
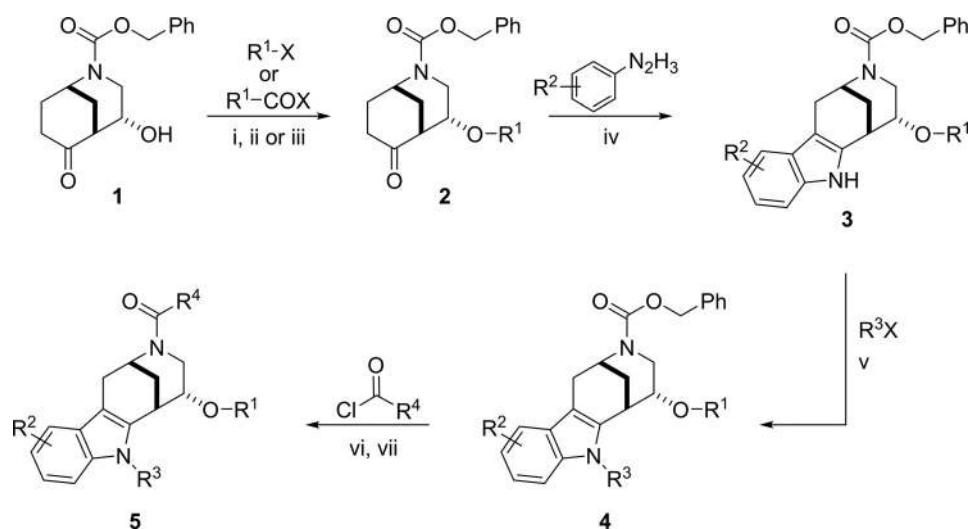


Figure 1. Design of the indomorphan pseudo-NP class based on the natural product fragments derived from indole (in blue) and morphan (in red) alkaloids.

contain characteristic structural fragments of the biosynthetically unrelated indole^[7] and morphan^[8] alkaloid classes, each of which is endowed with diverse and different bioactivities (Figure 1). A piperidine fragment is attached in an edge-on connection to an indole moiety in numerous polycyclic indole alkaloids,^[9] whereas a benzene ring is linked to a piperidine-containing fragment through a bridge in the scaffold of the morphinan alkaloids.^[9] In the indomorphans, these fragments are merged to a bridged, bicyclic indolyethylamine-containing novel pseudo-natural product class (Figure 1).

Biological investigation of the compound collection revealed that the indomorphans define a truly novel, structurally unprecedented glucose uptake inhibitor chemotype. The most potent inhibitor, termed Glupin, selectively blocks glucose uptake mediated by

GLUT-1 and GLUT-3 with low nanomolar potency, inhibits glycolysis, and efficiently suppresses the growth of various cancer cell lines.



Scheme 1. Synthesis of the indomorphan pseudo-NP class. i) MeI (5 equiv), Ag₂O (5 equiv), CH₂Cl₂, 72 h, rt; ii) AcCl, CH₂Cl₂/pyridine, 3 h, -10°C; iii) allyl ethylcarbonate (2 equiv) Pd(PPh₃)₄ (5 mol%), THF, 80°C, 2.5 h; iv) ArN₂H₃·HCl (1 equiv), AcOH, reflux, 1.5 h; v) R³Br (3 equiv), Cs₂CO₃ (3 equiv), DMF, 6 h, rt; vi) H₂, Pd/C, EtOH, 5 h, rt; vii) ArCOCl (1.0 equiv), NEt₃ (1.2 equiv), CH₂Cl₂, 12 h, rt.

Results

For the synthesis of the indomorphan library, bicyclic ketone **1**^[10] (Scheme 1) was derivatized to yield ethers **2**, which were then transformed into different indole derivatives **3** by means of the Fisher indole synthesis. Alkylation of the indole nitrogen atom (\rightarrow **4**), deprotection, and acylation of the morphan nitrogen atom yielded indomorphans **5**. All the transformations are preparatively simple and have wide scope. The synthesis yielded 43 indomorphans in five linear steps. In addition, 19 selected indomorphans were obtained commercially^[11] (see Table S1 in the Supporting Information).

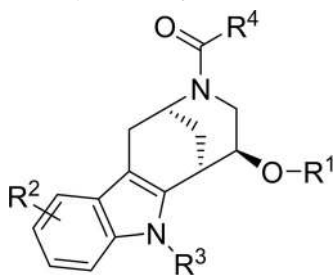
Calculation of the natural product score (NP-score) distribution^[12] of this pseudo-NP class, and comparison with the scores calculated for the NP set in ChEMBL,^[13] as well as for the set of marketed and experimental drugs listed in DrugBank,^[14] revealed that indomorphans exhibit a narrow distribution in a part of the NP-score graph that is not heavily populated by NPs but rather by synthetically accessible biologically relevant molecules (Figure S2a). Analysis of the principal moments of inertia (PMI),^[15] used as a crude measure of molecular shape, revealed that these pseudo-NPs possess a higher degree of three-dimensional character than common synthetically accessible compound collections, which reside typically on the rod-like to disc-like vertex of a PMI plot (Figure S2b).^[16] In addition, indomorphans display similar shape characteristics as indole- or morphan-containing NPs and non-naturally occurring biologically relevant compounds (see Figure S3 and Schemes S3 and S4). Further analysis using “Lipinski-Ro5” criteria, showed that 61 % of this collection is included within the limits of drug-like space (see Figure S2c and Table S2), thus indicating that de novo combinations of NP-derived fragments may result in compound collections with desirable properties.

