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26 Abstract

27 A series of picolyl amides of betulinic acid (3a-3c and 6a-6c) was prepared and subjected to the cytotoxicity screening tests. Structure-activity relationships studies resulted in finding 28 29 differences in biological activity in dependence on o_{-} , m_{-} and p_{-} substitution of the pyridine ring in the target amides, when cytotoxicity data of **3a–3c** and **6a–6c** were obtained and 30 compared. The amides **3b** and **3a** displayed cytotoxicity (given in the IC₅₀ values) in G-361 31 $(0.5 \pm 0.1 \,\mu\text{M} \text{ and } 2.4 \pm 0.0 \,\mu\text{M}, \text{ respectively}), \text{MCF7} (1.4 \pm 0.1 \,\mu\text{M} \text{ and } 2.2 \pm 0.2 \,\mu\text{M},$ 32 respectively), HeLa ($2.4 \pm 0.4 \mu$ M and $2.3 \pm 0.5 \mu$ M, respectively) and CEM ($6.5 \pm 1.5 \mu$ M 33 and $6.9 \pm 0.4 \,\mu$ M, respectively) tumor cell lines, and showed weak effect in the normal human 34 fibroblasts (BJ). Selectivity against all tested cancer cells was determined and compared to 35 normal cells with therapeutic index (TI) between 7 and 100 for compounds 3a and 3b. The 36 therapeutic index (TI = 100) was calculated for human malignant melanoma cell line (G-361) 37 38 *versus* normal human fibroblasts (BJ). The cytotoxicity of other target amides (3c and 6a–6c) revealed lower effects than 3a and 3b in the tested cancer cell lines. 39

40

41 **Key words**: betulinic acid; picolyl amine; amide; cytotoxicity; therapeutic index

42

43 **1. Introduction**

In the past years, we have published a paper dealing with the investigation of heteroaromatic amides of selected steryl hemiesters [1]. Several of those compounds were based on *o*-, *m*and *p*-picolyl amines, displaying only moderate cytotoxicity. Those compounds suffered from low polarity of the steryl skeleton. Nevertheless, even in that range of biological activity, difference in cytotoxicity values could be seen, based on *o*-, *m*- or *p*-substitution in picolyl amides. More recently, our attention has been focused at the investigation of triterpenoid acid and their derivatives, which mostly showed considerable cytotoxicity and high and sometimes

51	selective antimicrobial activity. The compounds completed a majority of ADME parameters
52	required for prediction of potential drugs, and - in several cases - displayed important
53	supramolecular characteristics, among which formation of supramolecular hydrogels was one
54	of the most important features [2,3].
55	Betulinic acid, (3β) -3-hydroxylup-20(29)-en-28-oic acid, is practically insoluble in water,
56	mainly due to the presence of large lipophilic backbone and inadequate number of hydrophilic
57	groups [4]. Even with this feature, which means, in fact, that this natural product does not
58	correspond to the Lipinski [5] and Ghose [6] rules describing potentially biologically active
59	compounds, it displays anti-HIV activity [7], antitumor [8,9] and antidiabetic activity [10,11].
60	Nature itself possesses natural mechanisms for increasing bioavailability of natural products
61	by forming their water-soluble conjugates (e.g. glycosides) [12]. To improve its
62	pharmacological characteristics in forms of the novel compounds has always been a challenge
63	for researchers, to find its new derivatives and related compounds [12]. Betulinic acid is a
64	plant product, to be found in genus Betulla, Diospiros, Paeonia, Syzigium or Ziziphus [13].
65	Concerning the antitumor effect, derivatives of betulinic acid have been used against a variety
66	of tumor cell lines: malignant brain tumor, primitive neuroectodermal tumor [14], human
67	chronic myelogenous leukemia, and against most of prevalent human cancer types, such as
68	cervical, prostate, breast, lung or colorectal cancer [12].
69	Many bioactive secondary metabolites of plants, including betulinic acid, induce apoptosis
70	pathway in cancer cells to exert their selective cytotoxic effects [15]. The cysteine-aspartic
71	acid protease (caspase) family proteins play a central role in the execution phase of apoptosis,
72	and the mechanism of initiation of apoptosis is mediated by the caspase cascade activation
73	[16]. Caspase-3 and caspase-7 are downstream factors activated by caspase-8 and caspase-9,
74	which – in turn – are activated predominantly by the extrinsic (death receptor) and intrinsic
75	(mitochondrial) pathways, respectively [17]. Caspase-3 is an executioner protease that results

in cleavage of poly-(ADP-ribose) polymerase (PARP), subsequent DNA degradation and 76 apoptotic death [18]. Previous studies showed that betulinic acid also displays anti-77 inflammatory, anti-HIV, anti-helminthic, anti-nociceptive activities [19]. Several studies were 78 done to explain the molecular mechanisms of betulinic acid mediated antitumor activity. The 79 process seems to be largely dependent on the ability of betulinic acid to trigger the 80 mitochondrial pathway of apoptosis in cancer cells. Betulinic acid shows direct effect on 81 mitochondria. Successive treatment of betulinic acid on cancer cells disrupts the 82 mitochondrial transmembrane potential, which facilitates the cells to undergo apoptosis. 83 Involvement of excessive amount of reactive oxygen species was found to be the most 84 important factor for loss of mitochondrial membrane integrity of melanoma cells treated with 85 betulinic acid. It directly targets to mitochondria, which - in turn - regulates the downstream 86 caspase activation and side by side overcomes resistance property. Thus, betulinic acid has 87 88 not developed resistance in cancer cells, and, thereby, it became permissible agents for future cancer therapy [15]. 89 90 Aminomethylpyridines (picolyl amines), and related heteroaromatic amines, including

numbers of their derivatives, and also the derived N-oxides, had already been investigated for 91 their pharmacological activity [1,20-22], as markers of solid tumors [23] or agents in 92 supramolecular self-assembly [24]. Lipophilic derivation of picolyl amines have resulted in 93 obtaining conjugates, bearing ester and amide bond in each molecule; that type of study was 94 made in our previous work [1]. The importance of such compounds consists in enabling 95 transportation of potentially biologically active compounds through biomembrane and they 96 often form cationic immune-stimulating complexes [25,26]. The derivatives of picolyl amines 97 have also been found to be involved in activation of the caspase cascade and promote cell 98 death [27]. 99

The objectives of the current research consist in (a) a synthesis of a series of picolyl amides of
betulinic acid both, at C(17)-COOH and at C(3)-OH (in the latter subseries using succinic
acid as a junction unit), and (b) investigation of cytotoxicity of the target compounds. A
comparison of the current series of compounds with those published earlier [1] will also be
discussed.

105

106 **2. Experimental part**

107 2.1. General

The ¹H NMR and the ¹³C NMR spectra were recorded on a Bruker AVANCE 600 MHz 108 spectrometer at 600.13 MHz and 150.90 MHz in deuterochloroform, using tetramethylsilane 109 $(\delta = 0.0)$ as internal reference. ¹H NMR data are presented in the following order: chemical 110 shift (δ) expressed in ppm, multiplicity (s, singlet; d, doublet; t, triplet; q, quartet; m, 111 112 multiplet), coupling constants in Hertz, number of protons. For unambiguous assignment of both ¹H and ¹³C signals 2D NMR ¹H, ¹³C gHSQC and gHMBC spectra were measured using 113 114 standard parameters sets and pulse programs delivered by producer of the spectrometer. 115 Copies of the NMR spectra are presented in the Supplementary material. Infrared spectra (IR) were measured with a Nicolet 205 FT-IR spectrometer. Mass spectra (MS) were measured 116 with a Waters ZMD mass spectrometer in a positive ESI mode. Optical rotation was measured 117 on an Autopol IV instrument (Rudolph Research Analytical, USA) at 589 nm wavelength, and 118 the value was corrected to 20°C. The PE 2400 Series II CHNS/O Analyzer (Perkin Elmer, 119 USA) was used for simultaneous determination of C, H, and N (accuracy of CHN 120 determination better than 0.30 % abs.). TLC was carried out on silica gel plates (Merck 121 60F₂₅₄) and the visualization was performed by the UV detection and by spraying with the 122 methanolic solution of phosphomolybdic acid (5%) followed by heating. For column 123 chromatography, silica gel 60 (0.063-0.200 mm) from Merck was used. All chemicals and 124

solvents were purchased from regular commercial sources in analytical grade and the solventswere purified by general methods before use.

127

128 2.2. (3β)-3-[(3-Carboxypropanoyl)oxy]lup-20(29)-en-28-oic acid (2)

129 Succinic anhydride (1.62 g; 16.2 mmol; 3.5 eq) and DMAP (170 mg; 1.39 mmol, 0.30 eq)

130 were added to a solution of betulinic acid (1; 2.1 g; 4.6 mmol) in dry pyridine (20 mL). The

131 reaction mixture was stirred at r.t. for 5 days. After stopping the reaction, the resulting

132 mixture was poured onto ice, hydrochloric acid was added to adjust pH = 7, the organic layer

133 was extracted with chloroform, and the extract was dried over sodium sulfate. Evaporation of

the solvent afforded a solid that was purified by column chromatography, yielding 1.93 g

135 (75%) of **2**.

¹H NMR: δ 0.79 (3H, s, H23), 0.79 (3H, s, H24), 0.80 (3H, s, H25), 0.87 (3H, s, H26), 0.94

137 (3H, s, H27), 1.64 (3H, dd, $J_1 = 0.7$ Hz, $J_2 = 1.4$ Hz, H29), 2.09-2.13 (2H, m, H16), 2.22 (1H,

138 ddd, $J_1 = 3.6$ Hz, $J_2 = 11.6$ Hz, $J_3 = 12.9$ Hz, H13), 2.44-2.52 (4H, m, H2'-H3'), 2.95 (1H, dt, J_1

139 = 4.8 Hz, J_2 = 11.0 Hz, J_3 = 11.0 Hz, H19), 4.38 (1H, dd, J_1 = 4.7 Hz, J_2 = 11.6 Hz, H3), 4.56

140 (1H, dq, $J_1 = 1.4$ Hz, $J_2 = 1.4$ Hz, $J_3 = 1.4$ Hz, $J_4 = 2.4$ Hz, H30), 4.69 (1H, dq, $J_1 = 0.7$ Hz, $J_2 = 0.7$

141 0.7 Hz, $J_3 = 0.7$ Hz, $J_4 = 2.4$ Hz, H30). ¹³C NMR: δ 14.31 (q, C27), 15.65 (q, C24), 15.81 (q,

142 C25), 16.35 (q, C26), 17.67 (t, C6), 18.90 (q, C29), 20.42 (t, C11), 23.27 (t, C2), 25.00 (t,

143 C12), 27.56 (q, C23), 28.77 (t, C2'), 29.15 (t, C21), 29.15 (t, C3'), 30.06 (t, C15), 31.64 (t,

144 C16), 33.68 (t, C22), 36.28 (t, C1), 36.60 (s, C10), 37.38 (s, C4), 37.54 (d, C13), 37.67 (t,

145 C7), 40.21 (s, C8), 41.99 (s, C14), 46.57 (d, C19), 48.50 (d, C18), 49.61 (d, C9), 54.62 (d,

146 C5), 55.36 (s, C17), 79.98 (d, C3), 109.58 (t, C30), 150.26 (s, C20), 171.56 (s, C1'), 173.32 (s,

147 C4'), 177.17 (s, C28). MS: *m/z* 555.2 [M-H]⁻. IR (KBr; cm⁻¹): 2946 (C–H val.), 2872 (–CH₃

148 val.), 1734 (–C=O val.). For $C_{34}H_{52}O_6$ (556.77) calculated C (73.34), H (9.41), found C

^{149 (73.37),} H (9.40). M.p. 244-246 °C.

