

SUPPORTING INFORMATION  
FOR

**Guanylation reactions for the rational design of  
cancer therapeutic agents.**

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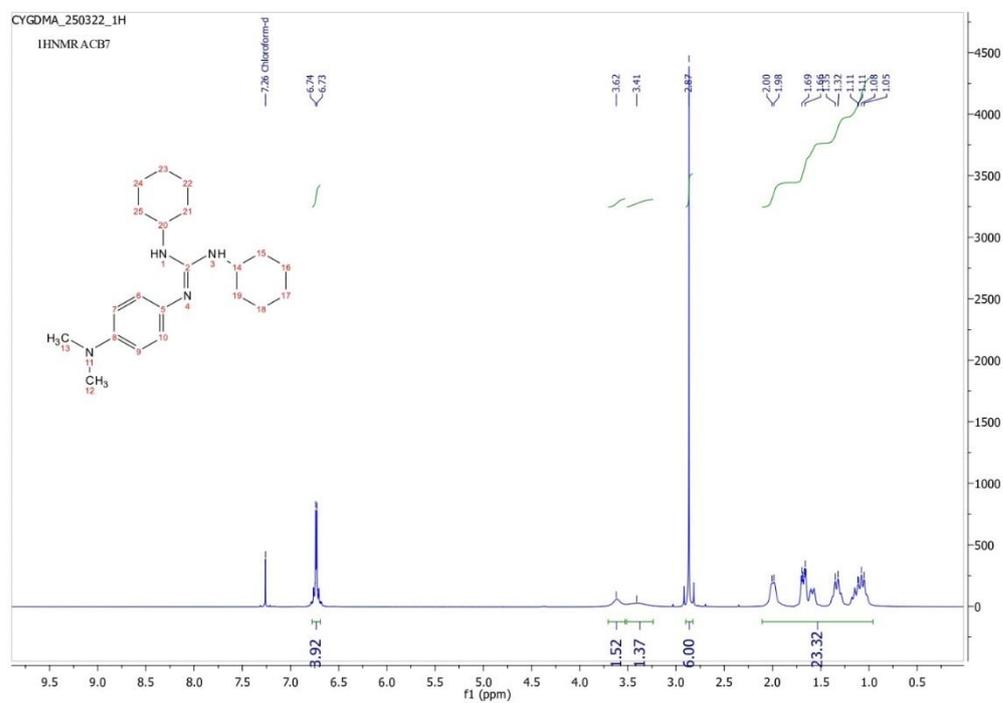
