

LYMPHOID NEOPLASIA

Growth differentiating factor 15 enhances the tumor-initiating and self-renewal potential of multiple myeloma cells

Toshihiko Tanno,¹ Yiting Lim,¹ Qiuju Wang,¹ Marta Chesi,² P. Leif Bergsagel,² Geoff Matthews,³ Ricky W. Johnstone,³ Nilanjan Ghosh,¹ Ivan Borrello,¹ Carol Ann Huff,¹ and William Matsui¹

¹Division of Hematologic Malignancies, Sidney Kimmel Comprehensive Cancer Center, Johns Hopkins University School of Medicine, Baltimore, MD;

²Comprehensive Cancer Center and Laboratory Medicine and Pathology, Mayo Clinic Arizona, Scottsdale, AZ; and ³Gene Regulation Laboratory, Cancer Therapeutics Program, Peter MacCallum Cancer Centre, Melbourne, VIC, Australia

Key Points

- The tumor microenvironment drives myeloma cell clonogenic growth and self-renewal through GDF15.

Disease relapse remains a major factor limiting the survival of cancer patients. In the plasma cell malignancy multiple myeloma (MM), nearly all patients ultimately succumb to disease relapse and progression despite new therapies that have improved remission rates. Tumor regrowth indicates that clonogenic growth potential is continually maintained, but the determinants of self-renewal in MM are not well understood. Normal stem cells are regulated by extrinsic niche factors, and the tumor microenvironment (TME) may similarly influence tumor cell clonogenic growth and self-renewal. Growth differentiating factor 15 (GDF15) is aberrantly secreted by bone marrow stromal cells (BMSCs) in MM. We found that GDF15 is produced by BMSCs after direct contact with plasma cells and enhances the tumor-initiating potential and self-renewal of MM cells in a protein kinase B- and SRY (sex-determining region Y)-box-dependent manner. Moreover, GDF15 induces the expansion of MM tumor-initiating cells (TICs), and changes in the serum levels of GDF15 were associated with changes in the frequency of clonogenic MM cells and the progression-free survival of MM patients. These findings demonstrate that GDF15 plays a critical role in mediating the interaction among mature tumor cells, the TME, and TICs, and strategies targeting GDF15 may affect long-term clinical outcomes in MM. (*Blood*. 2014;123(5):725-733)

ation factor 15 (GDF15) is aberrantly secreted by bone marrow stromal cells (BMSCs) in MM. We found that GDF15 is produced by BMSCs after direct contact with plasma cells and enhances the tumor-initiating potential and self-renewal of MM cells in a protein kinase B- and SRY (sex-determining region Y)-box-dependent manner. Moreover, GDF15 induces the expansion of MM tumor-initiating cells (TICs), and changes in the serum levels of GDF15 were associated with changes in the frequency of clonogenic MM cells and the progression-free survival of MM patients. These findings demonstrate that GDF15 plays a critical role in mediating the interaction among mature tumor cells, the TME, and TICs, and strategies targeting GDF15 may affect long-term clinical outcomes in MM. (*Blood*. 2014;123(5):725-733)

Introduction

Multiple myeloma (MM) is characterized by the clonal expansion of malignant plasma cells. Advances in MM treatment have significantly improved remission rates, but the vast majority of patients will eventually relapse and succumb to their disease.¹ The continuous risk of relapse suggests that therapy-resistant tumor cells are self-renewing and indefinitely maintain the potential for clonogenic growth. The factors influencing MM self-renewal are poorly understood, but normal stem cells are extrinsically regulated by accessory cells and extracellular matrix components within niches.^{2,3} Therefore specific factors within the tumor microenvironment (TME) may similarly influence MM cell clonogenic growth and self-renewal.

Bone marrow stromal cells (BMSCs) are a major component of the TME in MM and aberrantly secrete several cytokines including growth differentiation factor 15 (GDF15, also known as MIC-1, PTGF- β , PDF, PLAB, PL74, and NAG-1), a member of the transforming growth factor- β family.^{4,6} Elevated circulating levels of GDF15 may correlate with poor clinical outcomes in endometrial, prostate, pancreatic, and colorectal cancers.⁷⁻¹⁰ Similarly, increased GDF15 levels have correlated with disease stage and been associated with worse event-free survival and overall survival in MM patients.⁵ GDF15 may enhance the survival of MM cells in vitro.^{4,5} However, these effects are relatively modest, suggesting that GDF15 influences other properties such as self-renewal and clonogenic growth, which better explain the relationship between circulating cytokine levels and clinical outcomes.

