

1 Regulation of the ryanodine receptor by anti-apoptotic Bcl-2 is
2 independent of its BH3-domain-binding properties

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18

19 **Abstract**

20 The regulation of intracellular Ca^{2+} signaling is an important aspect of how anti-apoptotic B-cell
21 lymphoma 2 (Bcl-2) proteins regulate cell death and cell survival. At the endoplasmic reticulum (ER)
22 the Bcl-2 homology (BH) 4 domain of Bcl-2 is known to bind to and inhibit both inositol 1,4,5-
23 trisphosphate receptors (IP_3Rs) and ryanodine receptors (RyRs). Besides this, drugs that target the
24 hydrophobic cleft of Bcl-2 have been reported to deplete ER Ca^{2+} stores in an IP_3R - and RyR-
25 dependent way. This suggests that the hydrophobic cleft of Bcl-2 may also be involved in regulating
26 these ER-located Ca^{2+} -release channels. However, the contribution of the hydrophobic cleft on the
27 binding and regulatory properties of Bcl-2 to either IP_3Rs or RyRs has until now not been studied.
28 Here, the importance of the hydrophobic cleft of Bcl-2 in binding to and inhibiting the RyR was
29 assessed by using a genetic approach based on site-directed mutagenesis of Bcl-2's hydrophobic cleft
30 and a pharmacological approach based on the selective Bcl-2 hydrophobic cleft inhibitor, ABT-199.
31 Both binding assays and single-cell Ca^{2+} measurements indicated that RyR binding and the inhibition
32 of RyR-mediated Ca^{2+} release by Bcl-2 is independent of its hydrophobic cleft.

33 **Keywords:** B-cell lymphoma 2, ryanodine receptors, hydrophobic cleft, BH3 mimetic, ABT-199, P2A

34

35 **Abbreviations:** Bcl-2-associated death promoter (Bad), Bcl-2 homologous antagonist killer (Bak),
36 Bcl-2-associated X protein (Bax), B-cell lymphoma 2- (Bcl-2), Bcl-2 homology (BH), endoplasmic
37 reticulum (ER), erythropoietin (Epo), inositol 1,4,5-trisphosphate receptor (IP_3R), mammalian protein-
38 protein interaction trap (MAPPIT), ryanodine receptor (RyR), sarco-endoplasmic reticulum Ca^{2+}
39 ATPase (SERCA)

40 **Highlights**

- 41 • **Bcl-2's hydrophobic cleft is not involved in binding to and regulating RyRs**
- 42 • **The BH3 mimetic ABT-199 does not alter Bcl-2's ability to regulate RyRs**
- 43 • **Bcl-2-P2A-mCherry can efficiently be used in single-cell Ca^{2+} measurements**

44 **Introduction**

45 The modulation of intracellular Ca^{2+} signalling is an important aspect of how proteins of the B-cell
46 lymphoma 2 (Bcl-2)-family regulate cell death and cell survival [1, 2, 3, 4]. The three C-terminally
47 located Bcl-2 homology (BH) domains (BH1, BH2 and BH3) of anti-apoptotic Bcl-2 family members
48 (such as Bcl-2 and Bcl- X_L) form a hydrophobic cleft, which is used to bind to the BH3 domains
49 present in the pro-apoptotic Bcl-2-family members, thereby inhibiting their function [5]. The N-
50 terminally located BH4 domain also critically determines its anti-apoptotic function in at least two
51 ways: (i) by binding to and inhibiting the pro-apoptotic, multi-domain protein Bcl-2-associated X
52 protein (Bax), which upon activation oligomerizes and permeabilizes mitochondrial outer membranes
53 [6], and (ii) by binding to and inhibiting the inositol 1,4,5-trisphosphate (IP_3) receptor (IP_3R), an
54 endoplasmic reticulum (ER)-located Ca^{2+} -release channel, thereby preventing mitochondrial Ca^{2+}
55 overload [7].

56 We recently showed that, similarly to IP_3Rs , anti-apoptotic Bcl-2 via its BH4 domain also targets
57 another important ER-located family of Ca^{2+} -release channels: the ryanodine receptors (RyRs) [8].
58 The BH4 domain of Bcl-2 was sufficient to interact with and suppress RyR-mediated Ca^{2+} release. For
59 Bcl- X_L the situation was different. In contrast to the IP_3R , where the BH4 domain of Bcl- X_L does not
60 bind to and inhibit the channel [9], the BH4 domain of Bcl- X_L is able to bind to and inhibit RyRs [10].
61 However, in the full-length Bcl- X_L Lys87 located in the BH3 domain, which is part of the
62 hydrophobic cleft, also plays a role in targeting Bcl- X_L to the RyR. This suggests that the hydrophobic
63 cleft of the anti-apoptotic Bcl-2 proteins may also play a role in targeting them to the RyR. In addition
64 to this, drugs that target the hydrophobic cleft of Bcl-2 and thus compete with pro-apoptotic Bcl-2
65 family members have been reported to deplete the ER Ca^{2+} stores in pancreatic acinar cells [11]. As
66 these cells express both IP_3Rs and RyRs [12], it was suggested that dissociation of the pro-apoptotic
67 Bcl-2 family members from the anti-apoptotic family members by these Bcl-2 antagonists may
68 increase the sensitivity of the IP_3R and RyR to Ca^{2+} -induced Ca^{2+} release [11]. However, it was not
69 investigated whether these compounds effectively altered the binding of Bcl-2 to either of these ER-
70 located Ca^{2+} -release channels. Furthermore, the Bcl-2 antagonists used in this study are not selective

71 Bcl-2 inhibitors, thereby affecting other proteins. For instance, the Bcl-2 antagonist HA14-1 also
72 inhibits the activity of the sarco-/endoplasmic reticulum Ca^{2+} ATPase (SERCA), thereby causing ER
73 Ca^{2+} -store depletion [13, 14].