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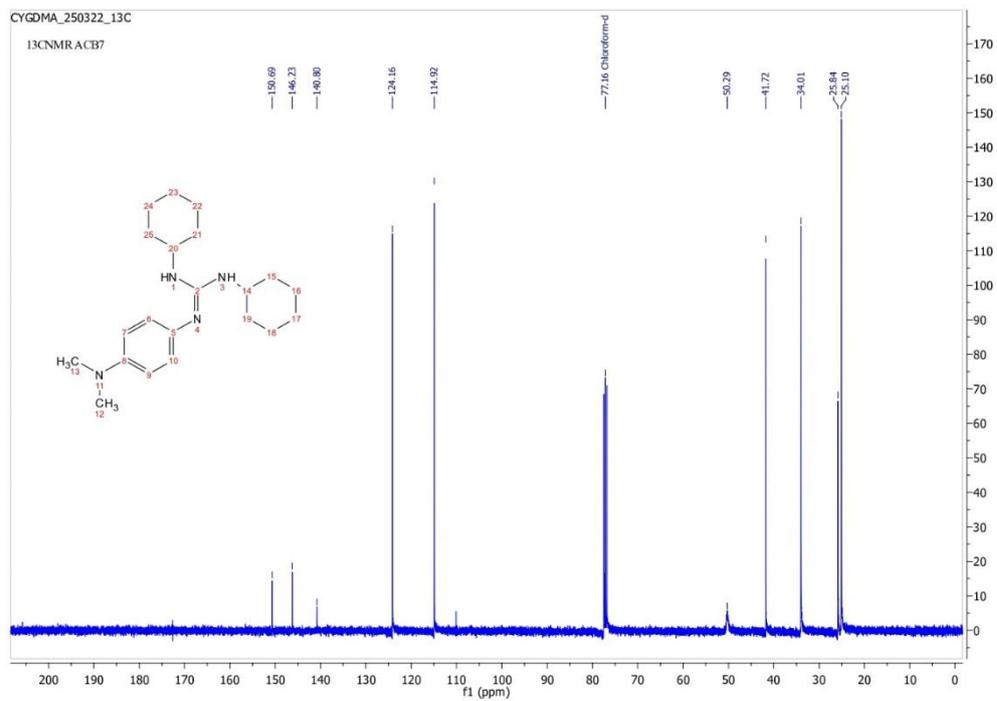
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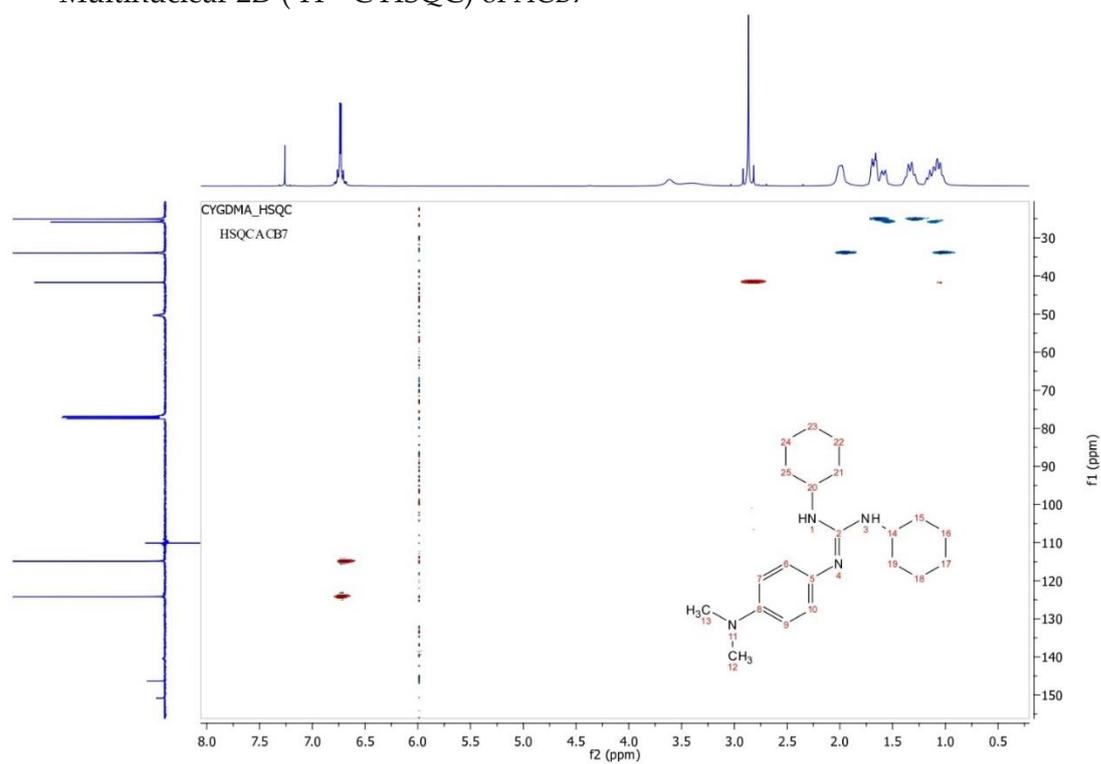
$^1\text{H-NMR}$  of ACB7 in  $\text{CDCl}_3$ .