We examined the effects of GDF15 on clonogenic MM growth and found that it increased both tumor cell colony formation in vitro and the engraftment of immunodeficient mice in a protein kinase B- and SRY (sex-determining region Y)-box (SOX2)-dependent manner. To evaluate self-renewal, we carried out serial transplantation studies and found that secondary MM engraftment was increased by the treatment of tumor cells with GDF15 and impaired by the loss of GDF15 within the bone marrow microenvironment. Moreover, the impact of GDF15 on the clonogenic growth and self-renewal of human MM was limited to phenotypically defined tumor-initiating cells (TICs) rather than bulk tumor cells. Finally, we studied the relationship between GDF15 and MM TICs in the clinical setting and found that changes in the serum levels of GDF15 were associated with changes in in vitro clonogenic MM growth and progression-free survival. Therefore GDF15 plays a novel role within the TME by enhancing the tumor-initiating potential and self-renewal of MM TICs, and the development of strategies targeting GDF15 may represent a novel approach for the treatment of MM.

Methods

Cell lines and clinical specimens

Human MM cell lines NCI-H929, RPMI 8226, U266, and MM1.S were obtained from the American Type Culture Collection (Manassas, VA) and

Submitted August 29, 2013; accepted December 2, 2013. Prepublished online as *Blood* First Edition paper, December 17, 2013; DOI 10.1182/blood-2013-08-524025.

The online version of this article contains a data supplement.

The publication costs of this article were defrayed in part by page charge payment. Therefore, and solely to indicate this fact, this article is hereby marked "advertisement" in accordance with 18 USC section 1734.

© 2014 by The American Society of Hematology

KMS11 cells from the Japanese Collection of Research Bioresources (National Institutes of Health Sciences, Japan). Cells were cultured in complete media (CM) as previously described.¹¹ Cells were incubated with human recombinant GDF15 (PeproTech, Rocky Hill, NJ), the Akt-1/2 inhibitor (124018; EMD Millipore, Billerica, MA), or a mouse anti-human GDF15 monoclonal antibody (R&D Systems, Minneapolis, MN) for the indicated doses and time periods. For long-term treatment with GDF15, cells were collected by centrifugation (300g) and resuspended in CM containing GDF15 twice per week.

Clinical bone marrow and peripheral blood samples were obtained from MM patients or healthy donors who gave informed consent in accordance with the Declaration of Helsinki, as approved by the Johns Hopkins Medical Institutes Institutional Review Board. Bone marrow mononuclear cells were isolated by density centrifugation (Ficoll-Paque; Pharmacia, Piscataway, NJ), and plasma cells were isolated using anti-CD138 magnetic microbeads (Miltenyi Biotec, Auburn, CA). Peripheral blood GDF15 levels were measured using the DuoSet enzyme-linked immunosorbent assay for human GDF15 (R&D Systems, Minneapolis, MN) according to the manufacturer's protocol.

Human primary BMSCs were obtained from 4 MM patients and 4 healthy volunteers by culturing bone marrow mononuclear cells (10^6 cells/mL) in 25 cm² tissue culture flasks coated with 0.1% gelatin (Sigma, St Louis, MO) and CM containing 10% horse serum. After 24 hours of incubation at 37°C and 5% CO₂, nonadherent cells were removed and the adherent fraction was cultured in fresh media. MM cell lines and plasma cells derived from clinical specimens (5×10^4 cells) were either combined with BMSCs (5×10^4 cells at passage 2-4) or cultured using cell culture inserts (0.8 μ m pores; Greiner Bio One, Monroe, NC) for 4 days and then evaluated for GDF15 production and tumor cell colony formation.

Tumor cell colony formation

After treatment, tumor cell colony formation in methylcellulose was used to quantify in vitro clonogenic growth according to our previously published methods.^{11,12} MM cell lines (1000 cells/mL) were washed twice in phosphate-buffered saline (PBS) after treatment then plated in triplicate into 35-mm² tissue culture dishes containing 1.2% methylcellulose, 30% fetal bovine serum, 1% bovine serum albumin, 10^{-4} M 2-mercaptoethanol, and 2 mM L-glutamine. For clinical specimens, mononuclear cells were isolated from primary clinical bone marrow aspirates, depleted of CD34⁺ and CD138⁺ cells using magnetic microbeads, treated with GDF15, and then plated (5×10^5 /mL) in methylcellulose cultures containing 10% lymphocyte conditioned media as a source of growth factors. After 10 to 21 days of culture at 37°C and 5% CO₂, tumor colonies consisting of >40 cells were quantified using an inverted microscope. For serial plating studies, tumor cell colonies were harvested, washed twice in CM, and resuspended (1000 cells/mL) in methylcellulose.

