

## Acquisition of Resistance toward HYD1 Correlates with a Reduction in Cleaved $\alpha 4$ Integrin Expression and a Compromised CAM-DR Phenotype

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### Abstract

We recently reported that the  $\beta 1$  integrin antagonist, referred to as HYD1, induces necrotic cell death in myeloma cell lines as a single agent using *in vitro* and *in vivo* models. In this article, we sought to delineate the determinants of sensitivity and resistance toward HYD1-induced cell death. To this end, we developed an HYD1 isogenic resistant myeloma cell line by chronically exposing H929 myeloma cells to increasing concentrations of HYD1. Our data indicate that the acquisition of resistance toward HYD1 correlates with reduced levels of the cleaved  $\alpha 4$  integrin subunit. Consistent with reduced VLA-4 ( $\alpha 4\beta 1$ ) expression, the resistant variant showed ablated functional binding to fibronectin, VCAM-1, and the bone marrow stroma cell line HS-5. The reduction in binding of the resistant cell line to HS-5 cells translated to a compromised cell adhesion-mediated drug resistant phenotype as shown by increased sensitivity to melphalan- and bortezomib-induced cell death in the bone marrow stroma coculture model of drug resistance. Importantly, we show that HYD1 is more potent in relapsed myeloma specimens than newly diagnosed patients, a finding that correlated with  $\alpha 4$  integrin expression. Collectively, these data indicate that this novel D-amino acid peptide may represent a good candidate for pursuing clinical trials in relapsed myeloma and in particular patients with high levels of  $\alpha 4$  integrin. Moreover, our data provide further rationale for continued preclinical development of HYD1 and analogues of HYD1 for the treatment of multiple myeloma and potentially other tumors that home and/or metastasize to the bone. *Mol Cancer Ther*; 10(12); 2257–66. ©2011 AACR.

### Introduction

Multiple myeloma is a disease characterized by the homing and uncontrolled growth of malignant plasma cells within the confines of the bone marrow (1, 2). Despite the recent advances in therapy, multiple myeloma remains an incurable disease. Fourteen thousand new cases of multiple myeloma are diagnosed each year in the United States with a 5-year survival rate of 37% (3). Although standard therapy will typically cause an initial response, myeloma patients ultimately develop drug resistance and become unresponsive to a variety of anti-

cancer agents, a phenomenon known as multidrug resistance. Clinical observations indicate that despite divergent genetic changes typical of myeloma, current therapy is not curative in any subset of patients.

We, as well as others, previously reported that adhesion of myeloma and leukemia cells to components of the extracellular matrix (ECM) is sufficient to cause drug resistance (4–13). We recently used a D-amino acid containing peptide (kikmvswkg), referred to as HYD1, known to block adhesion of prostate cells to ECMs (14, 15) and found that, in addition to blocking adhesion of multiple myeloma cells to fibronectin, HYD1 induced caspase-independent cell death in myeloma cell lines as a single agent *in vitro* and *in vivo* (16). Experimental evidence indicated that in prostate cancer cell lines, HYD1 interacts with  $\alpha 3$  and  $\alpha 6$  integrin (15). To delineate the molecular pathway of HYD1-induced caspase-independent cell death, we developed an acquired isogenic HYD1-resistant H929 myeloma cell line that we refer to as H929-60 cells. In this article, we show that the acquisition of resistance toward HYD1 does not result in a phenotype that is cross-resistant to other agents used to treat myeloma, including bortezomib and melphalan. Moreover, acquisition of resistance toward HYD1 occurs at a cost in overall fitness, as the resistant variant shows reduced binding to ECMs and is

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not resistant to melphalan- or bortezomib-induced cell death in the bone marrow coculture model system. Finally, in this article, we show that specimens obtained from relapsed myeloma patients were significantly more sensitive to HYD1-induced cell death than specimens obtained from newly diagnosed patients. Collectively, our data continue to support that HYD1 is an attractive agent for treating multiple myeloma patients and may be an important strategy for the treatment of relapsed disease.

## Materials and Methods

### Cell culture

NCI-H929, U266 and RPMI-8226, and HS-5 cells were obtained from the American Type Culture Collection. Melphalan-resistant 8226/LR and U266/LR6 were developed and characterized by Dr. William Dalton's laboratory, Moffitt Cancer Center, Tampa, FL (7, 17, 18). Myeloma cell lines were tested for secretion of  $\kappa$  (H929) or  $\lambda$  (RPMI-8226) levels by ELISA and mycoplasma every 4 months. Resistance levels of drug-selected cell lines are monitored every 4 months. 293FT cells were obtained from Invitrogen and grown in Iscove's Dulbecco's Modified Eagle's Medium (Cellgro) supplemented with 10% FBS. Normal bone marrow aspirate was purchased from Lonza Inc. Mesenchymal stroma cells (MSC) were generated by plastic adherence of the bone marrow aspirate. MSCs were confirmed by CD105, Stro-1, CD29, and CD73 positivity and CD34, CD33, and CD45 negativity (data not shown). MSCs were grown in minimum essential medium  $\alpha$ /GlutaMAX supplemented with 10% FBS qualified and 1% 100 $\times$  penicillin–streptomycin–glutamine (Invitrogen).

### Chemical reagents, antibodies, and peptides

Please refer to Supplemental Materials and Methods for sources of purchased materials.

### Selection of a drug-resistant cell line

NCI-H929 cells were exposed to increasing concentrations of HYD1 for 24 weeks. The emerging drug-resistant cell line was named H929-60 and is maintained in media containing 60  $\mu$ g/mL HYD1, once a week for 24 hours.

### Cell death analysis

After treatment with HYD1, cells were washed with PBS and incubated with 2 nmol/L TO-PRO-3 iodide for 45 minutes. The cells were analyzed for fluorescence intensity with the use of a FACSCalibur (BD Biosciences).

### Measurement of $\Delta\psi_m$

After treatment, cells were incubated for 15 minutes with 15 nmol/L of 3,3'-dihexyloxycarbocyanine iodide (Invitrogen). Cells were washed and resuspended in PBS, and the loss of mitochondrial membrane potential was measured with FACScan.

### ATP measurement

Treated and control cells were lysed in RIPA buffer, and ATP concentrations were measured by the ENLITEN ATP bioluminescence detection kit per manufacturer's instructions (Promega). ATP levels were normalized to the protein content of the lysates.

