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# The PI3K/AKT/mTOR pathway is involved in direct apoptosis of CLL cells induced by ROR1 monoclonal antibodies

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## 1 **Letter to the Editor**

### 2 **The PI3K/AKT/mTOR pathway is involved in direct apoptosis of CLL cells** 3 **induced by ROR1 monoclonal antibodies**

4  
5 PI3K/AKT/mTOR signaling is a central pathway regulating growth of malignant cells and is  
6 constitutively activated in several types of cancer. PI3K is a key regulator of survival in cancer cells and  
7 the downstream molecules AKT and mTOR have anti-apoptotic effects (1). The class IA PI3K p110  
8 (PI3K $\delta$ ) catalytic subunit is activated by the SRC homology 2 domain binding of p85 regulatory subunits  
9 to phosphorylated tyrosine motifs of receptor tyrosine kinases (RTKs). Mutations in the  
10 PI3K/AKT/mTOR pathway have been noted in different malignancies. In chronic lymphocytic leukemia  
11 (CLL), PI3K has been shown to be constitutively activated phosphorylating AKT and mTOR.

12 In an attempt to find new tumor antigens with a specific expression in CLL, we identified the RTK ROR1  
13 (2). ROR1 is a surface receptor participating in cellular processes, as signal transduction, cell-cell  
14 interaction, proliferation, metabolism and survival. ROR1 has been suggested to be a survival factor in  
15 CLL (3). Silencing of ROR1 in CLL induced downregulation of the ROR1 gene and protein as well as  
16 apoptosis of the leukemic cells (4). Patients with progressive CLL had a higher expression of ROR1 as  
17 compared to patients with non-progressive disease (5).

18 We have previously shown that an anti-CRD ROR1 mAb induced specific direct apoptosis of primary  
19 CLL cells and dephosphorylated the intracytoplasmic TK domain of the ROR1 molecule (5, 6). To further

20 understand the effects of our anti-CRD ROR1 mAb, we analysed effects on the PI3K/AKT/mTOR  
21 pathway as well as on SRC and CREB.

22 Surface staining of CLL cells and PBMC of healthy donors, preparations of cell lysates and Western blot  
23 analysis have been described previously (5). The following total and phosphorylated proteins were  
24 analysed: ROR1, SRC, PI3K, AKT, mTOR and CREB. Phosphoproteins were measured before and after  
25 2h of incubation with the mAbs. In cytotoxicity experiments, CLL cells and PBMC of healthy donors  
26 were incubated with mAbs against CRD (5), and Ig domains of ROR1 (Miltenyi Biotec, Bergisch  
27 Gladbach, Germany) for 24-72h and measured by the MTT assay.

28 The percentage of ROR1 positive PBMC of CLL patients analysed by the anti-CRD ROR1 mAb was  $82 \pm$   
29  $5\%$  (mean  $\pm$  SD) (range: 78-89%) (n=15). Normal PBMC was negative in PCR for ROR1 and  $< 0.5\%$   
30 cells were stained.

31 The frequency of apoptotic cells induced by different mAbs is shown in Fig 1. The anti-CRD ROR1 mAb  
32 was significantly more effective in inducing apoptosis than the anti-Ig ROR1 mAb at all time points (24h;  
33  $p < 0.0381$ , 48h;  $p < 0.002$ , 72h;  $p < 0.002$ ). There was no difference between the isotype control and the anti-  
34 Ig ROR1 mAb.

35 The anti-CRD ROR1 mAb induced dephosphorylation of ROR1 as well as of SRC in CLL cells (Fig 2).  
36 SRC proteins have been shown to be activated in lung, breast and pancreatic carcinoma cells and involved  
37 in survival, proliferation and invasion. Phosphorylated ROR1 can physically interact with and

38 phosphorylate SRC and suggested to be a critical component for multiple signaling pathways involved in  
39 tumorigenesis (7).

40 Treatment of CLL cells with the anti-CRD ROR1 mAb also decreased the level of phosphorylated AKT  
41 (Fig 2). ROR1 mediated SRC phosphorylation has been shown to trigger AKT activation in lung  
42 adenocarcinoma cells and ROR1 knockdown induced dephosphorylation of AKT as well as inhibited  
43 growth and induced apoptosis (7).

