

## SUPPLEMENTAL INFORMATION

### **The Dynamic Character of the *BCL2* Promoter i-Motif Provides a Mechanism for Modulation of Gene Expression by Compounds That Bind Selectively to the Alternative DNA Hairpin Structure**

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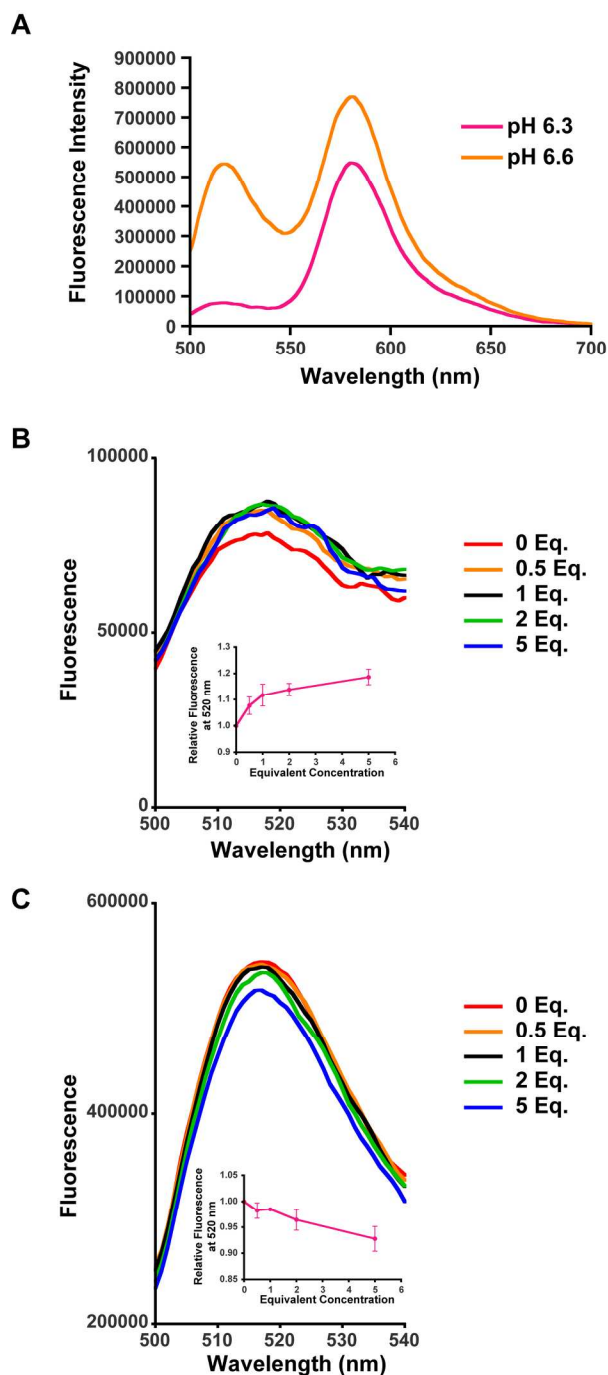
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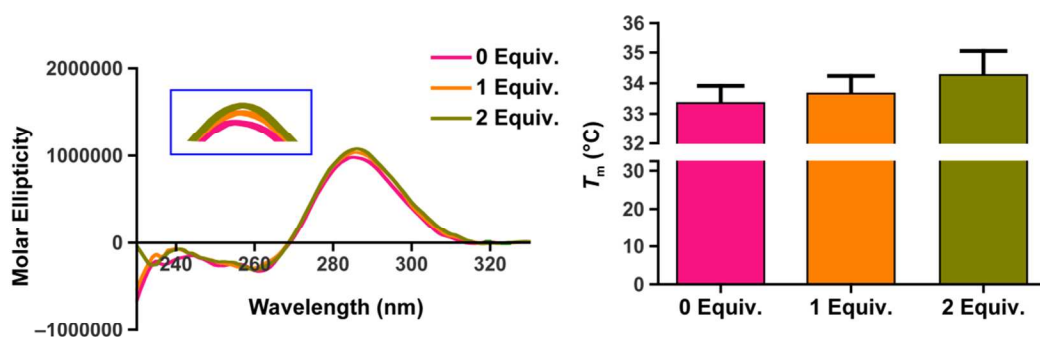
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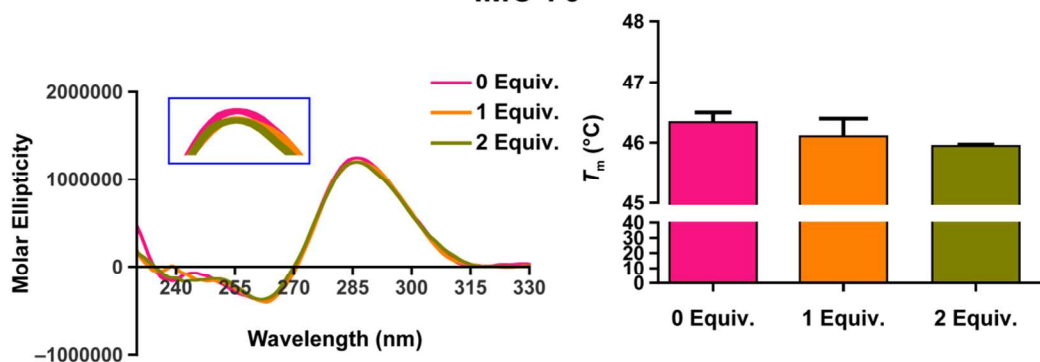