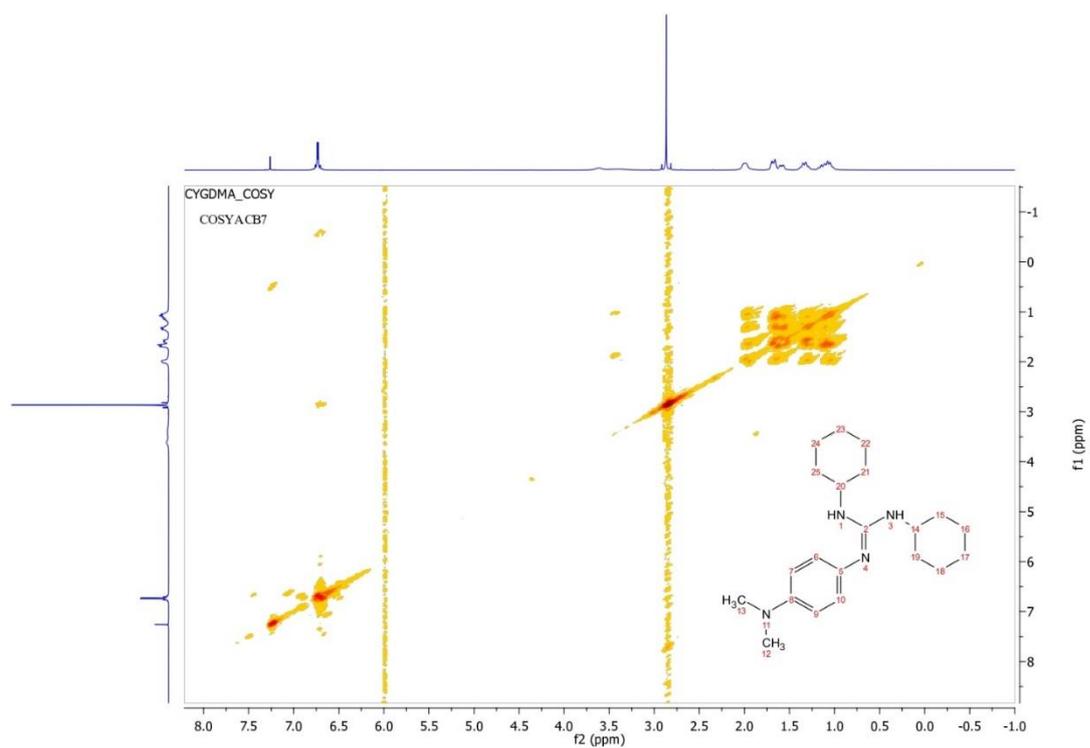
$^{13}\text{C-NMR}$  of ACB7 in  $\text{CDCl}_3$ .



