

Synthesis and screening of 6-alkoxy purine analogs as celltype selective apoptotic inducers in Jurkat cells

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Synthesis and screening of 6-alkoxy purine analogs as cell-type selective apoptotic inducers in Jurkat cells

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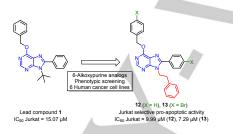
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A new series of 6-alkoxy purines derived from **1** was designed, synthesized and the pro-apoptotic activity of this chemotype evaluated on cancer cells. Using a phenotypic approach based on the cell cycle analysis, the compound library was screened against a 6-member panel of human cancer cell lines. The cell death induction mechanism and cell viability effects of hit compounds were also examined, showing **12** and **13** as T cell leukemia Jurkat-selective pro-apoptotic compounds.

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Abstract

Purines are ubiquitous structures in cell biology involved in a multitude of cellular processes. This has made substituted purines and analogs to be considered privileged scaffolds in drug design. In this work, we explored the key structural features of a purine-based pro-apoptotic hit, 8-*tert*-butyl-9-phenyl-6-benzyloxy-9*H*-purine (1), by making a library of 6-alkoxypurines with the aim to elucidate the structural requirements that govern its biological activity and to study the cell selectivity of this chemotype. This was addressed by a phenotypic screening approach based on cell cycle analysis of a panel of 6 human cancer cell lines, including T cell leukemia Jurkat cells. From this study, two derivatives (12 and 13) were identified as Jurkat-selective pro-apoptotic compounds, displaying superior potency and cell-selectivity than hit 1.

KEYWORDS: Anticancer drugs, Purine analogs, Phenotypic screening, Apoptosis, Leukemia

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1 INTRODUCTION

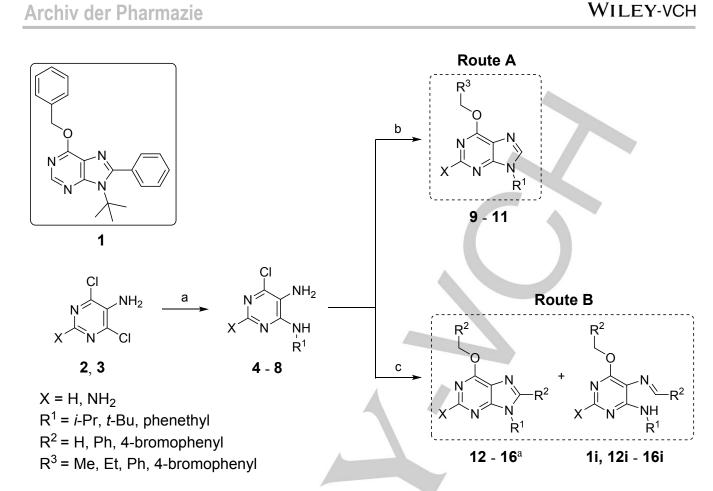
The purine ring system (imidazo[4,5-d]pyrimidine) is the most abundant nitrogen-based heterocycle in nature.^[1,2] This scaffold is contained in adenine and guanine, and therefore it is part of nucleic acids and nucleotides (AMP, GMP), essential molecules that regulate cellular energy and intracellular signaling processes (ATP, GTP) and enzyme cofactors (NADH, coenzyme A).^[3] The biological ubiquity of this heterocyclic ring makes it a privileged scaffold in drug design. Hence, several purine analogs have been developed for the treatment of different diseases such as leukemias, viral infections, cancer or immune disorders, and many others are currently being studied.^[4–6]

Typically, purine analogs are generated by the introduction of substituents at positions C2, C6, C8 and/or N9 using different synthetic methodologies.^[7,8] In this respect, our group developed a two-step synthetic methodology that allows the preparation of purine derivatives substituted at positions C6, C8 and N9.^[9] This is achieved by the use of different 6-chloro-4,5-diaminopyrimidines, alcohols and *N*,*N*-dimethylamides under strong basic conditions.^[10,11] Additionally, the compounds generated with this procedure present an ether group attached at C6, which is a less explored structural feature in purine libraries, where C6-aminated analogs prevail. The initial compound series generated with this strategy led to the identification of the purine **1**, an antiproliferative compound that exerts its activity by strong apoptotic induction on Jurkat (acute T cell leukemia) cells.^[12]

2 RESULTS AND DISCUSSION

2.1 Chemistry

Compounds 9 - 16, 1i, 12i - 16i were synthesized according to Scheme 1. Dichloropyrimidines 2 and 3 were substituted with different alkylamines at position C4 of the pyrimidine ring, providing intermediates 4 - 8. These chloropyrimidines were treated with the appropriate alcohol and an excess of the suitable *N*,*N*-dimethylamide under strong basic conditions to simultaneously carry out a displacement at C6 and an annulation reaction, affording purine analogs 9 - 15. This tandem reaction was controlled by the type of the *N*,*N*-dimethylamide used as solvent/reagent. Thus, DMF drove the synthesis through route A, generating C8 non-substituted purines, while sterically hindered *N*,*N*-dimethyl-amides, such as dimethylbenzamide (DMB), led the synthesis through route B, where the alcohol provided radicals at both C6 and C8. For compounds containing the 2,5-diamino pyrimidine core, microwave irradiation at higher temperatures was employed due to the lower reactivity of this core compared with the 5-amino-2-chloropyrimidine one. In addition to the proposed library, in those cases where DMB was used together with bulky alcohols (benzyl alcohol or 4-bromobenzyl alcohol), the intermediate imines 12i - 16i were also isolated (Table 1). Interestingly, in the preparation of the aminopurine 16 only the corresponding imine 16i was isolated.



SCHEME 1 Structure of compound 1 and synthesis of compounds 9 – 16, 1i and 12i – 16i. Reagents and conditions: (a) H₂N-R¹, NaHCO₃, THF, 55 °C, 18 h, or n-butanol, 190 °C (MW), 2 h, 46 – 84%; (b) NaH, DMF, 90 °C, 18 h or 150 °C (MW), 4 h, 7 – 45%; (c) DMB, BnOH or 4-bromobenzyl alcohol, NaH, 1,4-dioxane, 90 °C, 18 h or 150 °C (MW), 4h, 5 - 19%. a16 was not isolated.