151 2.3. (3β)-3-({4-Oxo-4-[(pyridin-n-ylmethyl)amino]butanoyl}oxy)lup-20(29)-en-28-oic acid
152 [3a (n=2), 3b (n=3) and 3c (n=4)]
153 *o*-, *m*- or *p*-Picolyl amine (20 μl; 0.198 mmol; 1.1 eq) and T3P (0.32 ml; 0.54 mmol; 3 eq)
154 were added to a solution of 2 (100 mg; 0.18 mmol) in dry pyridine (2 mL), and the reaction

mixture was stirred at r.t. for 1 day. After stopping the reaction, saturated solution of sodium

bicarbonate (4 mL) was added to reaction mixture, and stirring continued for several more

157 hours. The resulting mixture was extracted with chloroform, and dried over sodium sulfate.

158 Evaporation of the solvent afforded a solid, which was purified by column chromatography.

- 159 Yields: **3a** (85 %), **3b** (91 %), **3c** (89 %).
- **3a**: ¹H NMR: δ 0.77 (3H, s, H23), 0.78 (3H, s, H24), 0.80 (3H, s, H25), 0.87 (3H, s, H26),

161 0.95 (3H, s, H27), 1.65 (3H, dd, $J_1 = 0.7$ Hz, $J_2 = 1.3$ Hz, H29), 2.09-2.14 (2H, m, H16), 2.23

162 (1H, ddd, $J_1 = 4.7$ Hz, $J_2 = 11.7$ Hz, $J_3 = 12.9$ Hz, H13), 2.44-2.55 (4H, m, H2'-H3'), 2.95 (1H,

- 163 dt, $J_1 = 5.3$ Hz, $J_2 = 11.0$ Hz, $J_3 = 11.0$ Hz, H19), 4.31 (1H, dd, $J_1 = 6.0$ Hz, $J_2 = 15.9$ Hz, H5'),
- 164 4.35 (1H, dd, $J_1 = 6.0$ Hz, $J_2 = 15.9$ Hz, H5'), 4.38 (1H, dd, $J_1 = 4.8$ Hz, $J_2 = 11.6$ Hz, H3),
- 165 4.56 (1H, dq, $J_1 = 1.3$ Hz, $J_2 = 1.3$ Hz, $J_3 = 1.3$ Hz, $J_4 = 2.4$ Hz, H30), 4.69 (1H, bd, J = 2.4 Hz,

166 H30), 7.24 (1H, dddt, $J_1 = 0.6$ Hz, $J_2 = 0.6$ Hz, $J_3 = 1.2$ Hz, $J_4 = 4.8$ Hz, $J_5 = 7.5$ Hz, H8'), 7.26

167 (1H, dtt, $J_1 = 0.6$ Hz, $J_2 = 0.6$ Hz, $J_3 = 1.1$ Hz, $J_4 = 1.1$ Hz, $J_5 = 7.8$ Hz, H10'), 7.72 (1H, dt, J_1

168 = 1.8 Hz, $J_2 = 7.7$ Hz, $J_3 = 7.7$ Hz, H9'), 8.45 (1H, t, J = 6.0 Hz, NH), 8.48 (1H, ddd, $J_1 = 0.9$

169 Hz, $J_2 = 1.8$ Hz, $J_3 = 4.8$ Hz, H7'). ¹³C NMR: δ 14.32 (q, C27), 15.66 (q, C24), 15.82 (q, C25),

170 16.36 (q, C26), 17.67 (t, C6), 18.90 (q, C29), 20.43 (t, C11), 23.29 (t, C2), 25.01 (t, C12),

171 27.58 (q, C23), 29.16 (t, C21), 29.35 (t, C2'), 29.87 (t, C3'), 30.06 (t, C15), 31.66 (t, C16),

- 172 33.70 (t, C22), 36.29 (t, C1), 36.59 (s, C10), 37.38 (s, C4), 37.54 (d, C13), 37.69 (t, C7),
- 173 40.22 (s, C8), 41.99 (s, C14), 44.17 (t, C5'), 46.58 (d, C19), 48.50 (d, C18), 49.62 (d, C9),
- 174 54.63 (d, C5), 55.37 (s, C17), 79.87 (d, C3), 109.58 (t, C30), 120.88 (d, C10'), 121.97 (d, C8'),

136.56 (d, C9'), 148.71 (d, C7'), 150.28 (s, C20), 158.64 (s, C6'), 170.88 (s, C1'), 171.88 (s, 175 C4'), 177.18 (s, C28). MS: *m/z* 647.3 [M+H]⁺, 645.0 [M-H]⁻. IR (KBr; cm⁻¹): 3377 (N–H val. 176 amide), 2945 (C-H val.), 2871 (-CH₃ val.), 1728 (-C=O val. ester), 1659 (-C=O val. amide). 177 For C₄₀H₅₈N₂O₅ (646.90) calculated C (74.27), H (9.04), N (4.33), found C (74.29), H (9.03), 178 N (4.35). M.p. 121-124 °C. $[\alpha]_D^{20} = +15.5$ (*c* 0.323). 179 **3b**: ¹H NMR: δ 0.70-0.73 (1H, m, H5), 0.76 (3H, s, H24), 0.76 (3H, s, H23), 0.78 (3H, d, J = 180 0.6 Hz, H25), 0.88 (3H, s, H26), 0.92 (3H, d, J = 0.7 Hz, H27), 1.63 (3H, dd, $J_1 = 0.7$ Hz, $J_2 =$ 181 1.3 Hz, H29), 2.17 (1H, ddd, $J_1 = 3.7$ Hz, $J_2 = 11.6$ Hz, $J_3 = 12.9$ Hz, H13), 2.22 (2H, ddd, $J_1 =$ 182 2.9 Hz, *J*₂ = 4.0 Hz, *J*₃ = 13.1 Hz, H16), 2.46 (2H, t, *J* = 6.6 Hz, H3'), 2.58-2.66 (2H, m, H2'), 183 2.94 (1H, dt, $J_1 = 5.0$ Hz, $J_2 = 11.0$ Hz, $J_3 = 11.0$ Hz, H19), 4.39 (1H, dd, $J_1 = 5.9$ Hz, $J_2 = 14.9$ 184 Hz, H5'), 4.41 (1H, dd, *J*₁ = 5.7 Hz, *J*₂ = 14.9 Hz, H5'), 4.42 (1H, dd, *J*₁ = 6.2 Hz, *J*₂ = 10.1 Hz, 185 H3), 4.55 (1H, dq, $J_1 = 1.3$ Hz, $J_2 = 1.3$ Hz, $J_3 = 1.3$ Hz, $J_4 = 2.4$ Hz, H30), 4.68 (1H, dq, $J_1 = 1.3$ Hz, $J_2 = 1.3$ Hz, $J_3 = 1.3$ Hz, $J_4 = 2.4$ Hz, H30), 4.68 (1H, dq, $J_1 = 1.3$ Hz, $J_2 = 1.3$ Hz, $J_3 = 1.3$ Hz, $J_4 = 2.4$ Hz, H30), 4.68 (1H, dq, $J_1 = 1.3$ Hz, $J_2 = 1.3$ Hz, $J_3 = 1.3$ Hz, $J_4 = 2.4$ Hz, H30), 4.68 (1H, dq, $J_1 = 1.3$ Hz, $J_2 = 1.3$ Hz, $J_3 = 1.3$ Hz, $J_4 = 2.4$ Hz, H30), 4.68 (1H, dq, $J_1 = 1.3$ Hz, $J_2 = 1.3$ Hz, $J_3 = 1.3$ Hz, $J_4 = 2.4$ Hz, H30), 4.68 (1H, dq, J_1 = 1.3 Hz, $J_4 = 1.3$ Hz, $J_4 = 2.4$ Hz, H30), 4.68 (1H, dq, J_1 = 1.3 Hz, $J_4 = 1.3$ Hz, 186 187 $0.7 \text{ Hz}, J_2 = 0.7 \text{ Hz}, J_3 = 0.7 \text{ Hz}, J_4 = 2.3 \text{ Hz}, \text{H30}$, 6.20 (1H, bt, J = 5.8 Hz, NH), 7.23 (1H, ddd, $J_1 = 0.8$ Hz, $J_2 = 4.8$ Hz, $J_3 = 7.8$ Hz, H10'), 7.63 (1H, ddt, $J_1 = 0.7$ Hz, $J_2 = 0.7$ Hz, $J_3 = 0.7$ Hz, $J_3 = 0.7$ Hz, $J_2 = 0.7$ Hz, $J_3 = 0.7$ Hz, $J_4 = 0.7$ Hz, $J_5 = 0.7$ Hz, J188 189 1.8 Hz, $J_4 = 2.1$ Hz, $J_5 = 7.8$ Hz, H11'), 8.48 (1H, bdd, $J_1 = 1.8$ Hz, $J_2 = 4.8$ Hz, H9'), 8.50 (1H, bs, H7'). ¹³C NMR: δ 14.71 (q, C27), 16.13 (q, C24), 16.17 (q, C25), 16.51 (q, C26), 18.24 (t, 190 C6), 19.41 (q, C29), 20.99 (t, C11), 23.77 (t, C2), 25.60 (t, C12), 28.02 (q, C23), 29.78 (t, 191 C21), 29.78 (t, C2'), 30.72 (t, C3'), 31.25 (t, C15), 32.29 (t, C16), 34.38 (t, C22), 37.09 (t, C1), 192 193 37.23 (s, C10), 37.95 (s, C4), 38.49 (d, C13), 38.49 (t, C7), 40.85 (t, C5'), 41.09 (s, C8), 42.56 (s, C14), 47.01 (d, C19), 49.46 (d, C18), 50.57 (d, C9), 55.56 (d, C5), 56.38 (s, C17), 81.68 194 (d, C3), 109.64 (t, C30), 123.82 (d, C10'), 134.65 (s, C6'), 136.46 (d, C11'), 147.92 (d, C9'), 195 148.35 (d, C7'), 150.45 (s, C20), 171.74 (s, C1'), 172.73 (s, C4'), 179.99 (s, C28). MS: m/z 196 647.3 [M+H]⁺, 645.3 [M-H]⁻. IR (KBr; cm⁻¹): 3350 (N–H val. amide), 2946 (C–H val.), 2871 197 198 (-CH₃ val.), 1729 (-C=O val. ester), 1657 (-C=O val. amide). For C₄₀H₅₈N₂O₅ (646.90)