### Confocal microscopy

To assess whether (i) HYD1 bound the cell surface and (ii) whether binding of HYD1 was reduced in the resistant cell line, FAM-HYD1 was used to image peptide binding in the parental and resistant cell line by confocal microscopy. A 35-mm glass bottom microwell dishes (Mattek Cultureware) were plated with 10  $\mu$ L Cell-tak (BD Biosciences) per manufacturer's instructions. Media containing 1  $\mu$ mol/L Alexa Fluor 594 wheat germ agglutinin (WGA; Invitrogen) and 20  $\mu$ mol/L Hoechst 33342 (Invitrogen) was placed back in the plates, and the samples were incubated for 30 minutes. After 30 minutes, the cells were washed and treated with media containing 6.25  $\mu$ g/mL FAM-conjugated HYD1 for 10 minutes. Samples were immediately viewed with a Leica DMI6000 confocal microscope (Leica Microsystems). Gain, offset, and pinhole setting were identical for all samples within the treatment group.

### Reverse transcriptase polymerase chain reaction

Reverse transcriptase polymerase chain reaction (RT-PCR) was used to determine whether the decrease in  $\alpha$ 4 integrin protein levels in the resistant cell line was due to decreased transcription. RNA was extracted from log growth cells with RNeasy columns (Qiagen) per manufacturer's instructions. First-strand cDNA synthesis was carried out with the Quantitect Probe RT-PCR Kit (Qiagen) per manufacturer's instructions. Real-time PCR primers for  $\alpha$ 4 integrin were obtained from Applied Biosystems. The gene expression level was normalized by the endogenous control gene *GAPDH*. Real-time PCR reactions were carried out with ABI 7900 Sequence Detection System (Applied Biosystems).

### Cell adhesion to ECM proteins and stroma

Cell adhesion assays using ECMs were conducted as previously described (19). For stromal adhesion, 10,000 HS-5 cells or MSCs were incubated overnight on immunosorb 96-well plates (Nunc). H929 and H929-60 cells were incubated with 1  $\mu$ mol/L of 5-chloromethylfluorescein diacetate (CMFDA; Invitrogen) for 30 minutes, washed, and incubated for 45 minutes to allow unbound dye to diffuse out of the cells. Labeled cells were allowed to adhere for 2 hours and nonadherent cells were removed with 3 washes in PBS. Intensity was read on a fluorescence plate reader (9).

### Transfection of short hairpin RNAs

Short hairpin RNA (shRNA)-targeting strategies were used to determine whether  $\alpha$ 4 integrin expression was causally related to HYD1-induced cell death.  $\alpha$ 4 (TRCN0000029656) and  $\beta$ 1 (TRCN0000029645) shRNA and

nonsilencing clone sets were purchased from Open Biosystems and transfected into a lentivirus using the BLOCK-iT Lentiviral Pol II miR RNAi Expression System (Invitrogen). At 72 hours of infection, 1  $\mu\text{g}/\text{mL}$  puromycin (Invitrogen) was added to RPMI-8226 cells to allow for the selection of a stable population of cells. For the H929 cell line, transient infections were used to reduce integrin expression.

### Biotin-HYD1 pull-down assay

To identify HYD1-interacting proteins, biotin-conjugated HYD1 was used as bait as previously described (15). Briefly, membrane pellets were solubilized in AP buffer containing 0.2% NP40, and protein was quantified with BCA reagents (Pierce). Five hundred micrograms of biotin-HYD1 was bound to 30  $\mu\text{L}$  of UltraLink NeutraAvidin Plus beads (Pierce) for 1 hour in a buffer containing 0.5 mmol/L KCl, 0.3 mmol/L  $\text{KH}_2\text{PO}_4$ , 27.6 mmol/L NaCl, and 1.6 mmol/L  $\text{Na}_2\text{HPO}_4$  at pH 7.4. After 1 hour, the beads were washed twice in AP buffer and 50  $\mu\text{g}$  of membrane extracts were added to a total volume of 500  $\mu\text{L}$  and incubated with beads for 18 hours at 4 degrees. The beads were washed 3 times in AP buffer containing 0.2% NP40, and samples were suspended in SDS-PAGE sample buffer and bound proteins were resolved by SDS-PAGE.

### Isolation of CD138-positive and CD138-negative populations derived from multiple myeloma specimens

To determine whether HYD1 was active in primary patient specimens, 7 newly diagnosed and 7 relapsed

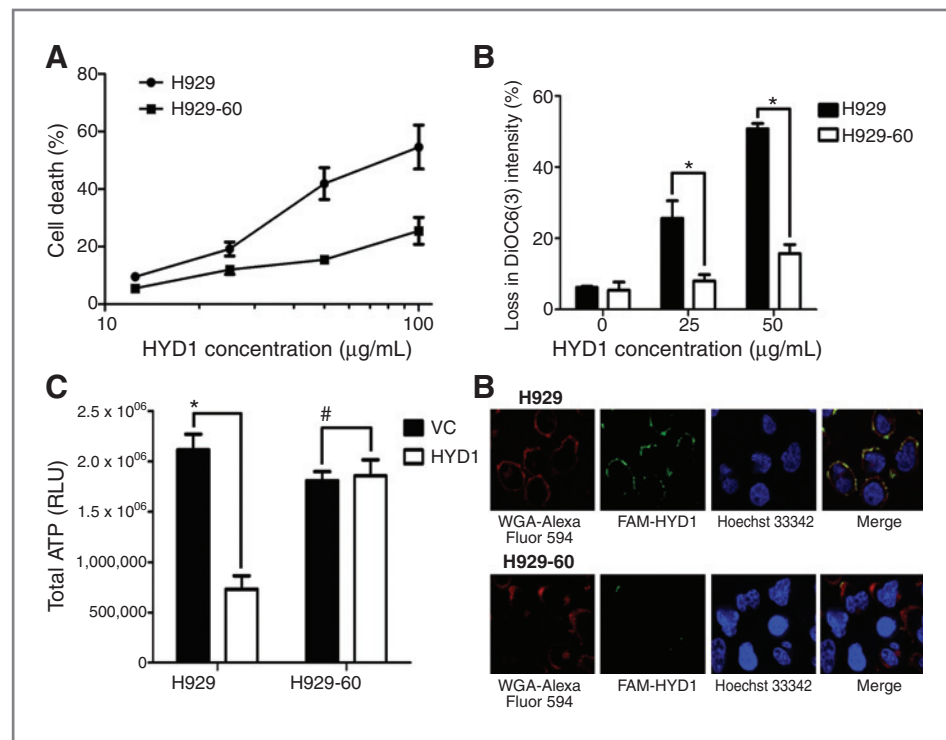
specimens were obtained. Myeloma patients were consented through the Total Cancer Care tissue banking protocol per Institutional Review Board regulations. Mononuclear cells were separated from human blood through the use of Ficoll-Paque PLUS (GE Healthcare). After separation, CD138-positive cells were sorted using 25 MS MACS Separation Columns (Miltenyi Biotec) and CD138 microbeads (Miltenyi) per manufacturer's instructions. For each specimen obtained,  $\alpha 4$  integrin surface expression was determined by fluorescence-activated cell sorting (FACS) analysis and HYD1-induced cell death was determined by TO-PRO-3 staining and FACS analysis in the CD138-positive and CD138-negative fraction.