**Supplemental Figure 1.** Fluorescence spectra analysis to confirm the conformational change of the *BCL2* i-motif produced by IMC-76 and IMC48. (A) pH-dependent FRET changes in the *BCL2* i-motif labeled with FAM at the 5'-end and TAMRA at the 3'-end of the oligomer. At low pH (pH 6.3), the  $I_{580}/I_{520}$  ratio is 7.45, showing a closer distance between two dyes by folding the i-motif, while at high pH (pH 6.6) the  $I_{580}/I_{520}$  ratio is 1.45, showing a longer distance between two dyes by unfolding the i-motif. (B) Dose-dependent spectra change by IMC-76 at pH 6.3,  $I_{580}$ . (C) Dose-dependent spectra change by IMC-48 at pH 6.6,  $I_{580}$ . Insets in (B) and (C) show a graph of the dose response from the spectral analysis at  $I_{580}$ . A PTI fluorometer was used to measure the spectra.

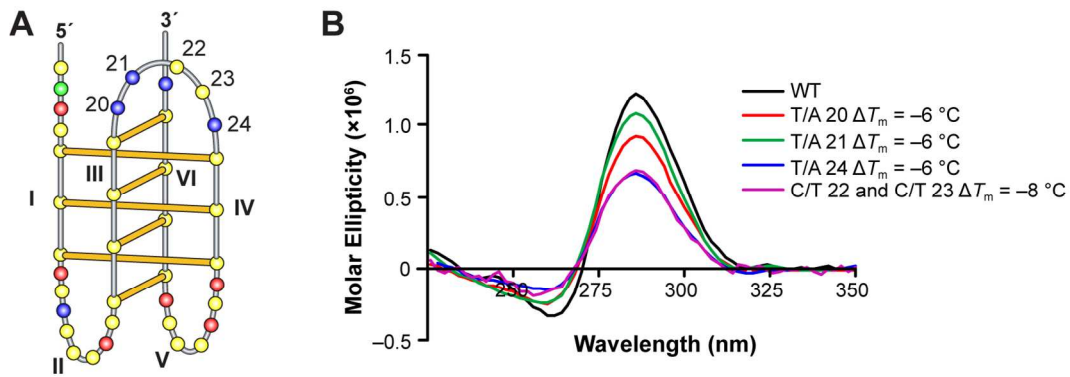
### IMC-48



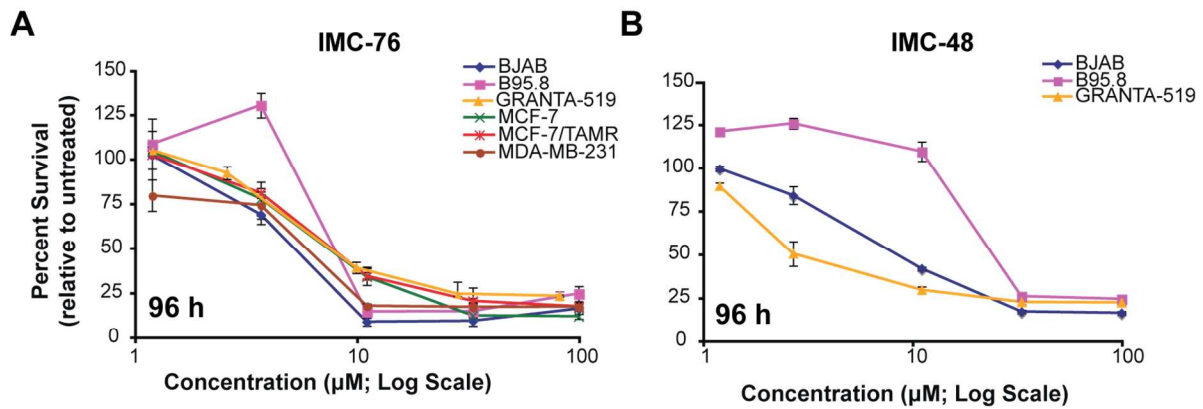
### IMC-76



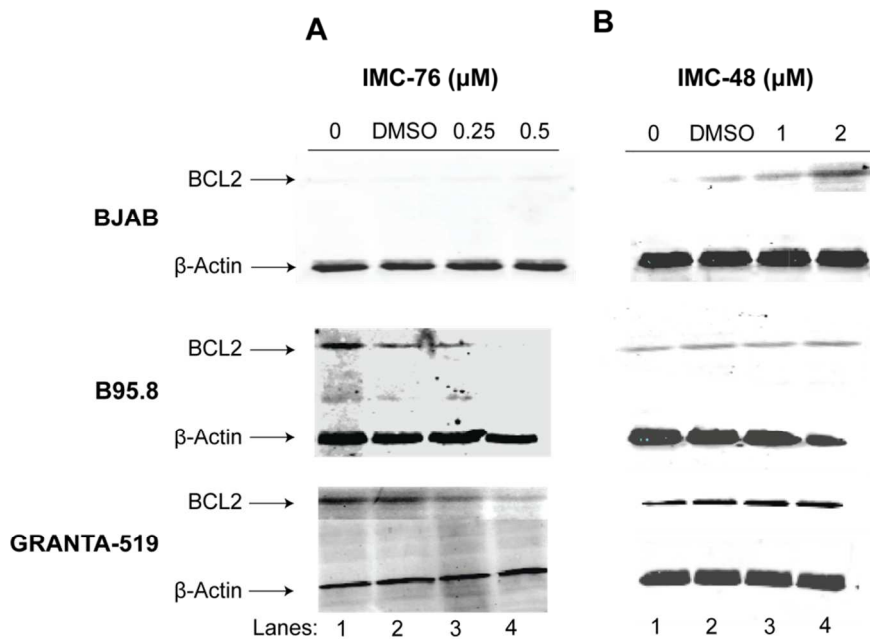
**Supplemental Figure 2.** CD spectra of the BCL2 i-motif with IMC-48 and IMC-76. IMC-48 and IMC-76 were incubated with the DNA at 50 mM Na cacodylate buffer (pH 6.6 and pH 6.3, respectively). The ellipticity at 286 nm was monitored with 1 °C/min of temperature increase for melting analysis, and then sigmoidal curve fitting was used to calculate  $T_m$ . DMSO was treated as a control.