Mouse transplantation

All animal studies were approved by the Johns Hopkins Medical Institutes Animal Care Committee. NCI-H929 and RPMI 8226 cells were injected via tail vein into NSG (NOD.Cg-Prkdcscid Il2rgtm1Wjl/SzJ) mice. For ex vivo GDF15 treatment, cells were washed twice in PBS before injection. After the development of hind limb paralysis, or after 6 months, engraftment was confirmed by enzyme-linked immunosorbent assay for serum human immunoglobulin (Ig) κ light chain (Bethyl Laboratories, Montgomery, TX) and human CD138⁺ cells within the bone marrow by flow cytometry. Secondary transplants were carried out by determining the concentration of human CD138⁺ cells within the bone marrow of primary recipients by flow cytometry, then injecting whole bone marrow cells containing 5×10^3 human CD138⁺ cells into secondary recipients. For Vk*My mouse tumor cells (1×10^4), *Gdf15* null or wild-type C57/B16 recipients were conditioned with 6 Gy whole-body irradiation before intravenous injection. Tumor engraftment was evaluated by the identification of monoclonal Ig production by serum protein electrophoresis (The Binding Site, San Diego, CA) and increased (>10%) bone marrow mouse CD138⁺ cells by flow cytometry. Secondary transplants were carried out by injecting whole bone marrow cells containing

1×10^4 mouse CD138⁺ cells into irradiated wild-type C57/B16 animals. TIC frequency was determined using ELDA software.¹³

Real-time polymerase chain reaction

RNA samples were extracted using the RNeasy mini plus kit (Qiagen, Valencia, CA). Quantitative real-time polymerase chain reaction (qRT-PCR) was carried out using the SYBR green system (Life Technologies, Grand Island, NY). All quantitative calculations were performed using the $\Delta\Delta$ Ct method, and primer sequences included: *GDF15*, 5'-CTCCAGATTCCGAG AGTTGC-3', 5'-AGAGATACGCAGGTGCAGGT-3'; *SOX2*, 5'-CCGGTA CGCTCAAAAAGAAA-3', 5'-AGTGTGGATGGGATTGGTGT-3'; and *ACTB*, 5'-ATCCACGAACTACCTTCACTCCAT-3', 5'-CATACTCTGCTTGC TGATCCACATC-3'.

Transient transfections

NCI-H929 cells were transfected with 300 nM *SOX2* or scrambled control siRNA (Santa Cruz Biotechnology, Santa Cruz, CA) using the Nucleofector system (Lonza, Basel, Switzerland), and 24 hours after transfection, the culture media were replaced with fresh media containing 100 ng/mL GDF15. *SOX2* knockdown was evaluated by qRT-PCR 3 days after transfection.

Immunoblot analysis

Western blot analysis was carried out on whole-cell lysates prepared from NCI-H929 cells, separated by 4% to 20% sodium dodecyl sulfate polyacrylamide gel electrophoresis, transferred to a polyvinylidene fluoride membrane (Life Technologies, Grand Island, NY), then incubated with antibodies against phospho-AKT, AKT, and BMI-1 (Cell Signaling Technology, Danvers, MA) as well as *SOX2* and β -ACTIN (EMD Millipore). Antigen detection was carried out using a goat anti-rabbit IgG secondary antibody conjugated to horseradish peroxidase and the Supersignal West Chemiluminescence reagent (Thermo Fisher Scientific).

Flow cytometry and cell isolation

Cells were stained with fluorescein isothiocyanate-conjugated mouse anti-human CD138, R-phycoerythrin rat anti-mouse CD138, or relevant isotypic control antibodies (BD Pharmingen, San Diego, CA) for 30 minutes at 4°C. Cells were stained for aldehyde dehydrogenase (ALDH) activity using the Aldefluor reagent (Stem Cell Technologies, Vancouver, BC, Canada) according to the manufacturer's instructions. Cells were subsequently washed and resuspended in PBS containing 5 μ M propidium iodide (PI; Sigma) and analyzed on a FACSCalibur flow cytometer (Becton Dickinson, Mountain View, CA) or isolated using a FACS Aria II cell sorter. Cells were initially gated to exclude PI⁺ cells and then analyzed for CD138 or ALDH expression. After cell sorting, flow cytometry demonstrated <5% contamination by relevant antigen-expressing cells.