Table 1 Compound library

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Purine analogs Scaffold	Х	R ¹	R ²	R ³	MW	cLogD ^[a]	Compd.
	⊢н	$\overset{\sim}{\prec}$	$\vdash \bigcirc$	$\vdash \bigcirc$	358.45	5.17	1
	⊢н	\sim	<mark>⊢</mark> н	−сн₃	268.32	2.73	9
	⊢н	\sim	Ļн	<ch<sup>3</ch<sup>	282.35	3.26	10
R ³	⊢н	$\sim $	$\vdash \bigcirc$	$\vdash \!\!\!\! \bigtriangledown$	406.49	6.13	12
$N \rightarrow R^2$	ŀн	\swarrow	⊢	⊢ →Br	564.28	7.66	13
X N N R ¹	-NH ₂	$\langle \mathbf{x} \rangle$	⊢н	$\vdash \!\!\!\! \bigcirc$	283.34	2.71	11
	-NH ₂	\swarrow	$\vdash \bigcirc$	$\vdash \bigcirc$	359.43	4.74	14
	-NH ₂	\swarrow	$\vdash \!\!\! \bigtriangledown$	$\vdash \!\!\!\! \bigtriangledown$	373.46	5.02	15
	-NH ₂	$\sim \bigcirc$	$\vdash \bigcirc$	$\vdash \!\!\! \bigcirc$	421.50	5.98	16 ^[b]
Pyrimidine Schiff b Scaffold	ase analo X	ngs R ¹	R ²	R ³	MW	cLogD ^[a]	Compd.
Scanolu		κ. 					
	Ļн	Ϋ́Υ	\vdash	\vdash	360.46	5.21	1i
R ³	⊢н	$\sim \bigcirc$	$\vdash \bigcirc$	$\vdash \bigcirc$	408.51	6.16	12i
	⊢н	$\sim \bigcirc$	⊢ ← Br	⊢	566.30	7.70	13i
	-NH ₂	\swarrow	$\vdash \bigcirc$	$\vdash \bigcirc$	361.45	4.69	14i
X N NH R ¹	-NH ₂	\swarrow	$\vdash \!\!\! \bigcirc$	$\vdash \!\!\!\! \bigcirc$	375.48	4.97	15i
	-NH ₂	\sim	$\vdash \bigcirc$	\mapsto	423.52	5.92	16i

[a] clogD values were calculated using Chemicalize by ChemAxon Ltd. at pH 7.4.^[13] [b] Not isolated.

2.2 Pharmacology/Biology

Cell cycle analysis was chosen as the primary screening output due to the capability of this method to identify active compounds and discriminate whether the new analogs maintain or not the pro-apoptotic activity of the lead molecule **1**. Thus, the final compound library was tested against a panel of six human cancer cell lines (hematopoietic: Jurkat, acute T cell leukemia; K562, chronic erythroleukemia; non-hematopoietic: HeLa, cervix adenocarcinoma; G361, malignant melanoma; MDA-MB-231, breast adenocarcinoma and HCT116, colorectal adenocarcinoma). Cells were treated with each compound for 48 h at a single dose (30 μ M). Cells treated with **1** (30 μ M) or DMSO (equivalent amount to that contained in the 30 μ M dose) were used as a reference and negative control, respectively, and the cell cycle profiles were determined by flow cytometry after staining of nuclear DNA with propidium iodide (PI) as previously described.^[12] The percentages of cells within the sub-G₁ region (apoptotic cells) were measured and values from negative controls (DMSO) subtracted to show the variation on the apoptotic populations.

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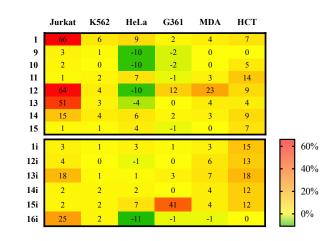


FIGURE 1 Phenotypic screening of compounds 9 - 15, 1i and 12i - 16i by cell cycle analysis. Cells were incubated for 48 h with the indicated compounds at 30 μ M, then stained with propidium iodide and cell cycles analyzed by flow cytometry. The heatmap represents changes in the percentage of cells within the sub-G₁ region after subtracting values from negative controls (DMSO) in each cell line. Compound 1 was used as reference.

Extending the length and bulkiness of the substituent at R^1 by the introduction of a phenethyl group afforded compounds with different cellular activities. When these analogs were not substituted in R^2 (=H) and also contained alkyloxy groups at C6 position (9 and 10), the purine analogs did not show any significant activity. In contrast, when both positions were occupied by lipophilic substituents such as phenyl or 4-bromophenyl group (12 and 13, respectively), a significant increment on the apoptotic populations in Jurkat cells was detected. Compound 12 showed a 64% increase in this cell line and a moderate activity over MDA-MB-231 cells (23%). Meanwhile, compound 13 showed a very selective activity over Jurkat cells with a sub-G₁ cell increment of 51% while being negligible against the rest of cancer cell lines. The structural similarities between these two compounds with 1 suggest that the presence of bulky aromatic groups at C6 and C8 positions drives the apoptotic activity of these compounds towards Jurkat cells.

Compounds **11**, **14** – **15** contained an amino group at C2 position. None of these compounds showed any significant effect in the cell cycle of the lines tested. Interestingly, the C2 aminated analog of **1**, compound **15** was completely inactive. This suggests that the presence of the amino group disrupts the biological interaction of **1** with its biological target.

The imines **1i** and **12i** – **16i** were tested next. These molecules were the pyrimidine Schiff base analogs of the corresponding compounds **1**, **12** – **16**. Among these, only the C2 aminated imine of **1**, compound **15i**, exhibited a notable activity, inducing a 41% increase in the sub-G₁ population of melanoma G361 cells, a phenotypic activity profile only seen with this derivative. Also, moderate activities were observed for compound **16i** on Jurkat cells. However, apart from these two imine derivatives, none of the rest showed significant activities, not even the corresponding Schiff base analogs of the highly active Jurkat-selective compounds **12**, **13** (**12i** and **13i**) or the parent compound **1** (**1i**), probably because of the different spatial arrangement and physicochemical properties of the substituents compared with the purines analogs. Based on the data from this phenotypic screening, compounds **12** and **13** were selected as the most active compounds and their activity were further analyzed.

A titration of the activity of the new compounds was done by culturing the cells with 30 μ M, 10 μ M and 3 μ M for 48 h followed by cell cycle analysis (**Figure 2**). Compound **12** showed a stronger capacity to induce cell death, as observed with the lowest dose of 3 μ M compared to the other two compounds.



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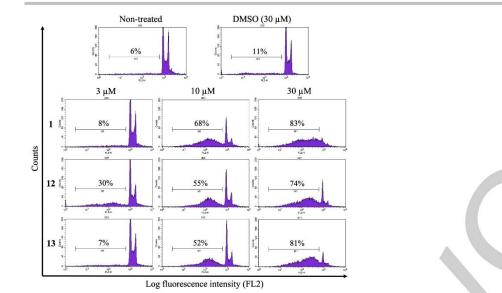


FIGURE 2 Titration of the proapoptotic activity of compounds 1, 12, and 13. Jurkat cells were treated for 48 h with the indicated doses and percentages of cells within the sub-G1 region measured by flow cytometry after staining with propidium iodide.