## Results

### H929-60 cells are resistant to HYD1-induced cell death but do not show cross-resistance to other active myeloma agents

As shown in Fig. 1A, the H929-60-acquired resistant cell line is significantly resistant to HYD1-induced cell death ( $P < 0.05$ , ANOVA) when compared with the parental H929 cells as measured by TO-PRO-3 positivity and FACS analysis. HYD1-induced cell death was previously characterized by the loss of mitochondrial membrane potential, ATP depletion, and an increase in reactive oxygen species (ROS). To determine whether the acquisition of resistance occurred upstream or downstream of mitochondria dysfunction, the mitochondria membrane potential, ATP levels, and ROS levels were compared following HYD1 treatment in the resistant (H929-60) and

**Figure 1.** H929-60 cells are resistant to HYD1-induced cell death and showed reduced binding of FAM-HYD1 to the cell surface. A, H929 and H929-60 cells were incubated with varying concentrations of HYD1 for 6 hours, and HYD1-induced cell death was determined by TO-PRO-3 staining and FACS analysis (ANOVA,  $P < 0.05$ ,  $n = 9$ ). B, loss of mitochondrial membrane permeability was determined by DiOC<sub>6</sub> staining following 2 hours of HYD1 treatment in H929 and H929-60 cells (\*,  $P < 0.05$ ;  $n = 9$ , Student *t* test). C, ATP levels were determined in the H929 and H929-60 cell lines following 6 hours of HYD1 (50  $\mu\text{g}/\text{mL}$ ) treatment. (\*,  $P < 0.05$  and #,  $P > 0.05$ ;  $n = 9$ , Student *t* test). D, H929 and H929-60 cells were stained with Alexa Fluor 594 WGA and Hoechst 33342 for 30 minutes. FAM-HYD1 (6.25  $\mu\text{g}/\text{mL}$ ) was added 10 minutes before analysis by confocal microscopy. The experiment was repeated 3 independent times and shown is a representative experiment.



sensitive parental cell line (H929). H929-60 cells were shown to be resistant to the loss of mitochondrial membrane potential (Fig. 1B), and ATP levels were not depleted following HYD1 treatment (Fig. 1C). Finally, ROS levels were reduced following HYD1 treatment in the resistant cell line compared with the parental cell line (see Supplementary Fig. S1). We used a FAM-conjugated HYD1 peptide to determine whether the H929-resistant variant showed a reduction in binding of FAM-HYD1 to the cell membrane. As shown in Fig. 1D, FAM-HYD1 localizes to the plasma membrane in the parental cell line. Furthermore, the localization of FAM-HYD1 was not evenly distributed across the cell membrane, but rather FAM-HYD1 showed punctuated staining in the parental cell line, suggesting potential clustering of the binding target. H929-60 cells treated with 6.25  $\mu\text{g}/\text{mL}$  FAM-HYD1, showed a 2.7-fold reduction in FAM-HYD1 binding relative to the parental cell line as determined by FACS analysis (see Supplementary Fig. S2). Collectively, these data indicate that the mechanism causative for resistance toward HYD1 occurs upstream of mitochondrial dysfunction and generation of ROS and that the resistant mechanism is likely the result of qualitative or quantitative changes in the HYD1-binding complex located on the cell membrane.

We hypothesized that because HYD1 induces necrotic cell death, selection with HYD1 would not result in a phenotype that conferred resistance to other agents commonly used to treat myeloma. To address this question, we compared the  $\text{IC}_{50}$  values of H929-60 and the parental H929 cells to the alkylating agent, melphalan, the topoisomerase II inhibitor, mitoxantrone, and the proteasome inhibitor, bortezomib. As predicted, H929-60 cells were not resistant to other classes of agents known to induce apoptosis commonly used to treat multiple myeloma (see Table 1). These data further support the potential advantage of targeting necrosis in combination with inducers of apoptosis for the treatment of multiple myeloma, as cross-resistance between these 2 agents is unlikely to occur during the course of drug treatment.

**Table 1.** H929-60 cells are not resistant to standard myeloma therapies

Compound	H929 $\text{IC}_{50}$ ( $n = 3$ )	H929-60 $\text{IC}_{50}$ ( $n = 3$ )
Melphalan	$11.3 \pm 2.5 \mu\text{mol}/\text{L}$	$13.1 \pm 3.7 \mu\text{mol}/\text{L}$
Mitoxantrone	$1.42 \pm 0.4 \mu\text{mol}/\text{L}$	$1.05 \pm 0.7 \mu\text{mol}/\text{L}$
Bortezomib	$11.3 \pm 5.1 \text{nmol}/\text{L}$	$12.2 \pm 3.9 \text{nmol}/\text{L}$

NOTE: H929 and H929-60 cells were treated with increasing concentrations of melphalan, mitoxantrone, or bortezomib for 24 hours. After 24 hours, cells were stained with annexin V-fluorescein isothiocyanate/propidium iodide for 45 minutes and acquired on FACScan.  $\text{IC}_{50}$  values were determined by linear regressions. Shown is the mean and SDs of 3 independent experiments.