Multinuclear 2D ( $^1\text{H}$ - $^{13}\text{C}$  HSQC) of ACB7

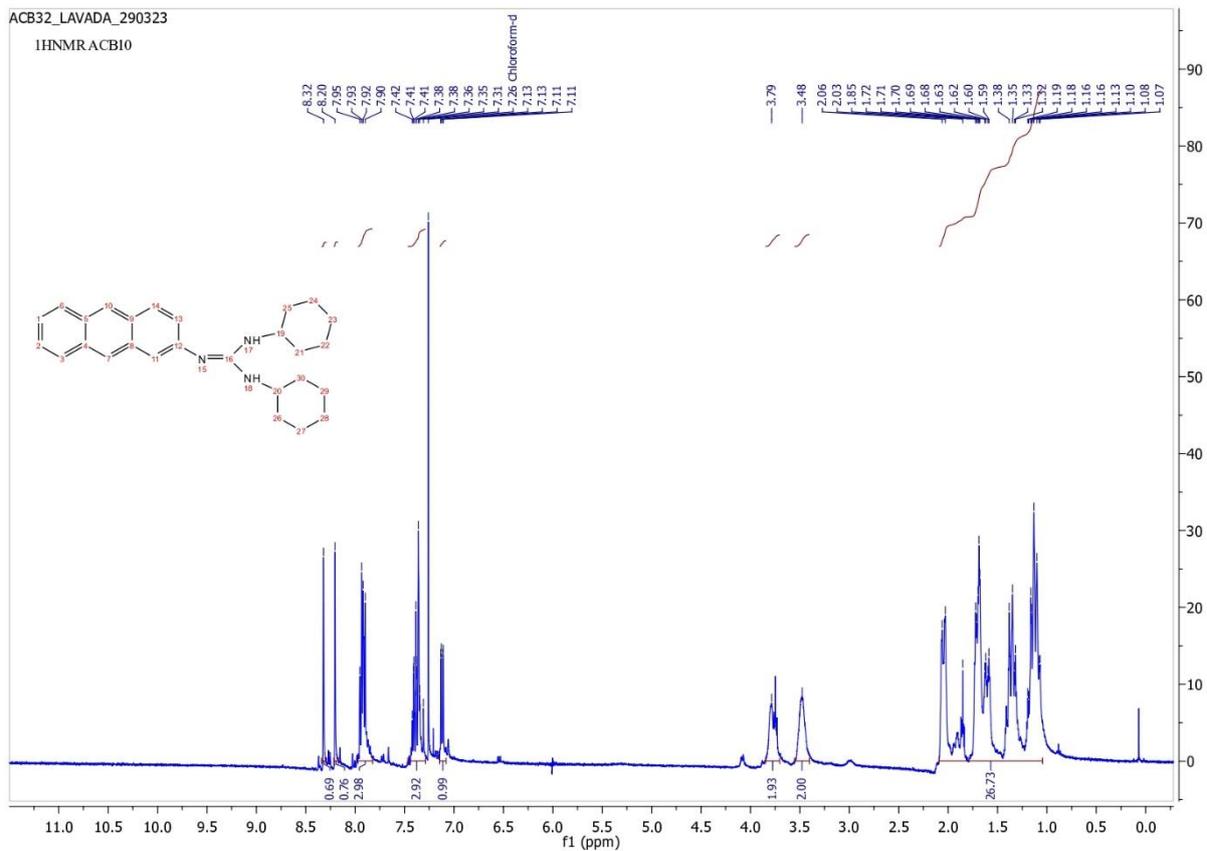


Multinuclear 2D ( $^1\text{H}$ - $^1\text{H}$  COSY) of ACB7

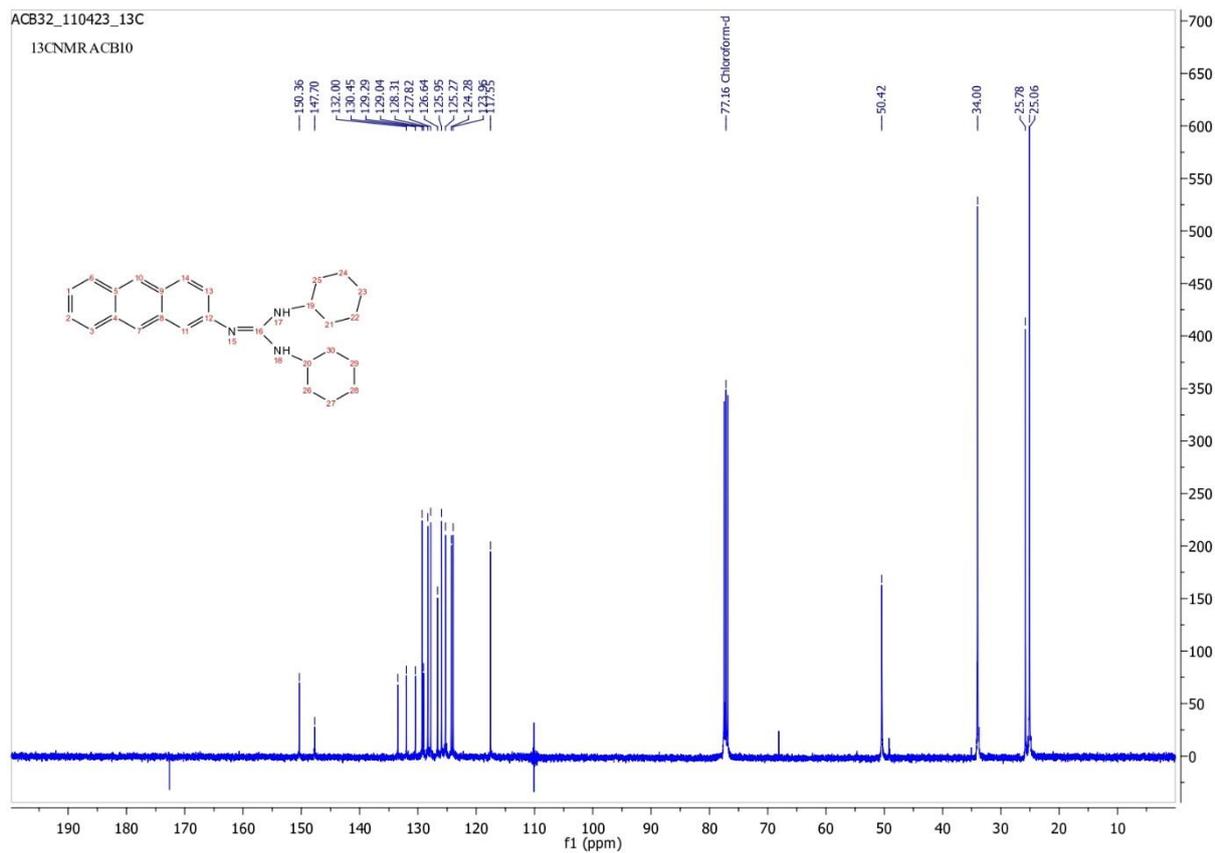