Data analysis

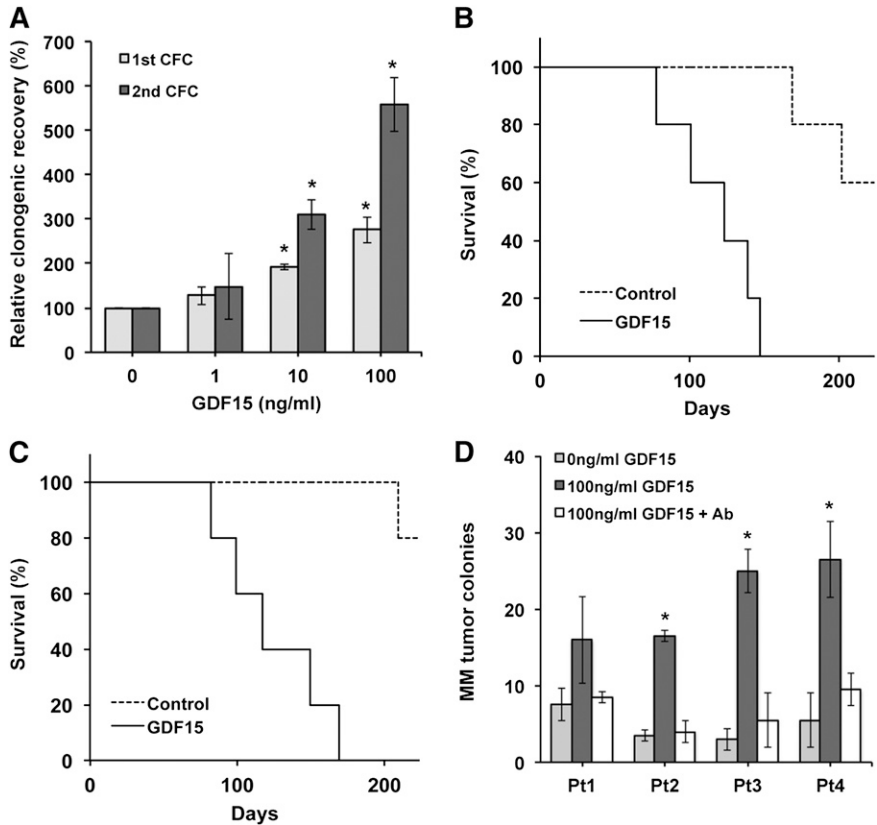
A Student *t* test was used to detect differences, and significance was defined as *P* < .05. Association between trends in serum GDF15 levels and MM-colony-forming unit (CFU) counts in 15 MM patients was determined using the Fisher exact test. Survival rate and progression-free survival were expressed as the percentage of surviving patients calculated using the Kaplan-Meier method. The log-rank test was used to test the difference between the study groups. These analyses were carried out in STATA version 12 (StataCorp, College Station, TX).

Results

GDF15 enhances clonogenic MM growth and self-renewal

To determine the effects of GDF 15 on clonogenic MM growth, we quantified the effects of GDF15 on the colony-forming potential of

Figure 1. GDF15 enhances the clonogenic growth of MM. (A) Primary and secondary colony formation (CFC) of NCI-H929 cells cultured with GDF15 for 7 days. **P* < .05 compared with 0 ng/mL GDF15. (B) Survival of NSG mice after injection of NCI-H929 cells treated ex vivo with or without GDF15 (100 ng/mL) for 14 days (*n* = 5 for each condition). (C) Survival of mice after secondary transplantation of GDF15 cells. (D) MM colony formation by primary clinical bone marrow specimens after treatment with GDF15 ± anti-GDF15 antibody (Ab) for 3 days. **P* < .05 compared with 0 ng/mL GDF15.