The indomorphans were investigated in several cell-based assays that monitor biological signaling, for example, the Wnt and Hedgehog pathway, or metabolic processes such as autophagy and glucose uptake. The indomorphans proved to be inhibitors of glucose uptake in HCT116 cells, with the most potent compounds showing IC_{50} values in the very low nanomolar range (see below). Cancer cells meet their increased demand for metabolic energy by increasing glucose uptake, accelerated glycolysis, and conversion of pyruvate into lactate, even in the presence of oxygen (this phenomenon is termed the Warburg effect).^[17] To this end, tumors frequently upregulate facilitative, high-affinity glucose transporters (GLUTs). Several small molecules have been reported to inhibit glucose transporters (for a review, see Granchi et al.^[18]). However, many of them displayed low potency or selectivity. Very recently BAY-876, a potent and GLUT-1-selective inhibitor was developed by Bayer.^[19] However, both GLUT-1 and GLUT-3 are upregulated in various cancers^[20] and GLUT-3 expression can increase upon radio-chemotherapy or primary surgery. Therefore, cells overexpressing GLUT-3 may possess a survival advantage and may be clonally selected.^[21] Moreover, induction of *GLUT3* expression has been reported to be a protective mechanism against hypoglycemia in neurons,^[22] and cancer cells respond sim-

ilarly to decreased glucose levels.^[23] Thus, efficient inhibition of glucose uptake in tumors and cancer cells may require simultaneous targeting of both GLUT-1 and GLUT-3, and we recently developed the first GLUT-1/GLUT-3-selective compounds.^[4b,24]

The glucose uptake assay monitored the inhibition of 2-deoxyglucose (2DG) uptake as described by Yamamoto et al.^[25] Briefly, 2DG uptake is determined by quantification of the 2DG-6-phosphate (2DG6P) that accumulates inside the cells by using the diaphorase-resazurin/-resorufin system (see Figure S7a). Several indomorphans inhibited 2DG uptake with IC_{50} values ranging from submicromolar to low micromolar levels (Table 1 and Table S1), with compound **5b** (Table 1, entry 2) being the most active with an IC_{50} value of 53 ± 23 nM.

Investigation of the structure-activity relationship (SAR) profile of the indomorphan scaffold revealed that potent analogues were obtained when small groups, such as hydrogen, methyl, or acetyl, were used at the R^1 position (Table 1, entries 1–4). Even a PEG-based linker introduced at R^1 afforded a highly potent compound (entry 5). Substitution at the benzene ring (R^2) strongly reduced the activity, particularly with substitution at position 5, but potency was maintained when a hydroxy group was introduced (entries 6–9). Alkylation of the indole nitrogen atom (R^3) was essential for activity (entries 10–17). Derivatives with an ester functional group embedded in the substituent yielded the most potent compounds, whilst amides or a carboxylic acid resulted in a loss of potency. A one-carbon linker to the ester group was found to be optimal (entries 16 and 17). These findings show that an ester group is best, although not absolutely necessary, for the highest activity. For the ethyl ester and the carboxylic acid we calculated the $\log P$ values to be 3.4 and 2.3, respectively. Moreover, calculation of the pharmacokinetic properties of the acid and the ethyl ester revealed similar predicted permeability in Caco2 cells, namely, 1.12 and 1.08 ($\log P_{app}$ values in $10^{-6} \text{ cm s}^{-1}$) using the pkCSM tool^[41] and 38.9 and 23.0 using the preADMET tool.^[42] These data indicate that cell penetration by both the ester and the carboxylic acid is similar, such that the activity of the acid, if at all, represents only to a minor extent the possible hydrolysis of the ethyl ester in cellulo. This finding is further supported by the activity recorded for the *tert*-butyl ester and the amide, both of which would be hydrolyzed only slowly or not at all. Nevertheless, we cannot fully rule out that the ethyl ester may represent a prodrug. To explore whether other substituents may be tolerated, furanymethyl derivative **5o** (Table 1, entry 15) was synthesized. This analogue inhibited glucose import with an IC_{50} value of 6.6 ± 2.4 μM . This heterocycle is not a full replacement of the ester, but the result indicates that other groups are tolerated at this site. Acylation of the morphan nitrogen atom (R^4) was essential for the inhibition of 2DG uptake (entries 18–25). Activity in the low micromolar range was observed for several heterocycles and more potent analogues carried a functionalized nicotinate group. However, over-functionalization of the pyridine ring was disadvantageous for the activity, as exemplified by the introduction of a CF_3 group at positions 4 and 5 (entries 24 and 25).

Table 1: Structure-activity relationship analysis for the indomorphan class (see Table S1 for further details).