199	calculated C (74.27), H (9.04), N (4.33), found C (74.30), H (9.02), N (4.35). M.p. 131-134
200	^o C. $[\alpha]_D^{20} = +12.8 \ (c \ 0.383).$
201	3c : ¹ H NMR: δ 0.70-0.74 (1H, m, H5), 0.76 (3H, s, H23), 0.76 (3H, s, H24), 0.78 (3H, s,
202	H25), 0.87 (3H, s, H26), 0.91 (3H, bd, H27), 0.98 (2H, dq, $J_1 = 4.0$ Hz, $J_2 = 13.0$ Hz, $J_3 = 13.0$
203	Hz, $J_4 = 13.0$ Hz, H12), 1.62 (3H, dd, $J_1 = 0.7$ Hz, $J_2 = 1.3$ Hz, H29), 2.16 (1H, ddd, $J_1 = 3.6$
204	Hz, $J_2 = 11.5$ Hz, $J_3 = 12.8$ Hz, H13), 2.21 (2H, ddd, $J_1 = 3.2$ Hz, $J_2 = 3.8$ Hz, $J_3 = 13.0$ Hz,
205	H16), 2.48 (1H, dt, $J_1 = 6.5$ Hz, $J_2 = 6.5$ Hz, $J_3 = 15.0$ Hz, H3'), 2.50 (1H, dt, $J_1 = 5.7$ Hz, $J_2 = 6.5$ Hz, $J_3 = 15.0$ Hz, H3'), 2.50 (1H, dt, $J_1 = 5.7$ Hz, $J_2 = 6.5$ Hz, $J_3 = 15.0$ Hz, H3'), 2.50 (1H, dt, $J_1 = 5.7$ Hz, $J_2 = 6.5$ Hz, $J_3 = 15.0$ Hz, H3'), 2.50 (1H, dt, $J_1 = 5.7$ Hz, $J_2 = 6.5$ Hz, $J_3 = 15.0$ Hz, H3'), 2.50 (1H, dt, $J_1 = 5.7$ Hz, $J_2 = 6.5$ Hz, $J_3 = 15.0$ Hz, H3'), 2.50 (1H, dt, $J_1 = 5.7$ Hz, $J_2 = 6.5$ Hz, $J_3 = 15.0$ Hz, H3'), 2.50 (1H, dt, $J_1 = 5.7$ Hz, $J_2 = 6.5$ Hz, $J_3 = 15.0$ Hz, $J_3 = 15.0$ Hz, $J_4 = 5.7$ Hz, $J_5 = 5.7$ Hz, J_5
206	5.7 Hz, $J_3 = 15.0$ Hz, H3'), 2.62 (1H, dt, $J_1 = 5.7$ Hz, $J_2 = 5.7$ Hz, $J_3 = 17.3$ Hz, H2'), 2.66 (1H,
207	dt, J_1 = 6.6 Hz, J_2 = 6.6 Hz, J_3 = 17.3 Hz, H2'), 2.95 (1H, dt, J_1 = 5.0 Hz, J_2 = 10.9 Hz, J_3 =
208	10.9 Hz, H19), 4.38 (1H, bdd, $J_1 = 6.1$ Hz, $J_2 = 16.0$ Hz, H5'), 4.42 (1H, bdd, $J_1 = 6.1$ Hz, $J_2 = 6.$
209	16.0 Hz, H5'), 4.43 (1H, dd, $J_1 = 5.5$ Hz, $J_2 = 11.0$ Hz, H3), 4.54 (1H, dq, $J_1 = 1.3$ Hz, $J_2 = 1.3$
210	Hz, $J_3 = 1.3$ Hz, $J_4 = 2.3$ Hz, H30), 4.67 (1H, dq, $J_1 = 0.7$ Hz, $J_2 = 0.7$ Hz, $J_3 = 0.7$ Hz, $J_4 = 2.3$
211	Hz, H30), 6.30 (1H, t, <i>J</i> = 6.1 Hz, NH), 7.16-7.18 (2H, m, H7'), 8.47-8.49 (2H, m, H8'). ¹³ C
212	NMR: δ 14.68 (q, C27), 16.06 (q, C24), 16.15 (q, C25), 16.49 (q, C26), 18.18 (t, C6), 19.37
213	(q, C29), 20.92 (t, C11), 23.75 (t, C2), 25.50 (t, C12), 28.01 (q, C23), 29.73 (t, C21), 29.94 (t,
214	C2'), 30.64 (t, C3'), 31.16 (t, C15), 32.26 (t, C16), 34.28 (t, C22), 37.08 (t, C1), 37.16 (s,
215	C10), 37.90 (s, C4), 38.38 (t, C7), 38.40 (d, C13), 40.75 (s, C8), 42.42 (s, C14), 42.48 (t, C5'),
216	46.97 (d, C19), 49.33 (d, C18), 50.47 (d, C9), 55.48 (d, C5), 56.34 (s, C17), 81.73 (d, C3),
217	109.63 (t, C30), 122.47 (d, C7'), 148.49 (s, C6'), 149.08 (d, C8'), 150.49 (s, C20), 171.98 (s,
218	C1'), 172.91 (s, C4'), 180.46 (s, C28). MS: <i>m/z</i> 647.2 [M+H] ⁺ , 645.3 [M-H] ⁻ . IR (KBr; cm ⁻¹):
219	3358 (N-H val. amide), 2946 (C-H val.), 2870 (-CH ₃ val.), 1729 (-C=O val. ester), 1663 (-
220	C=O val. amide). For $C_{40}H_{58}N_2O_5$ (646.90) calculated C (74.27), H (9.04), N (4.33), found C
221	(74.26), H (9.05), N (4.31). M.p. 145-147 °C. $[\alpha]_D^{20} = +12.8 \ (c \ 0.188).$
222	

223 2.4. (3β)-3-(Acetyloxy)lup-20(29)-en-28-oic acid (**4**)

224	Acetic anhydride (0.45 mL; 4.7 mmol; 1.44 eq), DMAP (52 mg; 0.43 mmol, 0.13 eq) and
225	EDIPA (1 mL) were added to a solution of betulinic acid (1; 1.5 g; 3.3 mmol) in dry THF (10
226	mL). The reaction mixture was refluxed for 3 h. After stopping the reaction, water (10 mL)
227	was added and the resulting mixture was stirred for 1 hour. The resulting mixture was
228	extracted with chloroform, and the extract was dried over sodium sulfate. Evaporation of the
229	solvent afforded a solid, which was purified by column chromatography, resulting finally in
230	1.55 g (95 %) of 4 .
231	¹ H NMR: δ 0.83 (3H, s, H23), 0.84 (3H, s, H25), 0.85 (3H, bs, H24), 0.93 (3H, s, H26), 0.97
232	(3H, bs, H27), 1.05 (2H, dq, J_1 = 4.9 Hz, J_2 = 13.0 Hz, J_3 = 13.0 Hz, J_4 = 13.0 Hz, H12), 1.19
233	(2H, dt, $J_1 = 3.2$ Hz, $J_2 = 3.2$ Hz, $J_3 = 13.6$ Hz, H21), 1.70 (3H, dd, $J_1 = 0.7$ Hz, $J_2 = 1.4$ Hz,
234	H29), 2.04 (3H, s, OAc), 2.13 (2H, ddd, J_1 = 3.7 Hz, J_2 = 11.6 Hz, J_3 = 12.8 Hz, H1), 2.27
235	(2H, dt, $J_1 = 3.8$ Hz, $J_2 = 3.8$ Hz, $J_3 = 13.1$ Hz, H16), 3.01 (1H, dt, $J_1 = 5.0$ Hz, $J_2 = 10.9$ Hz, $J_3 = 13.1$ Hz, H16), 3.01 (1H, dt, $J_1 = 5.0$ Hz, $J_2 = 10.9$ Hz, $J_3 = 13.1$ Hz, H16), 3.01 (1H, dt, $J_1 = 5.0$ Hz, $J_2 = 10.9$ Hz, $J_3 = 13.1$ Hz, H16), 3.01 (1H, dt, $J_1 = 5.0$ Hz, $J_2 = 10.9$ Hz, $J_3 = 13.1$ Hz, H16), 3.01 (1H, dt, $J_1 = 5.0$ Hz, $J_2 = 10.9$ Hz, $J_3 = 13.1$ Hz, H16), 3.01 (1H, dt, $J_1 = 5.0$ Hz, $J_2 = 10.9$ Hz, $J_3 = 10.1$ Hz, $J_3 = 10.1$ Hz, $J_4 = 10.1$ Hz, $J_5 = 10.1$ Hz, $J_$
236	= 10.9 Hz, H19), 4.47 (1H, dd, J_1 = 5.5 Hz, J_2 = 10.9 Hz, H3), 4.61 (1H, dq, J_1 = 1.4 Hz, J_2 =
237	1.4 Hz, $J_3 = 1.4$ Hz, $J_4 = 2.3$ Hz, H30), 4.74 (1H, dq, $J_1 = 0.7$ Hz, $J_2 = 0.7$ Hz, $J_3 = 0.7$ Hz, $J_4 = 0.7$ Hz, $J_5 = 0.7$ Hz, $J_7 = 0.7$ Hz, $J_8 = 0.7$ Hz,
238	2.3 Hz, H30). ¹³ C NMR: δ 14.64 (q, C27), 16.02 (q, C24), 16.17 (q, C25), 16.46 (q, C26),
239	18.15 (t, C6), 19.32 (q, OAc), 19.33 (q, C29), 20.84 (t, C11), 23.68 (t, C2), 25.43 (t, C12),
240	27.94 (q, C23), 29.68 (t, C21), 30.54 (t, C15), 32.13 (t, C16), 34.22 (t, C22), 37.03 (d, C13),
241	37.11 (s, C10), 37.79 (t, C7), 38.37 (s, C4), 38.39 (t, C1), 40.68 (s, C8), 42.41 (s, C14), 46.92
242	(d, C19), 49.25 (d, C18), 50.38 (d, C9), 55.40 (d, C5), 56.35 (s, C17), 80.94 (d, C3), 109.74 (t,
243	C30), 150.36 (s, C20), 171.06 (s, OAc), 181.47 (s, C28). MS: <i>m/z</i> 497.1 [M-H] ⁻ . IR (KBr; cm ⁻
244	¹): 2945 (C–H val.), 2871 (–CH ₃ val.), 1735 (–C=O val. ester), 1694 (–C=O val. conjugated
245	acid). For C ₃₂ H ₅₀ O ₄ (498.74) calculated C (77.06), H (10.10), found C (77.09), H (10.08).
246	M.p. 263-266 °C.

2.5. (3β)-28-Oxo-28-[(pyridin-n-ylmethyl)amino]lup-20(29)-en-3-yl acetate [5a (n=2), 5b 248 249 (n=3) and **5c** (n=4)] A solution of oxalyl chloride in dry dichloromethane (2 M; 1.4 mL; 2.8 mmol; 7 eq) was 250 added to a solution of 4 (200 mg; 0.4 mmol) in dry dichloromethane (4 mL), the reaction 251 mixture was stirred for 3 h, then evaporated and dissolved again in dry dichloromethane (4 252 mL). EDIPA (0.182 mL; 1.04 mmol; 2.6 eq) and o-, m- or p-picolyl amine (0.049 mL; 0.48 253 mmol; 1.2 eq) were added, and the reaction mixture was stirred at r.t. through night (for 12 h). 254 255 After stopping the reaction, evaporation of the solvent afforded a solid, which was purified by column chromatography. Yields: **5a** (86 %), **5b** (99 %), **5c** (87 %). 256 **5a**: ¹H NMR: δ 0.72 (1H, dd, $J_1 = 2.1$ Hz, $J_2 = 11.7$ Hz, H5), 0.75 (3H, s, H24), 0.77 (3H, s, 257 H25), 0.77 (3H, s, H23), 0.78 (3H, s, H26), 0.90 (3H, d, *J* = 0.6 Hz, H27), 1.08 (2H, dt, *J*₁ = 258 $3.2 \text{ Hz}, J_2 = 3.2 \text{ Hz}, J_3 = 13.4 \text{ Hz}, \text{H21}, 1.64 (3\text{H}, \text{dd}, J_1 = 0.7 \text{ Hz}, J_2 = 1.4 \text{ Hz}, \text{H29}), 1.99$ 259 260 $(3H, s, H2'), 2.04 (2H, dt, J_1 = 3.3 Hz, J_2 = 3.3 Hz, J_3 = 13.6 Hz, H16), 2.38 (1H, ddd, J_1 = 3.7 Hz)$ Hz, $J_2 = 11.5$ Hz, $J_3 = 12.9$ Hz, H13), 3.10 (1H, dt, $J_1 = 4.6$ Hz, $J_2 = 11.1$ Hz, $J_3 = 11.1$ Hz, 261 262 H19), 4.41 (1H, dd, $J_1 = 5.5$ Hz, $J_2 = 10.0$ Hz, H3), 4.46 (1H, dd, $J_1 = 5.2$ Hz, $J_2 = 15.9$ Hz, H3'), 4.52 (1H, dd, $J_1 = 5.2$ Hz, $J_2 = 15.9$ Hz, H3'), 4.54 (1H, dq, $J_1 = 1.4$ Hz, $J_2 = 1.4$ Hz, $J_3 = 1.4$ Hz, $J_3 = 1.4$ Hz, $J_2 = 1.4$ Hz, $J_3 = 1.4$ Hz, $J_4 = 1.4$ Hz, $J_5 = 1.4$ Hz, $J_7 = 1.4$ Hz, J_7 263 1.4 Hz, $J_4 = 2.4$ Hz, H30), 4.69 (1H, dq, $J_1 = 0.7$ Hz, $J_2 = 0.7$ Hz, $J_3 = 0.7$ Hz, $J_4 = 2.4$ Hz, 264 H30), 6.85 (1H, t, J = 5.2 Hz, NH), 7.17 (1H, m, $J_1 = 0.6$ Hz, $J_2 = 0.6$ Hz, $J_3 = 1.2$ Hz, $J_4 = 4.9$ 265 266 Hz, $J_5 = 7.5$ Hz, H7'), 7.29 (1H, m, $J_1 = 0.6$ Hz, $J_2 = 0.6$ Hz, $J_3 = 1.2$ Hz, $J_4 = 7.8$ Hz, H9'), 7.64 (1H, dt, $J_1 = 1.8$ Hz, $J_2 = 7.6$ Hz, $J_3 = 7.6$ Hz, H8'), 8.49 (1H, ddd, $J_1 = 0.8$ Hz, $J_2 = 1.8$ 267 Hz, $J_3 = 4.9$ Hz, H6'). ¹³C NMR: δ 14.58 (q, C27), 15.98 (q, C24), 16.17 (q, C25), 16.45 (q, 268 C26), 18.15 (t, C6), 19.42 (q, C29), 20.91 (t, C11), 21.31 (q, C2'), 23.67 (t, C2), 25.56 (t, 269 C12), 27.91 (q, C23), 29.41 (t, C21), 30.86 (t, C15), 33.65 (t, C16), 34.26 (t, C22), 37.08 (s, 270 C10), 37.71 (d, C13), 37.76 (s, C4), 38.33 (t, C1), 38.35 (t, C7), 40.70 (s, C8), 42.45 (s, C14), 271 44.17 (t, C3'), 46.78 (d, C19), 50.06 (d, C18), 50.49 (d, C9), 55.40 (d, C5), 55.76 (s, C17), 272