44 Phosphorylation of PI3K $\delta$  (Fig 2) and PI3K p85 but not the p55 isoform (data not shown) also decreased  
45 in CLL cells treated with the anti-CRD ROR1 mAb. In CLL and AML cells, increased AKT activity was  
46 shown to correlate with phosphorylation of PI3K $\delta$  which was the predominant isoform. Phosphorylation  
47 of PI3K $\delta$  was mediated by SRC and p85 recruitment increased the catalytic activity of the PI3K $\delta$  subunit.

48 Treatment of CLL cells with the anti-CRD ROR1 mAb also dephosphorylated mTOR. mTOR is  
49 important for the regulation of cell growth as well as metabolism and is activated in different tumor types  
50 translating proteins required for cell cycle progression from the G1 to S phase.

51 Furthermore, the anti-CRD ROR1 mAb induced dephosphorylation of the transcription factor CREB (Fig  
52 2). Oncogenic transcription factors play a central role in tumorigenesis. CREB is activated through  
53 phosphorylation by kinases, including AKT. CREB has been shown to be overexpressed and  
54 constitutively phosphorylated in AML and NSCLC and important for the pathogenesis of these diseases.

55 Apoptosis of CLL cells induced by the anti-CRD ROR1 mAb was preceded by dephosphorylation of  
56 ROR1, SRC, PI3K p85, PI3K $\delta$ , AKT, mTOR and CREB proteins. Binding of the anti-CRD ROR1 mAb

57 to ROR1 decreased phosphorylation of SRC which might lead to dephosphorylation of the PI3K p85  
58 isoform abrogating p85 recruitment and inactivation of the PI3K p110 catalytic subunit preventing signal  
59 transmission downstream of PI3K. Activation of CREB might occur via the PI3K/AKT/mTOR pathway  
60 which may enhance expression of genes augmenting resistance of tumor cells to apoptosis as well as  
61 promoting tumor cell growth. ROR1 has been shown to utilize distinct kinase dependent and independent  
62 mechanisms to sustain a favorable balance between PI3K/AKT mediated pro-survival signals and the pro-  
63 apoptotic p38 pathway (7). A significant association between the expression of ROR1 and activated  
64 AKT/CREB enhancing tumor cell growth in different tumor types has recently been reported (8).

65 The present study indicates that our anti-CRD ROR1 mAb inhibited the PI3K/AKT/mTOR pathway  
66 which is of major importance in tumorigenesis (1). This pathway is a validated target for e.g. EGFR and  
67 HER-2, two other receptor tyrosine kinases of the RTK families. ROR1 might be an interesting  
68 therapeutic target using mAbs for CLL and other cancers expressing ROR1. A support of a therapeutic  
69 effect of anti-ROR1 mAbs was recently reported using the anti-ROR1 D10 mAb in an animal CLL model  
70 (9). The D10 ROR1 mAb is directed against a CRD close epitope and induced direct apoptosis, while  
71 antibodies against the Ig-domain did not induce direct apoptosis (10), as confirmed in the present study.  
72 In addition, our anti-CRD ROR1 mAb also killed leukemic cells in ADCC and CDC (5). Thus, our  
73 specific anti-CRD ROR1 mAb seems to have multifunctional activities and might support a notion that  
74 the ROR1 binding site is of importance for the activity of cytotoxic ROR1 antibodies.

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77 **Author contributions**

78 Contribution: AHDM performed the experiments, analysed the data and wrote the manuscript. MHF  
79 performed experiments and reviewed the manuscript. AM, ASK and EM reviewed the manuscript. AÖ  
80 provided clinical material, analysed the data and wrote the manuscript. HM designed and supervised the  
81 study, provided clinical material, analysed the data and wrote the manuscript