To validate the proapoptotic character of these two molecules, we stained the treated cells with annexin-V, which selectively binds to the phosphatidyl serine translocated to the cell membrane during the process of apoptosis, followed by flow cytometric analysis as previously described.^[12] For this study, Jurkat cells were treated for 48 h with compounds **12** and **13** at 30 μ M, 10 μ M and 3 μ M. DMSO (30 μ M) and **1** (30 μ M, 10 μ M and 3 μ M) were used as negative and positive apoptotic controls, respectively (**Figure 3**). As expected, untreated Jurkat cells presented a negligible apoptotic population with the majority of cells gated within the viable cell region (R1). However, when cells were treated with compounds **12** and **13** at the indicated concentrations, a transition behavior from early (R2) to late apoptotic (R3) stages was observed. In addition, this effect was displayed in a dose-dependent manner, and these results are fully coherent with those shown in **Figure 2**, since compound **12** induced a significantly higher apoptosis as revealed in the percentages of R2 and R3 at the lowest concentration of 3 μ M in both assays. Finally, no necrotic populations were detected for any of these compounds **12** and **13** was equivalent to that shown by the parent compound **1**, suggesting that the three compounds share the same mechanism of action and that the introduction of the phenethyl group at N9 or 4-bromophenyl groups at C8 and C6 positions do not affect the biological behavior of this chemotype.

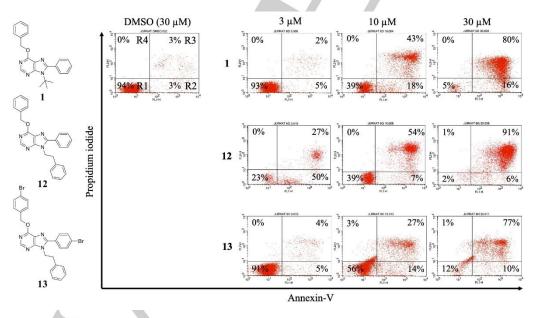
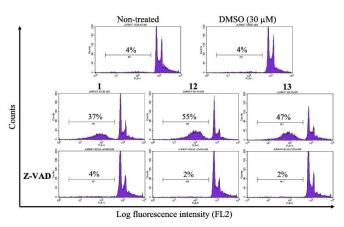


FIGURE 3 Membrane phosphatidylserine expression after treatment of Jurkat cells with compounds 12 and 13. Jurkat cells were treated for 48 h with the indicated doses and the percentages of apoptotic cells measured by flow cytometry after double staining with propidium iodide and Annexin-V-FLUOS. Untreated cells (DMSO, 30 µM) and cells treated with 1 were included as negative and positive controls of apoptosis, respectively. Cell populations (gating regions): R1, viable cells (Annexin-V⁻/PI⁻); R2, early apoptotic cells (Annexin-V⁺/PI⁻); R3, late apoptotic cells (Annexin-V⁺/PI⁺); R4, necrotic cells (Annexin-V⁺/PI⁺).

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To further demonstrate that the observed effect was due to the induction of apoptosis, we carried out assays where Jurkat cells were pre-treated for 1 h at 37 °C with the pan-caspase inhibitor Z-VAD-FMK (carbobenzoxy-valyl-alanyl-aspartyl-[O-methyl]-fluoromethylketone). Indicated compounds were then added for 24 h at the dose of 30 µM and subjected thereafter to cell cycle analysis (**Figure 4**). Whereas untreated cells showed a significant population contained within the sub-G1 region, this effect was completely abrogated in Z-VAD-FMK-pretreated cells. This indicates that induction of cell death mediated by the tested compounds is fully dependent on caspase activation, a hallmark of apoptosis pathways.



 μ M). [b] cLogD values were calculated using Chemicalize by ChemAxon Ltd. at pH 7.4.^[13]

FIGURE 3 Caspase-dependent Jurkat cell death induced by compounds 12 and 13. Cells were either pretreated (bottom panels) or not with the pan-caspase inhibitor Z-VAD-FMK for 1 h and thereafter treated with indicated compounds (30 µM) for 24h.

Additionally, to further compare the activities of compounds **12** and **13** with the reference molecule **1**, cell viability was measured on Jurkat cells using PrestoBlue^M reagent and analyzed by spectrofluorometry. Half-maximal inhibitory concentrations (IC₅₀) values were calculated using a 9-point half-log dose-response study (0.78 μ M – 200 μ M). **Table 2** shows the IC₅₀ values obtained and the theoretically calculated cLogD values.

The two new analogs exhibited superior activities than the parent compound **1** (15.07 μ M), exhibiting a direct correlation between the antiproliferative activity and their lipophilicity character. Thus, compound **12** showed an IC₅₀ of 9.99 μ M, while the value for its brominated analog (**13**) was 7.29 μ M. Therefore, even though compound **13** was found to be the most potent compound of the series on inhibition of proliferation, leading to a 2-fold potency increase over the parent compound **1**, compound **12** was the most active in inducing apoptosis, as demonstrated by cell cycle and annexin-V analysis. These effects were mostly restricted to Jurkat cells, showing a marginally superior selectivity over other cancer cell lines.

Compd.	IC ₅₀ (µM) ^[a]	cLogD ^[b]	
1	15.07	5.17	
12	9.99	6.13	
13	7.29	7.66	

3 CONCLUSION

In summary, we have prepared a library of 6-alkoxy purine analogs aiming to further explore the previously-reported pro-apoptotic activity of this chemotype over human cancer cells. Our screening campaign revealed that treatment with several of our analogs resulted in increments on the sub-G1 cell population on cell cycle, which were particularly conspicuous for compounds **12** and **13**. These effects, however, were restricted to T cells leukemia Jurkat cells, and were fully coherent with induction of apoptosis as demonstrated by annexin-V and Z-VAD-FMK assays. Additionally, this study has informed the design of subsequent analogs to further enhance the cellular activity of this chemotype.^[14] Follow-on studies will aim to identify the target of the analogs to facilitate future medicinal chemistry campaigns.

4 EXPERIMENTAL

4.1 Chemistry

4.1.1 General

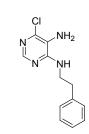
Unless otherwise noted, reagents and solvents were obtained from commercial suppliers and were used without further purification. ¹H and ¹³C NMR spectra were obtained by using a Varian Innova Unity (300 MHz), BRUKER Nanobay Avance III HD (400 MHz) or BRUKER Avance NEO (400 or 500 MHz) spectrometers, and chemical shifts are reported in parts per million (ppm, δ) downfield from tetramethylsilane (TMS). Coupling constants (*J*) are reported in Hz. Spin multiplicities are described as s (singlet), bs (broad singlet), d (doublet), t (triplet), q (quartet) and m (multiplet). High resolution mass spectra (HRMS) were obtained using a Waters LCT Premier XE Spectrometer. Analytical thin-layer chromatography (TLC) was performed using TLC Silica gel 60 F₂₅₄ plates and visualized under a UV lamp. Purifications were carried out by flash chromatography using a silica gel 220 – 440 mesh (Sigma-Aldrich). All compounds used in the biological experiments were >95% pure by ¹H/¹³C NMR. Stock solutions (100 mM) were prepared in dimethyl sulfoxide and stored at -20 °C until used.