- 273 80.93 (d, C3), 109.35 (t, C30), 122.34 (d, C7'), 122.64 (d, C9'), 137.12 (d, C8'), 148.50 (d,
- 274 C6'), 150.98 (s, C20), 157.08 (s, C4'), 171.01 (s, C1'), 176.39 (s, C28). MS: *m/z* 589.2
- 275 $[M+H]^+$, 587.2 $[M-H]^-$. IR (KBr; cm⁻¹): 2946 (C–H val.), 2870 (–CH₃ val.), 1734 (–C=O val.)
- ester), 1651 (-C=O val. amide). For C₃₈H₅₆N₂O₃ (588.86) calculated C (77.51), H (9.59), N
- 277 (4.76), found C (77.53), H (9.58), N (4.75). M.p. 110-115 °C.
- **5b**: ¹H NMR: δ 0.72 (1H, dd, $J_1 = 2.1$ Hz, $J_2 = 11.0$ Hz, H5), 0.77 (3H, s, H24), 0.78 (3H, d, J) 278 = 0.7 Hz, H25), 0.78 (3H, s, H23), 0.79 (3H, s, H26), 0.90 (3H, s, H27), 1.07 (2H, dt, J₁ = 3.3 279 Hz, $J_2 = 3.3$ Hz, $J_3 = 13.5$ Hz, H21), 1.64 (3H, dd, $J_1 = 0.7$ Hz, $J_2 = 1.4$ Hz, H29), 1.89 (2H, dt, 280 *J*₁ = 3.4 Hz, *J*₂ = 3.4 Hz, *J*₃ = 13.7 Hz, H16), 1.99 (3H, s, H2'), 2.39 (1H, ddd, *J*₁ = 3.7 Hz, *J*₂ = 281 11.5 Hz, $J_3 = 12.9$ Hz, H13), 3.09 (1H, dt, $J_1 = 4.5$ Hz, $J_2 = 11.2$ Hz, $J_3 = 11.2$ Hz, H19), 4.31 282 $(1H, dd, J_1 = 5.9 Hz, J_2 = 14.9 Hz, H3'), 4.41 (1H, dd, J_1 = 5.3 Hz, J_2 = 9.6 Hz, H3), 4.48 (1H, J_1 = 5.9 Hz, J_2 = 14.9 Hz, H3')$ 283 dd, $J_1 = 6.1$ Hz, $J_2 = 14.9$ Hz, H3'), 4.55 (1H, dq, $J_1 = 1.4$ Hz, $J_2 = 1.4$ Hz, $J_3 = 1.4$ Hz, $J_4 = 2.4$ 284 285 Hz, H30), 4.69 (1H, dq, $J_1 = 0.7$ Hz, $J_2 = 0.7$ Hz, $J_3 = 0.7$ Hz, $J_4 = 2.4$ Hz, H30), 6.08 (1H, t, J = 6.0 Hz, NH), 7.24 (1H, ddd, J_1 = 0.9 Hz, J_2 = 4.9 Hz, J_3 = 7.8 Hz, H8'), 7.66 (1H, bddd, J_1 = 286 287 1.8 Hz, *J*₂ = 2.2 Hz, *J*₃ = 7.8 Hz, H9'), 8.47 (1H, dd, *J*₁ = 1.8 Hz, *J*₂ = 4.9 Hz, H7'), 8.52 (1H, bd, J = 2.2 Hz, H5'). ¹³C NMR: δ 14.57 (q, C27), 16.01 (q, C24), 16.21 (q, C25), 16.45 (q, 288 C26), 18.14 (t, C6), 19.45 (q, C29), 20.91 (t, C11), 21.31 (q, C2'), 23.67 (t, C2), 25.56 (t, 289 C12), 27.91 (q, C23), 29.39 (t, C21), 30.81 (t, C15), 33.68 (t, C16), 34.29 (t, C22), 37.09 (s, 290 291 C10), 37.68 (d, C13), 37.77 (s, C4), 38.32 (t, C1), 38.37 (t, C7), 40.71 (t, C3'), 40.73 (s, C8), 42.44 (s, C14), 46.65 (d, C19), 50.10 (d, C18), 50.51 (d, C9), 55.43 (d, C5), 55.66 (s, C17), 292 80.91 (d, C3), 109.48 (t, C30), 123.66 (d, C8'), 135.15 (s, C4'), 136.22 (d, C9'), 148.24 (d, 293 C7'), 148.73 (d, C5'), 150.74 (s, C20), 171.03 (s, C1'), 176.26 (s, C28). MS: *m/z* 589.2 294 [M+H]⁺, 587.2 [M-H]⁻. IR (KBr; cm⁻¹): 3270 (N–H val. amide), 2941 (C–H val.), 2871 (–CH₃) 295 val.), 1725 (-C=O val. ester), 1659 (-C=O val. amide). For C₃₈H₅₆N₂O₃ (588.86) calculated C 296 (77.51), H (9.59), N (4.76), found C (77.50), H (9.60), N (4.78). M.p. 241-244 °C. 297

298	5c : ¹ H NMR: δ 0.72 (1H, dd, $J_1 = 2.1$ Hz, $J_2 = 11.1$ Hz, H5), 0.78 (3H, s, H24), 0.79 (3H, s,
299	H25), 0.79 (3H, s, H23), 0.82 (3H, s, H26), 0.91 (3H, s, H27), 1.11 (2H, dt, <i>J</i> ₁ = 3.4 Hz, <i>J</i> ₂ =
300	3.4 Hz, $J_3 = 13.5$ Hz, H21), 1.64 (3H, dd, $J_1 = 0.7$ Hz, $J_2 = 1.4$ Hz, H29), 1.91 (2H, dt, $J_1 = 3.3$
301	Hz, $J_2 = 3.3$ Hz, $J_3 = 13.9$ Hz, H16), 1.99 (3H, s, H2'), 2.39 (1H, ddd, $J_1 = 3.7$ Hz, $J_2 = 11.6$
302	Hz, $J_3 = 13.0$ Hz, H13), 3.08 (1H, dt, $J_1 = 4.5$ Hz, $J_2 = 11.1$ Hz, $J_3 = 11.1$ Hz, H19), 4.31 (1H,
303	dd, $J_1 = 6.0$ Hz, $J_2 = 15.6$ Hz, H3'), 4.42 (1H, dd, $J_1 = 5.9$ Hz, $J_2 = 10.7$ Hz, H3), 4.45 (1H, dd,
304	$J_1 = 6.1$ Hz, $J_2 = 15.6$ Hz, H3'), 4.55 (1H, dq, $J_1 = 1.4$ Hz, $J_2 = 1.4$ Hz, $J_3 = 1.4$ Hz, $J_4 = 2.4$ Hz,
305	H30), 4.69 (1H, dq, $J_1 = 0.7$ Hz, $J_2 = 0.7$ Hz, $J_3 = 0.7$ Hz, $J_4 = 2.4$ Hz, H30), 6.10 (1H, t, $J = 0.7$ Hz, $J_2 = 0.7$ Hz, $J_3 = 0.7$ Hz, $J_4 = 2.4$ Hz, H30), 6.10 (1H, t, $J = 0.7$ Hz, $J_4 = 0.7$ Hz, $J_5 = 0.7$ Hz, $J_7 = 0.7$ Hz, $J_8 = $
306	6.1 Hz, NH), 7.18-7.21 (2H, m, H5'), 8.48-8.51 (2H, m, H6'). ¹³ C NMR: δ 14.58 (q, C27),
307	16.09 (q, C24), 16.22 (q, C25), 16.45 (q, C26), 18.15 (t, C6), 19.45 (q, C29), 20.91 (t, C11),
308	21.31 (q, C2'), 23.66 (t, C2), 25.55 (t, C12), 27.91 (q, C23), 29.46 (t, C21), 30.80 (t, C15),
309	33.72 (t, C16), 34.34 (t, C22), 37.09 (s, C10), 37.68 (d, C13), 37.77 (s, C4), 38.38 (t, C7),
310	38.38 (t, C1), 40.77 (s, C8), 42.24 (s, C14), 42.46 (t, C3'), 46.60 (d, C19), 50.10 (d, C18),
311	50.52 (d, C9), 55.44 (d, C5), 55.72 (s, C17), 80.91 (d, C3), 109.55 (t, C30), 122.57 (d, C5'),
312	148.98 (s, C4'), 149.41 (d, C6'), 150.65 (s, C20), 171.04 (s, C1'), 176.38 (s, C28). MS: <i>m/z</i>
313	589.2 [M+H] ⁺ , 587.2 [M-H] ⁻ . IR (KBr; cm ⁻¹): 3378 (N–H val. amide), 2946 (C–H val.), 2869
314	(-CH ₃ val.), 1734 (-C=O val. ester), 1642 (-C=O val. amide). For C ₃₈ H ₅₆ N ₂ O ₃ (588.86)
315	calculated C (77.51), H (9.59), N (4.76), found C (77.49), H (9.57), N (4.78). M.p. 105-108
316	°C.

- 317
- 318 2.6. (3β)-3-Hydroxy-*N*-(pyridin-n-ylmethyl)lup-20(29)-en-28-amide [6a (n=2), 6b (n=3) and
 319 6c (n=4)]

LiOH·H₂O (54 mg; 1.3 mmol; 5 eq) was added to a solution of **5a**, **5b** or **5c** (151 mg; 0.26 mmol) in methanol (15 mL), and the reaction mixture was refluxed for 5 h. After stopping the reaction, the solvent was evaporated, and the residue was dissolved in chloroform and filtered.