74 Therefore, we here assessed the involvement of the hydrophobic cleft of Bcl-2 in binding to and
75 inhibiting the RyR by using both site-directed mutagenesis and a novel specific Bcl-2 inhibitor, ABT-
76 199. This analysis shows that RyR binding and inhibition by Bcl-2 is independent of its hydrophobic
77 cleft properties.

78 **Materials and methods**

79 **Chemicals, antibodies and peptides**

80 Unless otherwise mentioned all chemicals were obtained from Sigma-Aldrich. The antibodies used in
81 this study include: HRP conjugated anti-FLAG M2 antibody (Sigma-Aldrich), mouse monoclonal
82 anti-RyR antibody 34C (Thermo Scientific, Rockford, IL, USA, or Developmental Studies Hybridoma
83 Bank, University of Iowa, USA) and rabbit polyclonal anti-Bax (Santa Cruz). The peptide used for
84 elution of the FLAG pull-downs was MDYKDHDGDYKDHDIDYKDDDDK (Life Tein).

85

86 **Plasmids and constructs**

87 The used restriction enzymes were obtained from New England BioLabs. 3XFLAG-tagged Bcl-2
88 (3XFLAG-Bcl-2) was obtained as described previously [9]. PCR site-directed mutagenesis using the
89 3XFLAG-Bcl-2 construct as template (primers forward:
90 5'ATCCCCATGGCAGCAGTAGATCAAGCGCTGAGGGAGGCA3', and reverse:
91 5'TGCCTCCCTCAGCGCTTGATCTACTGCTGCCATGGGGAT3') was performed to obtain the
92 3XFLAG-Bcl-2 G145A R146A (3XFLAG-Bcl-2^{GR/AA}) double mutant. The pCMV24 vector encoding
93 the P2A-mCherry or 3XFLAG-Bcl-2-P2A-mCherry were obtained as follows. A P2A-encoding DNA
94 sequence (corresponding to amino acid sequence GSGATNFSLLKQAGDVEENPGP) [15] was
95 obtained as a duplex with overhanging ends corresponding to cleaved EcoRI and BglII restriction sites
96 forward:

97 5'AATTCGGATCCGGAGCCACGAACTTCTCTCTGTAAAGCAAGCAGGAGACGTGGAAGA
98 AAACCCTGGTCCTA3', and reverse
99 5'GATCTAGGACCAGGGTTTTCTCCACGTCTCCTGCTTTAACAGAGAGAAGTTCGT
100 GGCTCCGGATCCG3'. The duplex was ligated into the empty pCMV24 vector utilizing EcoRI and
101 BglII as restriction sites. After removing the BglII site from the mCherry construct via site-directed
102 mutagenesis (primers forward: 5'GGACGAGCTGTACAAGAGGTCTCGAATCACAAGTTTG3',
103 and reverse: 5'CAAACCTTGTGATTCGAGACCTCTTGACAGCTCGTCC3'), mCherry was cloned

104 at the 3' side of the P2A sequence (primers forward:
105 5'GCGGCGAGATCTGTGAGCAAGGGCGAGGAGGAC3', and reverse:
106 5'GCGGCGGTCGACTTACGTTTCTCGTTCAGC3') utilizing the BglII and SalI restriction
107 enzymes. At the 5' side of the P2A sequence, Bcl-2 was cloned (primers forward:
108 5'GCGGCGGCGGCCGCAGCGCACGCTGGGAGAAC3', and reverse:
109 5'GCGGCGGAATTCCTTGTGGCCCAGATAGGCAC3') utilizing the NotI and EcoRI restriction
110 enzymes and the 3XFLAG-Bcl-2 plasmid as template. One nucleotide was added just before the
111 multiple cloning site in order to put the pCMV24-P2A-mCherry construct in the correct open reading
112 frame (primers forward: 5'GGATGACGATGAGCAAGCTTGCGGC3', and reverse:
113 5'GCCGCAAGCTTGCTCATCGTCATCC3').

114

115 **Cell culture and transfections**

116 All media used in this study were obtained from Life Technologies. HEK293T cells, and HEK293
117 cells stably overexpressing RyR3 (HEK RyR3) were cultured as previously [10].

118 One day after seeding, the 3XFLAG-Bcl-2 or 3XFLAG-Bcl-2^{GR/AA} construct were introduced into
119 HEK RyR3 cells utilizing the X-tremeGENE HP DNA transfection reagent (Roche) according to the
120 manufacturer's protocol. For single-cell Ca²⁺ measurements two different methods for introducing
121 mCherry, as selection marker for transfected cells, into the cells were used. First, 48 hours after
122 seeding the pCMV24 vector (negative control), 3XFLAG-Bcl-2 or the 3XFLAG-Bcl-2^{GR/AA} construct
123 were transfected together with a pcDNA 3.1(-) mCherry expressing vector as a selection marker at a
124 3:1 ratio as previously described [10]. Second, 48 hours after seeding, 0.25 µg of a pCMV24 vector
125 encoding the P2A-mCherry or a pCMV24 vector encoding 3XFLAG-Bcl-2-P2A-mCherry was
126 introduced into the HEK RyR3 cells utilizing the X-tremeGENE HP DNA transfection reagent. All
127 single-cell Ca²⁺ measurements were performed two days after the transfection.