4.1.2 General procedure for the synthesis of compounds 4 and 5.

To a stirred 0.48 M solution of 5-amino-4,6-dichloropyrimidine (2) (1 equiv.) in dry THF was added NaHCO₃ (2 equiv.). The mixture was then heated to 55 °C and at this temperature, a 1.6 M solution of the appropriate amine (2 equiv.) in THF was added dropwise. The mixture was heated at reflux for 18 h and then cooled to room temperature. The mixture was diluted with dichloromethane, filtered, the solvent removed under reduced pressure and the crude material was purified by silica gel column chromatography.



5-Amino-4-(*tert***-butylamino)-6-chloropyrimidine (4).** The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 25%) to provide the tittle compound as a white solid (851.2 mg, 4.233 mmol, 46% yield). ¹H-NMR (400 MHz, CDCl₃): δ 8.05 (1H, s, CH), 4.82 (2H, bs, NH₂), 1.92 (1H, bs, NH), 1.46 (9H, s, 3 x CH₃). ¹³C-NMR (101 MHz, CDCl₃): δ 158.11 (C), 152.49 (CH), 135.25 (C), 101.82 (C), 51.47 (C), 29.86 (3 x CH₃). HRMS (ES + ve), C₈H₁₄N₄Cl (M + H)⁺: Calculated 201.0907. Obtained 201.0893.



5-Amino-6-chloro-4-(phenethylamino)pyrimidine (5). The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 33%) to provide the tittle compound as a yellow solid (1.13 g, 4.533 mmol, 50% yield). ¹H-NMR (400 MHz, CDCl₃): δ 8.13 (1H, s, CH), 7.37 (2H, t, *J* = 7.2 Hz, 2 x CH), 7.34 – 7.24 (3H, m, 3 x CH), 5.19 (1H, bs, NH), 3.88 – 3.78 (2H, m, CH₂), 3.25 (2H, bs, NH₂), 3.01 (2H, t, *J* = 7.0 Hz, CH₂). ¹³C-NMR (101 MHz, CDCl₃): δ 155.09 (C), 149.60 (CH), 142.57 (C), 138.95 (C), 128.91 (2 x CH), 128.80 (2 x CH), 126.72 (CH), 122.04 (C), 42.78 (CH₂), 35.54 (CH₂). HRMS (ES + ve), C₁₂H₁₄N₄Cl (M + H)⁺: Calculated 249.0907. Obtained 249.0902.

4.1.3 General procedure for the synthesis of compounds 6 – 8.

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A microwave vial was charged with a 0.4 M solution of 4,6-dichloropyrimidine-2,5-diamine (3) (1 equiv.) in n-butanol, NaHCO₃ (2.5 equiv.) and the appropriate amine (5 equiv.). Then the mixture was heated at 190 °C for 2 h using microwave radiation. The mixture was diluted with dichloromethane, filtered, the solvent removed under reduced pressure and the crude material was purified by silica gel column chromatography.



2,5-Diamino-6-chloro-4-(isopropylamino)pyrimidine (6). The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 66%) to provide the tittle compound as an off-white solid (189.7 mg, 0.978 mmol, 58% yield). ¹H-NMR (400 MHz, CDCl₃): δ 5.39 (1H, s, CH), 4.79 (2H, bs, NH₂), 4.28 – 4.07 (1H, m, CH), 2.66 (2H, bs, NH₂), 1.23 (6H, d, *J* = 6.5 Hz, 2 x CH₃). ¹³C-NMR (126 MHz, CDCl₃): δ 158.45 (C), 157.37 (C), 146.61 (C), 111.73 (C), 42.92 (CH), 22.96 (2 x CH₃). HRMS (ES + ve), C₇H₁₃N₅Cl (M + H)⁺: Calculated 202.0859. Obtained 202.0849.



2,5-Diamino-4-(*tert***-butylamino)-6-chloropyrimidine (7).** The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 20 – 33%) to provide the tittle compound as a dark green solid (187 mg, 0.865 mmol, 47 yield%). ¹H-NMR (400 MHz, CDCl₃): δ 5.37 (1H, bs, NH), 4.67 (2H, bs, NH₂), 2.73 (2H, bs, NH₂), 1.41 (9H, s, 3 x CH₃). ¹³C-NMR (126 MHz, CDCl₃): δ 158.91 (C), 157.27 (C), 147.53 (C), 111.80 (C), 51.98 (C), 28.97 (3 x CH₃). HRMS (ES + ve), C₈H₁₅N₅Cl (M + H)⁺: Calculated 216.1016. Obtained 216.1009.



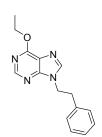
2,5-Diamino-6-chloro-4-(phenethylamino)pyrimidine (8). The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 33%) to provide the tittle compound as a yellow solid (406.8 mg, 1.54 mmol, 84% yield). ¹H-NMR (400 MHz, CDCl₃): δ 7.32 (2H, t, *J* = 7.4 Hz, 2 x CH), 7.28 – 7.14 (3H, m, 3 x CH), 5.42 (1H, bs, NH), 4.65 (2H, bs, NH₂), 3.66 (2H, q, *J* = 6.7 Hz, CH₂), 2.90 (2H, t, *J* = 6.7 Hz, CH₂), 2.65 (2H, bs, NH₂). ¹³C-NMR (101 MHz, CDCl₃): δ 159.21 (C), 157.91 (C), 148.36 (C), 139.20 (C), 128.91 (2 x CH), 128.76 (2 x CH), 126.63 (CH), 111.75 (C), 42.37 (CH₂), 35.76 (CH₂). HRMS (ES + ve), C₁₂H₁₅N₅Cl (M + H)⁺: Calculated 264.1016. Obtained 264.1022.

4.1.3 General procedure for the synthesis of compounds 1i, 9, 10, 12, 12i, 13 and 13i.

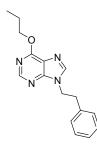
A suspension of NaH (60%) in mineral oil (10 equiv.) was dissolved in a mixture cooled at 0 °C constituted both from the appropriate alcohol (10 equiv.) and amide (10 equiv.). When *N*,*N*-dimethylbenzamide was used as the amide source dioxane was employed as a solvent. The solution was stirred at room temperature for 30 min and then at 90 °C for the same time. At 90 °C, the corresponding diaminopyrimidine (**4**, **5**) (1 equiv.) dissolved in the appropriate amide (50 equiv.) was added dropwise and the mixture heated for 18 h. The reaction mixture was quenched to pH 7 with saturated aqueous NH₄Cl, extracted with dichloromethane and washed with saturated brine. The combined organic extracts were dried over anhydrous Na₂SO₄, filtered and the solvent was removed under reduced pressure. The residue was purified by silica gel column chromatography.