- 323 Evaporation of the solvent afforded a solid, which was purified by column chromatography.
- 324 Yields: **6a** (82 %), **6b** (84 %), **6c** (83 %).
- **6a**: ¹H NMR: δ 0.62 (1H, dd, J_1 = 2.3 Hz, J_2 = 11.5 Hz, H5), 0.70 (3H, s, H24), 0.74 (3H, d, J
- 326 = 0.8 Hz, H25), 0.75 (3H, s, H23), 0.90 (3H, s, H26), 0.91 (3H, d, *J* = 0.6 Hz, H27), 1.08 (2H,
- 327 dt, $J_1 = 3.1$ Hz, $J_2 = 3.1$ Hz, $J_3 = 13.5$ Hz, H21), 1.64 (3H, dd, $J_1 = 0.7$ Hz, $J_2 = 1.4$ Hz, H29),
- 328 2.04 (2H, dt, $J_1 = 3.3$ Hz, $J_2 = 3.3$ Hz, $J_3 = 13.5$ Hz, H16), 2.38 (1H, ddd, $J_1 = 3.6$ Hz, $J_2 = 11.6$
- 329 Hz, $J_3 = 13.1$ Hz, H13), 3.10 (1H, dt, $J_1 = 4.6$ Hz, $J_2 = 11.1$ Hz, $J_3 = 11.1$ Hz, H19), 3.13 (1H,
- 330 dd, $J_1 = 4.9$ Hz, $J_2 = 11.5$ Hz, H3), 4.46 (1H, dd, $J_1 = 5.1$ Hz, $J_2 = 12.0$ Hz, H1'), 4.53 (1H, dd,
- 331 $J_1 = 5.1$ Hz, $J_2 = 12.0$ Hz, H1'), 4.54 (1H, dq, $J_1 = 1.4$ Hz, $J_2 = 1.4$ Hz, $J_3 = 1.4$ Hz, $J_4 = 2.4$ Hz,
- 332 H30), 4.69 (1H, dq, $J_1 = 0.7$ Hz, $J_2 = 0.7$ Hz, $J_3 = 0.7$ Hz, $J_4 = 2.4$ Hz, H30), 6.85 (1H, bt, J = 0.7 Hz, $J_4 = 0.7$ Hz, $J_5 = 0.7$ Hz, $J_7 = 0.7$ Hz, $J_8 = 0.7$ Hz, $J_$
- 333 5.1 Hz, NH), 7.17 (1H, ddd, $J_1 = 1.3$ Hz, $J_2 = 5.0$ Hz, $J_3 = 7.6$ Hz, H5'), 7.29 (1H, bdt, $J_1 = 1.0$
- 334 Hz, $J_2 = 1.0$ Hz, $J_3 = 7.7$ Hz, H7'), 7.64 (1H, dt, $J_1 = 1.8$ Hz, $J_2 = 7.6$ Hz, $J_3 = 7.6$ Hz, H6'),
- 335 8.49 (1H, ddd, J_1 = 1.0 Hz, J_2 = 1.8 Hz, J_3 = 5.0 Hz, H4'). ¹³C NMR: δ 14.63 (q, C27), 15.34
- 336 (q, C24), 15.98 (q, C25), 16.12 (q, C26), 18.28 (t, C6), 19.46 (q, C29), 20.90 (t, C11), 25.61
- 337 (t, C12), 27.40 (t, C2), 27.99 (q, C23), 29.43 (t, C15), 33.67 (t, C16), 33.88 (t, C21), 34.34 (t,
- 338 C22), 37.17 (s, C10), 37.74 (d, C13), 38.34 (t, C7), 38.69 (t, C1), 38.83 (s, C4), 40.69 (s, C8),
- 339 42.46 (s, C14), 44.21 (t, C1'), 46.77 (d, C19), 50.07 (d, C18), 50.59 (d, C9), 55.34 (d, C5),
- 340 55.77 (s, C17), 78.95 (d, C3), 109.30 (t, C30), 122.33 (d, C4'), 122.61 (d, C7'), 137.05 (d,
- 341 C6'), 148.58 (d, C3'), 151.02 (s, C20), 157.12 (s, C2'), 176.36 (s, C28). MS: *m/z*
- 342 547.3[M+H]⁺, 545.3 [M-H]⁻. IR (KBr; cm⁻¹): 3368 (N–H val. amide), 2942 (C–H val.), 2868
- 343 (-CH₃ val.), 1649 (-C=O val. amide). For $C_{36}H_{54}N_2O_2$ (546.83) calculated C (79.07), H
- 344 (9.95), N (5.12), found C (79.09), H (9.93), N (5.14). M.p. 210-211 °C. $[\alpha]_D^{20} = +10.3$ (c
- 345 0.350).
- **6b**: ¹H NMR: δ 0.61 (1H, dd, J_1 = 2.3 Hz, J_2 = 11.0 Hz, H5), 0.70 (3H, s, H24), 0.76 (3H, d, J
- 347 = 0.7 Hz, H25), 0.79 (3H, s, H23), 0.91 (3H, s, H26), 0.91 (3H, d, *J* = 0.8 Hz, H27), 1.07 (2H,

348	dt, $J_1 = 3.2$ Hz, $J_2 = 3.2$ Hz, $J_3 = 13.4$ Hz, H21), 1.63 (3H, dd, $J_1 = 0.7$ Hz, $J_2 = 1.4$ Hz, H29),
349	1.88 (2H, dt, $J_1 = 3.3$ Hz, $J_2 = 3.3$ Hz, $J_3 = 13.5$ Hz, H16), 2.39 (1H, ddd, $J_1 = 3.7$ Hz, $J_2 = 11.5$
350	Hz, $J_3 = 13.0$ Hz, H13), 3.09 (1H, dt, $J_1 = 4.4$ Hz, $J_2 = 11.0$ Hz, $J_3 = 11.0$ Hz, H19), 3.12 (1H,
351	dd, $J_1 = 4.9$ Hz, $J_2 = 11.4$ Hz, H3), 4.31 (1H, dd, $J_1 = 5.8$ Hz, $J_2 = 15.0$ Hz, H1'), 4.47 (1H, dd,
352	$J_1 = 6.1$ Hz, $J_2 = 15.0$ Hz, H1'), 4.55 (1H, dq, $J_1 = 1.4$ Hz, $J_2 = 1.4$ Hz, $J_3 = 1.4$ Hz, $J_4 = 2.4$ Hz,
353	H30), 4.69 (1H, dq, $J_1 = 0.7$ Hz, $J_2 = 0.7$ Hz, $J_3 = 0.7$ Hz, $J_4 = 2.4$ Hz, H30), 6.06 (1H, bt, $J = 0.7$ Hz, $J_2 = 0.7$ Hz, $J_3 = 0.7$ Hz, $J_4 = 2.4$ Hz, H30), 6.06 (1H, bt, $J = 0.7$ Hz, $J_4 = 0.7$ Hz, $J_5 = 0.7$ Hz, $J_7 = 0.7$ Hz, $J_8 $
354	6.0 Hz, NH), 7.22 (1H, ddd, $J_1 = 0.9$ Hz, $J_2 = 4.9$ Hz, $J_3 = 7.9$ Hz, H6'), 7.64 (1H, bddd, $J_1 = 0.9$ Hz, $J_2 = 4.9$ Hz, $J_3 = 7.9$ Hz, H6'), 7.64 (1H, bddd, $J_1 = 0.9$ Hz, $J_2 = 4.9$ Hz, $J_3 = 7.9$ Hz, H6'), 7.64 (1H, bddd, $J_1 = 0.9$ Hz, $J_2 = 4.9$ Hz, $J_3 = 7.9$ Hz, H6'), 7.64 (1H, bddd, $J_1 = 0.9$ Hz, $J_2 = 4.9$ Hz, $J_3 = 7.9$ Hz, H6'), 7.64 (1H, bddd, $J_1 = 0.9$ Hz, $J_2 = 4.9$ Hz, $J_3 = 7.9$ Hz, H6'), 7.64 (1H, bddd, $J_1 = 0.9$ Hz, $J_2 = 4.9$ Hz, $J_3 = 7.9$ Hz, H6'), 7.64 (1H, bddd, $J_1 = 0.9$ Hz, $J_2 = 4.9$ Hz, $J_3 = 7.9$ Hz, H6'), 7.64 (1H, bddd, $J_1 = 0.9$ Hz, $J_2 = 4.9$ Hz, $J_3 = 7.9$ Hz, H6'), 7.64 (1H, bddd, $J_1 = 0.9$ Hz, $J_2 = 0.9$ Hz, $J_3 = 7.9$ Hz, H6'), 7.64 (1H, bddd, $J_1 = 0.9$ Hz, $J_2 = 0.9$ Hz, $J_3 = 7.9$ Hz, H6'), 7.64 (1H, bddd, $J_1 = 0.9$ Hz, $J_2 = 0.9$ Hz, $J_3 = 7.9$ Hz, H6'), 7.64 (1H, bddd, J_1 = 0.9
355	1.7 Hz, $J_2 = 2.3$ Hz, $J_3 = 7.6$ Hz, H7'), 8.49 (1H, bdd, $J_1 = 1.7$ Hz, $J_2 = 4.9$ Hz, H5'), 8.50 (1H,
356	bdq, J_1 = 0.7 Hz, J_2 = 0.7 Hz, J_3 = 0.7 Hz, J_4 = 2.3 Hz, H3'). ¹³ C NMR: δ 14.62 (q, C27), 15.34
357	(q, C24), 16.02 (q, C25), 16.15 (q, C26), 18.26 (t, C6), 19.49 (q, C29), 20.90 (t, C11), 25.60
358	(t, C12), 27.39 (t, C2), 27.96 (q, C23), 29.41 (t, C15), 30.83 (t, C21), 33.70 (t, C16), 34.37 (t,
359	C22), 37.17 (s, C10), 37.71 (d, C13), 38.33 (t, C7), 38.70 (t, C1), 38.83 (s, C4), 40.42 (s, C8),
360	40.72 (t, C1'), 42.45 (s, C14), 46.63 (d, C19), 50.12 (d, C18), 50.61 (d, C9), 55.35 (d, C5),
361	55.67 (s, C17), 78.94 (d, C3), 109.42 (t, C30), 123.62 (d, C6'), 135.04 (s, C2'), 136.06 (d,
362	C7'), 148.41 (d, C5'), 148.83 (d, C3'), 150.77 (s, C20), 176.23 (s, C28). MS: <i>m</i> / <i>z</i>
363	547.3[M+H] ⁺ , 545.3 [M-H] ⁻ . IR (KBr; cm ⁻¹): 3284 (N–H val. amide), 2933 (C–H val.), 2867
364	(-CH ₃ val.), 1640 (-C=O val. amide). For C ₃₆ H ₅₄ N ₂ O ₂ (546.83) calculated C (79.07), H
365	(9.95), N (5.12), found C (79.05), H (9.96), N (5.11). M.p. 254-256 °C. $[\alpha]_D^{20} = +11.1 (c$
366	0.307).
367	6c : ¹ H NMR: δ 0.62 (1H, dd, $J_1 = 2.2$ Hz, $J_2 = 11.1$ Hz, H5), 0.71 (3H, s, H24), 0.77 (3H, bs,
368	H25), 0.82 (3H, s, H23), 0.92 (3H, bs, H27), 0.92 (3H, s, H26), 1.12 (2H, dt, <i>J</i> ₁ = 3.5 Hz, <i>J</i> ₂ =
369	3.5 Hz, $J_3 = 13.5$ Hz, H21), 1.64 (3H, dd, $J_1 = 0.7$ Hz, $J_2 = 1.4$ Hz, H29), 1.74 (2H, ddd, $J_1 =$