Entry	Compound	R ¹	R ²	R ³	R ⁴	IC ₅₀ [μM] ^[a]
1	(±)-Glupin (5a)	Me	H	CH ₂ CO ₂ Et	2-(methylthio)pyridin-3-yl	0.055 ± 0.017
2	5b	Ac	H	CH ₂ CO ₂ Et	2-(methylthio)pyridin-3-yl	0.053 ± 0.023
3	5c	H	H	CH ₂ CO ₂ Et	2-(methylthio)pyridin-3-yl	0.13 ± 0.05
4	5d	ⁿ Pr	H	CH ₂ CO ₂ Et	2-(methylthio)pyridin-3-yl	0.087 ± 0.015
5	5e	PEG ^{∗[b]}	H	CH ₂ CO ₂ Et	2-(methylthio)pyridin-3-yl	0.083 ± 0.026
6	5f	Me	5-OH	CH ₂ CO ₂ Et	2-(methylthio)pyridin-3-yl	0.10 ± 0.06
7	5g	Me	5-Cl	CH ₂ CO ₂ Et	2-(methylthio)pyridin-3-yl	1.4 ± 0.2
8	5h	Me	7-Cl	CH ₂ CO ₂ Et	2-(methylthio)pyridin-3-yl	0.21 ± 0.09
9	5i	Me	5-CO ₂ Et	CH ₂ CO ₂ Et	2-(methylthio)pyridin-3-yl	> 30
10	5j	Me	H	H	2-(methylthio)pyridin-3-yl	> 30
11	5k	Me	H	CH ₂ CO ₂ ^t Bu	2-(methylthio)pyridin-3-yl	0.19 ± 0.06
12	5l	Me	H	CH ₂ CO ₂ ⁿ Pr	2-(methylthio)pyridin-3-yl	0.093 ± 0.035
13	5m	Me	H	CH ₂ CONHMe	2-(methylthio)pyridin-3-yl	1.45 ± 0.6
14	5n	Me	H	CH ₂ CO ₂ H	2-(methylthio)pyridin-3-yl	10.0 ± 4.0
15	5o	Me	H	furan-2-ylmethyl	2-(methylthio)pyridin-3-yl	6.7 ± 2.4
16	5p	Me	H	CO ₂ Et	2-(methylthio)pyridin-3-yl	28 ± 2
17	5q	Me	H	(CH ₂) ₂ CO ₂ Et	2-(methylthio)pyridin-3-yl	8.4 ± 3.6
18	5r	Me	H	CH ₂ CO ₂ Et	<i>N</i> -morpholino	3.16 ± 0.6
19	5s	Me	H	CH ₂ CO ₂ Et	2-furyl	3.4 ± 0.3
20	5t	Me	H	CH ₂ CO ₂ Et	pyridine-3-yl	2.3 ± 0.8
21	5u	Me	H	CH ₂ CO ₂ Et	2-fluoropyridin-3-yl	0.54 ± 0.16
22	5v	Me	H	CH ₂ CO ₂ Et	2-methoxypyridin-3-yl	0.23 ± 0.05
23	5w	Me	H	CH ₂ CO ₂ Et	2-trifluoromethylpyridin-3-yl	0.12 ± 0.03
24	5x	Me	H	CH ₂ CO ₂ Et	2-(methylthio)-4-(trifluoromethyl)pyridin-3-yl	6.7 ± 1.6
25	5y	Me	H	CH ₂ CO ₂ Et	2-(methylthio)-5-(trifluoromethyl)pyridin-3-yl	14 ± 4

[a] IC₅₀ values determined for the inhibition of 2DG uptake in HCT116 cells. Data are mean values ($N \geq 3$ independent experiments, $n \geq 3$ independent replicates). Error represents mean ± S.D. [b] PEG[∗]: (2,2-dimethyl-4-oxo-3,9,12,15-tetraoxa-5-azaocadecan-18-yl)carbamic acid.

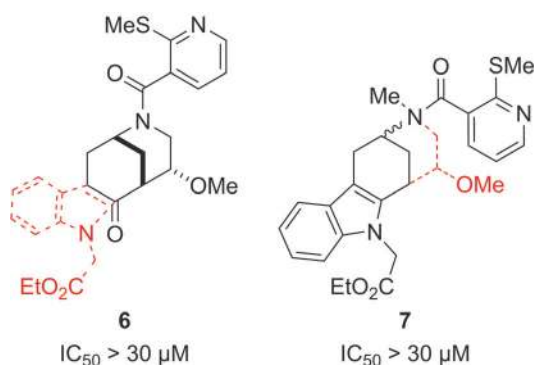


Figure 2. Structure and activity of the Glupin-based fragment analogues **6** and **7**. Red dashed lines show the fragments missing from the indomorphan core, with R¹ to R⁴ overall shown as in Glupin.

To investigate the contribution of the indole and morphan fragments on the bioactivity and to determine whether the novel fragment combination is required, we prepared indomorphan fragments **6** and **7** (Figure 2) which were decorated

with the same functional groups as the active compound **5a** (Table 1, entry 1). Notably, the inhibiting activity of 2-DG uptake was not shared by the indolyethyl fragment, which lacks the morphan scaffold, or by the bicyclic ketomorphan fragment, which lacks the indole ring. These results suggest that the observed biological activity of the indomorphan scaffold is not shared by the individual NP fragments and is, therefore, a property of the new pseudo-NP scaffold.