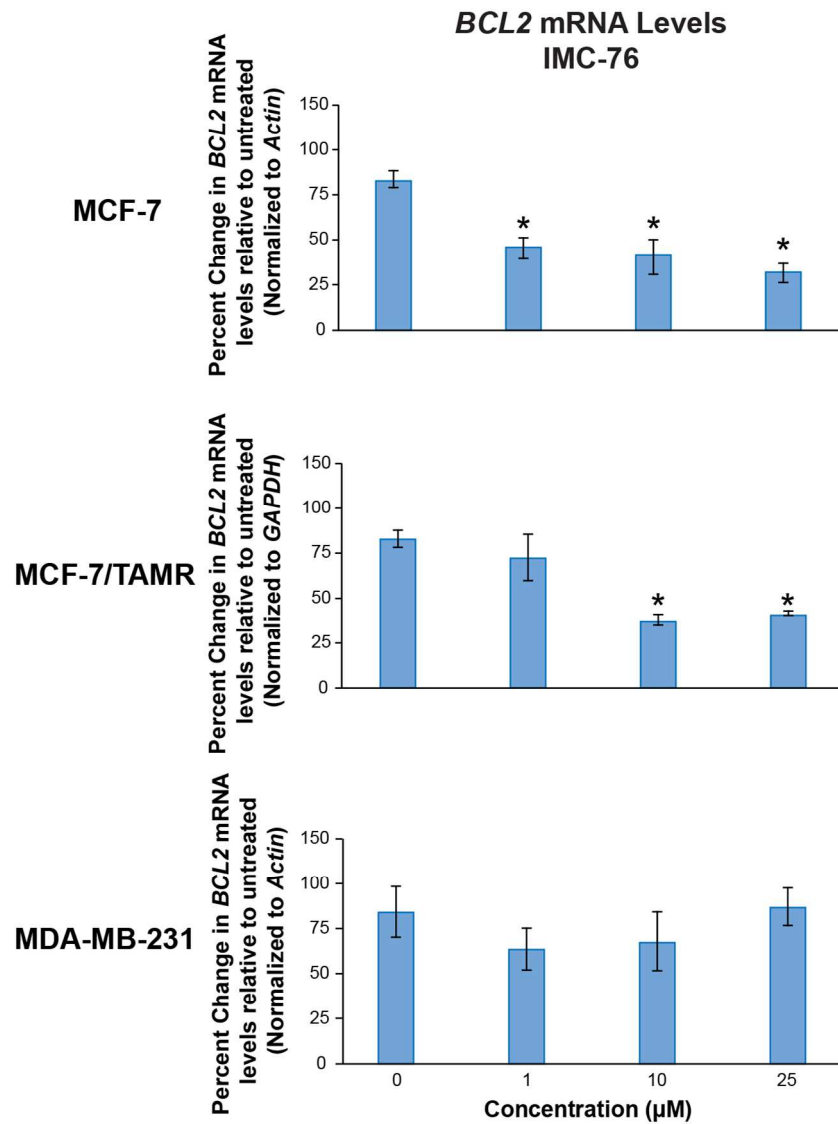
**Supplemental Figure 3.** CD spectral and thermal stability analyses of capping mutants. (A) Folding pattern of the *BCL-2* i-motif with the capping structure nucleotides labeled. (B) CD spectra of the *BCL-2* i-motif capping mutants C22, C23, T20, T21, and T24 in comparison to the wild-type sequence.



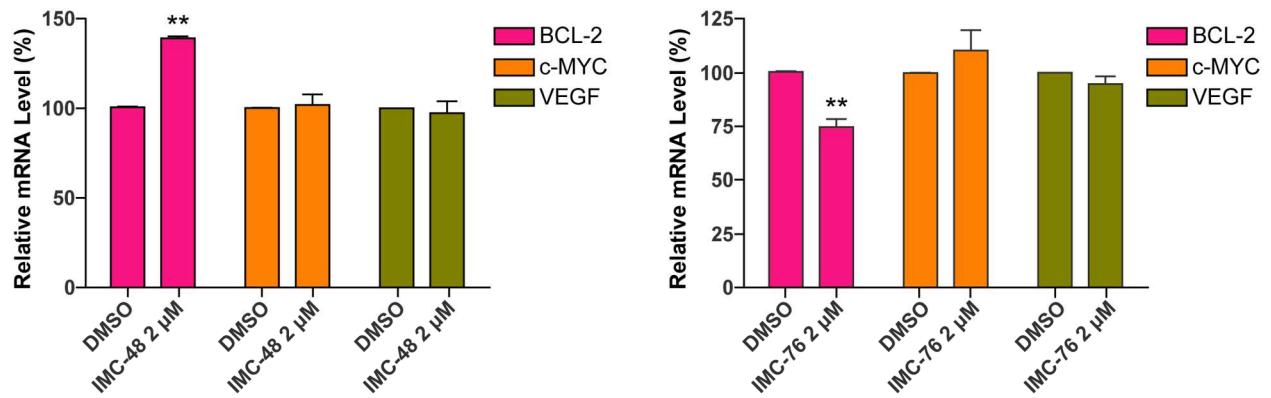
**Supplemental Figure 4.** Effect of IMC-76 and IMC-48 on cell toxicity in lymphoma and breast cancer cell lines. Percent survival determined by the MTS cytotoxicity assay in response to treatment with IMC-76 (A) for BJAB, B95.8, and GRANTA-519 lymphoma and MCF-7, MCF-7/TAMR, and MDA-MB-231 breast cancer cell lines or IMC-48 (B) for BJAB and B95.8 lymphoma cell lines at 96 h. Percent survival was calculated relative to untreated controls from three independent experiments.