- 370 0.9 Hz, $J_2 = 8.1$ Hz, $J_3 = 12.3$ Hz, H7), 2.39 (1H, ddd, $J_1 = 3.6$ Hz, $J_2 = 11.6$ Hz, $J_3 = 12.8$ Hz,
- 371 H13), 3.09 (1H, dt, J_1 = 4.6 Hz, J_2 = 11.1 Hz, J_3 = 11.1 Hz, H19), 3.13 (1H, dd, J_1 = 4.9 Hz, J_2
- 372 = 11.4 Hz, H3), 4.32 (1H, dd, J_1 = 6.0 Hz, J_2 = 15.6 Hz, H1'), 4.45 (1H, dd, J_1 = 6.1 Hz, J_2 =

373 15.6 Hz, H1'), 4.55 (1H, dq, $J_1 = 1.4$ Hz, $J_2 = 1.4$ Hz, $J_3 = 1.4$ Hz, $J_4 = 2.4$ Hz, H30), 4.69 (1H, dq, $J_1 = 0.7$ Hz, $J_2 = 0.7$ Hz, $J_3 = 0.7$ Hz, $J_4 = 2.4$ Hz, H30), 6.05 (1H, bt, J = 6.0 Hz, NH), 374 7.16-7.18 (2H, m, H4'), 8.49-8.51 (2H, m, H3'). ¹³C NMR: δ 14.63 (q, C27), 15.33 (q, C24), 375 16.10 (q, C25), 16.16 (q, C26), 18.27 (t, C6), 19.49 (q, C29), 20.90 (t, C11), 25.60 (t, C12), 376 27.39 (t, C2), 27.96 (q, C23), 29.47 (t, C15), 30.82 (t, C21), 33.75 (t, C16), 34.42 (t, C22), 377 37.18 (s, C10), 37.71 (d, C13), 38.39 (t, C7), 38.71 (t, C1), 38.84 (s, C4), 40.76 (s, C8), 42.21 378 (t, C1'), 42.48 (s, C14), 46.59 (d, C19), 50.12 (d, C18), 50.61 (d, C9), 55.37 (d, C5), 55.73 (s, 379 C17), 78.96 (d, C3), 109.49 (t, C30), 122.47 (d, C3'), 148.49 (s, C2'), 149.78 (d, C4'), 150.70 380 (s, C20), 176.23 (s, C28). MS: m/z 547.3[M+H]⁺, 545.3 [M-H]⁻. IR (KBr: cm⁻¹): 3352 (N-H 381 val. amide), 2942 (C-H val.), 2871 (-CH₃ val.), 1670 (-C=O val. amide). For C₃₆H₅₄N₂O₂ 382 (546.83) calculated C (79.07), H (9.95), N (5.12), found C (79.06), H (9.94), N (5.15). M.p. 383 246-248 °C. $[\alpha]_{D}^{20} = +3.7$ (*c* 0.454). 384

385

386 2.7. Cell Cultures

The screening cell lines: T-lymphoblastic leukemia CEM; breast carcinoma MCF7; cervical 387 carcinoma HeLa; malignant melanoma G-361; and human foreskin fibroblasts BJ were 388 obtained from the American Type Culture Collection (Manassas, VA, USA). Cells were 389 cultured in DMEM (Dulbecco's Modified Eagle Medium, Sigma, MO, USA). Media used 390 were supplemented with 10% fetal bovine serum, 2 mM L-glutamine, and 1% penicillin-391 streptomycin. The cell lines were maintained under standard cell culture conditions at 37 °C 392 and 5% CO₂ in humid environment. Cells were subcultured twice or three times a week using 393 the standard trypsinization procedure. 394

395

396 2.8. Cytotoxicity screening tests

397 Description of the experimental procedure used in cytotoxicity test was already published

398 [1,2]. IC₅₀ values obtained with the compounds 3a-3c and 6a-6c are shown in Table 1.

399

400 2.9. Flow cytometry analysis

G-361 cancer cells were trypsinized, seeded in 100 mm culture dishes, and after 24 h of 401 stabilization incubated with the tested compounds. After additional 24 h, the cells were again 402 detached with trypsin, washed, fixed and stained in 0.1% [v/v] Triton X-100, 0.2 mg.mL⁻¹ 403 RNase A and 10 µg.mL⁻¹ propidium iodide in PBS. Their DNA contents were then assessed 404 by a flow cytometer (FACS VerseTM, Becton Dickinson, NJ, USA), and the distribution of 405 cells in subG₁ ("apoptotic cells"), G_0/G_1 , S and G_2/M peaks were quantified by histogram 406 analysis using BD FACSuite software (Becton Dickinson, NJ, USA). The experiments were 407 repeated three times. The differences between control and treated cells were analyzed by one-408 way analysis of variance ANOVA using Microsoft Excel 2010. Signification was marked 409 410 with asterisk (p < 0.05).

411

412 2.10. SDS–polyacrylamide gel electrophoresis and immunoblotting

Cells (G-361) were seeded into culture medium in 100-mm culture dishes at a density of $1.5 \times$ 413 10⁶ cells.mL⁻¹ and after 24 h of stabilization treated with tested compounds. DMSO was used 414 as a vehicle for controls. After another 24 h of treatment, the cells were washed three times 415 with cold PBS (10 mM, pH 7.4) and lysed in ice-cold RIPA protein extraction buffer (20 mM 416 Tris-HCl, pH 7.4, 5 mM EDTA, 2 mM EGTA, 100 mM NaCl, 2 mM NaF, 0.2% Nonidet P-417 40, 30 mM PMSF, 1 mM DTT, 10 mg.mL⁻¹ of aprotinin and leupeptin). The lysate was 418 collected into a microfuge tube and incubated on ice for 1 h. It was then cleared by 419 centrifugation at 18,000 × g for 30 min at 4 °C and the supernatant was collected. Proteins in 420 lysates were quantified by the Bradford method and diluted with Laemmli electrophoresis 421 buffer. The proteins were then separated on 10% or 12% SDS-polyacrylamide gels, 422

transferred to nitrocellulose membranes (Bio-Rad Laboratories, CA, USA) and stained with 423 Ponceau S (Sigma Aldrich, St. Louis, MO, USA) to check equal protein loading. The 424 membranes were blocked with 5% (w/v) non-fat dry milk and 0.1% Tween-20 in PBS for 2 h 425 and probed with specific primary antibodies (Cell Signaling Technology, Danvers, MA, USA; 426 Santa Cruz Biotechnology, CA, USA) overnight. Anti-caspase-3 and anti-caspase-7 427 recognized zymogen and also its fragment. After washing in PBS and PBS with 0.1% Tween-428 20, the membranes were probed with horseradish peroxidase-conjugated secondary antibodies 429 and visualized with West Pico Supersignal chemiluminescent detection reagent (Thermo 430 Fisher Scientific, Rockford, USA). Signal was detected by a CCD camera (Fujifilm, Tokio, 431 Japan). To confirm equal protein loading, immunodetection was performed with anti-β-actin 432 monoclonal antibody (Santa Cruz Biotechnology, CA, USA). The experiments were repeated 433 three times. The protein expression in treated cells was compared to untreated controls. 434 435

436 2.11. Activities of caspase 3/7

Treated G-361 cells were harvested by centrifugation and homogenized in an extraction buffer 437 (10 mM KCl, 5 mM HEPES, 1 mM EDTA, 1 mM EGTA, 0.2% CHAPS, plus protease 438 inhibitors: aprotinin, leupeptin, PMSF; pH 7.4) on ice for 20 min. The resulting homogenates 439 were clarified by centrifugation at $18,000 \times g$ for 20 min at 4°C. The protein contents of the 440 samples were quantified by the Bradford method and they were diluted to equivalent protein 441 concentrations. Lysates were then incubated for 1 h with 100 mM Ac-DEVD-AMC as a 442 substrate (Sigma, MO, USA) in an assay buffer (25 mM PIPES, 2 mM EGTA, 2 mM MgCl₂, 443 5 mM DTT, pH 7.3). For negative controls, the lysates were supplemented with 100 mM Ac-444 DEVD-CHO (Sigma, MO, USA) as a caspase-3/7 inhibitor. The fluorescence of the product 445 was measured using a Fluoroskan Ascent microplate reader (Labsystems, Vantaa, Finland) at 446 346/442 nm (ex/em). Experiments were repeated three times in triplicates. The differences 447

- between control and treated cells were analyzed by one-way analysis of variance ANOVA using Microsoft Excel 2010. Signification was marked with asterisk (p < 0.05).
- 450

451 **3. Results and Discussion**

452 3.1. Synthetic protocol

The synthesis of both series of the target picolyl amides (3a-3c and 6a-6c) is efficient and 453 easy. To get **3a–3c** (Scheme 1), betulinic acid (1) was first esterified by succinic anhydride 454 [2,26] to prepare its hemiester 2, which was then subjected to a formation of an amide bond 455 with o-, m- and p-picolyl amines, using T3P as condensation agent [2,27]. To get 6a–6c 456 (Scheme 2), protecting of the C(3)-OH group as acetate ester was required, and achieved by 457 reacting betulinic acid (1) with acetic anhydride in dry THF, using EDIPA as a base and 458 DMAP as reaction promotor [2]. Free carboxylic group of **1** was then converted to betulinic 459 460 acyl chloride by means of oxalyl chloride in dichloromethane [2,28], and used subsequently without isolation in a formation of amide with o-, m- and p-picolyl amines, using again 461 462 EDIPA as a base [27], affording 5a-5c. Protecting acetate group was removed by LiOH.H₂O in methanol [2], yielding the required amides **6a–6c**. 463 *Cytotoxicity*: The experimental data obtained (Table 1) revealed high importance of **3b** and 464 3a, while 3c and 6a–6c displayed only moderate cytotoxicity. The cytotoxicity data of 3a–3c 465 clearly demonstrate the importance of substitution of pyridine ring in structure-activity 466 relationship studies: while o- and m-substitution of the pyridine ring resulted in highly 467 cytotoxic compounds **3a** and **3b**, *p*-substitution revealed its disadvantage in **3c**. Both cytotoxic 468 compounds, **3a** and **3b**, showed very low cytotoxicity in normal fibroblasts, and became 469 potential candidates for cytotoxic agents against G-361 ($2.4 \pm 0.0 \mu$ M and $0.5 \pm 0.1 \mu$ M, 470

- 471 respectively), MCF7 ($2.2 \pm 0.2 \mu$ M and $1.4 \pm 0.1 \mu$ M, respectively), HeLa ($2.3 \pm 0.5 \mu$ M and
- 472 $2.4 \pm 0.4 \,\mu$ M, respectively) and CEM ($6.9 \pm 0.4 \,\mu$ M and $6.5 \pm 1.5 \,\mu$ M, respectively) cancer

473	cell lines. Moreover, 3b was the most active in melanoma cancer cell line G-361 (IC $_{50}$ 0.5 \pm
474	0.1 μ M) compared to other tested cancer cell lines with weak activity in normal human cells
475	(BJ) that is important for wide therapeutic window. We determined selectivity against all
476	tested cancer cells compared to normal cells with therapeutic index between 7 and 100 for
477	compounds 3a and 3b . The highest therapeutic index ($TI = 100$) was calculated for human
478	malignant melanoma cell line (G-361) versus normal human fibroblasts (BJ) with 3b.
479	Difluoroderivatives of dihydrobetulinic acid showed selectivity with therapeutic index
480	between 2 and 8 against cancer lines [29]. In another study, therapeutic index of 9.1 was
481	measured for conjugate of (3β) -3-O-acetylbetulinic acid with a triazole ring further
482	substituted by benzaldehyde [30]. Based on the literature search, no derivative of betulinic
483	acid was as highly active in melanoma cancer cell line G-361 as 3b . In a future, we would like
484	to design other compounds with higher selectivity, mainly based on the structure of 3b .
485	Together with obtained results, these derivatives could be promising for in vivo experiments
486	for development new anticancer drugs.
487	Concerning 6a–6c , their lower cytotoxicity has been expected due to the substitution of
488	carboxyl group in these derivatives of betulinic acid. However, the great disadvantage of $6a$ -
489	6c is their cytotoxicity in normal fibroblasts, which is sometimes even higher than that in the
490	cancer cell lines. The cytotoxicity of 6a–6c in CEM, MCF7 and HeLa cancer cell lines is
491	comparable with the cytotoxicity of the compounds investigated earlier [1]. The equal
492	moderate cytotoxicity with no selectivity as for 6a–6c was also shown for saponins of
493	betulinic acid in different cancer cell lines [33].
494	