Based on the potency and structure, compound **5a**, termed (±)-Glupin (from glucose uptake inhibitor; Table 1, entry 1), was selected for further analysis. (±)-Glupin showed an IC₅₀ value of 5 ± 1 nM in separate, non-automated determinations on HCT116 cells and also reduced the uptake of 2DG in the highly glycolytic human breast cancer cell line MDA-MB-231 (IC₅₀ = 17 ± 5 nM; Figure 3a).^[26] This finding was further confirmed by the inhibition of tritium-radiolabeled 2DG uptake by (±)-Glupin with an IC₅₀ value of 8 ± 3 nM (Figure 3b). (±)-Glupin also potentially suppressed 2DG uptake in HeLa (IC₅₀ = 22 ± 2 nM) and CHO cells (IC₅₀ = 4 ± 3 nM). (±)-Glupin enantiomers were separated by preparative chiral HPLC, and the absolute configuration was assigned

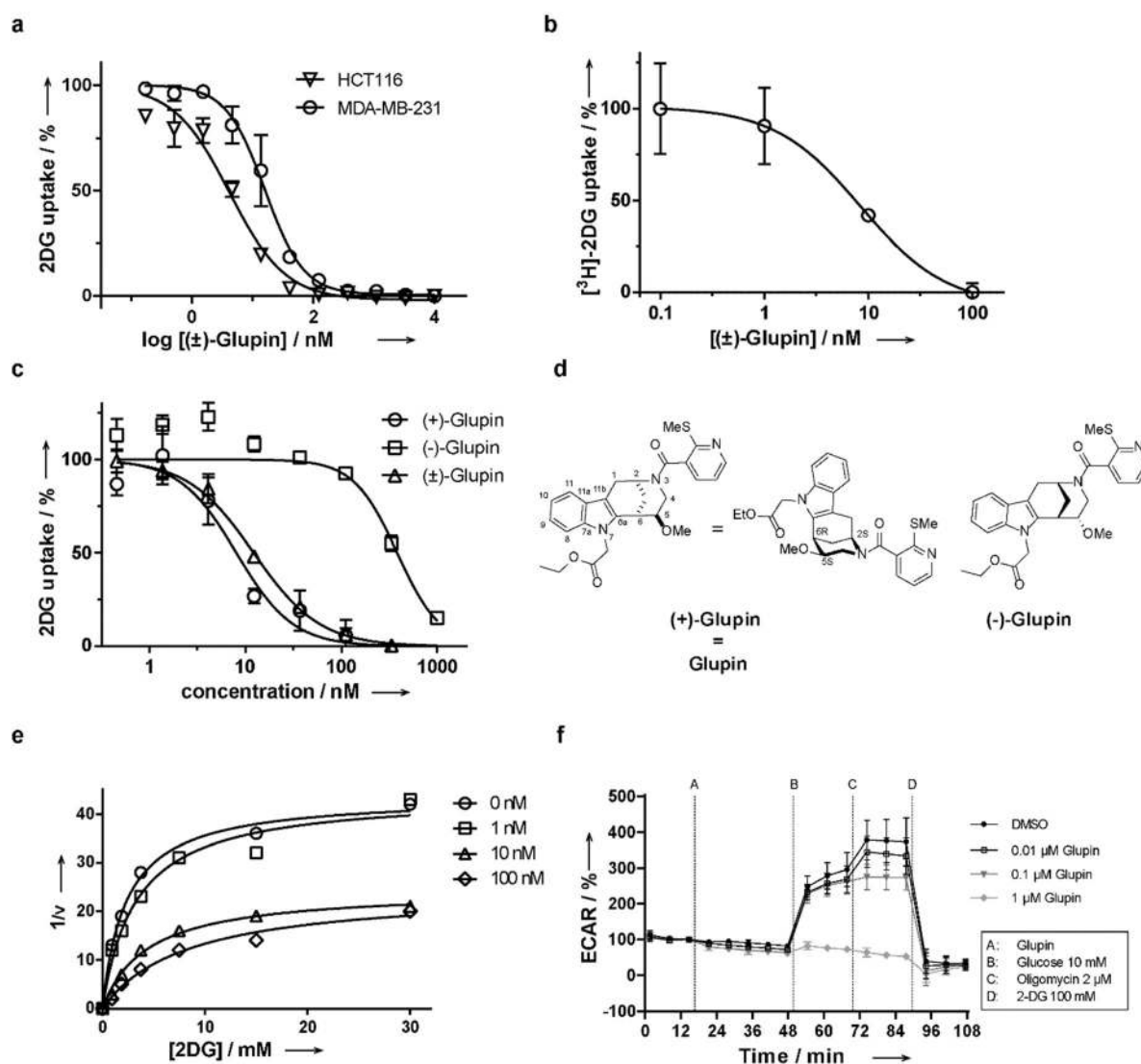


Figure 3. Inhibition of 2DG uptake by Glupin. a) (\pm)-Glupin inhibition of 2DG uptake in MDA-MB-231 and in HCT116. Data are mean values \pm SD ($N=3$, $n=3$). b) (\pm)-Glupin inhibition of ^3H -2DG uptake in MDA-MB-231 cells. Cells were treated with compound or DMSO and $1\ \mu\text{Ci}$ ^3H -2DG for more than 30 min. Data are mean values \pm SD ($N=3$, $n=3$). c) Glupin is the more potent enantiomer. 2DG uptake was measured in MDA-MB-231 cells. Data are mean values \pm SEM ($n=3$). d) Structures of (+)-Glupin (=Glupin) and the (-)-enantiomer. e) The 2DG uptake in the presence of different amounts of Glupin was measured in a time- and concentration-dependent manner in MDA-MB-231 cells and then analyzed by nonlinear regression using the Michaelis–Menten equation to determine the V_{max} and $K_{\text{M,app}}$ values. f) Influence of Glupin on glycolysis. Glucose-starved MDA-MB-231 cells were treated with Glupin and the extracellular acidification rate (ECAR) was measured over time in a Seahorse XFp Extracellular flux analyzer. Data are mean values \pm SD ($n=3$).

