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# JNK1 AS A SIGNALING NODE IN VDR-BRAF INDUCTION OF CELL DEATH IN AML

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# Abstract

Numerous clinical studies of vitamin D, its derivatives or analogs, have failed to clearly demonstrate sustained benefits when used for the treatment of human malignant diseases. However, given the strong preclinical evidence of anti-neoplastic activity and the epidemiological associations suggesting that vitamin D compounds may have a place in cancer therapy, attempts are continuing to devise new approaches to their therapeutic use. This laboratory has developed a strategy to enhance the effectiveness of the currently standard therapy of Acute Myeloid Leukemia (AML) by the immediate addition of the vitamin D2 analog Doxercalciferol combined with the plant polyphenol-derived Carnosic acid to AML cells previously treated with Cytarabine (AraC). Enhancement of AML cell death was noted to be dependent on VDR and BRAF kinase. Here we document that the stress-related kinase JNK is an important additional component of cell death enhancement of AraC-induced cell death, and we show that JNK signaling to the apoptosis regulator BIM and Caspase executioners of cell death are downstream of VDR and BRAF. A clear understanding of the molecular basis for the increased efficacy of AraC in the therapy of AML is expected to bring this regimen to a clinical trial.

# **Graphical Abstract**

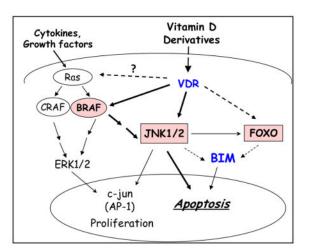
Potential signals downstream from VDR-BRAF pathway in the context of DNA damage.

Disclosure of conflict of interest

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In this cell context, solid lines show proven signaling; bolded if shown here, and interrupted lines show signaling that needs further study.

#### Keywords

Vitamin D; JNK; BRAF; FOXO; Acute Myeloid Leukemia; Carnosic acid

# INTRODUCTION

Standard therapy for Acute Myeloid Leukemia (AML), with cytotoxic agents such as Cytarabine (AraC) typically fails as curative therapy, especially in the elderly or relapsed patients (1–3). Not surprisingly, this has motivated numerous attempts to provide more than only generally cytotoxic therapeutic regimens to AML patients. Using molecularly targeted approaches, some initial clinical success has recently been reported in several subsets of patients, including patients with IDH1/2 mutations (4–6), and with the upregulation of HOXA9 transcription factor (7), or in AML with BCL2 dependence (8). Preclinical studies also suggest that chromatin regulators such as MLL1 and DOT1L can be therapeutic targets in NPM1-mutated AML (9–11).

The above studies generally depended on the use of small synthetic molecules which target cellular enzymes or transcription factors. In contrast, we have recently provided pre-clinical evidence that the small molecules related to natural compounds can increase the potentially therapeutic efficacy of AraC studied in AML blasts in ex vivo culture (12). These compounds, an analog of vitamin D2 Doxercalciferol (D2) and a plant-derived antioxidant carnosic acid (CA), have a powerful differentiating effect when combined, and this increases the cytotoxicity of AraC to AML blasts. Importantly, this action is selective for neoplastic cells, as the enhancement of cell death by this differentiation agent combination has no observable effect on the normal bone marrow cells (13, 14).

The initial investigation of the mechanism responsible for the enhancement of AraC-induced cell death by treating cells that have damaged DNA with a differentiation agent combination (D2/CA) has revealed a paradoxical tumor suppressor-like activity of BRAF. We found that

reducing the level or inhibiting the kinase activity of BRAF diminishes the ability of D2/CA to enhance ALM cell death (13). Also, D2/CA increases BRAF expression in a VDR-dependent manner, which correlates with the increased levels of the DNA damage marker H2AX (14). We now provide data supporting the role of JNK as a transmitter of these signals to the cell death machinery, downstream from VDR-BRAF.