495 3.2. Picolyl amides of betulinic acid caused apoptosis

496 We have observed whether compounds 3a-3c influence the cell cycle and cause apoptosis in

497 human malignant melanoma cells G-361. Cells were treated with 0.5, 2.5, 12.5 or 25 μ M of

three substances (3a-3c) for 24 h. Then the cells were harvested, stained and analyzed by 498 flow cytometry, western blotting or caspase activation assay. Flow cytometric analysis 499 showed an increase in the subG₁ phase of the cell cycle (apoptotic cells) in G-361 cell line 500 after treatment with **3a** and **3b**. In turn, **3c** did not enhance the number of subG₁ cells and 501 caused no changes in the cell cycle distribution. In these cells, treatment with **3a** and **3b** 502 increased the proportion of cells in the G_0/G_1 phase and decreased their proportion in the 503 concomitant phases, mainly G₂/M (Figure 1). Apoptotic markers were detected by western 504 505 blotting to examine the antiproliferative activities of tested compounds 3a-3c. The initiation of apoptosis is mediated by caspase cascade activation [34]. Caspase-3 and caspase-7 are 506 executioner proteases that result in cleavage of poly-(ADP-ribose) polymerase (PARP) and 507 subsequent DNA degradation and apoptotic death [35]. The western blots showed degradation 508 of caspase-7 into cleaved fragments (a part of the apoptotic cascade) after treatments with 3a 509 510 and **3b** in G-361 cells (Figure 2). PARP cleavage, a final step of caspase activation, was observed in human malignant melanoma cells treated by 3a and 3b (Figure 2). pRb was 511 512 downregulated by 12.5 µM and 25 µM of 3a and 3b, but total Rb also decreased in those 513 treatments. It is probably due to reaching the cytotoxicity towards human melanoma cells. No changes in the expressions of anti-apoptotic proteins Bcl-2 and Mcl-1 and also p53 were 514 observed. Downregulation of phospho-p44/42 MAPK (Erk 1/2) (Thr202/Tyr204) was 515 516 detected after the treatment with 12.5 and 25 µM of **3a** and **3b** for 24 h. Total level of p44/42 was not changed. This kinase is responsible for proliferation, differentiation, motility, and 517 death of cells. The compounds **3a** and **3b** inhibited those activities mediated by Erk 1/2. These 518 data were supplemented by estimation of caspase-3/7 activity in G-361 treated cells using the 519 fluorogenic substrate Ac-DEVD-AMC and/or caspase-3/7 inhibitor Ac-DEVD-CHO. 520 521 Compound **3a** increased activity of caspase-3/7 after the treatment with 25 μ M up to eighteen fold. Compound **3b** induced the activity of caspase-3/7; after 24 h treatment with 522

concentration 25 μM, a twelve fold increase in the effector caspases was observed compared
to the untreated control (Figure 3). Unlike 3a and 3b, compound 3c did not affect the activity
of caspases-3/7 after 24 h. These results show that 3a and 3b can induce apoptosis by caspase3 activation, and target proliferation and differentiation in G-361 cells.

527

528 3.3. Physico-chemical parameters

To support rational design of the target compounds investigated in this work, their physico-529 chemical parameters have been calculated using ACD/iLabs software and databases [36]. The 530 parameters calculated for **3a–3c** and **6a–6c** were compared with the Lipinski [5] rule of five 531 and with the Ghose [6] rules. The rules describe molecular properties important for a small 532 molecule drug pharmacokinetics in the human body, including their absorption, distribution, 533 metabolism and excretion (known as ADME parameters). However, the rules do not predict 534 535 displaying of the pharmacological activity. Lipinski [5] rule of five considers partition coefficient (log P, range -0.4 to +5.6), molar refractivity (range 40 to 130), molecular weight 536 537 (range 180 to 500), number of atoms in the molecule (20 to 70) and polar surface area (up to 14 nm). The data for comparison are presented in Table 2. However, the target compounds 538 **3a–3c** and **6a–6c** do not correspond to the range given for any of the above Lipinski and 539 Ghose rules [5,6], except of the parameter giving the number of atoms in the molecule. 540 Despite those facts, **3a–3c** and **6a–6c** display cytotoxicity. Exceptions are already known, 541 where pharmacologically active compounds do not correspond to all rules, we have observed 542 such result as well [1,2]. 543

The most important molecular descriptors are the partition coefficient (log *P*) and the distribution coefficient (log *D*) [37]. In chemical and pharmaceutical sciences, both, log *P* and log *D* are measures of hydrophilicity or hydrophobicity of the studied compound, and are useful for estimating distribution of a drug within the body, where log *D* shows the

dependence on the pH of the matrix. Hydrophobic drugs are then preferentially distributed to 548 hydrophobic compartments (e.g., lipid bilayer of cells), while hydrophilic drugs are 549 preferentially distributed to hydrophilic compartments (e.g., blood serum). The distribution 550 coefficient is a pH dependent value, and, therefore, the value at pH = 7.4 (the physiological 551 pH value of blood serum) is of particular importance (see also Table 2). Thus, log *P* expresses 552 a ratio of concentrations of non-ionized compound between two phases, non-polar (octanol) 553 and polar (water), while log D expresses the ratio of the sum of the concentrations of all forms 554 555 of the compound (ionized and non-ionized) in each of the two phases. In pharmacology, $\log P$ and log D indicate how easily the drug can reach its intended target in the body, how strong 556 its effect will be once it reaches its target and how long it will remain in the body in an active 557 form. The log P values calculated for either **3a–3c** or **6a–6c** are identical, and they exceed the 558 values given by the Lipinski and Ghose rules [5,6] (Table 2). Another supportive parameter is 559 the predicted aqueous solubility, $\log S$. S (in mol dm⁻³) is the concentration of the solute in a 560 saturated solution that is in equilibrium with the crystalline solid, and it is a pH dependent 561 parameter. Standard range for log S at pH 7.4 is -6.5/+0.5. Among the prepared compounds, 562 only **3a–3c** show values for log *S* in this range (Table 2). 563

564

565 3.4. ADME parameters:

The importance of some of the ADME parameters for evaluation of pharmacokinetic properties of the prepared compounds is summarized in Table 2. The blood brain barrier (BBB) and plasma protein binding are two of the important factors affecting distribution of the compound in the human body. Several parameters assist in evaluation of each potential drug for its BBB transport [38] The rate of brain penetration, log *PS*, is a logarithm of the permeability-surface area coefficient that measures the ability of a drug to cross the BBB and to move into brain tissue over time. It is one of the relevant parameters for evaluation of the

573	rate of BBB penetration. The extent of brain penetration parameter, log <i>PB</i> , indicates if the
574	drug might be active or inactive on the central nervous system (CNS). The typical values for
575	log <i>PB</i> are $-1.5/+1.5$. Most of the current drugs show the log PB value up to +2 (active on
576	CNS) or down to -2 (inactive on CNS). The log <i>PB</i> values calculated for 3a–3c and 6a–6c
577	indicate that these compounds will be inactive on CNS (Table 2). However, the complex of
578	the BBB parameters is completed by the brain/plasma equilibration rate, the parameter
579	expressed as log $PS * f_{u, brain}$ that is a mathematical modeling parameter based on time
580	required for reaching brain equilibrium. It is dependent on the brain unbound fraction $(f_{u, brain})$.
581	This value indicates if the drug may potentially be active on CNS together with the log BB
582	parameter, the predicted brain/blood partition coefficient (cf. Supplementary material). The
583	parameter log BB is a hybrid parameter determined by permeability, plasma and brain tissue
584	binding, and active transport mechanism, standard range $-3.0/+1.2$. The log BB values
585	calculated for 3a–3c and 6a–6c appear in the required range of this parameter. Nevertheless,
586	there are numerous exceptions already known. A supportive parameter, plasma protein
587	binding (PPB) gives calculated quantity of a drug bound to a protein.
588	Another important ADME parameter is bioavailability. Among the prepared compounds, the
589	calculated bioavailability for 3a–3c is lower than that for 6a–6c . However, the experimental
590	data show that 3a–3c are generally more cytotoxic to the tested cancer cell lines than 6a–6c .
591	The rules [5,6] describe molecular properties important for a small molecule drug
592	pharmacokinetics in the human body, including their absorption, distribution, metabolism and
593	excretion (ADME parameters). However, the rules do not predict whether a compound will
594	display pharmacological activity. Nevertheless, the importance of several ADME parameters,
595	namely blood-brain barrier and plasma protein binding, reflects the distribution of the
596	compound in the human body [39]. A comparison of the calculated physico-chemical
597	properties with the measured cytotoxicity shows that every time a new class of compound is

being investigated, no available experimental screening may be intentionally omitted (Tables1 and 2).

600

3.5. A comparison of **3a–3c** and **6a–6c** with the compounds prepared earlier [1, 2] 601 Earlier prepared compound, bearing picolyl amine moiety in the molecule, derived either 602 from lanosterol, cholesterol or situaterol, displayed cytotoxicity in the range of $IC_{50} = 25-45$ 603 μ M, or were inactive (IC₅₀ > 50 μ M). The earlier published data also show that the physico-604 chemical and ADME parameters calculated for those compounds, corresponded to the 605 606 Lipinski and Ghose rules [5,6] partially only. The current compounds **3a–3c** and **6a–6c** show generally higher cytotoxicity than the earlier series of compounds. Only **3a** and **3b** display 607 considerable cytotoxicity, and **3b** show the IC₅₀ value on G361 in the range IC₅₀ = 0.5 ± 0.1 608 μ M, with the therapeutic index TI = 100. This finding augments the importance of this 609 particular compound **3b**. In turn, even **3a** and **3b** corresponded to a limited number of 610 calculated physico-chemical and ADME parameters. Any calculation of those parameters is 611 useful and often desired in the drug design. However, the conclusions of the calculations 612 should always be supported by the experimental data for confirmation or correction. 613

614

615 **4. Conclusion**

Amides **3b** and **3a** represent the most important members of this series of compounds because of their high cytotoxicity against human melanoma cell line G-361 (Table 1). The former of this couple of amides (**3b**) is active in concentration $0.5 \pm 0.1 \mu$ M, and, subsequently it does not damage healthy cells. It also displays high cytotoxicity in other tested cell lines. Investigating structure-activity relationship in dependence on substitution of pyridine ring, the conclusions are different for each subseries. In the former subseries (**3a–3c**), the cytotoxicity of amides bearing *m*- and *o*-substituted pyridine ring (**3b** and **3a**) showed high cytotoxicity in

623	all tested cell lines, while p -substitution of the pyridine ring (3c) resulted in weak cytotoxicity
624	(Table 1). Our hypothesis is that this result might be connected with different mechanism of
625	action of 3b and 3a in comparison with 3c due to the modifications in their structures. This
626	hypothesis will be focused in a future investigation, because its confirmation may be
627	important for better understanding of structure-activity relationship. However, in the latter
628	subseries ($6a-6c$), amides bearing <i>p</i> - and <i>o</i> -substituted pyridine ring showed either
629	comparable cytotoxicity, or p -substituted amide was more active, and the m -substituted
630	derivative displayed the lowest cytotoxicity (Table 1). The most important finding with 6a–6c
631	was their high activity in the normal human cells. Due to that finding, 6a–6c are much less
632	suitable anti-cancer agents than $3a-3c$ that are, in turn, potential anti-cancer candidate agents.
633	
634	Supplementary material is available on the web page of the journal.
635	
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771	Abbreviations: CEM, cells of human T-lymphoblastic leukemia; MCF7, cells of human breast
772	adenocarcinoma; HeLa, cells of human cervical cancer; G-361, cells of human malignant
773	melanoma; BJ, normal human fibroblasts; T3P, propylphosphonic anhydride; EDIPA, ethyl
774	diisopropyl amine.
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796 [Scheme and Figure Captions]

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798 Scheme 1

i: succinic anhydride, DMAP, dry pyridine; ii: 2-, 3- or 4-aminomethylpyridine, T3P, dry

800 pyridine

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Scheme 2

i: acetanhydride, EDIPA, DMAP, THF; ii: oxalyl chloride, dichloromethane; iii: 2-, 3- or 4-

aminomethylpyridine, EDIPA, dichloromethane; iv: LiOH.H₂O, methanol

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806 Figure 1

Histograms with the distributions of G-361 cells in the G_0/G_1 , S, and G_2/M cell cycle phases

(A), and the subG₁ fraction of cells (**B**) measured by flow cytometric analysis, after 24 h

treatment with **3a**, **3b** and **3c** relative to untreated controls. Data indicate the percentages (%)

of the number of cells in respective phases. Experiments were repeated three times in

811 triplicates. Error bars are omitted for clarity. (A) Analysis of variance (one-way ANOVA)

between control and treated cells failed due to high biological variability in replicates. (B)

B13 Differences between control and treated cells among subG₁ phase were significant (p < 0.05)

814 *.