**Figure S1.**  $^1\text{H}$ -NMR,  $^{13}\text{C}$ -NMR, multinuclear 2D ( $^1\text{H}$ - $^{13}\text{C}$  HSQC) and multinuclear 2D ( $^1\text{H}$ - $^1\text{H}$  COSY) NMR spectra of ACB7

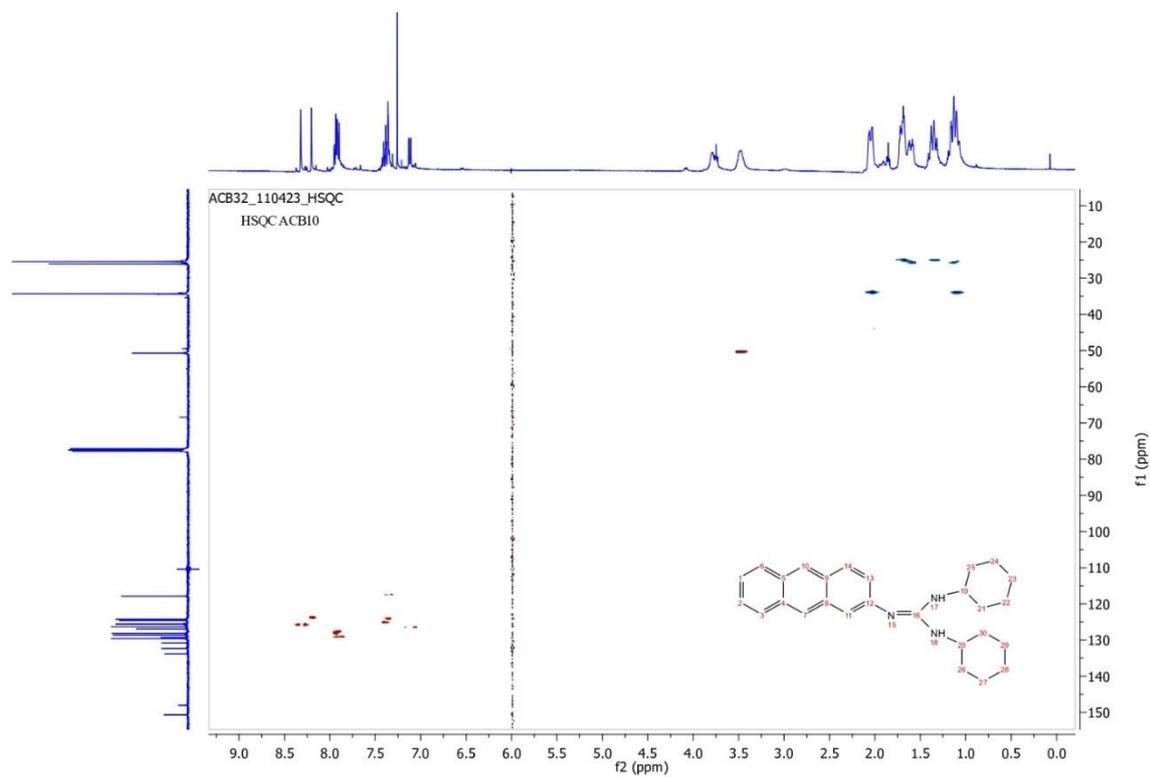
# <sup>1</sup>H-NMR of ACB10 in CDCl<sub>3</sub>.