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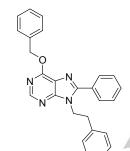
5-(Benzylideneamino)-6-(benzyloxy)-4-(*tert***-butylamino)pyrimidine (1i).** The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 10 – 33%) to provide the tittle compound as a yellow solid (15.6 mg, 0.043 mmol, 6% yield). ¹H-NMR (300 MHz, CDCl₃): δ 9.17 (1H, s, CH), 8.15 (1H, s, CH), 7.81 – 7.71 (2H, m, 2 x CH), 7.49 – 7.33 (8H, m, 8 x CH), 6.28 (1H, bs, NH), 5.50 (2H, s, OCH₂), 1.54 (9H, s, 3 x CH₃). HRMS (ES + ve), C₂₂H₂₅N₄O (M + H)⁺: Calculated 361.2028. Obtained 361.2041.



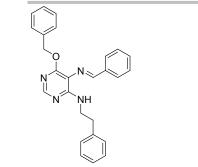
6-Ethoxy-9-phenethyl-9*H***-purine (9).** The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 33 - 100%) to provide the title compound as a yellow solid (50.4 mg, 0.187 mmol, 45% yield). ¹H-NMR (400 MHz, CDCl₃): δ 8.60 (1H, s, CH), 7.58 (1H, s, CH), 7.37 – 7.24 (3H, m, 3 x CH), 7.13 – 7.07 (2H, m, 2 x CH), 4.72 (2H, q, *J* = 7.1 Hz, OCH₂), 4.54 (2H, t, *J* = 7.0 Hz, CH₂), 3.24 (2H, t, *J* = 7.0 Hz, CH₂), 1.57 (3H, t, *J* = 7.1 Hz, CH₃). ¹³C-NMR (101 MHz, CDCl₃): δ 160.95 (C), 152.16 (CH), 152.05 (C), 142.15 (CH), 137.34 (C), 128.95 (2 x CH), 128.77 (2 x CH), 127.17 (CH), 121.57 (C), 63.21 (CH₂), 45.73 (CH₂), 36.25 (CH₂), 14.61 (CH₃). HRMS (ES + ve), C₁₅H₁₇N₄O (M + H)⁺: Calculated 269.1402. Obtained 269.1402.



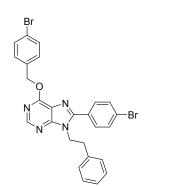
9-Phenethyl-6-propoxy-9H-purine (10). The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 50 – 66%) to provide the tittle compound as a brown solid (24.9 mg, 0.088 mmol, 19% yield). ¹H-NMR (400 MHz, CDCl₃): δ 8.54 (1H, s, CH), 7.54 (1H, s, CH), 7.31 – 7.18 (3H, m, 3 x CH), 7.08 – 7.01 (2H, m, 2 x CH), 4.56 (2H, t, *J* = 7.0 Hz, OCH₂), 4.48 (2H, t, *J* = 7.0 Hz, CH₂), 3.18 (2H, t, *J* = 7.0 Hz, CH₂), 2.00 – 1.86 (2H, m, CH₂), 1.08 (3H, t, *J* = 7.5 Hz, CH₃). ¹³C-NMR (101 MHz, CDCl₃): δ 161.18 (C), 152.23 (CH), 152.08 (C), 142.18 (CH), 137.36 (C), 129.02 (2 x CH), 128.84 (2 x CH), 127.26 (CH), 121.51 (C), 69.01 (CH₂), 45.83 (CH₂), 36.32 (CH₂), 22.36 (CH₂), 10.57 (CH₃). HRMS (ES + ve), C₁₆H₁₉N₄O (M + H)⁺: Calculated 283.1559.



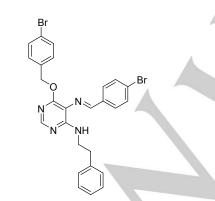
6-(Benzyloxy)-9-phenethyl-8-phenyl-9H-purine (12). The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 25%) to provide the tittle compound as a white solid (7.6 mg, 0.019 mmol, 5% yield). ¹H-NMR (400 MHz, CDCl₃): δ 8.58 (1H, s, CH), 7.60 – 7.53 (2H, m, 2 x CH), 7.52 – 7.41 (3H, m, 3 x CH), 7.39 – 7.24 (5H, m, 5 x CH), 7.21 – 7.13 (3H, m, 3 x CH), 6.96 – 6.87 (2H, m, 2 x CH), 5.71 (2H, s, OCH₂), 4.56 (2H, t, *J* = 7.4 Hz, CH₂), 3.10 (2H, t, *J* = 7.4 Hz, CH₂). ¹³C-NMR (126 MHz, CDCl₃): δ 161.23 (C), 158.80 (C), 154.40 (CH), 139.32 (C), 137.42 (C), 136.96 (C), 131.17 (CH), 129.89 (C), 129.12 (2 x CH), 128.81 (2 x CH), 128.78 (2 x CH), 128.66 (2 x CH), 128.34 (2 x CH), 128.13 (CH), 127.85 (2 x CH), 126.59 (CH), 111.10 (C), 68.25 (CH₂), 42.36 (CH₂), 36.18 (CH₂). HRMS (ES + ve), C₂₆H₂₃N₄O (M + H)⁺: Calculated 407.1872. Obtained 407.1848.



5-(Benzylideneamino)-6-(benzyloxy)-4-(phenethyl-amino)pyrimidine (12i). The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 20%) to provide the tittle compound as a brown solid (30.1 mg, 0.074 mmol, 19% yield). ¹H-NMR (500 MHz, CDCl₃): δ 9.17 (1H, s, CH), 8.19 (1H, s, CH), 7.68 – 7.61 (2H, m, 2 x CH), 7.47 – 7.40 (7H, m, 7 x CH), 7.40 – 7.35 (3H, m, 3 x CH), 7.34 – 7.27 (3H, m, 3 x CH), 6.30 (1H, bs, NH), 5.51 (2H, s, OCH₂), 3.83 (2H, q, *J* = 6.5 Hz, CH₂), 2.97 (2H, t, *J* = 6.8 Hz, CH₂). ¹³C-NMR (126 MHz, CDCl₃): δ 161.22 (CH), 158.79 (C), 154.40 (CH), 139.32 (C), 137.42 (C), 136.95 (C), 131.17 (CH), 129.89 (C), 129.12 (2 x CH), 128.81 (2 x CH), 128.77 (2 x CH), 128.66 (2 x CH), 128.34 (2 x CH), 128.12 (CH), 127.84 (2 x CH), 126.58 (CH), 111.09 (C), 68.25 (CH₂), 42.37 (CH₂), 36.18 (CH₂). HRMS (ES + ve), C₂₆H₂₅N₄O (M + H)⁺: Calculated 409.2028. Obtained 409.2039.