### Acquisition toward resistance to HYD1 results in reduced expression of $\alpha 4$ , $\beta 1$ integrin and ablated functional binding to fibronectin and VCAM-1

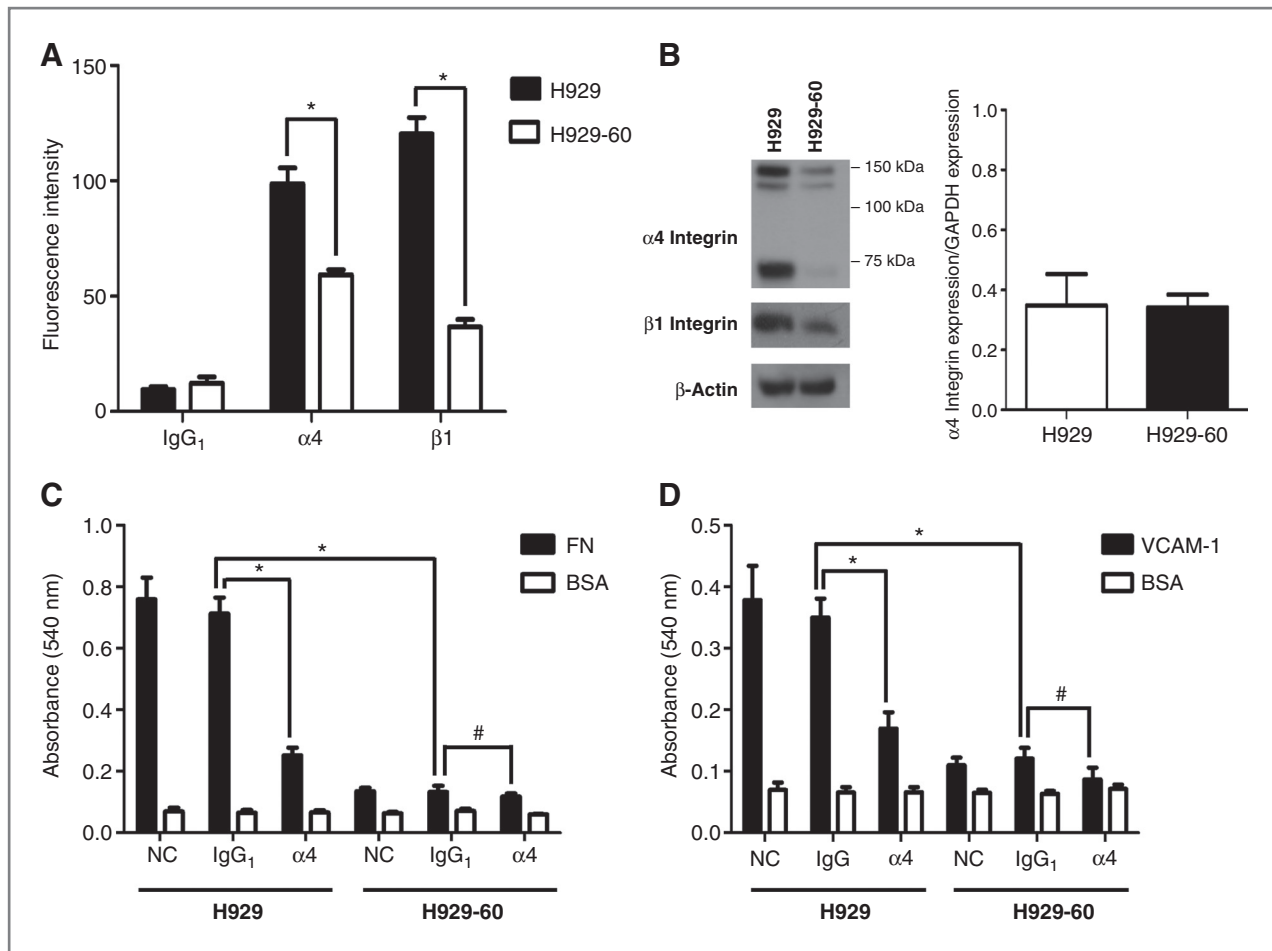
To determine whether the acquisition of resistance correlated with quantitative changes in integrin expression, we screened multiple  $\alpha$  integrin subunits ( $\alpha 4$ ,  $\alpha 5$ ,  $\alpha V$ , and  $\alpha 6$  integrin; data not shown) that are commonly expressed in hematopoietic cells and determined that  $\alpha 4$  integrin was the most abundant integrin expressed on the parental H929 cell line and expression was reduced in the resistant cell line (see Fig. 2A). Using FACS analysis, we determined that  $\alpha 4$  integrin was found to be reduced by 1.8-fold. As shown in Fig. 2B, the reduction at the cell surface corresponded to reduced protein expression with a whole-cell lysate preparation. Interestingly, when examining the whole-cell lysate, the most dramatic decrease was the cleaved form of  $\alpha 4$  integrin. In some cell types, the mature 150 kDa  $\alpha 4$  integrin is cleaved into a nondisulfide linked 80 and 70 kDa fragment. For example, activation of T cells was previously reported to correlate with increased cleavage of the mature  $\alpha 4$  integrin (20). However, the cleavage of  $\alpha 4$  integrin was found to not alter the adhesive properties of VLA-4 integrin to fibronectin or VCAM1 (21). The attenuation of protein expression is posttranscriptionally regulated as the parental and resistant cell line showed equal levels of  $\alpha 4$  mRNA (Fig. 2B). Further studies are warranted to delineate the posttranscriptional regulation of cleaved  $\alpha 4$  integrin in the resistant cell line.

We next sought to determine whether a reduction in the expression of VLA-4 integrin resulted in a functional reduction in adhesion to extracellular matrices. To this end, we compared the levels of adhesion of the sensitive and resistant cell line to fibronectin and the more specific ligand for  $\alpha 4$  integrin, VCAM-1. In Fig. 2C and D, H929-60 cells showed a dramatic reduction in the binding to fibronectin and VCAM-1. An  $\alpha 4$  integrin blocking was used as a positive control for blocking the parental cell line to fibronectin and VCAM-1.

### Reducing the expression of $\alpha 4$ and $\beta 1$ integrins is causative for resistance to HYD1-induced cell death in H929 and 8226 multiple myeloma cells.