128

129 **Co-immunoprecipitations, FLAG pull-downs and immunoblots**

130 Co-immunoprecipitation experiments were performed utilizing a co-immunoprecipitation kit (Thermo
131 Scientific) in HEK RyR3 cells as described previously [10]. The negative control IgG was obtained
132 from Thermo Scientific. Cell lysis and wash steps were performed with a CHAPS-based buffer (50
133 mM Tris-HCl pH 7.5, 100 mM NaCl, 2 mM EDTA, 50 mM NaF, 1 mM Na₃VO₄, 1% CHAPS and
134 protease inhibitor tablets (Roche)). When indicated, 3 μM of ABT-199 (Active Biochem) was added
135 during the overnight incubation. In the control co-immunoprecipitations an equal amount of the
136 vehicle (DMSO) was included. For FLAG pull-downs, 300 μg of HEK RyR3 cells transfected with the
137 empty vector, 3XFLAG-Bcl-2 or 3XFLAG-Bcl-2^{GR/AA} were lysed (10 mM HEPES pH 7.5, 0.25 %
138 Nonidet P-40, 142 mM KCl, 5 mM MgCl₂, 2 mM EDTA, 1 mM EGTA and protease inhibitor tablets)
139 and incubated for 2 hours at 4°C with 30 μl of anti-DYKDDDDK (L5)-tag agarose (Biolegend). After
140 extensive washing (50 mM Tris pH 8, 150 mM NaCl, 1.0% NP-40, 0.5% sodium deoxycholate, 0.1%
141 sodium dodecyl sulfate, and protease inhibitor tablets) elution was performed by incubating the resin
142 for 30 min at 15°C with FLAG elution peptide (250 μg/ml 50 mM Tris-HCl pH 7.4, 150 mM NaCl).
143 After the elution, the eluate was collected and 4X LDS (Life Technologies) supplemented with 1/200
144 β-mercaptoethanol. Samples were boiled (95°C) for 5 min just before SDS-PAGE on either NuPAGE
145 4-12% Bis-Tris gels or NuPAGE 3-8% Tris-Acetate gels (both from Life Technologies) when
146 RyRs were visualized. gels. Immunoblot analysis was performed as previously [10].

147

148 **Mammalian protein-protein interaction trap (MAPPIT)**

149 Bcl-2-associated death promoter (Bad), and Bcl-2 were cloned in the pSEL+2L bait vector, TRIP13
150 and Bcl-2 lacking its transmembrane domain were cloned in the pMG1 or pMG2 prey vector,
151 respectively [16]. Bad and TRIP13 entry clones are from the ORFeome8.1 collection received from
152 the Vidal lab (CCSB, Boston, USA), and were cloned as bait or prey, respectively using the Gateway
153 recombination technology as described by the manufacturer (Life Technologies). Using the pCAGGS-
154 Bcl-2 plasmid, obtained from BCCM/LMBP (Gent, Belgium) as template, Bcl-2 was amplified by
155 PCR using the following primers, forward: 5'GCGGAATTCATGGCGCACGCTGGGAGA3', and

156 reverse: 5'CGCGCGGCCGCTCACTTGTGGCCCAGATAGG3', and was cloned in the pMG2 prey
157 vector using the restriction enzymes EcoRI and NotI. Bcl-2 was cloned in the bait vector, by EcoRI
158 digestion of the above described Bcl-2 prey vector, made blunt by nucleotide fill-in, and followed by a
159 NotI restriction digest, and SalI digestion of the pSEL+2L bait vector, made blunt by nucleotide fill-in
160 and subsequent NotI digestion. The Bcl2 prey lacking its transmembrane domain was obtained by
161 PCR amplification on the Bcl-2 prey construct using the same forward primer as described above and
162 the following reverse primer: 5'CGCGCGGCCGCTCAGGAGAAATCAAACAGAGGCC3', and was
163 cloned in the pMG2 prey vector using the restriction enzymes EcoRI and NotI.

164 MAPPIT experiments were performed as previously [10], with minor changes. Briefly, HEK293T
165 cells were seeded in 96-well plates. Six wells per condition were transfected with the different
166 combinations of bait, prey and reporter plasmid (rPAP1-luci) using the calcium phosphate method.
167 The next day, a serial dilution of the ABT-199 compound was added and 4 hours later half of the wells
168 were stimulated with 5 ng/ml erythropoietin (Epo) while the other half were left untreated. One day
169 later the cells were lysed and after the addition of substrate the luciferase activity was determined
170 using a luminometer. The fold induction was obtained by dividing the average value of the stimulated
171 cells by the average value of the non-stimulated cells.