**Supplemental Figure 5.** Effect of IMC-76 and IMC-48 on BCL-2 protein levels in lymphoma cell lines. Western blot analysis of BCL-2 protein levels in BJAB, B95.8, and GRANTA-519 cell lines following 24 h treatment with IMC-76 (A) or IMC-48 (B). In both (A) and (B), lanes 1 and 2, represent untreated and DMSO vehicle cell lysates, respectively. In (A), lanes 3 and 4 contain lysates from cells treated with 0.25 and 0.5 μM IMC-76, respectively. In (B), lanes 3 and 4 contain lysates from cells treated with 1 and 2 μM IMC-48, respectively. The western blots are representative of three independent experiments with β-actin as a loading control.

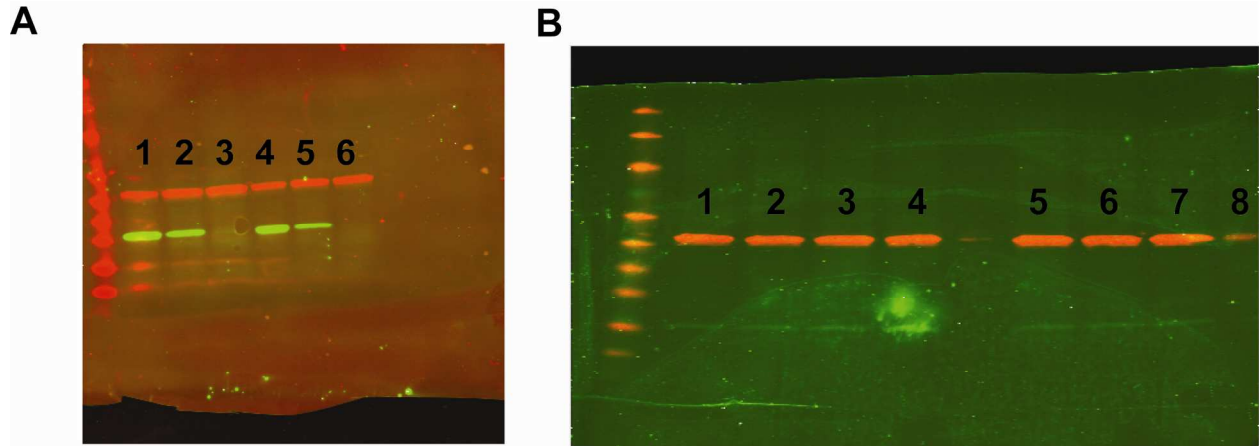


**Supplemental Figure 6.** Effect of IMC-76 on *BCL2* mRNA levels in breast carcinoma cell lines. The *BCL2* mRNA levels as determined by qPCR for MCF-7, MCF-7/TAMR and MDA-MB-231 breast carcinoma cell lines following 24 h treatment with IMC-76. Percent change in mRNA levels were calculated relative to untreated controls. MCF-7 and MDA-MB-231 *BCL-2* mRNA levels were normalized to  $\beta$ -actin and MCF-7/TAMR levels were normalized to *GAPDH*. \* $P \leq 0.02$ .



**Supplemental Figure 7.** Effect of IMC-48 and IMC-76 on the mRNA levels of BCL2, c-MYC, and VEGF. A BJAB cell line with a low basal level of BCL2 mRNA (left panel) and an MCF-7 cell line with a high basal level of BCL2 mRNA (right panel) were used for IMC-48 and IMC-76, respectively. Cells were treated for 24 h with compounds and DMSO served as a control. Relative mRNA level was determined by normalization of Ct values to GAPDH and the DMSO by qPCR assay.



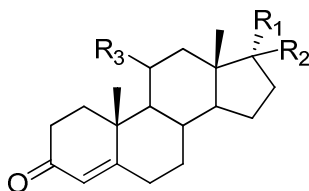


**Supplemental Figure 8.** Uncut western blot gels. (A) Western blot gel of basal BCL2 protein levels from GRANTA-519 (lanes 1 and 4), B95.8 (lanes 2 and 5), and BJAB (lanes 3 and 6) cell line lysates shown in Figure 5B. (B) Western blot gel of BJAB cells following 24 h treatment with IMC-76 from Supplemental Figure 5. Lanes 1 and 5 and lanes 2 and 6 represent untreated and DMSO vehicle cell lysates, respectively. Lanes 3 and 7 and lanes 4 and 8 contain lysates from cells treated with 0.25 and 0.5  $\mu$ M IMC-76, respectively. The lysates represent two of three independent experiments with  $\beta$ -actin as a loading control.