82 **Conflict of interest**

83 The authors declare no conflict of interest.

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101 **References**

- 102 1. Arcaro A, Guerreiro AS. The phosphoinositide 3-kinase pathway in human cancer: genetic  
103 alterations and therapeutic implications. *Curr Genomics*. 2007;8(5):271-306.
- 104 2. Daneshmanesh AH, Mikaelsson E, Jeddi-Tehrani M, Bayat AA, Ghods R, Ostadkarampour  
105 M, et al. Ror1, a cell surface receptor tyrosine kinase is expressed in chronic lymphocytic leukemia and  
106 may serve as a putative target for therapy. *Int J Cancer*. 2008;123(5):1190-5.
- 107 3. Fukuda T, Chen L, Endo T, Tang L, Lu D, Castro JE, et al. Antisera induced by infusions of  
108 autologous Ad-CD154-leukemia B cells identify ROR1 as an oncofetal antigen and receptor for Wnt5a.  
109 *Proceedings of the National Academy of Sciences of the United States of America*. 2008;105(8):3047-52.
- 110 4. Choudhury A, Derkow K, Daneshmanesh AH, Mikaelsson E, Kiaii S, Kokhaei P, et al.  
111 Silencing of ROR1 and FMOD with siRNA results in apoptosis of CLL cells. *Br J Haematol*.  
112 2010;151(4):327-35.
- 113 5. Daneshmanesh AH, Hojjat-Farsangi M, Khan AS, Jeddi-Tehrani M, Akhondi MM, Bayat AA,  
114 et al. Monoclonal antibodies against ROR1 induce apoptosis of chronic lymphocytic leukemia (CLL) cells.  
115 *Leukemia*. 2012;26(6):1348-55.
- 116 6. Hojjat-Farsangi M, Khan AS, Daneshmanesh AH, Moshfegh A, Sandin A, Mansouri L, et al.  
117 The tyrosine kinase receptor ROR1 is constitutively phosphorylated in chronic lymphocytic leukemia  
118 (CLL) cells. *PloS one*. 2013;8(10):e78339.
- 119 7. Yamaguchi T, Yanagisawa K, Sugiyama R, Hosono Y, Shimada Y, Arima C, et al. NKX2-  
120 1/TTF1/TTF-1-Induced ROR1 is required to sustain EGFR survival signaling in lung adenocarcinoma.  
121 *Cancer Cell*. 2012;21(3):348-61.
- 122 8. Zhang S, Chen L, Wang-Rodriguez J, Zhang L, Cui B, Frankel W, et al. The onco-embryonic  
123 antigen ROR1 is expressed by a variety of human cancers. *The American journal of pathology*.  
124 2012;181(6):1903-10.
- 125 9. Widhopf GF, 2nd, Cui B, Ghia EM, Chen L, Messer K, Shen Z, et al. ROR1 can interact with  
126 TCL1 and enhance leukemogenesis in Emu-TCL1 transgenic mice. *Proceedings of the National Academy  
127 of Sciences of the United States of America*. 2014;111(2):793-8.
- 128 10. Yang J, Baskar S, Kwong KY, Kennedy MG, Wiestner A, Rader C. Therapeutic potential and  
129 challenges of targeting receptor tyrosine kinase ROR1 with monoclonal antibodies in B-cell malignancies.  
130 *PloS one*. 2011;6(6):e21018.

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139 **Legends to figures**

140 **Figure 1.**

141 Apoptosis (%) (mean  $\pm$  SD) time response curves induced by the anti-CRD ROR1 mAb in CLL cells  
142 (n=7) (A) and PBMC of healthy donors (n=5) (B). For comparison a mouse mAb against the Ig domain of  
143 ROR1 as well as an isotype control mAb was included. Direct apoptosis induced by anti-CRD ROR1  
144 mAb was significantly higher compared to the anti-Ig ROR1 and the isotype control mAbs (anti-CRD  
145 ROR1 vs anti-Ig ROR1 at 24h;  $p < 0.0381$ , at 48h;  $p < 0.002$  and at 72h;  $p < 0.002$ ) (anti-CRD ROR1 vs  
146 control isotype control at 24h;  $p < 0.002$ , at 48h;  $p < 0.002$  and at 72h;  $p < 0.002$ ). Spontaneous apoptosis at  
147 each time point was deducted. At 72h the spontaneous apoptosis for CLL cells was  $27 \pm 5\%$  and for  
148 PBMC of healthy donors  $26 \pm 6\%$  (mean  $\pm$  SD). There were no statistically significant differences in  
149 apoptosis comparing the anti-CRD ROR1, the anti-Ig ROR1 and the isotype control mAbs using normal  
150 PBMC as targets.