172

173 **Single-cell Ca²⁺ measurements and fluorescence microscopy**

174 Single-cell Ca²⁺ measurements were performed as previously [8]. mCherry was used as selection
175 marker for the transfected cells. When used, ABT-199 or an equal volume of the vehicle (DMSO) was
176 included during the loading and de-esterification (1 hour in total) of the Fura2-AM in the HEK RyR3
177 cells. A Zeiss Axio Observer Z1 Inverted Microscope equipped with a 20x air objective and a high-
178 speed digital camera (Axiocam Hsm, Zeiss) was used for imaging mCherry expression.

179

180 **Statistical analysis**

181 Repeated measure ANOVA tests were performed. * indicates significantly different results (p<0.05).

182 Exact p-values are indicated in the figure legends. NS indicates non-significant.

183 **Results and discussion**

184 The hydrophobic cleft of Bcl-2 has been extensively characterized with respect to its binding to BH3
185 domains of the pro-apoptotic Bcl-2 family members. Mutations disrupting the hydrophobic cleft of
186 Bcl-2 and thereby abolishing binding of Bcl-2 to Bax and Bcl-2 homologous antagonist killer (Bak)
187 have been described [17]. Using a Bcl-2 hydrophobic cleft double mutant (G145A and R146A (Bcl-
188 2^{GR/AA})), we aimed to elucidate whether or not this region is involved in regulating RyRs.

189 We first validated that the Bcl-2^{GR/AA} mutant was unable to bind the pro-apoptotic protein Bax by
190 performing FLAG-pull-down assays in HEK RyR3 cells transiently overexpressing the empty vector
191 (pCMV24), 3XFLAG-Bcl-2 or the 3XFLAG-Bcl-2^{GR/AA} mutant (Fig. 1A). The 3XFLAG-tagged
192 constructs were pulled down (Fig 1A, top) after which the presence of endogenous Bax was assessed.
193 Bax specifically co-immunoprecipitated with 3XFLAG-Bcl-2, while it was absent in empty vector
194 control samples. Furthermore, in contrast to wild-type 3XFLAG-Bcl-2 the 3XFLAG-Bcl-2^{GR/AA}
195 mutant could not pull-down endogenous Bax (Fig. 1A, bottom), indicating that these mutations disrupt
196 the hydrophobic cleft as reported. Next, co-immunoprecipitation experiments were set up in order to
197 verify whether the hydrophobic cleft of Bcl-2 is involved in binding to RyR3. RyR3 was
198 immunoprecipitated from HEK RyR3 cells (Fig. 1B, top) transiently overexpressing 3XFLAG-Bcl-2
199 or 3XFLAG-Bcl-2^{GR/AA}. Strikingly, mutating the hydrophobic cleft of Bcl-2 resulted in an increased
200 binding to RyR3 (Fig. 1B, bottom). This might indicate that more of 3XFLAG-Bcl-2^{GR/AA} may be
201 available to bind to RyR3 due to its impaired recruitment to pro-apoptotic Bcl-2-family members.

202 Finally, Fura2-AM single-cell Ca²⁺ measurements were set up to assess whether disruption of Bcl-2's
203 hydrophobic cleft affected its ability to inhibit RyR3-mediated Ca²⁺ release. In these experiments, the
204 empty vector (pCMV24), 3XFLAG-Bcl-2 or the 3XFLAG-Bcl-2^{GR/AA} mutant were transfected in HEK
205 RyR3 cells at a 3:1 ratio with mCherry. Only mCherry-positive cells were included in these
206 measurements. RyR-mediated Ca²⁺ release was evoked with caffeine (1.5 mM) after chelating
207 extracellular Ca²⁺ with BAPTA (3 mM). An average Ca²⁺ trace of a typical experiment is shown in
208 Fig. 2A whereas the summary of all performed Ca²⁺ measurements is presented in Fig. 2B. From this it
209 was clear that, Bcl-2 mutations that disrupt the hydrophobic cleft did not affect the ability of Bcl-2 to
210 inhibit RyR3. This suggests that the hydrophobic cleft of Bcl-2 is not involved in targeting RyR3.

211 To further substantiate our findings, an independent pharmacological approach was performed,
212 utilizing a BH3-mimetic drug which occupies the hydrophobic cleft of Bcl-2. In recent years, several
213 advances have been made in generating highly specific BH3-mimetic drugs [18]. One of the most
214 recent compounds, ABT-199, was designed to have a high specificity for Bcl-2 over Bcl-X_L thereby
215 limiting side effects generated by unwanted Bcl-X_L inhibition [18]. In this study we used this ABT-
216 199 compound in order to independently assess the importance of the hydrophobic cleft of Bcl-2 for
217 RyR interaction and regulation by Bcl-2.

218 First, we assessed the efficiency of the compound in disrupting the binding of Bcl-2 to the pro-
219 apoptotic Bcl-2 family members. For this we utilized an *in cellulo* protein-protein interaction assay,
220 MAPPIT [10, 19]. In this technique, functional complementation of a chimeric cytokine receptor
221 serves as a read out for the binding of two proteins. The first binding partner, Bad (bait) was fused to
222 the membrane bound chimeric cytokine receptor consisting of the extracellular domain of the Epo
223 receptor fused to the transmembrane and mutated cytosolic part of the leptin receptor. The second
224 binding partner, Bcl-2 lacking its transmembrane domain (prey) was fused to a part of the glycoprotein
225 130 receptor which can trigger Epo-dependent STAT signaling when binding to the bait occurs. The
226 latter is monitored via a luciferase reporter assay driven by a STAT-sensitive promoter. As expected,
227 Bad could bind to Bcl-2 which could be inhibited by ABT-199 in a concentration-dependent way
228 (EC₅₀~20 nM) (Fig 3A). A positive control, consisting of Bcl-2 as bait and TRIP13 as prey, which
229 triggers STAT signaling independently from binding to the bait, showed no decreased binding in the
230 presence of ABT-199 indicating that the compound disrupts the Bad/Bcl-2 interaction specifically.