6-(4-Bromobenzyloxy)-8-(4-bromophenyl)-9-phenethyl-9H-purine (13). The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 25%) to provide the tittle compound as an orange solid (15.1 mg, 0.027 mmol, 7% yield). ¹H-NMR (500 MHz, CDCl₃): δ 8.56 (1H, s, CH), 7.58 – 7.54 (1H, m, CH), 7.53 – 7.41 (7H, m, 7 x CH), 7.31 – 7.27 (2H, m, 2 x CH), 7.20 – 7.14 (2H, m, 2 x CH), 6.97 – 6.89 (1H, m, 1 x CH), 5.02 (2H, s, OCH₂), 4.56 (2H, t, *J* = 6.8 Hz, 2 x CH), 3.10 (2H, t, *J* = 6.8 Hz, 2 x CH). ¹³C-NMR (126 MHz, CDCl₃): δ 159.93 (C), 158.10 (C), 151.48 (CH), 137.12 (C), 136.01 (C), 135.37 (C), 133.68 (C), 131.72 (2 x CH), 130.19 (C), 130.14 (2 x CH), 129.48 (C), 129.20 (2 x CH), 129.02 (2 x CH), 128.66 (2 x CH), 126.89 (CH), 121.86 (C), 114.93 (2 x CH), 69.28 (CH₂), 45.52 (CH₂), 35.58 (CH₂). HRMS (ES + ve), C₂₆H₂₁N₄OBr₂ (M + H)⁺: Calculated 563.0082. Obtained 563.0092.

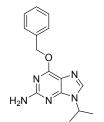


6-(4-Bromobenzyloxy)-5-[(4-bromobenzylidene)-amino]-4-(phenethylamino)pyrimidine (13i). The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 25%) to provide the tittle compound as an orange solid (9.7 mg, 0.017 mmol, 4% yield). ¹H-NMR (400 MHz, CDCl₃): δ 9.11 (1H, s, CH), 9.06 (1H, s, CH), 8.17 (1H, s, CH), 7.69 – 7.62 (1H, m, CH), 7.58 – 7.41 (5H, m, 5 x CH), 7.38 – 7.27 (6H, m, 6 x CH), 5.45 (2H, s, OCH₂), 3.83 (2H, q, *J* = 6.5 Hz, CH₂), 2.96 (2H, t, *J* = 6.5 Hz, CH₂). ¹³C-NMR (126 MHz, CDCl₃): δ 159.66 (C), 158.66 (CH), 154.57 (CH), 139.21 (C), 137.24 (C), 136.21 (C), 135.90 (C), 132.09 (2 x CH), 131.81 (2 x CH), 129.90 (C), 129.61 (2 x CH), 129.12 (2 x CH), 128.84 (2 x CH), 128.38 (2 x CH), 126.66 (CH), 122.18 (C), 110.83 (C), 67.50 (CH₂), 42.29 (CH₂), 36.09 (CH₂). HRMS (ES + ve), C₂₆H₂₃N₄OBr₂ (M + H)⁺: Calculated 565.0239. Obtained 565.0245.

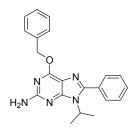
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4.1.4 General procedure for the synthesis of compounds 11, 14, 14i, 15, 15i, 16 and 16i.

A microwave vial was charged with a suspension of NaH (60%) in mineral oil (10 equiv.), *N*,*N*-dimethylformamide (40 equiv.) and the appropriate alcohol (10 equiv.). Then the corresponding triaminopyrimidine (6 - 8) (1 equiv.) dissolved in DMF (80 equiv.) was added dropwise and the mixture heated at 150 °C for 4 h using microwave radiation. When DMB was used as the amide source dioxane was employed as a solvent. The reaction mixture was quenched to pH 7 with saturated aqueous NH₄Cl, extracted with dichloromethane and washed with saturated brine. The combined organic extracts were dried over anhydrous Na₂SO₄, filtered and the solvent was removed under reduced pressure. The residue was purified by silica gel column chromatography.



2-Amino-6-(benzyloxy)-9-isopropyl-9H-purine (11). The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 50 – 100%) to provide the tittle compound as a red solid (8.5 mg, 0.03 mmol, 7% yield). ¹H-NMR (500 MHz, CDCl₃): δ 7.68 (1H, s, CH), 7.53 – 7.47 (2H, m, 2 x CH), 7.37 – 7.27 (3H, m, 3 x CH), 5.56 (2H, s, OCH₂), 4.84 (2H, bs, NH₂), 4.73 – 4.63 (1H, m, CH), 1.54 (6H, d, *J* = 6.8 Hz, 2 x CH₃). ¹³C-NMR (126 MHz, CDCl₃): δ 161.14 (C), 159.00 (C), 153.78 (C), 137.08 (CH), 136.71 (C), 128.49 (2 x CH), 128.38 (2 x CH), 128.06 (CH), 116.10, 68.10 (CH₂), 46.66 (CH), 22.75 (2 x CH₃). HRMS (ES + ve), C₁₅H₁₈N₅O (M + H)⁺: Calculated 284.1511. Obtained 284.1510.

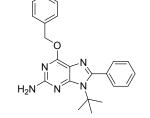


2-Amino-6-(benzyloxy)-9-isopropyl-8-phenyl-9H-purine (14). The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 33%) to provide the tittle compound as a brown solid (11 mg, 0.031 mmol, 6% yield). ¹H-NMR (400 MHz, CDCl₃): δ 7.64 – 7.56 (2H, m, 2 x CH), 7.58 – 7.42 (5H, m, 5 x CH), 7.39 – 7.31 (3H, m, 3 x CH), 5.58 (2H, s, OCH₂), 4.84 (2H, bs, NH₂), 4.68 – 4.55 (1H, m, CH), 1.64 (6H, d, *J* = 6.8 Hz, 2 x CH₃). ¹³C-NMR (126 MHz, CDCl₃): δ 160.95 (C), 158.20 (C), 150.34 (C), 136.81 (C), 131.74 (C), 129.74 (CH), 129.61 (2 x CH), 128.68 (2 x CH), 128.61 (2 x CH), 128.45 (2 x CH), 128.05 (CH), 127.13(C), 116.15 (C), 68.02 (CH₂), 49.31 (CH), 21.16 (2 x CH₃). HRMS (ES + ve), C₂₁H₂₂N₅O (M + H)⁺: Calculated 360.1824. Obtained 360.1820.

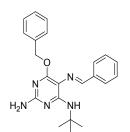
NH

2-Amino-5-(benzylideneamino)-6-(benzyloxy)-4-(isopropylamino)pyrimidine (14i). The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 33%) to provide the tittle compound as a yellow solid (29.9 mg, 0.083 mmol, 16% yield). ¹H-NMR (400 MHz, CDCl₃): δ 9.10 (1H, s, CH), 7.73 – 7.65 (1H, m, CH), 7.47 – 7.27 (9H, m, 9 x CH), 6.22 (1H, bs, NH), 5.43 (2H, s, OCH₂), 4.84 (2H, bs, NH₂), 4.32 – 4.19 (1H, m, CH), 1.27 (6H, d, *J* = 6.5 Hz, 2 x CH₃). ¹³C-NMR (126 MHz, CDCl₃): δ 160.51 (C), 155.24 (CH), 141.04 (C), 138.48 (C), 137.29 (C), 134.60 (C), 129.94 (CH), 128.68 (2 x CH), 128.58 (CH), 127.96 (CH), 127.82 (CH), 127.76 (CH), 127.55 (CH), 127.11 (2 x CH), 103.59 (C), 67.78 (CH₂), 42.56 (CH), 23.38 (2 x CH₃). HRMS (ES + ve), C₂₁H₂₄N₅O (M + H)⁺: Calculated 362.1981. Obtained 362.1978.