# MATERIALS AND METHODS

#### Reagents

Arabinocytosine (AraC) and Doxercalciferol (1α-hydroxyvitamin D<sub>2</sub>; 1-D2) were purchased from Sigma-Aldrich (St. Louis, MO). Carnosic acid (CA) was purchased from Enzo Life Sciences, Inc. (Farmingdale, NY). JNK inhibitor SP600125 was from Cell Signaling Inc (Danvers, MA). The antibodies that used for Western blotting as following: BRAF (#9433), CRAF (#9422), Phospho-JNK (Thr183/Tyr185, #9255), JNK (#9252), Phospho-p38 MAPK (Thr180/Tyr182, #4511), Phospho-C-JUN (Ser63, #2631), C-JUN (#9165), Phospho-BIM (Ser69, #4585), Phospho-H2AX (Ser139, #9718), FOXO3a (#2497), cleaved Caspase 9 (#9505), cleaved caspase 3 (#9661) and HRP-linked anti-rabbit (#7074) antibodies were purchased from Cell Signaling Technologies. BIM (sc-8625), VDR (sc-1008) and Crk-L (sc-319) were obtained from Santa Cruz Biotechnology (Dallas, TX).

### Cell culture

HL60 is an AML cell line isolated from a patient with acute promyeloblastic leukemia (15). The search of the database in cBioPortal for Cancer Genomics shows no mutations of BRAF and JNKs in AML. However, it has been reported that BRAF mutations have been found in a small proportion of secondary AML (16), which is not the case in the specimen used in this study. The cells were cultured in RPMI 1640 medium supplemented with 10% heat-inactivated bovine calf serum (BCS) at 37°C in a 5% CO2 atmosphere. Cells were passaged twice a week to maintain log phase growth. For experiments, the cells were seeded in 6-well plates at  $1.5 \times 10^5$  cells/mL followed by the addition of the various agents for the indicated times. Cell viability was routinely determined by using the Trypan blue exclusion assay with a Neubauer hemocytometer.

### Isolation of mononuclear cells from a patient blood sample

A blood sample was obtained during a diagnostic procedure from a patient volunteer diagnosed with FAB M5 subtype of AML, according to the Institutional Review Board protocol. Mononuclear cells were isolated from the specimen by using Histopaque-1077 (Sigma-Aldrich) gradient centrifugation as described previously (17).

### Knockdown of proteins by siRNA oligonucleotides

HL60 cells were transfected with 100 nM of SignalSilence JNK siRNA, VDR siRNA, siBRAF, siCRAF or scrambled Control siRNA (Cell Signaling Inc), using Endo-Porter delivery reagent from Gene Tools Inc (Philomath, OR) for 48–72 hours before exposure to other agents for the indicated times. The cells were then examined to determine the extent of knockdown of JNK or VDR expression at both mRNA and protein levels.

## Annexin V FITC assay

Following the indicated exposure of either HL60 cells or ex vivo AML blasts to AraC followed by D2/CA combination, the cells were collected, washed twice with 1xPBS, then re-suspended in the binding buffer, and stained using an Annexin V-FITC assay Kit (Fisher Scientific, Pittsburgh, PA). The cells were incubated with 50 µg/ml Annexin V and 10 µg/ml propidium iodide in 1x binding buffer at room temperature in the dark for 15 min, and immediately analyzed by flow cytometry (EPICS XL). Annexin V-positive/PI-negative cells were considered to be early apoptotic, cells with both Annexin V and PI positive to be late apoptotic, and Annexin V negative but PI positive to be "necrotic"(18).

# **Real time quantitative PCR**

Total RNA was extracted by using Trizol (Invitrogen, CA) according to manufacturer's protocol. Quantitative RT-PCR for JNK1/2 was carried out using an ABI 7500 Real-Time PCR System as described before (14). Fold changes of mRNA levels in target gene relative to the RNA polymerase II (RPII) control were calculated by using relative quantification analysis. Primers used for JNK1 were: forward 5'-CCCTGACAAGCAGTT AGATGAA-3', reverse 5'-CTGTCTGTATCAGAGGCCAAAG-3'; JNK2, forward 5'-ATGGAGCTGGATCATGAAAGAA-3', reverse 5'-GGTCAG TGCCTTGGAATATCA-3', for RP II, forward 5'-GCACCACGTCCAATGA CAT-3', reverse 5'-GTGCGG CTGCTTCCATAA-3'. The quality of all PCR products was monitored by using post-PCR melting curve analysis.