Considering H929-60 cells were resistant to HYD1-induced cell death and showed reduced surface expression of the  $\alpha 4$  integrin subunit, we next determined whether reducing the expression of  $\alpha 4$  integrins using shRNA-targeting strategies in myeloma cell lines was sufficient to induce resistance toward HYD1-induced cell death. As shown in Fig. 3A and B, reducing the expression of  $\alpha 4$  integrin conferred resistance to HYD1-induced cell death ( $P < 0.05$ , Student  $t$  test) in both 8226 and H929 cells.

Because  $\alpha 4$  integrin can heterodimerize with either  $\beta 1$  or  $\beta 7$  integrin, we tested whether reducing  $\beta 1$  integrin was sufficient to induce resistance to myeloma cell lines. As shown in Fig. 3C and D, reducing  $\beta 1$  integrins rendered H929 and 8226 cells resistant to HYD1-induced



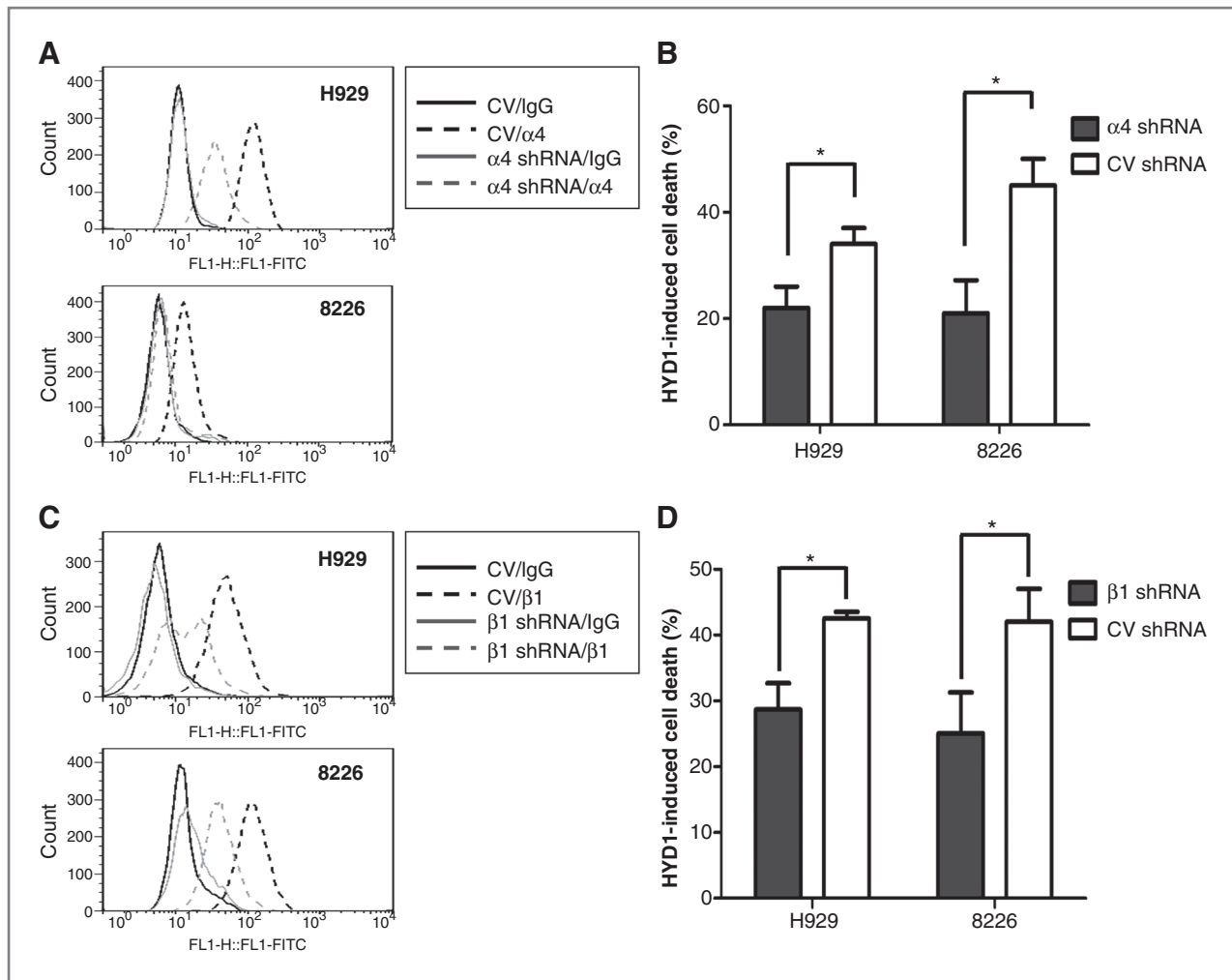
**Figure 2.** H929-60 cells have reduced  $\alpha 4$  integrin protein levels and reduced adhesion to fibronectin (FN) and VCAM-1. A, surface expression of  $\alpha 4$  and  $\beta 1$  integrin on H929 and H929-60 cells was determined by FACS analysis (\*,  $P < 0.05$ ;  $n = 9$ , Student  $t$  test). B, whole-cell lysates of H929 and H929-60 cells were probed for  $\alpha 4$  integrin,  $\beta 1$  integrin, and  $\beta$ -actin by Western blot analysis.  $\alpha 4$  Integrin mRNA levels were determined by real-time PCR.  $\alpha 4$  Integrin mRNA expression levels were normalized by dividing with GAPDH levels ( $P > 0.05$ ,  $n = 3$ ). C and D, H929 and H929-60 cells were incubated with  $\alpha 4$ -blocking antibody or IgG control antibody for 30 minutes and subsequently adhered to fibronectin (40  $\mu\text{g}/\text{mL}$ ; C)- or VCAM-1 (10  $\mu\text{g}/\text{mL}$ ; D)-coated plates for 2 hours. A representative of each experiment is shown (\*,  $P < 0.05$  or #,  $P > 0.05$ , Student  $t$  test). GAPDH, glyceraldehyde-3-phosphate dehydrogenase; IgG<sub>1</sub>, immunoglobulin G<sub>1</sub>.

cell death ( $P < 0.05$ , Student  $t$  test).  $\beta 1$  integrin can potentially heterodimerize with 11 different  $\alpha$  subunits. The observation that reducing  $\alpha 4$  or  $\beta 1$  integrin gave similar levels of protection indicates that the  $\alpha 4\beta 1$  integrin is the predominant  $\beta 1$  integrin partner associated with HYD1-induced cell death in myeloma cells. The observation that reducing integrin expression afforded only partial resistance may be due to (i) residual levels of  $\alpha 4\beta 1$  integrin remaining on the cell surface or (ii)  $\alpha 4\beta 1$  represents only one component of the binding complex required for HYD1-induced cell death.