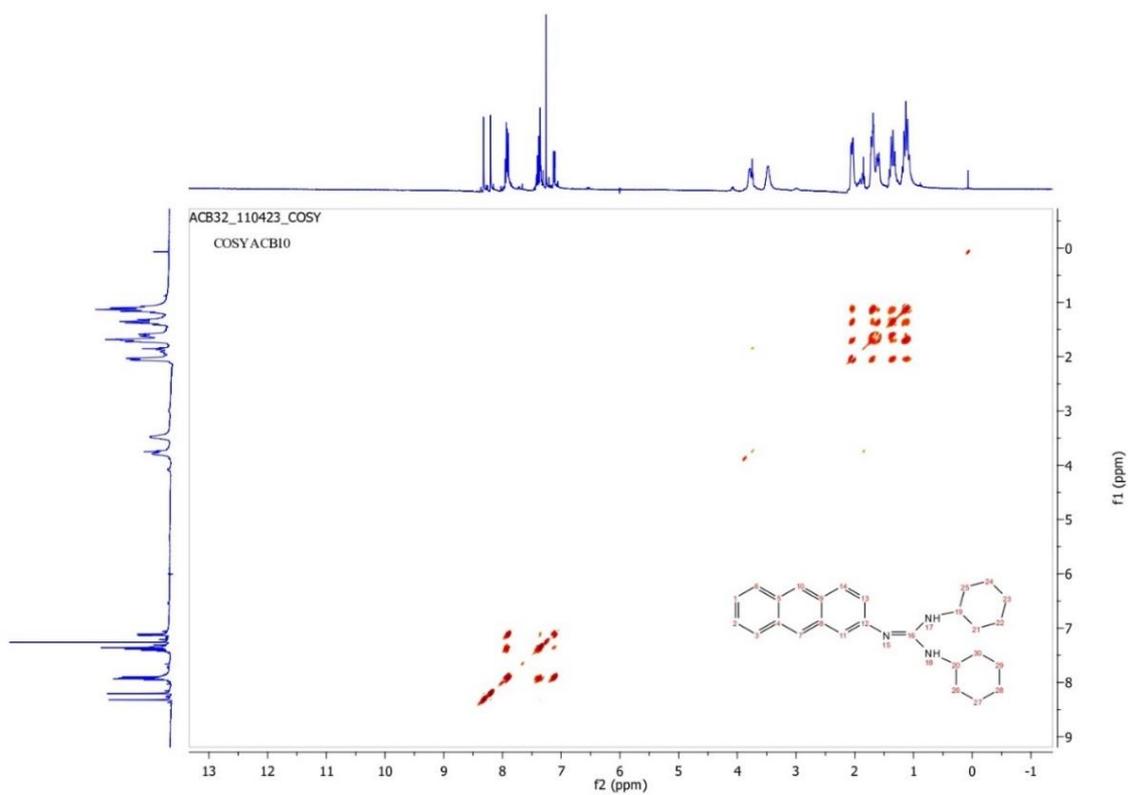
# <sup>13</sup>C-NMR of ACB10 in CDCl<sub>3</sub>