## Western blotting

Western blotting was performed using 40  $\mu$ g of whole cell extracts as described before (19). Briefly, membranes were incubated with primary antibodies for 1–2 hours according to the sensitivity of each antibody, and then blotted with a HRP-linked secondary antibody for 1 hour. The bands of protein were visualized using a chemiluminescence detection system (Pierce Biotechnology, Rockford, IL). Each membrane was stripped and reprobed for Crk-L, which was used as an internal loading control. The optical density of each band was quantified using ImageQuant 5.0 (Molecular Dynamics, Sunnyvale, CA).

### Statistical analysis

Each experiment was repeated at least 3 times. The results are presented as the Mean  $\pm$  Standard Deviation. Significance of the differences between mean values of two treatment groups was assessed by a Student's T-test using Microsoft EXCEL program. Statistical significance of multiple treatment groups were assessed by one way ANOVA followed by the Tukey HSD test. p < 0.05 was considered statistically significant.

# RESULTS

# 1. Phosphorylation of JNK1/2 is increased by D2/CA in AraC-treated cells and is required for the upregulation of BIM and the enhancement of cell death

HL60 cells were treated with siRNA to JNK1/2 to reduce the expression of this stresssignaling kinase, and Table 1 shows that this treatment abrogated the D2/CA-induced

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increases in JNK1 mRNA, and reduced the increases in JNK2 mRNA after AraC or D2/CA exposure. Western blotting confirmed the marked knock-down (KD) of JNK protein isoforms, and resulted in a robust reduction of JNK activation by phosphorylation, as shown in Fig 1. While the expression of p38 MAPK was unaffected by the JNK KD, showing its specificity, the reduction of JNK activation was further documented by the reduced phosphorylation of C-JUN, a target of activated JNK (Fig 1). The decrease in P-JNK levels was also accompanied by the reduced expression of the pro-apoptotic protein BIM, and less evidence of DNA damage as shown by the P-H2AX levels (Fig 1). These signaling changes were associated with markedly reduced D2/CA-induced enhancement of the AraC-induced cell death in these cells (Table 2A). Similar treatment of a small blood sample of AML cells obtained from a patient with AML also resulted in increase in cell death induced by AraC and its enhancement by D2/CA, which was reduced in the cells with reduced levels of JNK that resulted from an exposure to siJNK (Table 2B).

Thus, the reduction in JNK levels results in a decrease in signals for and the execution of cell death in AML cells with DNA damage.

# 2. Inhibition of JNK activation by SP600125 reduces the expression of FOXO3a and inhibits the activation of the apoptotic cascade

We also sought to confirm the role of JNK as a component of cell death signaling in this system by inhibiting the kinase activity of JNK with the selective pharmacological inhibitor SP600125. Indeed, the Westerns illustrated in Fig 2A show that in cells treated with AraC the enhancement of JNK phosphorylation by D2/CA was abrogated by this compound. This series of experiments also demonstrated that in this system the FOXO3a transcription factor is upregulated in a JNK-dependent manner by D2/CA. Since it is known that FOXO3a can control apoptosis by transcriptional regulation of BIM (20), the parallel effects of D2/CA and SP600125, observed here on FOXO3a, BIM, and on the activation by cleavage of Caspase 9 and Caspase 3, suggest that JNK is an upstream activator of a cascade of events that lead to apoptosis. Indeed, the effect of SP6000125 on cell death correlated well with its inhibition of JNK1 (Fig 2B).