231 Next, we wanted to assess the effect of ABT-199 on the binding of Bcl-2 to full size RyR3. For this,
232 similar co-immunoprecipitations as in Fig. 1B were set up. In the next experiments 3 μM of ABT-199
233 was used as this concentration was more than sufficient to completely disrupt binding of Bad and Bcl-
234 2 and we wanted to maximize the possible effects of the compound on the RyR/Bcl-2 interaction.
235 RyR3 was immunoprecipitated from lysates from HEK RyR3 cells overexpressing 3XFLAG-Bcl-2
236 after overnight incubation with ABT-199 or the vehicle (DMSO) (Fig. 3B, top). Similar amounts of
237 3XFLAG-Bcl-2 could be co-immunoprecipitated under these two conditions (Fig. 3B, bottom)
238 underpinning the data obtained with the Bcl-2 hydrophobic cleft mutant (Fig. 1B).

239 Next, we assessed the effect of the ABT-199 compound on the ability of Bcl-2 to inhibit RyR-
240 mediated Ca^{2+} release. For this we optimized the use of a different plasmid combining the 3XFLAG-
241 Bcl-2 construct with the mCherry selection marker in the same vector. The previously described virus-
242 derived P2A sequence was used for this purpose [15]. This sequence, once translated, is known to
243 cleave itself without any involvement of the host cell. The P2A sequence was introduced in the
244 pCMV24 vector flanked on its N-terminus by 3XFLAG-Bcl-2 and at the C-terminus by mCherry
245 (3XFLAG-Bcl-2-P2A-mCherry). A pCMV24 vector encoding P2A-mCherry was also made as
246 control. Before proceeding we verified whether the P2A sequence produced two separate proteins.
247 First, lysates from HEK RyR3 cells expressing either 3XFLAG-Bcl-2-P2A-mCherry or 3XFLAG-Bcl-
248 2 were immunoblotted. These showed in both cells the presence of 3XFLAG-Bcl-2 (Fig. 4A). In the
249 3XFLAG-Bcl-2-P2A-mCherry expressing cells this migrated at a slightly higher molecular mass
250 because part of the P2A sequence remains attached to the protein preceding the P2A sequence. As
251 reported previously, a weaker second product was also recognized at a slightly lower molecular mass
252 [15, 20]. Since this is also recognized by the anti-FLAG antibody, this is likely also 3XFLAG-tagged-
253 Bcl-2 protein lacking the P2A sequence. This band migrates slightly lower than the 3XFLAG-Bcl-2
254 next to it because the Myc-tag, present at the C-terminal end of the multiple cloning site of the
255 pCMV24 vector, is in this case attached to the mCherry protein and not to the 3XFLAG-Bcl-2.
256 Importantly, no FLAG-reactive signal could be detected at higher molecular weights. Since the
257 3XFLAG tag allows for a very sensitive detection, this result indicates that the vast majority of the
258 P2A-containing constructs was cleaved, resulting in the overexpression of two separate proteins. Next,
259 the two transfection protocols for single-cell Ca^{2+} measurements utilized in present study were
260 compared for their mCherry expression (Fig. 4B). Both methods resulted in a similar number of
261 mCherry-positive cells, indicating that the P2A-containing constructs can be used as selection marker
262 in the single-cell Ca^{2+} measurements. Fura2-AM single-cell Ca^{2+} measurements were performed in
263 HEK RyR3 cells overexpressing either P2A-mCherry as a control or 3XFLAG-Bcl-2-P2A-mCherry.
264 During the loading and de-esterification steps (1 hour in total), ABT-199 (3 μM) or the vehicle control
265 (DMSO) was added to the cells. The Ca^{2+} measurements were similarly performed as above.
266 Introducing 3XFLAG-Bcl-2 into these HEK RyR3 cells via the 3XFLAG-Bcl-2-P2A-mCherry

267 construct resulted in about 50% decrease in caffeine-induced Ca^{2+} release (Fig. 4 C, D). ABT-199 did
268 not impact the ability of Bcl-2 to inhibit caffeine-induced RyR-mediated Ca^{2+} release. It should be
269 noted that ABT-199 slightly and non-significantly reduced caffeine-induced Ca^{2+} release in control
270 conditions, which may suggest a minor direct impact of these drugs on RyRs.

271 The data obtained with the ABT-199 compound support the results using the Bcl-2 hydrophobic cleft
272 mutant and suggest that the ABT-199 BH3 mimetics do not alter Bcl-2's ability to inhibit RyRs
273 thereby decreasing the risk of potential side effects of these drugs.

274 In conclusion, disrupting the binding between the hydrophobic cleft of Bcl-2 and the pro-apoptotic
275 family members, does not alter Bcl-2-mediated regulation of the RyR.