#### **Biotin-HYD1 interacts with $\alpha 4$ integrin, and reduced binding was observed in the acquired drug resistant cell line.**

Previous data indicated that biotin-HYD1 associated with  $\alpha 3$  and  $\alpha 6$  on prostate cancer cell lines (15). We used a

similar strategy to determine whether biotin-HYD1 interacted with an  $\alpha 4$  integrin containing complex and whether that interaction was attenuated in the resistant cell line. The total membrane lysate was used as a control for detection of  $\alpha 4$  integrin. Again the reduction in the cleaved  $\alpha 4$  integrin subunit in membrane extracts (although not as dramatic as the whole-cell lysate) was most prominent in the resistant cell line compared with the mature  $\alpha 4$  integrin subunit. In addition, as shown in Fig. 4, using biotin-HYD1, we show that the resistant cell line shows preferentially decreased binding for the cleaved  $\alpha 4$  subunit. Our data indicate that the cleaved form may be important with respect to functional consequences of HYD1-induced cell death. Further studies are warranted to fully understand the significance of the cleaved  $\alpha 4$  integrin subunit in mediating HYD1-induced cell death and the cell adhesion-mediated drug resistant (CAM-DR) phenotype.

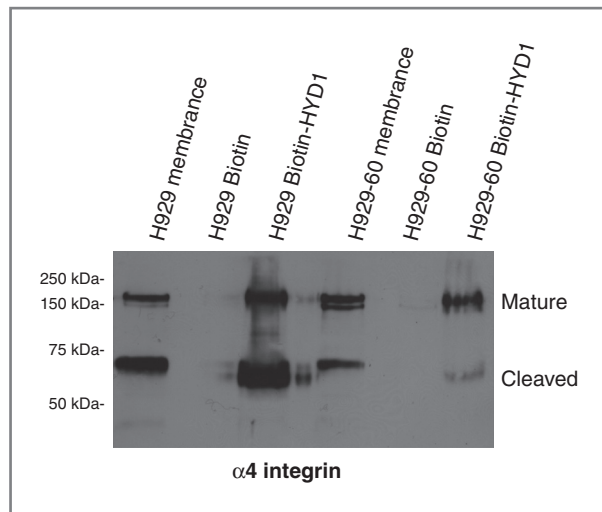


**Figure 3.** Reducing the expression of  $\alpha$ 4 and  $\beta$ 1 integrins caused partial resistance to HYD1-induced cell death in H929 and 8226 cells. A and C, in H929 cells,  $\alpha$ 4 and  $\beta$ 1 integrin expression was reduced via transient infection. A, at 72 hours postinfection of shRNA,  $\alpha$ 4 and  $\beta$ 1 integrin expression was determined by FACS analysis. 8226 cells were stably infected with  $\alpha$ 4 integrin shRNA,  $\beta$ 1 integrin shRNA, or control vector shRNA. FACS analysis was used to determine expression of  $\alpha$ 4 and  $\beta$ 1 integrin. Shown is a representative histogram from one experiment. B and D, H929 and 8226 cells were treated with 50  $\mu$ g/mL and 100  $\mu$ g/mL of HYD1, respectively, for 6 hours. After 6 hours, cell death was analyzed by TO-PRO-3 staining and FACS analysis. (\*,  $P < 0.05$ ;  $n = 9$ , Student  $t$  test).

### H929-60 cells displayed reduced binding to bone marrow stromal cells and show a compromised CAM-DR phenotype

In addition to attenuated adhesion to fibronectin and VCAM-1, H929-60 cells also exhibit a reduction in adhesion to the HS-5 stromal cell line and MSCs (Fig. 5A and B). We reasoned that acquisition of resistance toward HYD1, which correlated with reduced functional binding to fibronectin, VCAM-1, HS-5 stromal cells, and MSCs would likely result in a compromised CAM-DR phenotype. To test this premise, we used the HS-5 coculture model of drug resistance. H929- and the HYD1-resistant variant H929-60 cells were treated with either melphalan or bortezomib in the presence or absence of HS-5/GFP bone marrow stromal cells. As shown in Fig. 5C and D, the HYD1-resistant cell line was not resistant in the coculture

bone marrow stroma model to melphalan or bortezomib, respectively. There is some discrepancy in the literature whether coculturing myeloma cells with stroma cells causes resistance to bortezomib (22). The apparent discrepancy may be the result of endpoints that measure growth compared with cell death, dose of bortezomib, or the scheduling of how long myeloma cells interact with the stroma before drug exposure. In addition, exposure to bortezomib for 24 hours was shown to downregulate  $\alpha$ 4 integrin levels and thus drug scheduling could also impact observed results (23). However, our results are consistent with recent reports showing that myeloma cells are resistant to bortezomib in MSC coculture models (5). Together, these data indicate that as myeloma cells are selected for resistance to HYD1, they are losing resistance to standard chemotherapy in the context of the bone



**Figure 4.** Biotin–HYD1 interacts with  $\alpha 4$  integrin and binding to cleaved  $\alpha 4$  integrin is attenuated in the resistant H929-60 cells. Fifty micrograms of H929 and H929-60 protein obtained from membrane fractions was added to biotin or biotin–HYD1-bound NeutrAvidin beads for 18 hours as described in Materials and Methods. Biotin–HYD1 and biotin-bound proteins were denatured in SDS sample buffer, and  $\alpha 4$  integrin levels were detected by Western blot analysis. The experiment was repeated twice and a representative experiment is shown.