# 3. VDR is required for the optimal JNK upregulation

Given that the cell death enhancement by D2/CA combination involves the vitamin D2 analog Doxercalciferol, we tested whether vitamin D receptor (VDR) is required JNK activation. Indeed, the KD of VDR with a silencing construct previously shown to be effective (13) reduced JNK activating phosphorylation in parallel with cell death reduction, but did not reduce JNK protein levels (Fig 3). This shows that the activation of JNK by phosphorylation does require optimal levels of VDR.

### 4. JNK is downstream of BRAF, but not of CRAF

We have previously shown that the enhancement of cell death by D2/CA requires BRAF (14), but the possibility of an involvement of JNKs was not examined in that study. Using a pool of siBRAF construct oligonucleotides we now show that BRAF is required for JNK phosphorylation in this system, and JNK activation is also demonstrated by the dependence on BRAF of C-JUN phosphorylation, which is a target of JNK (Fig 4A). In contrast, KD of

# DISCUSSION

The JNK pathway is a well-recognized transmitter of stress and genotoxicity-related cell death in many cell types (21, 22). We, and others, have shown that JNK is activated during VDD-induced differentiation of AML cells (23–26), but whether this is a biomarker of this process, or has a causative role is not clear. JNK can have a protective role in some cell types (27) and some specific cell context in AML blasts (28). However, it has been reported that anthracycline resistance of AML patients can be due to the failure of JNK activation and apoptosis (29), and our evidence suggests that in HL60 cells and in ex vivo AML blasts JNK depletion reduces both AraC cell death and its enhancement (Table 2).

In our system, cell death signals from JNK appear to be transmitted by FOXO3a, at least in part. As pointed out above, FOXO3a controls apoptosis by transcriptional regulation of BIM in some cells (20, 30, 31), and JNK is one of FOXO3a upstream regulators.

On one hand, JNK can trigger apoptosis by the upregulation of genes which promote cell death such as BIM, PUMA and BNIP3, at least in part by modulating the nuclear localization, and thus the levels of FOXO3a protein in a number of different cell types (32–34). Specifically, phosphorylation of FOXO3a by JNK diminishes the interaction of FOXO3a with 14-3-3 protein, which protects FOXO3a from proteosomal degradation in the cytoplasm (35, 36). Nuclear localization of FOXO3a allows this transcription factor to upregulate a variety of genes, including BIM.

On the other hand, AKT promotes cell survival by phosphorylating FOXOs at three regulatory sites (Thr32, Ser253, and Ser315), inactivating FOXO3a by its re-localization from the nucleus to the cytoplasm (37). Therefore, an additional mechanism for the observed increase in AML cell death by D2/CA can be a decrease in AKT activity.

Although the specificity of JNK knock-down by siBRAF shown in Fig 4 was confirmed by CRISPR/Cas9 (data not shown), currently it is not clear why BRAF is required for D2/CA-induced cell death enhancement. A signaling connection between BRAF and AKT has been reported (38), so this connection will be investigated in future studies of death enhancement in AML cells with damaged DNA. Additionally, the Apoptosis Signaling Protein 1 (ASK1) is known to bind to both RAF and to JNK, so this may be another potential signaling cross-talk that leads to BIM upregulation in this cell system.

It therefore appears that there seems to be a clear signaling path from JNK to caspase activation and cell death, although the participation of AKT in FOXO modulation resulting in BIM upregulation still remains to be excluded. In contrast, how BRAF signals downstream has multiple potential scenarios discussed above, and these are currently under active investigation.