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332

334 **Figure legends**

335 **Figure 1: Mutational disruption of the hydrophobic cleft of Bcl-2 does not disrupt Bcl-2/RyR**
336 **complexes.**

337 (A) Immunoblot of FLAG pull-downs performed on lysates from HEK RyR3 cells transiently
338 overexpressing the empty vector (pCMV24), 3XFLAG-Bcl-2 or the 3XFLAG-Bcl-2^{GR/AA} mutant. The
339 FLAG-tagged proteins were pulled down using agarose coupled to a FLAG-recognizing antibody. The
340 empty vector (pCMV24) was used as negative control. Immunoblots were stained with an anti-FLAG-
341 HRP-conjugated antibody (top) or an anti-Bax antibody (bottom). All experiments were performed
342 independently at least three times. (B) Immunoblot of the performed co-immunoprecipitation
343 experiments using HEK RyR3 lysates overexpressing 3XFLAG-Bcl-2 or the 3XFLAG-Bcl-2^{GR/AA}
344 mutant. RyR3 was immunoprecipitated using a RyR antibody recognizing all RyR isoforms (top). Co-
345 immunoprecipitations with non-specific IgG antibodies were used for every condition as negative
346 control. The presence of the co-immunoprecipitated FLAG-tagged proteins was assessed with an anti-
347 FLAG HRP-conjugated antibody (bottom). The experiments were performed independently at least
348 three times. For clarity reasons, a longer exposure of the input sample lanes obtained from the same
349 gel/blot is shown.

350 **Figure 2: Mutational disruption of the hydrophobic cleft of Bcl-2 does not alter Bcl-2-mediated**
351 **inhibition of RyR-mediated Ca²⁺ release.**

352 (A) Average calibrated trace (15-20 cells) of a typical single-cell Ca²⁺ measurement utilizing Fura-
353 2AM as Ca²⁺ indicator. The empty vector (pCMV24), 3XFLAG Bcl-2 or 3XFLAG-Bcl2^{GR/AA} was
354 cotransfected with mCherry (3:1 ratio) to identify transfected cells. One minute after chelating
355 extracellular Ca²⁺ with BAPTA (3 mM), caffeine was added (1.5 mM) as indicated by the arrows. (B)
356 Summary of the performed single-cell Ca²⁺ measurement experiments. The data points indicate the
357 average caffeine-induced Ca²⁺ release ($[\text{Ca}^{2+}]_{\text{peak}} - [\text{Ca}^{2+}]_{\text{before caffeine}}$) of all traces performed on the same
358 day. All experiments were repeated three times independently and experiments performed on the same

359 day are indicated by the same symbol in the same color (N>80 cells/condition). The average \pm S.E.M.
360 is indicated in gray (p=0.0091).

361 **Figure 3: The BH3 mimetic ABT-199 does not disrupt Bcl-2/RyR complexes.**

362 (A) A representative example of the performed MAPPIT experiments. The binding is shown as fold
363 induction value, calculated by dividing the average luciferase activity of Epo-stimulated cells by the
364 average of non-stimulated cells. Binding of the positive control or of Bad to Bcl-2 is shown in the
365 presence of increasing concentrations of ABT-199. (B) Immunoblot of the performed co-
366 immunoprecipitation experiments in HEK RyR3 cells overexpressing 3XFLAG Bcl-2 as performed in
367 Figure 1B (top). ABT-199 (3 μ M) or an equal volume of DMSO was included during the overnight
368 incubation step. Quantification of the performed experiments. Non-specific 3XFLAG-Bcl-2 binding to
369 the corresponding negative control (IgG) was subtracted from the amount of 3XFLAG-Bcl-2 co-
370 immunoprecipitated with RyR3 and expressed relative to the DMSO control. Values depict average \pm
371 SD. Experiments were repeated three times independently (bottom).

372 **Figure 4: The BH3 mimetic ABT-199 does not alter Bcl-2-mediated inhibition of RyR-mediated**
373 **Ca²⁺ release.**

374 (A) Immunoblot stained with anti-FLAG HRP-conjugated antibody of HEK RyR3 lysates transiently
375 transfected with 3XFLAG-Bcl-2 or 3XFLAG-Bcl-2-P2A-mCherry. The * indicates a byproduct of the
376 P2A cleavage process also recognized by the anti-FLAG antibody. (B) Fluorescence and bright field
377 overlay image, comparing the mCherry signal after transfection with 3XFLAG-Bcl-2 at 3:1 ratio with
378 mCherry (top) or the 3XFLAG-Bcl-2-P2A-mCherry construct (bottom). (C) Average calibrated trace
379 (15-20 cells) of single cell Ca²⁺ measurements, performed as in Figure 2, in HEK RyR3 cells
380 transfected with the pCMV24 containing P2A-mCherry (control) or the 3XFLAG-Bcl-2-P2A-
381 mCherry construct. ABT-199 (3 μ M) or an equal volume of DMSO was added to the cells during the
382 loading and de-esterification (one hour in total) of Fura2-AM. (D) Summary of the performed single-
383 cell Ca²⁺ measurements. Experiments performed during the same day are indicated by the same
384 symbol and color. The data points indicate average caffeine-induced Ca²⁺ release ($[Ca^{2+}]_{peak} - [Ca^{2+}]_{before}$)

385 caffeine) of all traces performed on the same day. All experiments were performed three times
386 independently (N>80 cells/condition). The average \pm S.E.M. is indicated in gray (p=0.0015).

