Supporting information

Ent-beyerane diterpenes as a key platform for the development of ArnT-mediated colistin resistance inhibitors

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1.¹H and ¹³C NMR spectra of compounds 11-12, 14, 4H, 15



Figure S1. ¹H NMR (CD₃OD, 400 MHz) spectrum of compound 11.



Figure S2. ¹³ C NMR (CD₃OD, 100 MHz) spectrum of compound **11**.



Figure S3. ¹H NMR (CD₃OD, 400 MHz) spectrum of compound **12**.





Figure S5. 1H NMR (CDCl₃, 400 MHz) spectrum of compound 14.



S4



Figure S7. ¹H NMR (CDCl₃, 400 MHz) spectrum of compound 4H.





Figure S8. ¹³ C NMR (CDCl₃, 100 MHz) spectrum of compound 4H.



Figure S9. ¹H NMR (CDCl₃, 400 MHz) spectrum of compound 15.



Figure S10. ¹³ C NMR (CDCl₃, 100 MHz) spectrum of compound 15.

2. Biological activity



2.1. Effect of compounds on bacterial growth

Figure S11. Dose-dependent effect of compounds **1**, **2**, **3**, **10**, **12**, **14** and **15** on PA14 col^R 5 growth after 24 hours at 37°C in MH (without colistin). Growth values are expressed as percentage relative to the cultures treated with equivalent concentrations of DMSO and represent the mean (±SD) of three independent experiments.



Figure S12. Dose-dependent effect of compounds **12**, **14** and **15** on the growth of the two clinical isolates *P. aeruginosa* MG75 and ND76 after 24 hours at 37°C in MH (without colistin). Growth values are expressed as percentage relative to the cultures treated with equivalent concentrations of DMSO and represent the mean (±SD) of three independent experiments.







Figure S13. Viability of 16HBE and CFBE epithelial cells exposed to the compounds 1, 2, 3, 12, 14 and 15 at the indicated concentrations, or DMSO at equivalent concentrations, for 18 hours. Cell viability was assessed through the MTT assay and expressed as percentage relative to untreated cells. Data are the mean (±SD) of three independent experiments.

μΜ	1	2	3	12	14	15
125	P=0.5175	P=0.2991	P=0.0374	P=0.0127	P>0.9999	P=0.8276
62.5	P>0.9999	P=0.7566	P=0.0942	P=0.1647	P>0.9999	P>0.9999
31.25	P=0.7531	P>0.9999	P=0.0072	P=0.0283	P>0.9999	P=0.7359
15,62	P>0.9999	P>0.9999	P=0.1173	P=0.3854	P>0.9999	P>0.9999
7.81	P>0.9999	P>0.9999	P=0.0669	P=0.2039	P=0.9870	P=0.4300
3.9	P>0.9999	P>0,.9999	P>0.9999	P>0.9999	P=0.8629	P>0.9999
1.95	P=0.4499	P>0.9999	P>0.9999	P=0.8883	P>0.9999	P=0.7828

Table S1. P values of compound- respect to DMSO-treated 16HBE cells by two-way ANOVA.

Table S2. P values of compound- respect to DMSO-treated CFBE cells by two-way ANOVA.

μM	1	2	3	12	14	15
125	P=0.0922	P=0.6880	P>0.9999	P=0.3081	P>0.9999	P=0.0733
62.5	P=0.0401	P=0.5943	P>0.9999	P>0.9999	P=0.6961	P>0.9999
31.25	P=0.2250	P>0.9999	P>0.9999	P=0.7167	P>0.9999	P=0.2424
15,62	P=0.0215	P>0.9999	P>0.9999	P=0.7139	P>0.9999	P=0.6526
7.81	P>0.9999	P>0.9999	P>0.9999	P>0.9999	P=0.3976	P=0.0814
3.9	P>0.9999	P>0.9999	P>0.9999	P=0.6697	P>0.9999	P=0.5528
1.95	P>0.9999	P>0.9999	P>0.9999	P=0.5700	P>0.9999	P>0.9999

3. Molecular modeling

Compound	Chemgauss4 ^a	XSCORE ^b
1	-3.56	6.35
2	-5.14	6.09
3	-5.79	6.31
4	-2.25	6.29
5	-0.22	6.28
6	-2.17	6.22
7	-1.42	6.15
8	-0.46	6.15
9	-2.99	6.20
10	-4.30	6.26
11	1.33	6.20
12	-5.51	6.32
13	-2.49	6.15
14	-6.20	6.63
15	-3.21	6.38

 Table S3. XSCORE rescoring results on docking poses predicted with FRED.

^a – the lower, the better ^b – expressed as -pK_d, the higher the better



Figure S14. Structural overimposition between the predicted binding mode of compounds **15** (A), **14** (B), and **12** (C) with the crystallographic pose of undecaprenyl phosphate (magenta sticks). The crystallographic structure of ArnT coded by PDB ID: 5F15 is shown as green lines and cartoon. Small molecules studied in this work are shown as cyan sticks.