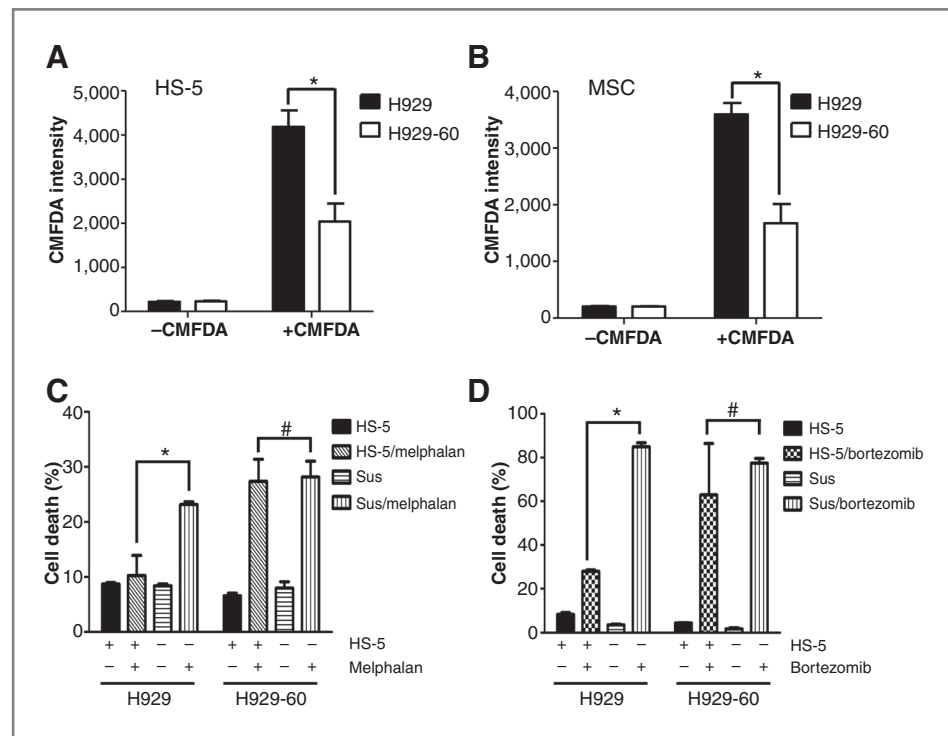
marrow microenvironment because of reduced capacity to adhere to bone marrow stroma cells. Interestingly, the inverse seems to be true because myeloma cell lines selected for resistance to melphalan (8226/LR5 and U266/LR6; refs. 7, 17, 18) show increased sensitivity to

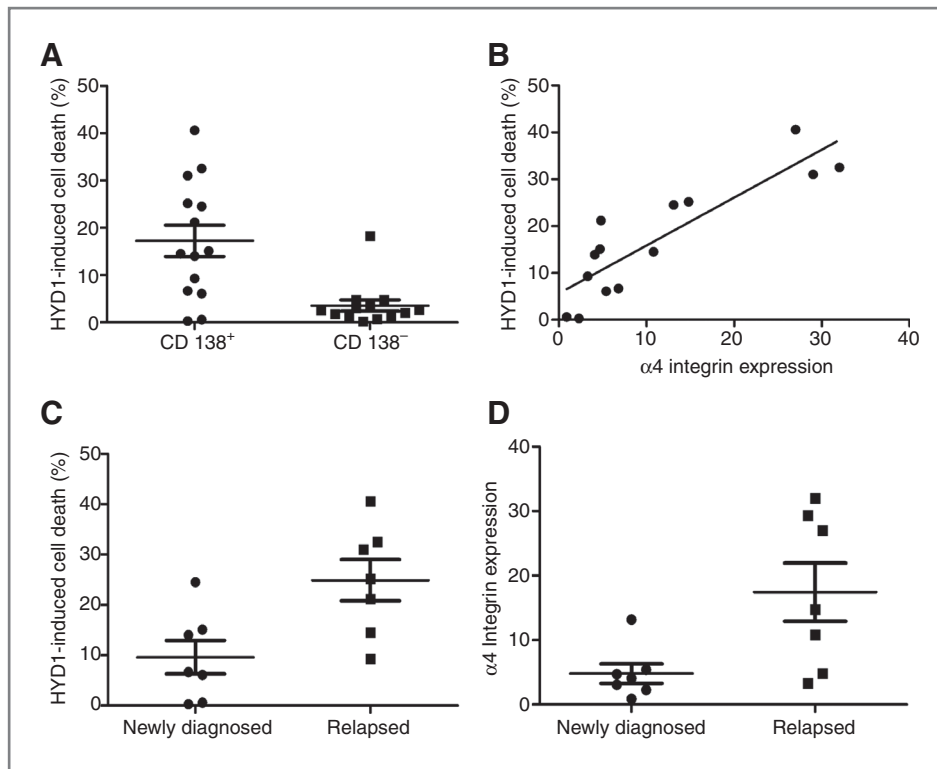
HYD1-induced cell death (see Supplementary Table S1). This correlated with increased  $\alpha 4$  expression and is consistent with previous reports that selection with melphalan coincides with increased  $\alpha 4$  integrin expression (6). The difference in sensitivity toward HYD1 or the comprised CAM-DR phenotype can not be attributed to changes in growth kinetics as the sensitive and resistant cell line show similar rates of proliferation (see Supplementary Fig. S3).

### HYD1-induced cell death is increased in relapsed myeloma patient specimens compared with newly diagnosed specimens and correlates with $\alpha 4$ integrin expression

To determine whether primary myeloma specimens are sensitive to HYD1-induced cell death, we collected 7 newly diagnosed and 7 relapsed primary myeloma specimens. Immunomagnetic beads were used to enrich for CD138-positive malignant plasma cell fractions. As shown in Fig. 6A, the CD138-positive tumor population was more sensitive to HYD1-induced cell death than the CD138-negative population. In addition to measuring cell death, we determined by FACS analysis the levels of  $\alpha 4$  integrin expression in the CD138-positive cells. As shown in Fig. 6B, HYD1-induced cell death positively correlates with  $\alpha 4$  integrin expression. These data obtained in patient specimens correlating  $\alpha 4$  integrin expression and sensitivity to HYD1-induced cell death is consistent with data generated in cell lines (see Table 1). Importantly, we observed that HYD1 was significantly ( $P < 0.05$ , Student *t* test) more active in relapsed patients

**Figure 5.** H929-60 cells display reduced binding to HS-5 cells and MSCs and are not resistant to bortezomib or melphalan in the stroma coculture model. A and B, H929 and H929-60 cells preincubated with CMFDA dye were adhered to 10,000 HS-5 cells or MSCs for 2 hours. After 2 hours, unadherent cells were removed, and the fluorescence intensity was measured on a fluorescent plate reader. C and D, H929 and H929-60 cells were adhered to HS-5/GFP cells for 24 hours. After 24 hours, samples were treated with 15  $\mu\text{mol/L}$  melphalan or 16 nmol/L bortezomib for 24 hours. Cell death was analyzed by a FACScalibur with the GFP-expressing HS-5 cells being excluded from the analysis. In A and C, a representative of 3 independent experiments is shown. (\*,  $P < 0.05$ ;  $n = 9$  or #,  $P > 0.05$ ;  $n = 9$  Student *t* test).