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Figure 2

817 Western blot analysis of apoptotic proteins (PARP, pRb, Rb, zymogen and fragment of

818 caspase-7, pAKT, AKT, pp44/42 MAPK (Erk 1/2) (Thr202/Tyr204), p44/42 MAPK (Erk 1/2)

in human malignant melanoma cells G-361 treated with amides. The expression of proteins in

cells treated with 0.5, 2.5, 12.5 and 25 μ M of compounds **3a**, **3b** and **3c** for 24 h were

821	compared with the protein expression of untreated control cells (0+ means control with
822	DMSO). The expression of β -actin was used as a protein loading marker.
823	
824	Figure 3
825	Activity of caspase-3/7 in malignant melanoma cells G-361 treated with 0.5, 2.5, 12.5 and 25
826	μ M amides 3a , 3b and 3c compared with untreated control cells for 24 h. The data indicate
827	the relative increase in caspase-3/7 activity. Experiments were repeated three times in
828	triplicates. * Differences between control and treated cells were significant ($p < 0.05$).
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846	Table 1. IC ₅₀ [µM]	values in four	cancer cell lines	and normal human	fibroblasts for 3a-3c
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Compound Cytotoxicity (IC ₅₀ [µM])															
	CEM ^b	MCF7 ^c	HeLa ^d	G-361 ^e	BJ ^f										
3a	6.9 ± 0.4	2.2 ± 0.2	2.3 ± 0.5	2.4 ± 0.0	46.2 ± 2.8	19.3									
3b	6.5 ± 1.5	1.4 ± 0.1	2.4 ± 0.4	0.5 ± 0.1	50.0 ± 0.0	100.0									
3c	22.6 ± 5.9	>50	40.9 ± 4.7	32.1 ± 2.1	>50	ND ^g									
6a	18.6 ± 1.0	27.0 ± 5.5	14.7 ± 0.2	16.4 ± 1.6	15.8 ± 1.4	1.0									
6b	25.3 ± 6.9	38.7 ± 1.0	23.6 ± 2.3	18.1 ± 0.1	17.5 ± 2.7	1.0									
6c	18.6 ± 3.7	21.1 ± 4.3	14.8 ± 0.2	11.6 ± 1.6	11.2 ± 1.4	1.0									
human fibro	blasts; ^g ND =	not determin	ed.												
	Ç														
	A C														
	P C														
	P C														

847 and **6a–6c** after 72 h

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Compd.	<mark>MW</mark>	Physico-chemical and ADME parameters ^a									
<mark>or ref.</mark>		log P	log D	log S (pH	<mark>bioav.</mark>	log PS *	log PS	log PB	log BB	PPB [%]	H _{acc} /
<mark>No.</mark>			(pH 7.4)	<mark>7.4)</mark>	<mark>[%]</mark>	<mark>fu, brain</mark>					H _{don}
) C				<mark>n.m.</mark>
<mark>3a</mark>	<mark>646.90</mark>	<mark>7.56</mark>	<mark>4.78</mark>	<mark>-5.01</mark>	<mark>< 30</mark>	<mark>-5.8</mark>	<mark>-4.0</mark>	<mark>-0.89</mark>	<mark>-0.0</mark>	<mark>99.80</mark>	<mark>7/2/</mark> 9
<mark>3b</mark>	<mark>646.90</mark>	<mark>7.56</mark>	<mark>5.26</mark>	<mark>-5.04</mark>	<mark>< 30</mark>	<mark>-5.8</mark>	<mark>-3.9</mark>	-0.89	<mark>-0.0</mark>	<mark>99.80</mark>	<mark>7/2/</mark> 9
<mark>3c</mark>	<mark>646.90</mark>	<mark>7.56</mark>	<mark>5.07</mark>	<mark>-5.48</mark>	<mark>< 30</mark>	<mark>-5.8</mark>	<mark>-3.9</mark>	<mark>-0.88</mark>	<mark>-0.0</mark>	<mark>99.80</mark>	<mark>7/2/</mark> 9
<mark>6a</mark>	<mark>546.83</mark>	<mark>7.00</mark>	<mark>7.00</mark>	<mark>-6.76</mark>	<mark>30-70</mark>	<mark>-4.1</mark>	<mark>-1.8</mark>	<mark>-0.30</mark>	<mark>-0.0</mark>	<mark>99.75</mark>	<mark>4/2/</mark> -
<mark>6b</mark>	<mark>546.83</mark>	<mark>7.00</mark>	<mark>7.00</mark>	<mark>-6.61</mark>	<mark>30-70</mark>	<mark>-4.1</mark>	<mark>-1.8</mark>	<mark>-0.30</mark>	<mark>-0.0</mark>	<mark>99.75</mark>	<mark>4/2/</mark> 4
<mark>6c</mark>	<mark>546.83</mark>	7.00	<mark>6.99</mark>	<mark>-6.58</mark>	<mark>30-70</mark>	<mark>-4.1</mark>	<mark>-1.8</mark>	<mark>-0.30</mark>	<mark>-0.0</mark>	<mark>99.75</mark>	<mark>4/2/</mark> 4
ref. [32]	max. 500	max. 5.0									max
					>						<mark>max</mark>
ref. [33]	Max. 500	Max. 5.6									

Table 2. Physico-chemical and ADME parameters of the target compounds **3a–3c** and **6a–6c** calculated using the ACD/iLabs software [36]

865	^a log P – partition coefficient; log D – distribution coefficient; log S – predicted aqueous solubility; bioav. = bioavailability – the degree of
866	availability of a chemical by the target tissue; $\log PS * f_{u, brain}$ – the brain/plasma equilibration rate, the parameter that is a mathematical modeling
867	parameter based on time required for reaching brain equilibrium; $\log PS - \log a$ the permeability-surface area coefficient; $\log PB - $ the
868	extent of brain penetration parameter; log BB – a hybrid parameter determined by permeability, plasma and brain tissue binding, and active
869	transport mechanism; PPB – plasma protein binding; $H_{acc} / H_{don} / n.m.b. =$ number of H-bond acceptors / number of H-bond donors / number of
870	movable bonds.
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881 Scheme 1



893 Scheme 2





914 Figure 2



932 Figure 3



950 Graphical Abstract



Compound		TI ^a				
	CEM ^b	MCF7 ^c	HeLa ^d	G-361 ^e	BJ ^f	
3 a	6.9 ± 0.4	2.2 ± 0.2	2.3 ± 0.5	2.4 ± 0.0	46.2 ± 2.8	19.3
3b	6.5 ± 1.5	1.4 ± 0.1	2.4 ± 0.4	0.5 ± 0.1	50.0 ± 0.0	100.0
3c	22.6 ± 5.9	>50	40.9 ± 4.7	32.1 ± 2.1	>50	ND ^g
6a	18.6 ± 1.0	27.0 ± 5.5	14.7 ± 0.2	16.4 ± 1.6	15.8 ± 1.4	1.0
6b	25.3 ± 6.9	38.7 ± 1.0	23.6 ± 2.3	18.1 ± 0.1	17.5 ± 2.7	1.0
6с	18.6 ± 3.7	21.1 ± 4.3	14.8 ± 0.2	11.6 ± 1.6	11.2 ± 1.4	1.0

Table 1. IC₅₀ [μ M] values in four cancer cell lines and normal human fibroblasts for **3a–3c** and **6a–6c** after 72 h

^a Therapeutic index (TI) calculated for G-361 line *versus* fibroblasts BJ; ^b CEM, cells of human T-lymphoblastic leukemia; ^c MCF7, cells of human breast adenocarcinoma; ^d HeLa, cells of human cervical cancer; ^e G-361, human malignant melanoma cell line ^f BJ, normal human fibroblasts; ^g ND = not determined

Compd.	MW	Physico-chemical and ADME parameters ^a									
or ref.		$\log P$	$\log D$	log S (pH	bioav.	$\log PS *$	log PS	log PB	log BB	PPB [%]	H _{acc} /
No.			(pH 7.4)	7.4)	[%]	$f_{u,\ brain}$					H_{don} /
							CY				n.m.b.
3a	646.90	7.56	4.78	-5.01	< 30	-5.8	-4.0	-0.89	-0.0	99.80	7/2/9
3b	646.90	7.56	5.26	-5.04	< 30	-5.8	-3.9	-0.89	-0.0	99.80	7/2/9
3c	646.90	7.56	5.07	-5.48	< 30	-5.8	-3.9	-0.88	-0.0	99.80	7/2/9
6a	546.83	7.00	7.00	-6.76	30-70	-4.1	-1.8	-0.30	-0.0	99.75	4/2/4
6b	546.83	7.00	7.00	-6.61	30-70	-4.1	-1.8	-0.30	-0.0	99.75	4/2/4
6c	546.83	7.00	6.99	-6.58	30-70	-4.1	-1.8	-0.30	-0.0	99.75	4/2/4
ref. [32]	max. 500	max. 5.0									max. 10 /
											max. 5 / -
ref. [33]	Max. 500	Max. 5.6	7	\mathcal{O}							
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Table 2. Physico-chemical and ADME parameters of the target compounds **3a–3c** and **6a–6c** calculated using the ACD/iLabs software [36]

^a log P – partition coefficient; log D – distribution coefficient; log S – predicted aqueous solubility; bioav. = bioavailability – the degree of availability of a chemical by the target tissue; $\log PS * f_{u, brain}$ – the brain/plasma equilibration rate, the parameter that is a mathematical modeling parameter based on time required for reaching brain equilibrium; $\log PS - \log arithm of the permeability-surface area coefficient; <math>\log PB - the$ extent of brain penetration parameter; $\log BB - a$ hybrid parameter determined by permeability, plasma and brain tissue binding, and active transport mechanism; PPB – plasma protein binding; $H_{acc} / H_{don} / n.m.b. =$ number of H-bond acceptors / number of H-bond donors / number of - nun movable bonds.













Highlights

Picolyl amides of betulinic acid as antitumor agents causing tumor cell apoptosis

Uladzimir Bildziukevich, Lucie Rárová, David Šaman, and Zdeněk Wimmer

- Picolyl amides of betulinic acid cause human malignant melanoma cell apoptosis;
- Comparison of substitution at the C(3)-OH and C(17)-COOH on cytotoxicity studied;
- Facilitating cellular uptake and enhancing antitumor properties;
- Cytotoxicity of one of the target compounds on G-361 found with TI = 100.

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