by means of chiral derivatization and independent synthesis of reference compounds (see the Supporting Information for details). The more potent (+)-(2*S*, 5*S*, 6*R*)-Glupin enantiomer inhibits 2DG uptake in MDA-MB-231 cells with $\text{IC}_{50} = 4 \pm 2\ \text{nM}$, while (-)-(2*R*, 5*R*, 6*S*)-Glupin displays an IC_{50} value of $335 \pm 47\ \text{nM}$ (Figure 3c,d). (+)-Glupin (hereafter referred to as “Glupin”) did not inhibit the downstream hexokinase, thus demonstrating direct interference with glucose uptake (see Figure S7b).

Kinetic measurements in MDA-MB-231 cells showed that the maximum rate (V_{max}) decreased, whereas the K_{M} value increased as the concentration of Glupin increased, thus indicating a mixed type of inhibition (Figure 3e). Investiga-

tion of the influence of Glupin on the extracellular acidification rate (as a measure of glycolysis) and the oxygen consumption rate (as a measure of oxidative phosphorylation) of MDA-MB-231 cells revealed that $1\ \mu\text{M}$ Glupin inhibited glycolysis to the level of glucose-starved cells (Figure 3f) and increased the oxygen consumption rate (Figure S8a). Moreover, treatment of MOLT16 cells with $50\ \text{nM}$ (\pm)-Glupin reduced the levels of metabolites linked to glycolysis (glycerol-3-phosphate, lactic acid), the tricarboxylic acid (TCA) cycle (citric/isocitric acid), the pentose phosphate pathway (ribitol/arabitol), and the hexosamine biosynthetic pathway (UDP-*N*-acetylglucosamine), which all utilize glucose-6-phosphate (Table S3). The level of aspartate, which