MM cell lines. Compared with vehicle-treated cells, GDF15 significantly increased the colony formation of NCI-H929 cells in a dose-dependent manner (Figure 1A; *P* < .01). Moreover, this increase was maintained through secondary colony formation despite no further treatment with GDF15 (Figure 1A; *P* < .01). We similarly found that GDF increased the colony formation of RPMI 8226, U266, KMS12, and MM1.S MM cells as well (supplemental Figure 1A-B, available on the *Blood* Web site).

To determine the impact of GDF15 on tumor initiation in vivo, we injected NCI-H929 cells into NSG mice after 14 days of ex vivo treatment. GDF15 treatment significantly increased tumor initiation because all mice receiving GDF15-treated cells were symptomatic and had detectable serum human IgG κ light chain and bone marrow plasma cells by 6 months compared with 40% of animals injected with the same number of vehicle-treated cells (Figure 1B; *P* < .0004). We also carried out limiting dilution studies and found that GDF15 treatment significantly increased the frequency of TICs more than tenfold over control-treated cells (Table 1; *P* < .0041). To assess the effects of GDF15 on MM self-renewal, we carried out serial transplants and found that all secondary recipients receiving cells originally treated ex vivo with GDF15 engrafted with MM compared with 20% of mice

transplanted with bone marrow cells from mice initially injected with control cells (Figure 1C; *P* < .0002).

We also examined the effects of GDF15 primary MM clinical specimens and found similar effects on tumor colony formation for all of the samples we studied (Figure 1D). In addition, this increase in clonogenic growth was specific for GDF15 because it could be blocked using an anti-GDF15 neutralizing antibody. Collectively, these results demonstrate that GDF15 acts in a specific manner to enhance clonogenic MM growth and self-renewal both in vitro and in vivo.

AKT-SOX2 signaling is required for the effects of GDF15 on MM

To examine the cellular processes responsible for the effects of GDF15 on clonogenic growth, we examined the AKT signaling pathway because it is activated by GDF15 in both normal and malignant tissues.^{5,14-16} GDF15 induced AKT activation in NCI-H929 cells as evidenced by increased levels of phosphorylated AKT (Figure 2A), and the AKT inhibitor (Akti)-1/2 (124018) significantly blocked the ability of GDF15 to enhance colony formation (Figure 2B, first plating; 100 nM, *P* = .020, 1 μ M, *P* = .0066, second plating; 100 nM, *P* = .0089, 1 μ M, *P* = .0037). We also examined the transcription factor SOX2 because it can be activated by AKT and is involved in the pluripotency and self-renewal of ES cells and induced pluripotent stem cells.¹⁷⁻¹⁹ Moreover, SOX2 expression increases during the transition from asymptomatic monoclonal gammopathy of unknown significance (MGUS) to overt MM.²⁰ GDF15 increased the expression of SOX2 in NCI-H929 MM cells (Figure 2A), and this induction could be inhibited by treatment with Akti-1/2 (Figure 2C). Furthermore, decreased SOX2 expression using siRNA significantly limited the GDF15-induced enhancement of colony formation (Figure 2D). In

Table 1. TIC frequency of control and GDF15-treated MM cells

| | Tumor formation (cells) | | | | TIC frequency ⁻¹ (95% CI) | P value |
|---------------------|-------------------------|-----------------|-----------------|-----------------|--------------------------------------|---------|
| | 10 ⁶ | 10 ⁵ | 10 ⁴ | 10 ³ | | |
| Control cells | 5/5 | 5/5 | 0/3 | 0/3 | 36 000 (12 000-110 000) | .0041 |
| GDF15-treated cells | 5/5 | 5/5 | 3/3 | 1/3 | 2 000 (500-10 000) | |

95% CI, 95% confidence interval.