# Acknowledgments

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# **Abbreviations list**

AP-1	Activator protein 1, c-Jun and dimerization partners				
AKT	Serine/threonine protein kinase B (PKB)				
AML	Acute myeloid leukemia				
AraC	Cytarabine/Arabinocytosine				
ASK1	Apoptosis signal regulating kinase 1				
BCL2	B-cell lymphoma 2				
BIM	Bcl-2-like protein 11				
BNIP3	Bcl2 interacting protein 3				
BRAF	B-Raf proto-oncogene				
CA	Carnosic acid				
CRAF	Raf-1 proto-oncogene				
Crk-L	CRK-like proto-oncogene				
D2	Doxercalciferol (1a-hydroxyvitamin D2; 1-D2)				
DOT1L	DOT like histone lysine methyltransferase				
FAB	French-American British classification of AML subtypes				
FOXO3	Forkhead box protein 3				
H2AX	H2A histone family member X				
HOXA9	Homeobox protein Hox-A9				
IDH1/2	Isocitrate dehydrogenase 1/2				
JNK	c-Jun N-terminal kinase				
JUN	Jun proto-oncogene				
MLL1	Myeloid/lymphoid or mixed-lineage leukemia 1 (MLL1)				
NPM1	Nucleophosmin				
p38MAPK	p38 mitogen-activated protein kinase				
VDD	Vitamin D Derivative				
VDR	Vitamin D Receptor				

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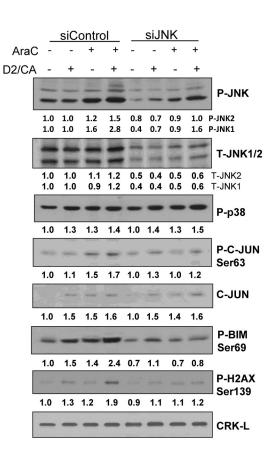
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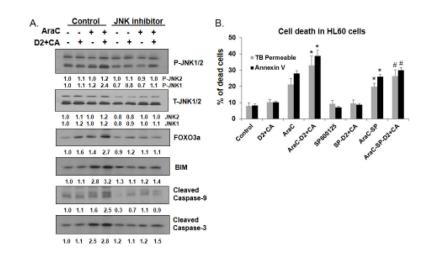
# Highlights

- In AML cells with DNA damage attempted differentiation results in apoptosis instead.
- Enhanced cell death is facilitated by an enhanced expression of VDR and BRAF.
- The execution of cell death in this setting may be signaled by JNK activation.



## Figure 1.

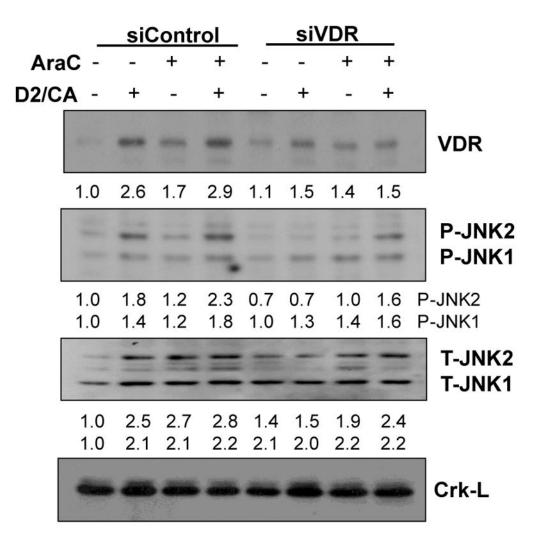
Western blots showing that phosphorylation of JNK1/2 and their downstream target C-JUN is increased by D2/CA in AraC-treated cells, and is required for the upregulation of BIM, and the DNA damage marker P-H2AX. The lack of change in P-p38 levels indicates the specificity of siJNK. CRK-L was used as an internal control for loading.



# Figure 2.

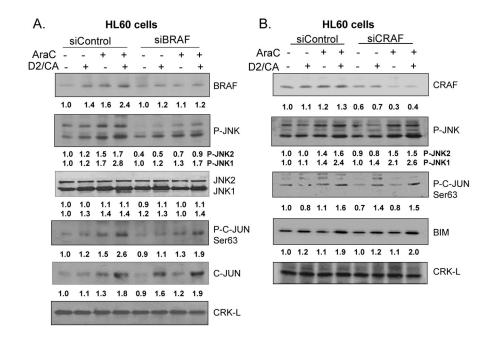
**A.** Inhibition of JNK activity by SP600125 reduces protein levels of FOXO3a and BIM in cells with AraC-induced cell damage treated with D2/CA. The levels of cleaved Caspases 9 and 3 are also reduced in these cells. **B**. The reduction of the enhancement of the cell death by D2/CA by inhibition of JNK phosphorylation parallels the molecular changes shown in panel A. \*= p < 0.05, when compared with AraC treated group. # = p < 0.05, when compared with corresponding control group. N=3

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#### Figure 3.