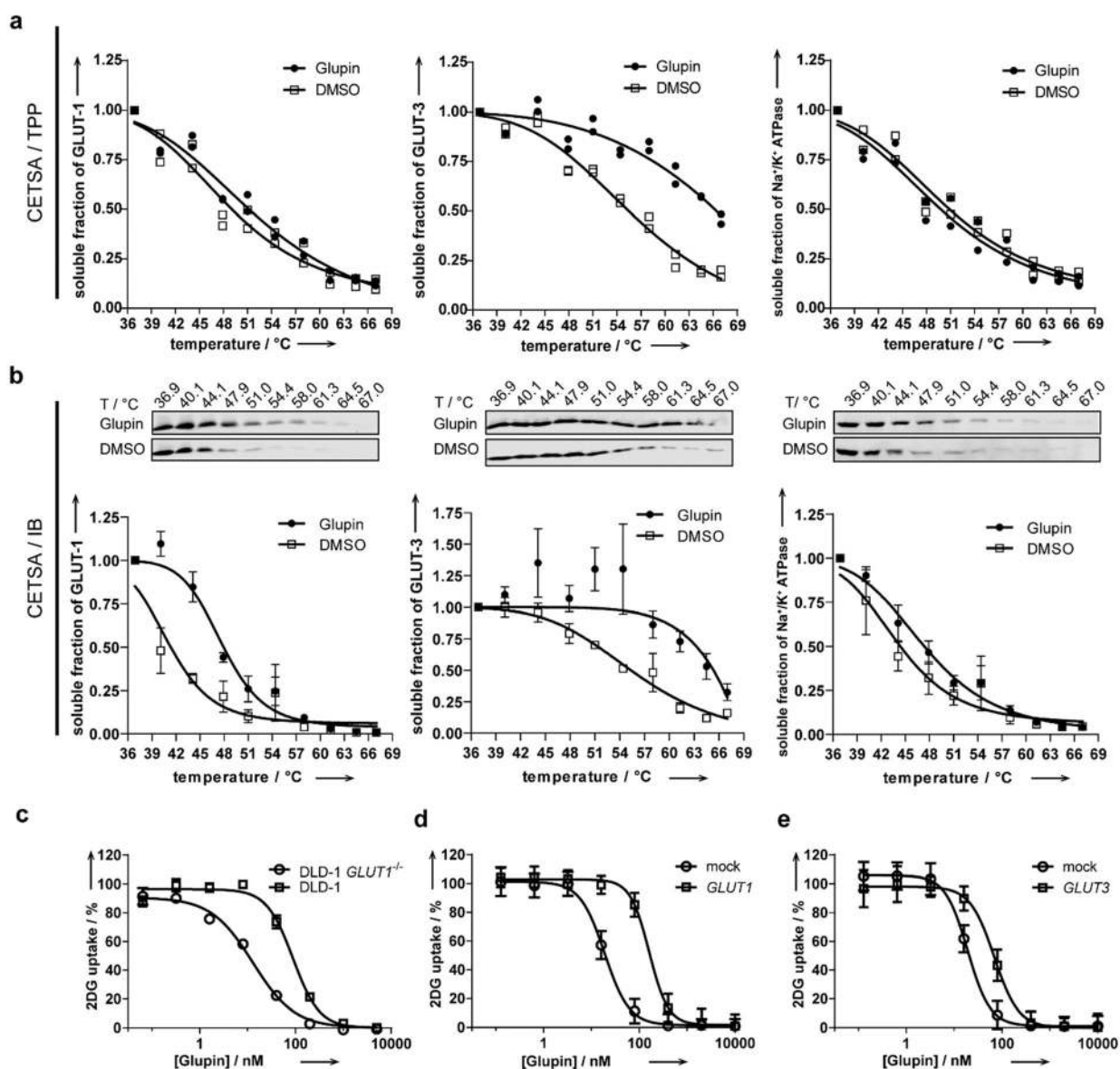


Figure 4. Target engagement of GLUT-1 and GLUT-3 by Glupin. The cellular thermal shift assay was performed with lysates from SW480 cells in the presence of 1 μM Glupin or DMSO using mass spectrometry (a) or immunoblotting (b) to detect GLUT-1 or GLUT-3 and Na⁺/K⁺-ATPase as a control. Data of two biological replicates (a) or mean values \pm SD and representative of $n=3$ (b) are shown. c) Uptake of 2DG in DLD-1 GLUT1^{-/-} cells after incubation with the compound for 30 min. d) Uptake of 2DG in CHO cells that ectopically express GLUT-1 compared to cells transfected with an empty vector (mock). e) Uptake of 2DG in CHO cells that ectopically express GLUT-3 compared to cells transfected with an empty vector (mock). All data are mean values \pm SD ($n=3$).

can fuel the TCA cycle, was elevated upon treatment with Glupin, most likely as a compensatory mechanism.^[27] These results are in agreement with inhibition of glucose uptake by Glupin and, consequently, the synthesis of glucose-6-phosphate.

In the absence of glucose, cells may utilize alternative energy sources and building blocks to sustain cell growth, in particular fatty acids, and altered lipid storage and metabolism is strongly connected with cancer malignancy.^[28] Treatment of MDA-MB-231 cells with Glupin in the presence of 400 μM oleic acid markedly reduced the number of lipid droplets with an IC₅₀ value of 84.9 nM (Figure S8b). Pre-

formed lipid droplets were not affected, thus suggesting that Glupin has no stimulating effect on lipolysis (Figure S8c).

Target engagement was demonstrated by means of a proteome-wide cellular thermal shift assay (CETSA, thermal proteome profiling, TPP)^[29] using SW480 cells, which revealed thermal stabilization of GLUT-3 ($\Delta T_m = 7.1 \pm 0.4^\circ\text{C}$) and GLUT-1 ($\Delta T_m = 1.9 \pm 0.3^\circ\text{C}$) in the presence of 1 μM Glupin (Figure 4a). No further members of the GLUT family could be detected. A separate CETSA experiment using immunoblotting (Figure 4b) confirmed this finding and showed thermal stabilization of GLUT-1 ($\Delta T_m = 6.5 \pm 0.6^\circ\text{C}$) and GLUT-3 ($\Delta T_m > 12^\circ\text{C}$).