**Figure 6.** HYD1-induced cell death in CD138<sup>+</sup> patient samples correlates with  $\alpha 4$  integrin expression. A, CD138<sup>+</sup> and CD138<sup>-</sup> cells were treated with 100  $\mu$ g/mL HYD1 for 24 hours. After 24 hours, cell death was measured by TO-PRO-3 staining and FACS analysis. ( $P < 0.05$ , Student  $t$ -test). B,  $\alpha 4$  integrin expression was compared with HYD1-induced cell death by a Pearson correlation coefficient. The CD138<sup>+</sup> population showed a significant correlation between  $\alpha 4$  integrin expression and HYD1-induced cell death ( $P < 0.05$ ). C and D, specimens were separated into 2 groups depending on the clinical diagnosis of either newly diagnosed or specimens obtained from patients that were considered relapsed. C, CD138<sup>+</sup> cells were treated with 100  $\mu$ g/mL HYD1 for 24 hours. After 24 hours, cell death was measured by TO-PRO-3 staining and FACS analysis (Student  $t$  test,  $P < 0.05$ ). D, CD138<sup>+</sup> cells were used to analyze for  $\alpha 4$  integrin expression by FACS analysis. (Student  $t$  test,  $P < 0.05$ ).

than newly diagnosed patients (see Fig. 6C). Finally as shown in Fig. 6D, we show that  $\alpha 4$  integrin levels are increased in CD138-positive cells isolated from relapsed myeloma patients compared with newly diagnosed patients ( $P < 0.05$ , Student  $t$  test). Together, these data suggest that  $\alpha 4$  integrin expression is selected for over the course of drug treatment and may contribute to the eventual failure to therapeutically manage multiple myeloma. In addition, patients with high levels of  $\alpha 4$  expression may benefit from combination strategies that include targeting this specific integrin complex such as Natalizumab (humanized  $\alpha 4$  antibody; ref. 24) or HYD1.

## Discussion

The development of an isogenic resistant cell line has been used by many investigators as model systems for rapidly delineating molecular determinants of response for novel as well as clinically approved agents (25–29). Selection with HYD1 resulted in a cell line with compromised adhesive interactions and failure to display a resistant phenotype in a bone marrow stroma coculture model of drug resistance. We propose that to ensure adequate designs of trials, it is essential to define biomarkers of response with patient specimens in early phases of drug development, as response markers can take time to validate and often lag behind the design of early clinical trials (30). To this end, we show that  $\alpha 4$  integrin expression positively correlated with HYD1-induced cell death. In addition, HYD1 was more potent

in specimens obtained from relapsed myeloma patients than newly diagnosed myeloma patients. These data suggest that HYD1 will likely be most effective in relapsed patients showing high levels of  $\alpha 4$  expression. It is attractive to speculate that agents that disrupt cell adhesion may indeed be a good strategy for comprising the overall fitness of cells in the context of the bone marrow microenvironment. Agents that disrupt cell adhesion may be critical to test the development of evolutionary based double bind strategies in which resistance to a drug is predicted to occur at the cost of fitness within the niche. The use of double bind strategies is a unique concept recently proposed by Gatenby and colleagues for delaying the emergence of resistant variants in the treatment of cancer and indeed HYD1 may fit the criteria for an agent to test this unique therapeutic strategy (31).

Our data indicate that reducing  $\alpha 4\beta 1$  integrin expression was sufficient to confer partial drug resistance. However, because only partial resistance was observed, we predict that additional components contained within the HYD1-binding complex may contribute to HYD1-induced cell death. Further studies are warranted for delineating the entire HYD1-interacting complex with unbiased approaches, which is currently a focus of our laboratory. In the drug-resistant variant cell line, we observed a predominant reduction in the cleavage of  $\alpha 4$  integrin compared with the mature form. It is intriguing to speculate whether the cleaved fragment of  $\alpha 4$  integrin is required for HYD1-induced cell death and whether the



cleaved form has any significance with respect to disease progression in multiple myeloma.

Recent data show that targeting  $\alpha 4$  integrin in a syngeneic mouse 5TGM1 model via monoclonal antibody treatment reduced the tumor burden in the bone marrow, spleen, and liver (32). Moreover, the VCAM-1/VLA-4 axis increases MIP-1  $\alpha$  and  $\beta$  levels and also increases the ability of myeloma cells to support osteoclastogenesis (33). On the basis of these findings, it will be important to determine whether HYD1 inhibits the ability of myeloma cells to disrupt bone homeostasis by either inhibiting the activation of osteoclasts or disrupting the ability of myeloma cells to inhibit the differentiation of osteoblasts (34–36). Another strategy for targeting integrins is to inhibit pathways required for inside-out activation of VLA-4 integrins. Thus, VLA-4 can be modulated by regulating the affinity for ligand as well as clustering or avidity of the integrin heterodimer (37, 38). A potential strategy could include targeting Rap1 that is known to be required for inside-out activation of VLA-4 (39). Another approach is inhibition of CXCR4, where recently, Azab and colleagues showed that AMD3100 inhibits adhesion of myeloma cells to stroma and sensitized myeloma cells to chemotherapy in coculture models of drug resistance (5). However, HYD1 is unique to our knowledge, as in addition to

blocking cell adhesion, HYD1 induces necrotic cell death directly on the myeloma cell, a finding that was not observed with  $\alpha 4$ -blocking antibody or RGD-containing peptides (data not shown). Historically, drug development has focused on aberrations in signaling intrinsic to the tumor cells using unicellular models. However, we and others have argued that tumors evolve in the context of the microenvironment, and thus it is feasible that some phenotypes observed in tumors such as drug resistance and metastasis will only be expressed in the context of cues derived from the microenvironment (22, 40–42).

### Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

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## Acquisition of Resistance toward HYD1 Correlates with a Reduction in Cleaved $\alpha 4$ Integrin Expression and a Compromised CAM-DR Phenotype

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