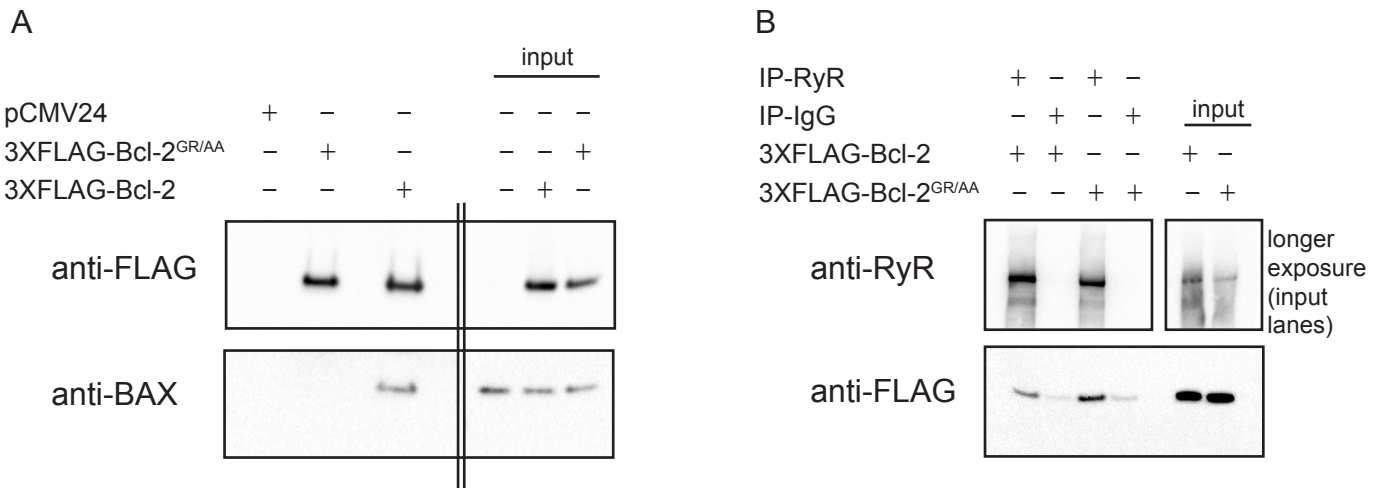
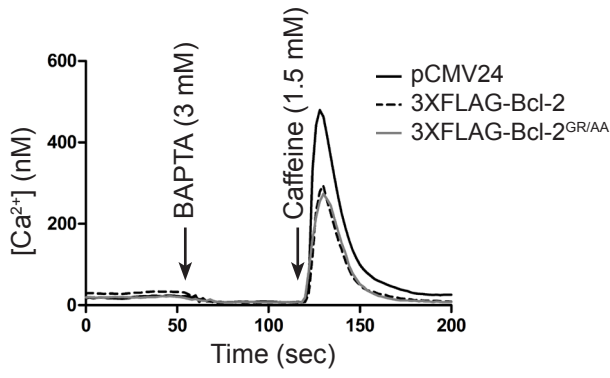