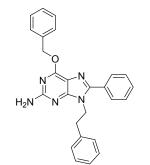




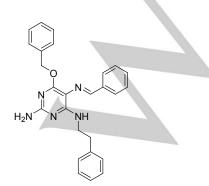
2-Amino-6-(benzyloxy)-9-(*tert***-butyl)-8-phenyl-9***H***-purine (15).** The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 20%) to provide the tittle compound as a brown solid (10.7 mg, 0.029 mmol, 7% yield). ¹H-NMR (400 MHz, CDCl₃): δ 7.58 – 7.25 (10H, m, 10 x CH), 5.54 (2H, s, OCH₂), 4.79 (2H, bs, NH₂), 1.58 (9H, s, 3 x CH₃). ¹³C-NMR (101 MHz, CDCl₃): δ 161.04 (C), 157.69 (C), 150.44 (C), 136.76 (C), 135.52 (C), 130.20 (2 x CH), 129.22 (CH), 128.73 (2 x CH), 128.42 (2 x CH), 128.04 (CH), 127.88 (2 x CH), 127.14 (C), 115.92 (C), 68.00 (CH₂), 60.10 (C), 30.88 (3 x CH₃). HRMS (ES + ve), C₂₂H₂₄N₅O (M + H)⁺: Calculated 374.1981. Obtained 374.1965.



2-Amino-5-(benzylideneamino)-6-(benzyloxy)-4-(*tert***-butylamino)pyrimidine (15i).** The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 20%) to provide the tittle compound as a yellow solid (7.3 mg, 0.019 mmol, 5% yield). ¹H-NMR (500 MHz, CDCl₃): δ 9.11 (1H, s, CH), 7.72 – 7.66 (2H, m, 2 x CH), 7.48 – 7.30 (8H, m, 8 x CH), 6.50 (1H, bs, NH), 5.45 (2H, s, OCH₂), 4.69 (2H, bs, NH₂), 1.53 (9H, s, 3 x CH₃). ¹³C-NMR (126 MHz, CDCl₃): δ 161.77 (C), 160.22 (C), 159.22 (C), 154.67 (CH), 138.60 (C), 137.49 (C), 129.79 (CH), 128.66 (2 x CH), 128.54 (2 x CH), 127.86 (CH), 127.77 (2 x CH), 127.45 (2 x CH), 103.94 (C), 67.49 (CH₂), 51.41 (C), 29.47 (3 x CH₃). HRMS (ES + ve), C₂₂H₂₆N₅O (M + H)⁺: Calculated 376.2137. Obtained 376.2152.



2-Amino-6-(benzyloxy)-9-phenethyl-8-phenyl-9H-purine (16). The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 25%) to provide the tittle compound as an orange solid (traces). ¹H-NMR (400 MHz, CDCl₃): δ 7.51 – 7.47 (3H, m, 3 x CH), 7.41 (2H, d, *J* = 7.2 Hz, 2 x CH), 7.38 – 7.32 (4H, m, 4 x CH), 7.31 – 7.27 (2H, m, 2 x CH), 7.20 – 7.17 (2H, m, *J* = 5.1, 2.0 Hz, 2 x CH), 6.97 – 6.93 (2H, m, 2 x CH), 5.58 (2H, s, OCH₂), 4.70 (2H, bs, NH₂), 4.36 (2H, t, *J* = 7.6 Hz, CH₂), 3.04 (2H, t, *J* = 7.6 Hz, CH₂). ¹³C-NMR (101 MHz, CDCl₃): δ 160.68 (C), 158.73 (C), 156.00 (C), 137.88 (C), 136.94 (C), 130.41 (C), 129.57 (CH), 129.16 (2 x CH), 128.87 (2 x CH), 128.70 (2 x CH), 128.66 (CH), 128.62 (2 x CH), 128.54 (2 x CH), 128.44 (2 x CH), 127.68 (CH), 126.81 (C), 115.22 (C), 68.00 (CH₂), 46.27 (CH₂), 45.23 (CH₂). HRMS (ES + ve), C₂₆H₂₃N₅ONa (M + Na)⁺: Calculated 444.1800. Obtained 444.1814.



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2-Amino-5-(benzylideneamino)-6-(benzyloxy)-4-(phenethylamino)pyrimidine (16i). The crude product was purified via silica gel column chromatography (Ethyl acetate/Hexane 25%) to provide the tittle compound as a yellow solid (21.1 mg, 0.05 mmol, 14% yield). ¹H-NMR (400 MHz, CDCl₃): δ 9.08 (1H, s, CH), 7.62 – 7.54 (2H, m, 2 x CH), 7.45 – 7.25 (12H, m, 12 x CH), 7.21 – 7.13 (1H, m, CH), 6.47 (1H, bs, NH), 5.44 (2H, s, OCH₂), 5.20 (2H, bs, NH₂), 3.75 (2H, q, *J* = 6.7 Hz, CH₂), 2.92 (2H, t, *J* = 6.7 Hz, CH₂). ¹³C-NMR (101 MHz, CDCl₃): δ 160.62 (C), 158.99 (C), 153.98 (CH), 140.13 (C), 139.73 (C), 138.73 (C), 137.56 (C), 129.65 (CH), 129.11 (2 x CH), 128.75 (2 x CH), 127.85 (2 x CH), 127.70 (2 x CH), 127.50 (2 x CH), 127.18 (CH), 126.43 (CH), 103.47 (C), 67.58 (CH₂), 45.85 (CH₂), 42.27 (CH₂). HRMS (ES + ve), C₂₆H₂₆N₅O (M + H)⁺: Calculated 424.2137. Obtained 424.2140.

4.2 Pharmacological/biological assays

Cell subculture. Jurkat (acute T cell leukemia) and K562 (chronic myelogenous leukemia) cell lines were grown in T75 Flasks in RPMI 1640 medium (BioWhittaker, Verviers, Belgium) with 10% fetal bovine serum (Gibco, Auckland, New Zealand), 1% L-glutamine (Gibco) and incubated at 37 °C under 5% CO₂. HeLa (cervical cancer), G361 (malignant melanoma), MDA-MB-231 (breast cancer) and HCT116 (colon cancer) cell lines were grown in T75 Flasks in Dulbecco's Modified Eagle Medium (DMEM) with 10% fetal bovine serum (Gibco, Auckland, New Zealand) and 1% L-Glutamine (Gibco) and incubated at 37 °C under 5% CO₂. Cell cultures were split at 60% – 80% confluence. Suspension cells were centrifuged (1500 rpm, 5 min). Cell pellets were resuspended in supplemented RPMI growth media, seeded at dilution factor 1:10 (T75 – 15 mL) and incubated until 60% – 80% confluence was achieved. Adherent cells were washed with PBS (T75 – 10 mL) and trypsinized (Trypsin: T75 – 1 mL, 5 min, 37 °C). Trypsin was inactivated with supplemented DMEM growth media (9 mL) and centrifuged (1500 rpm, 5 min). Cell pellets were resuspended in supplemented DMEM growth media, seeded at dilution factor 1:10, unless otherwise stated, in same media (T75 – 15 mL) and incubated until 60% – 80% confluence was achieved.