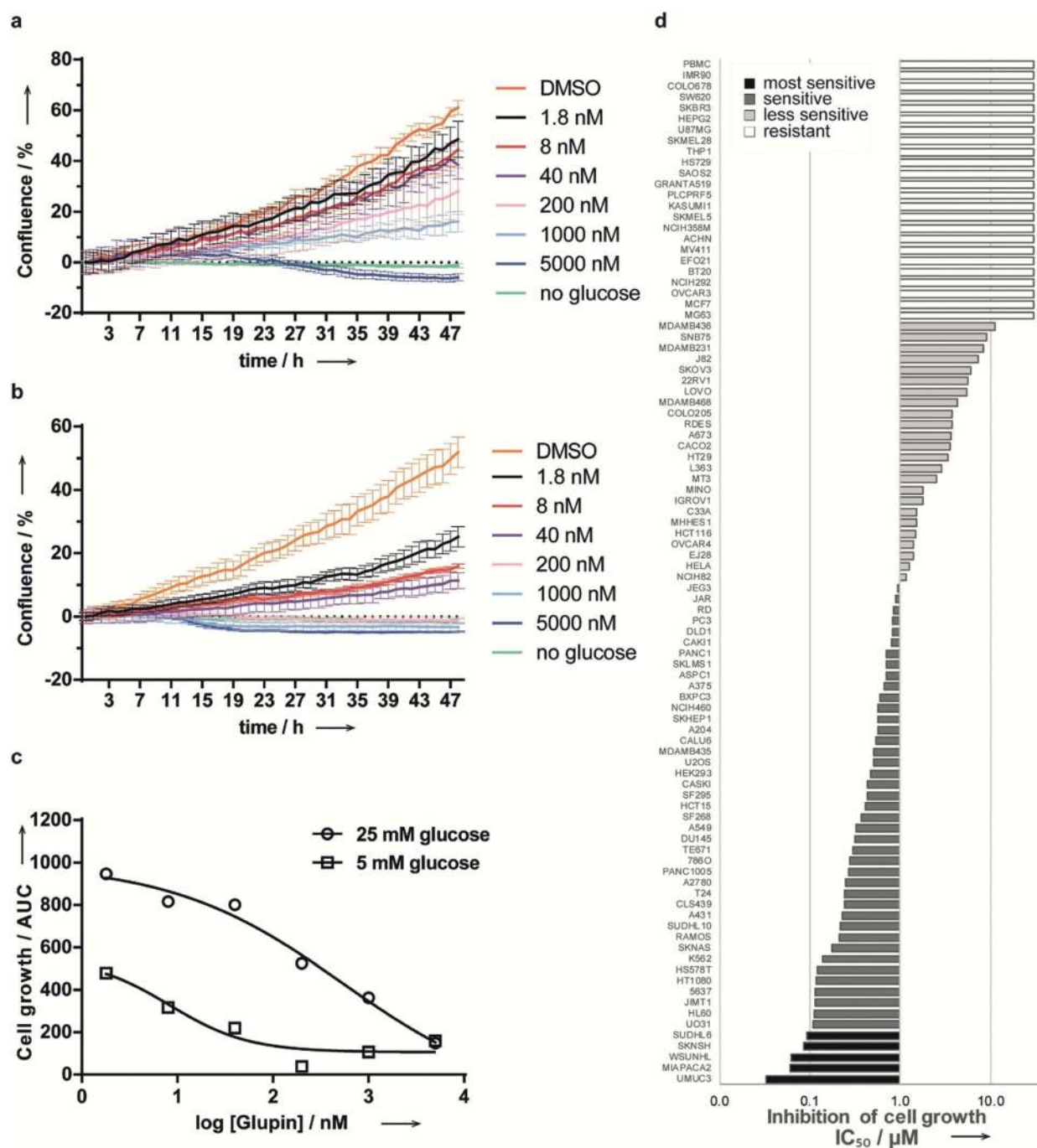


Figure 5. Inhibition of cancer cell growth by Glupin. a–c) The growth of MDA-MB-231 cells in 25 mM (a) or 5 mM glucose (b) was monitored in the presence of Glupin or DMSO by means of live-cell kinetic analysis and confluence as a measure. Data are mean values ($N=3$) \pm SD and representative of $n=3$. c) Comparison of cell growth at 5 and 25 mM glucose. The area under the curve (AUC) was calculated for the curves in (a) and (b). d) Glupin inhibits the growth of various cancer cell lines. 94 cell lines were treated with different concentrations of Glupin for 72 h followed by the sulforhodamine B assay and IC_{50} determination.

DLD-1 *GLUT1* ($-/-$) cells do not express GLUT-1 but exhibit substantial expression of GLUT-3 (Figure S9).^[4b] In contrast, the isogenic parental cell line DLD-1 expresses mainly GLUT-1 (Supporting Figure S9).^[4b,30] Glupin inhibited the uptake of 2DG in both cell lines with IC_{50} values of 59.6 ± 8.4 nM for DLD-1 and 11.4 ± 1.6 nM for DLD-1 *GLUT1* ($-/-$; Figure 4c). This finding confirms that Glupin targets both GLUT-1 and GLUT-3 and indicates that Glupin inhibits

GLUT-3 more strongly than GLUT-1. Ectopic expression and, thereby, increased abundance of GLUT-1 or GLUT-3 in CHO cells^[4b] (Figure S10a,c), led to a partial rescue of the Glupin-mediated inhibition of glucose uptake as detected by the increase in the IC_{50} values from 19.2 ± 2.1 nM to 162 ± 14 nM for GLUT-1 overexpression (Figure 4d) and from 19.1 ± 2.9 nM to 68 ± 7.2 nM for GLUT-3 (Figure 4e). In contrast, overexpression of GLUT-2 or GLUT-4 did not have

any influence on the activity of the compound (Figures S10b,d and S11 as well as Table S4). These results suggest that Glupin interacts with GLUT-1 and GLUT-3 but not with GLUT-2 and GLUT-4.

Real-time analysis of the growth of the MDA-MB-231 cell line in the presence of Glupin showed a dose-dependent suppression of cell growth in plain medium containing 25 mM glucose (Figure 5a). This effect was much stronger at a physiological glucose concentration (5 mM; Figure 5b,c). Exposure of the compound to more than 90 cancer cell lines and IMR-90 and PBMC cells as noncancerous controls (see Figure 5d and Table S5) revealed that the growth of the bladder cancer cell line UM-UC-3 ($IC_{50}=32$ nM), the pancreas cancer cell line MIA PaCa-2 ($IC_{50}=61$ nM), the lymphoma cell lines WSU-NHL and SU-DHL-6 ($IC_{50}=62$ nM and 92 nM, respectively), and the neuroblastoma cell line SK-N-SH ($IC_{50}=84$ nM), which depend on glucose for cell growth,^[31] was potently inhibited. IC_{50} values lower than 300 nM were obtained for more than 20 cell lines, whereas 24 cell lines were not affected by the compound, among them the peripheral blood mononuclear control cells (PBMCs). Sensitivity to Glupin was detected across various tissue types (Table S6). These results indicate that targeting glucose import by GLUT-1 and GLUT-3 may be a viable approach to inhibit the growth of cancer cells.