Figure 1

A



B

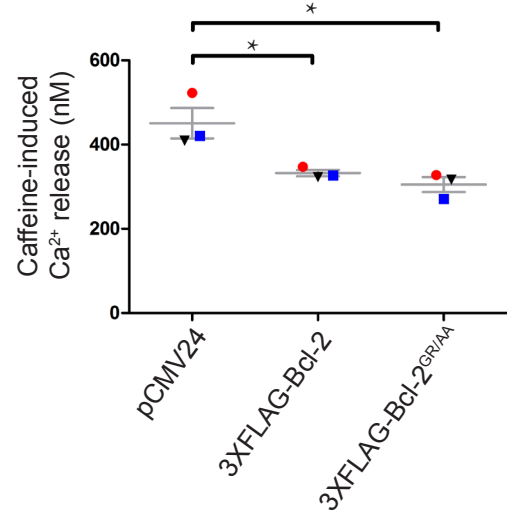


Figure 2

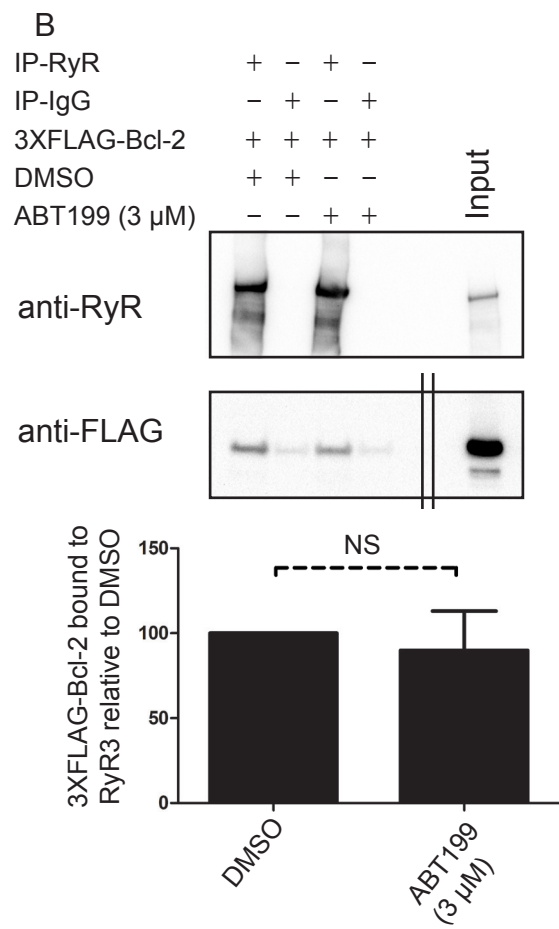
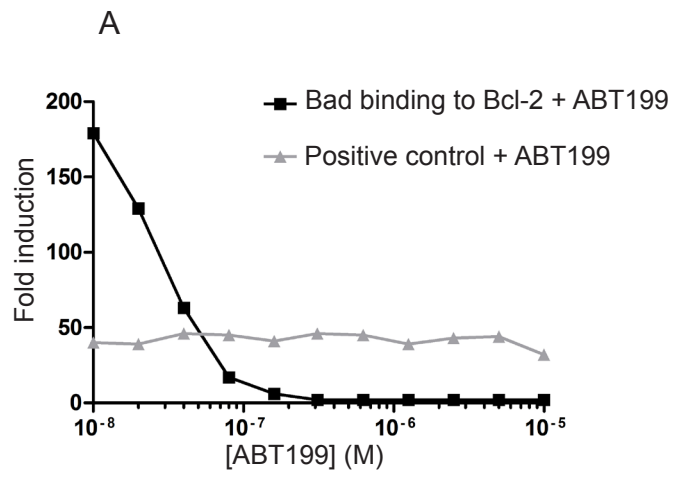


Figure 3

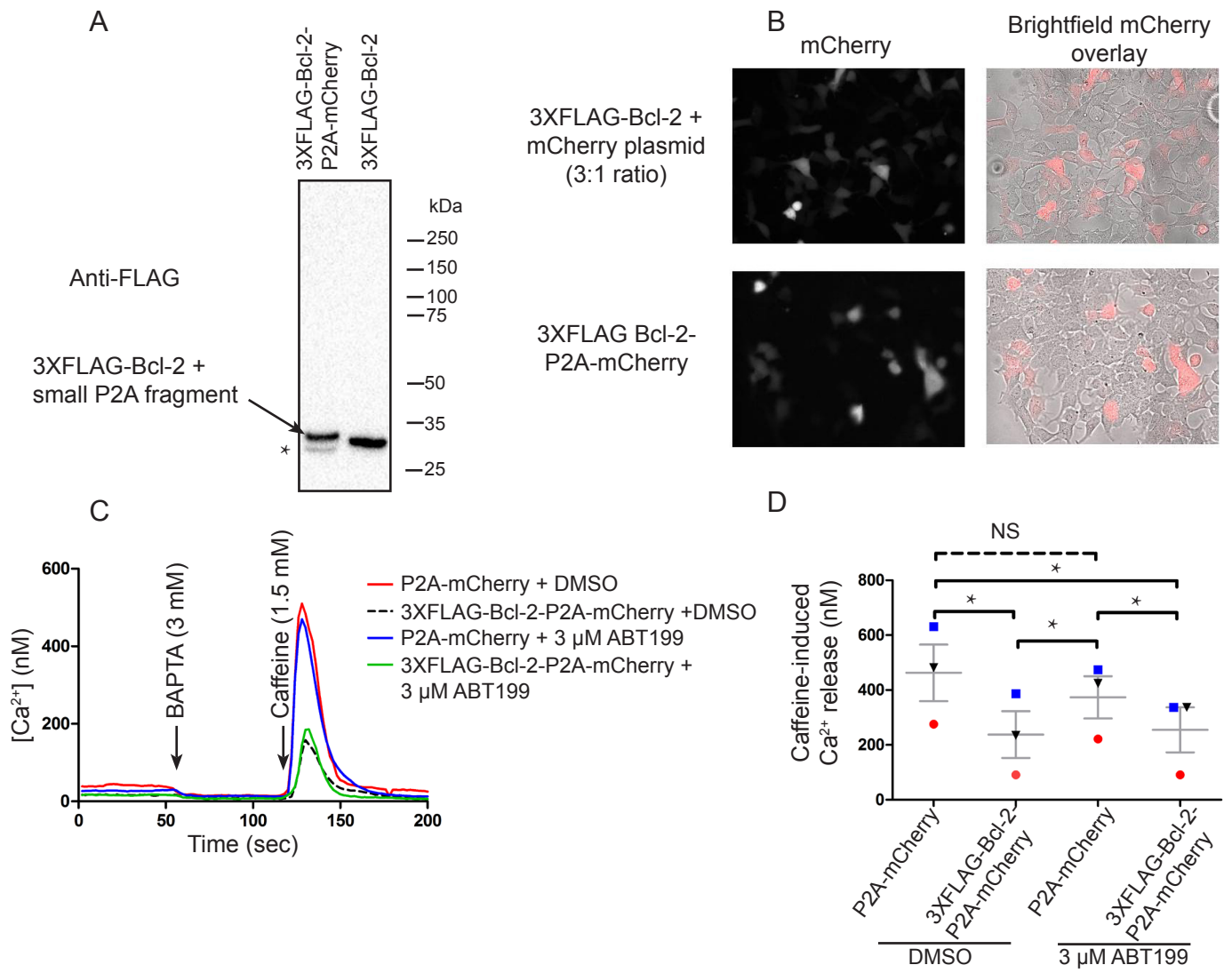


Figure 4