**Supplemental Table 1.** Sequences of *BCL2* FRET and NdU probes.

<b>DNA Probe</b>	<b>Sequence (5' → 3')</b>
<i>BCL2</i> i-motif for HTS	FAM-CAG <b>CCCCGCTCCCGCCCCCTTCCTCCCGCGCCCGCCCCT</b> - BHQ
<i>BCL2</i> i-motif	FAM-CAG <b>CCCCGCTCCCGCCCCCTTCCTCCCGCGCCCGCCCCT</b> - TAMRA
<i>BCL2</i> i-motif mutant	FAM -CAG <b>TTTTGCTCCCGCTTTC</b> TTCCT <b>TTT</b> GCG <b>CCCGCCCCT</b> - TAMRA
<i>BCL2</i> G-quadruplex	FAM-AG <b>GGGCGGGCGCGGGAGGAAGGGGGCGGGAGCGGGGCTG</b> - TAMRA
<i>c-MYC</i> i-motif	FAM- TCCCCACCTTCCCCACCCTCCCCACCCTCCCCA-TAMRA
<i>VEGF</i> i-motif	FAM-CTCCG <b>CCCCGCCGGGACCCCGCCCCCGGCCCGCCCC</b> -TAMRA
20T	CAG <b>CCCCGCTCCCGCCCCC</b> Nd <b>UTCCTCCCGCGCCCGCCCCT</b>
21T	CAG <b>CCCCGCTCCCGCCCCCT</b> Nd <b>UCCTCCCGCGCCCGCCCCT</b>
24T	CAG <b>CCCCGCTCCCGCCCCCTTCC</b> Nd <b>UCCCGCGCCCGCCCCT</b>
39T	CAG <b>CCCCGCTCCCGCCCCCTTCCTCCCGCGCCCGCCCC</b> Nd <b>U</b>

**Supplemental Table 2.** Structures and FRET values for steroidal compounds from the NCI Diversity Set and ChemDiv library.



Compound #	R <sub>1</sub>	R <sub>2</sub>	R <sub>3</sub>	% Increase in FRET value compared to probe (set to 100 %) (10 equivalent compared to DNA)
IMC-76	-	-	-	270
IMC-48	-	-	-	52
5829-8948	-OH	-H	-H	200.3
N050-0008	-OCOCH <sub>3</sub>	-H	-H	163.0
N050-0013	-OH	-COCH <sub>3</sub>	-H	213.0
N050-0014	-OH	-COCH <sub>2</sub> OH	-H	177.3
5776-0002	-OCOCH <sub>3</sub>	-COCH <sub>2</sub> OCOCH <sub>3</sub>	-H	234.6
N050-0017	-OH	-COCH <sub>2</sub> OH	=O	190.4
N056-0002	-OCOCH <sub>3</sub>	-COCH <sub>2</sub> OH	-OH	183.0
N056-0003	-OCOCH <sub>3</sub>	-COCH <sub>2</sub> OCOCH <sub>3</sub>	-OH	181.0

**Analysis of structure and FRET values:** IMC-76 was identified from the NCI Diversity Set and shows a FRET value of 270. It is a steroidal molecule predominantly substituted on ring D at C17. Screening of similar steroidal molecules from the ChemDiv collection resulted in identification of compounds that exert a similar effect on FRET: an increase indicating destabilization of the BCL2 i-motif. Although definite conclusions cannot be drawn about structure–activity relationships (SAR), activity trend was observed for steroidal compounds with C17 substitution. A detailed SAR would require design, synthesis, and testing of analogs; hence this would be a part of future studies on these compounds.

**Supplemental Table 3.** IMC-76 and IMC-48 IC<sub>50</sub> values at 96 h for breast carcinoma and lymphoma cell lines as determined by the MTS cytotoxicity assay.

Cell Line	IC <sub>50</sub> (μM)	
	IMC-76 (96 h)	IMC-48 (96 h)
MCF-7	9.4	ND
MCF-7/TAMR	9.4	ND
MDA-MB-231	5.8	ND
BJAB	5.2	8.4
B95.8	7.9	24.2
GRANTA-519	8.4	2.7

ND: not determined