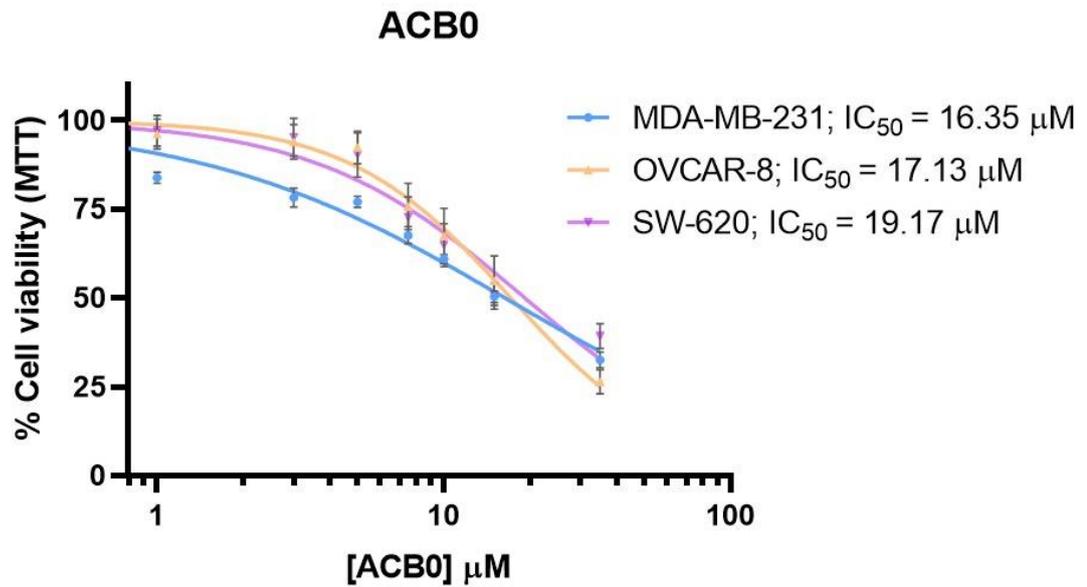
Multinuclear 2D ( $^1\text{H}$ - $^{13}\text{C}$  HSQC) of *ACB10*.



Multinuclear 2D ( $^1\text{H}$ - $^1\text{H}$  COSY) of *ACB10*



**Figure S2.**  $^1\text{H}$ -NMR,  $^{13}\text{C}$ -NMR, multinuclear 2D ( $^1\text{H}$ - $^{13}\text{C}$  HSQC) and multinuclear 2D ( $^1\text{H}$ - $^1\text{H}$  COSY) NMR spectra of *ACB10*.

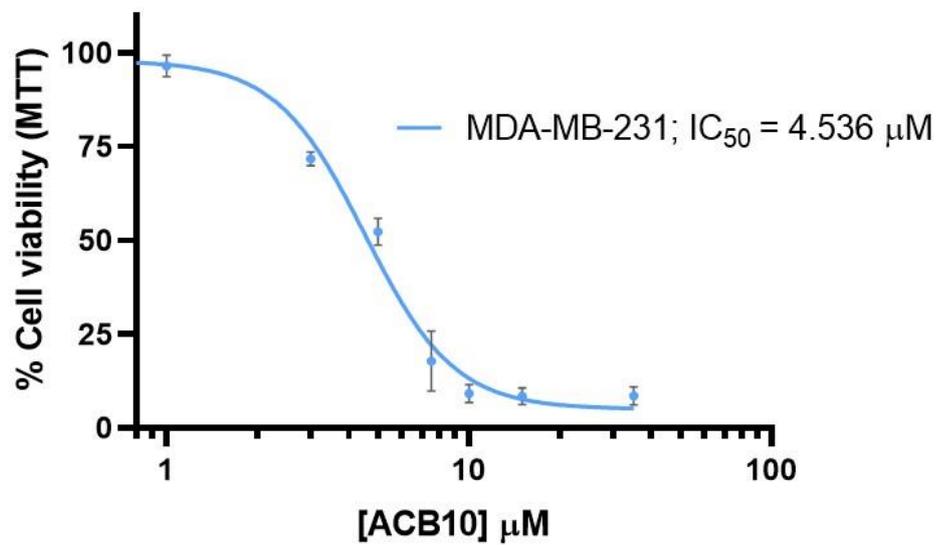


**Figure S3.** Antiproliferative effect induced by different concentrations of *ACB0* in the indicated cell lines at 72 h tested by MTT assay. Calculation of the concentrations  $IC_{50}$  based on a nonlinear regression curve fit (log(inhibitor) vs. normalized response-variable slope). The mean of three independent experiments (each in triplicate) is plotted together with the standard error (SEM).



Figure S4: Correlations between ADME parameters.

## ACB10



**Figure S5:** Antiproliferative effect induced by different concentrations of *ACB10* fluorescent guanidine in the MDA-MB-231 cell line at 72 h tested by MTT assay. The inhibitory concentration 50 (IC<sub>50</sub>) for MDA-MB-231 cell line is shown. Calculation of the concentrations IC<sub>50</sub> based on a nonlinear regression curve fit (log(inhibitor) vs. response-variable slope). The mean of three independent experiments (each in triplicate) is plotted together with the standard error (SEM).