Since glucose depletion may lead to increased expression of GLUT-1 and GLUT-3,^[22,23,32] we analyzed the influence of Glupin on the expression of *GLUT1* (*SLC2A1*) and *GLUT3* (*SLC2A3*). The application of Glupin for 48 h induced only a slight increase in the expression of both genes at the mRNA level (Figure S12). However, treatment with 0.5 μ M Glupin significantly increased GLUT-1 levels after 24 h of treatment (Figure S13 a,b) and of GLUT-3 levels after 48 h of treatment (Figure S13 c,d). These results indicate that Glupin-mediated inhibition of glucose uptake increases the levels of both GLUT-1 and GLUT-3 in DLD-1 cells.

Discussion

Our results demonstrate that the combination of natural product (NP) fragments in unprecedented arrangements may yield biologically relevant pseudo-NP collections with both novel NP-inspired structure and novel bioactivity. The indomorphan pseudo-NPs retain the characteristic structural elements of the guiding NPs but reside in an area of chemical space which is inaccessible to nature, reflected by the fact that they are not obtainable by known biosynthetic pathways. This novel, and structurally complex scaffold may be endowed by design with advantageous physiochemical properties, as the pseudo-NP collection displays a NP-score distribution closer to that of approved drugs.

The term “pseudo-natural product” and reasoning related to the concept have inspired synthesis efforts before. Thus, Suga and co-workers used the term to describe in vitro synthesized cyclic peptides embodying non-natural amino acids,^[33] and Oshima and co-workers qualified alkaloid-like compounds obtained by the interception of biosynthetic pathways as pseudo-NPs.^[34] Tietze et al.^[35] reviewed the

syntheses of natural product hybrids, that is, NP-like compounds which are composed of different NP substructures and may be considered pseudo-NPs if they represent pseudo-NP combinations that match our concept.^[4b,5] In attempts to generate potential alkaloid types which might not have been identified in nature at the time, Morrison et al.^[36] replaced 3,4-dihydroxyphenylalanine by tryptophan in a synthesis following the biogenetic route leading to the morphine scaffold. Bosch et al. (see Ref. [37] and references therein) developed heteromorphans, that is, compounds in which the benzene ring in 6,7-benzomorphan is replaced by a heterocyclic ring, including an indomorphan analogue. “Unnatural” polyketide natural products have been obtained by means of synthetic biology approaches, for example, through precursor-directed biosynthesis, mutasynthesis, or combinatorial biosynthesis (for reviews see Ref. [38]), and natural and non-natural structural elements have been combined in polyketide syntheses to yield natural product analogues not accessible by biosynthetic pathways (see for example, Ref. [39]).

The indomorphan pseudo-NPs define a novel glucose uptake inhibitor class with unprecedented chemotype. The most potent compound Glupin selectively targets glucose transporters GLUT-1 and -3 and potently inhibits the growth of cancer cells. Our results demonstrate that inhibition of GLUT-1 and GLUT-3 may be necessary to efficiently inhibit the growth of cancer cells since both glucose transporters GLUT-1 and GLUT-3 may be upregulated upon withdrawal of glucose.^[22,23,32] Dual GLUT-1- and GLUT-3-selective compounds such as Glupin would inhibit both transporters. Indeed, treatment with Glupin increased the expression of both GLUT-1 and GLUT-3 in DLD-1 cells, which primarily express GLUT-1. Upregulation of GLUT-3 upon glucose deprivation also ensures the uptake of glucose at lower glucose concentrations in the cell environment compared to GLUT-1 because of its higher affinity for glucose.^[40] Thus, in particular, the overexpression of GLUT-3 may pose a hurdle for the efficacy of GLUT-1-selective agents and emphasizes the need to simultaneously target both proteins. These findings may inspire new drug discovery programs aimed at the modulation of tumor metabolism.

Acknowledgements

Research at the Max Planck Institute of Molecular Physiology was supported by the Max Planck Society and the European Research Council under the European Union's Seventh Framework Programme (FP7/2007–2013)/ERC Grant agreement no. 268309. M.S., J.C., E.S.R., and J.W. acknowledge the International Max-Planck Research School for a doctoral scholarship. G.K. and L.L. received funding from the Alexander von Humboldt Stiftung for a postdoctoral fellowship; M.C.C. received funding from the Fonds der chemischen Industrie (Kekulé fellowship) and the Swiss National Science Foundation (Doc.mobility). M. Sellstedt received funding from the Swedish Research Council, grant number 350-2012-6611. The compound management and screening center (COMAS) Dortmund is acknowledged for performing the high-throughput screening and the orthogonal and counter

screens. The Lead Discovery Center (LDC) in Dortmund is acknowledged for performing the cell panel analysis. We are grateful to Andreas Brockmeyer (Max Planck Institute of Molecular Physiology) for assistance during MS/MS measurements. The Swedish Metabolomics Centre's Open Access Lab is acknowledged for access to, and assistance with, mass spectrometers for determinations of metabolite levels.

Conflict of interest

The development of glucose transport inhibitors is an actively pursued drug discovery project at the Lead Discovery Center GmbH (LDC) of the Max Planck Gesellschaft, sponsored by H.W.

Keywords: antitumor agents · glucose transporters · inhibitors · natural products · pseudo-natural products

How to cite: *Angew. Chem. Int. Ed.* **2019**, *58*, 17016–17025
Angew. Chem. **2019**, *131*, 17172–17181

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Manuscript received: July 29, 2019

Revised manuscript received: August 12, 2019

Accepted manuscript online: August 30, 2019

Version of record online: October 7, 2019