Cell cycle analysis. For the analysis of cell cycle, 250,000 cells/well (Jurkat, K562), 200,000 cells/well (G361), 150,000 cells/well (MDA-MB-231, HCT116) in 24-well plates; or 50,000 cells/well (HeLa) in 6-well plates were cultured in 500 µL (24-well) or 1.5 ml (6-well) of the appropriate media and incubated with the chosen compounds at 30 µM for 48 h. Untreated cells were incubated with the maximum DMSO concentration used. Cell cycles were analyzed by staining of cells with propidium iodide (PI) staining. Thus, after treatment with the corresponding compounds, the cells were collected and washed with 2 ml phosphate-buffered saline (PBS) at 4 °C and fixed with 100 µL of PBS and 900 µL of 70% ethanol, on ice for 5 minutes. After washing with PBS, cells were resuspended in 250 µL of PBS and another 250 µL of a DNA extraction buffer (0.2M Na₂HPO₄, 0.1M C₆H₈O₇, pH 7.8) and incubated at 37 °C for 10 minutes. The supernatant was removed and 200 µL of the staining solution [8 µL propidium iodide (1mg / mL) and 2 µL RNAse 100 (µg / mL)] were added and the samples incubated for 10 min at 37 °C in the dark. Fluorescence was measured in the FL2 detector of the FACScalibur cytometer (Becton Dickinson & Co., NJ, USA) and the analysis of the sub-G1 population (apoptotic cells) was done using the Cell Quest software (BD, Biosciences).

Annexin-V staining. For the analysis of phosphatidylserine exposure, Jurkat cells (250,000 cells/well) were cultured in 24-well plates in 500 µl of supplemented RPMI medium and incubated with the chosen compounds at the indicated doses for 48 h at 37 °C and 5% CO₂. Etoposide was used as positive control and untreated cells were used as controls of live cells. After treatment, cells were collected and washed with annexin V binding buffer (10 mM HEPES pH 7.4, 140 mM NaCl, 2.5 mM CaCl₂) and re-suspended in 100 µl of this buffer. Then, 2.5 µl of Annexin-V-FLUOS solution (Roche, Mannheim, Germany) was added to each sample, followed by incubation for 20 min in the dark at room temperature. After a final wash, the pellet was resuspended in annexin V binding buffer containing 10 µl of Pl at 10 µg/ml. Stained samples were immediately analyzed by flow cytometry using a FACSCalibur flow cytometer and analyzed with the Cell Quest software.

Z-VAD-FMK treatment. For the study of the induction of caspase-dependent cell death, Jurkat cells (200,000 cells/well) were cultured in 24-well plates in 500 µl of supplemented RPMI medium and incubated with Z-VAD-FMK (50 µM) for 1 h at 37 °C and 5% CO₂. Thereafter, cells were treated with the chosen compounds (30 µM) and incubated for 24 h at 37 °C and 5% CO₂. **6d** and DMSO were used as positive and negative controls, respectively. Cell cycles were analyzed by staining of cells with propidium iodide (PI) staining. Thus, after treatment with the corresponding compounds, the cells were collected and washed with 2 ml phosphate-buffered saline (PBS) at 4 °C and fixed with 100 µL of PBS and 900 µL of 70% ethanol, on ice for 5 minutes. After washing with PBS, cells were resuspended in 250 µL of PBS and another 250 µL of a DNA extraction buffer (0.2M Na₂HPO₄, 0.1M C₆H₈O₇, pH 7.8) and incubated at 37 °C for 10 minutes. The supernatant was removed and 200 µL of the staining solution [8 µL propidium iodide (1mg / mL) and 2 µL RNAse 100 (µg / mL)] were added and the samples incubated for 10 min at 37 °C in the dark. Fluorescence was measured in the FL2 detector of the FACScalibur cytometer (Becton Dickinson & Co., NJ, USA) and the analysis of the sub-G1 population (apoptotic cells) was done using the Cell Quest software (BD, Biosciences).

Dose-response cell viability assays.

- Hematopoietic cells (Jurkat and K562). Compounds, including DMSO, were prepared as intermediate solutions at 2x in supplemented RPMI medium. The compounds were then added in serial dilution in 96-well flat-bottomed plates to give a final volume of 50 μL/well. Then 50 μL of a previously prepared supplemented RPMI medium at 1000 cells/well (Jurkat, K562) were added to each well and the plates incubated for 5 days in an incubator at 37 °C and 5% CO₂. Final compound concentration range among 0.05 μM and 200 μM. Untreated cells were incubated with the maximum DMSO concentration used (0.2% v/v). After 5 days, PrestoBlue™ cell viability reagent (10 μl) was added to each well and the plates incubated for 90 120 min. Fluorescence emission was detected using a Synergy™ HTX Multi-Mode Microplate Reader (excitation 530/20 nm, emission 590/35 nm).
- Non-hematopoietic (G361, MDA-MB-231, HCT116, HeLa). Cells were seeded in 96-well flat-bottomed plates at 500 cells/well (G361 and HCT116 cells), 1,000 cells/well (HeLa cells) and 2,000 cells/well (MDA-MB-231 cells) in 50 μL of supplemented

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medium and incubated for 24 h in an incubator at 37 °C and 5% CO₂. After 24 h, the media was aspirated from each well and replaced with 100 µL of fresh medium. Compounds, including DMSO, were prepared as intermediate solutions at 2x in supplemented DMEM medium. Then compounds were added in serial dilution to give a final volume of 100 µL/well. Final compound concentration range among 0.05 µM and 200 µM. Untreated cells were incubated with the maximum DMSO concentration used (0.2% v/v). After 5 days, PrestoBlue™ cell viability reagent (10 µl) was added to each well and the plates incubated for 90 – 120 min. Fluorescence emission was detected using an Envision[®] fluorescence plate reader (excitation 530/20 nm, emission 590/35 nm).

All conditions were normalized to the untreated cells (100%) and curves were fitted using a four-parameter logistic fit with minimum value constrained to zero using GraphPad Prism software, to calculate half-maximal inhibitory concentration (IC_{50}) values. IC_{50} values are expressed as mean of three independent experiments.

4.3 cLogD determinations.

Calculated distribution coefficients (clogD) were determined using Chemicalize by ChemAxon Ltd.^[13] at the pH of 7.4.

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CONFLICT OF INTEREST

The authors declare no competing financial interest.

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