Knockdown of VDR with siVDR reduces JNK phosphorylation, but does not reduce JNK protein levels. CRK-L was used as a loading control.



# Figure 4.

Downstream targets of RAF signaling. A: Knockdown of BRAF with siBRAF shows that BRAF is required for optimal phosphorylation of JNK and its target C-JUN. B. Knockdown of CRAF with siCRAF showed little if any effect on the phosphorylation of JNK or C-JUN, and the levels of its downstream target BIM.

# Table 1

Knockdown of JNK1 and JNK2 with siJNK diminishes the AraC-D2/CA-induced increases in JNK1/2 mRNA

Table 1A	mKNA	mRNA vs siCTL	vs AraC	No CATUS	r value
09TH	JNK1	p value	p value	p value	vs corresp group
siControl-Control	96.0				
siControl-D2/CA	1.26	0.180			
siControl-AraC	0.82	0.038			
siControl-AraC-D2/CA	1.46	0.179	0.040	0.261	
siJNK-Control	0.89				0.157
siJNK-D2/CA	0.87	0.788			0.147
siJNK-AraC	0.85	0.462			0.656
siJNK-AraC-D2/CA	0.81	0.326	0.126	0.327	0.105
Table 1B	mRNA	vs siCTL	vs AraC	vs CA+D2	P value
	6211VI			-	
HL60	2NNL	p value	p value	p value	vs corresp group
siControl-Control	1.13				
siControl-D2/CA	1.75	0.011			
siControl-AraC	2.48	0.020			
siControl-AraC-D2/CA	2.96	0.007	0.018	0.050	
siJNK-Control	0.80				0.003
siJNK-D2/CA	1.09	0.069			0.010
siJNK-AraC	1.33	0.024			0.061
siJNK-AraC-D2/CA	1.92	0.006	0.042	0.047	0.033

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## Table 2

Reduction of AraC-induced cell death by D2/CA when JNK expression is knocked down by a siJNK construct.

	HL60		AML ex vivo blasts	
	TB Permeable	Annexin V	TB Permeable	Annexin V
	Cell death %	Cell death %	Cell death %	Cell death %
siControl-Control	$8.3 \pm 1.8$	$10.4\pm1.8$	$13.3\pm3.6$	$15.6\pm2.8$
siControl-D2/CA	$11.2\pm2.5$	$15.5\pm2.6$	$15.7\pm2.9$	$16.7\pm3.5$
siControl-AraC	$\textbf{33.0} \pm \textbf{4.1}$	$40.5\pm3.6$	$42.3\pm4.4$	$40.2\pm3.9$
siControl-AraC-D2/CA	$49.2\pm5.9$	$56.9 \pm 4.6$	$55.3\pm6.2$	$58.9 \pm 6.1$
siJNK-Control	$9.7\pm2.8$	$11.7\pm3.0$	$12.0\pm3.1$	$14.4\pm3.5$
siJNK-D2/CA	$12.3\pm2.9$	$16.3\pm2.9$	$14.7\pm2.6$	$16.4\pm3.2$
siJNK-AraC	$31.0\pm4.5$	$\textbf{33.6} \pm \textbf{4.8}$	$\textbf{38.0} \pm \textbf{4.8}$	$35.7\pm4.2$
siJNK-AraC-D2/CA	<b>39.0</b> ± <b>4.7</b>	42.6 ± 4.4	44.3 ± 4.9	45.5 ± 4.9

There was statistically significant enhancement of cell death (p < 0.05, n=3) in siControl- treated cells (both HL60 and AML ex vivo), but not in siJNK-treated cells.