151 **Figure 2.**

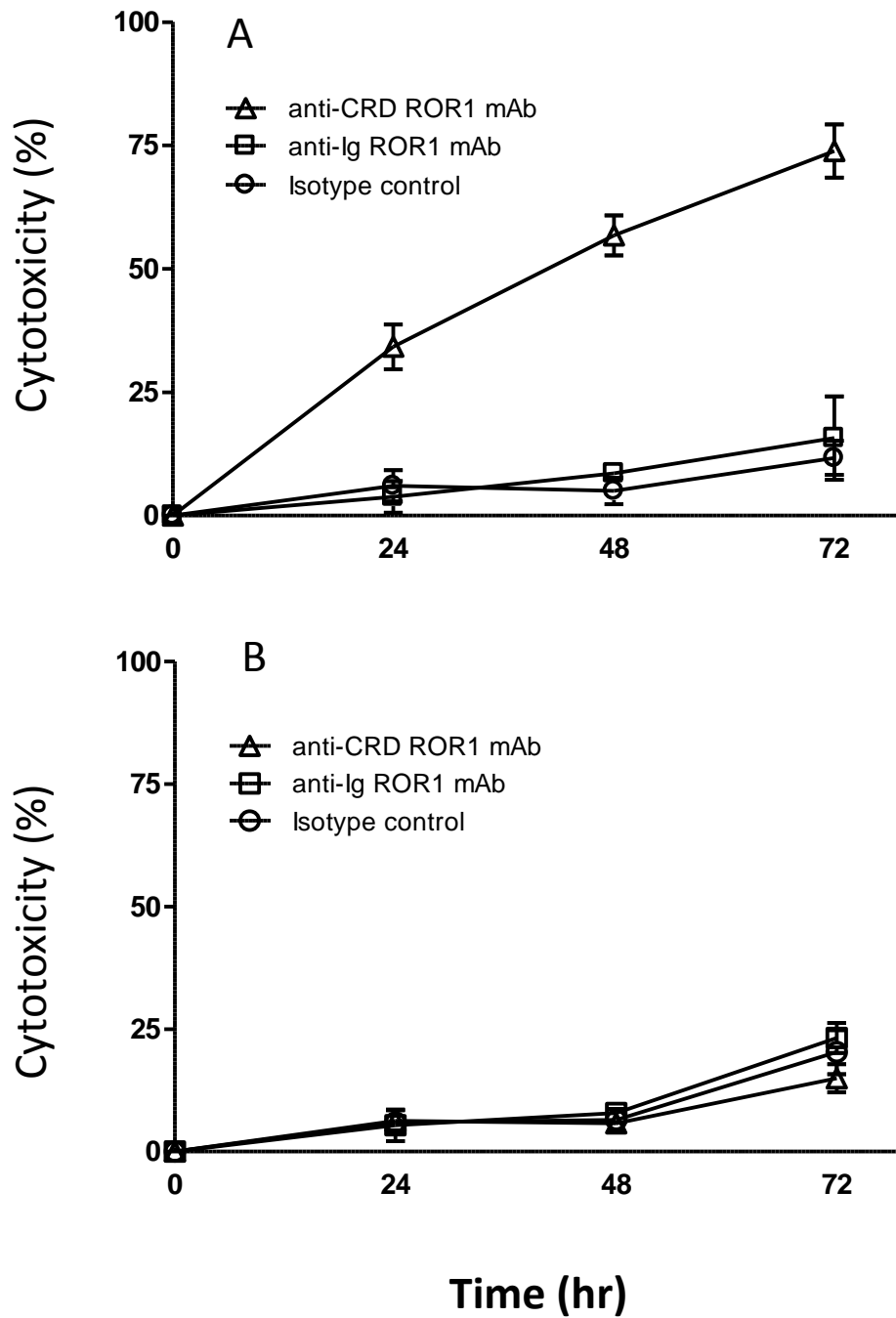
152 **A)** Representative experiments of five CLL patients showing dephosphorylation of ROR1, SRC, PI3K $\delta$ ,  
153 AKT, mTOR (signaling molecules) and CREB (transcription factor) within 2h of incubation of CLL cells  
154 with the anti-CRD ROR1 mAb (+) and an isotype control mAb (-). Time kinetics experiments showed  
155 that 2h was the optimal time point to achieve maximum dephosphorylation of the signaling proteins (data  
156 not shown). The Western blot methods as well as the anti-CRD ROR1 mAb and the rabbit polyclonal  
157 pROR1 antibody have been described previously (5, 6). The other antibodies were monoclonal antibodies  
158 purchased from Cell Signaling Technology (Danvers, MA, USA) and Santa Cruz Biotechnology (Dallas,  
159 TX, USA). **B)** Relative intensity of phosphorylated proteins to total proteins was calculated after 2h of



160 incubation without ( □ ) and with ( ■ ) the anti-CRD ROR1 mAb. Statistically significant levels are  
161 shown at the top. Intensity was measured by ImageJ software (National Institutes of Health (NIH),  
162 Bethesda, MD, USA), as previously described (6).

163

Figure 1



**Figure 2**