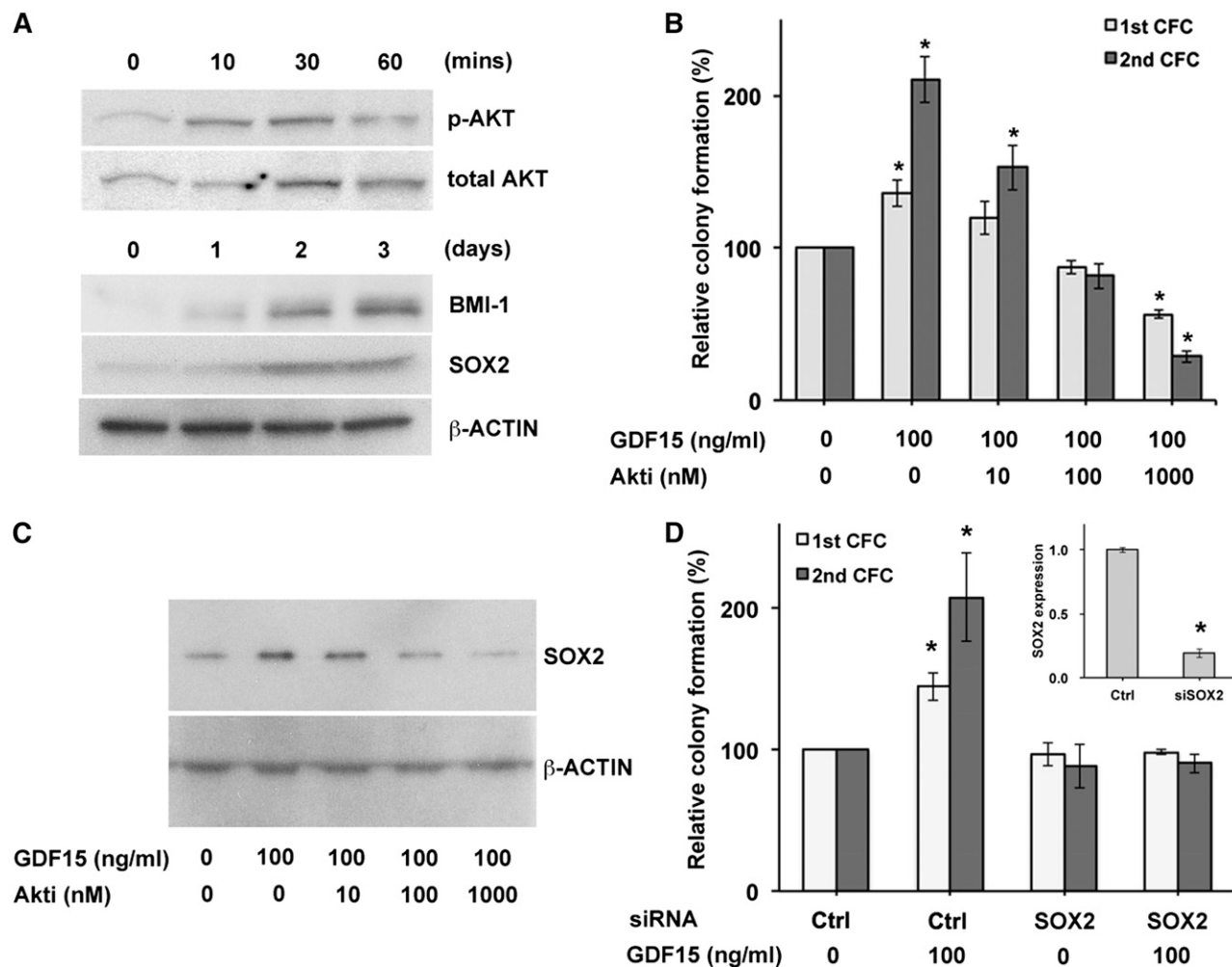


Figure 2. GDF15 activity is dependent on AKT and SOX2. (A) Western blot detection of phosphorylated AKT, BMI-1, and SOX2 expression by NCI-H929 cells after treatment with GDF (100 ng/mL). (B) Serial colony formation after treatment with GDF15 and the Akti 124018 for 3 days. * $P < .05$ compared with 0 ng/mL GDF15. (C) Western blot analysis of SOX2 after treatment with GDF15 and AKT inhibitor. (D) Serial colony formation of NCI-H929 cells after transfection of *SOX2* or control short interfering RNA (siRNA) and treatment with GDF15 for 3 days. * $P < .05$ compared with scramble control and 0 ng/mL GDF15. The inset shows *SOX2* expression by qRT-PCR 3 days after transfection.

addition, AKT and SOX2 inhibition similarly inhibited the clonogenic growth of RPMI 8226 cells (supplemental Figure 2). We also examined the effects of GDF15 on the expression of BMI-1, a member of the polycomb-repressive complex 1 required for the self-renewal of normal adult stem cells.^{21,22} In MM, BMI-1 is similarly required for clonogenic MM growth,²³ and we found that its expression was induced by GDF15 (Figure 2A). Taken together, these results indicate that GDF15 enhances clonogenic MM growth through key factors regulating self-renewal.

BMSC-derived GDF15 enhances clonogenic myeloma growth

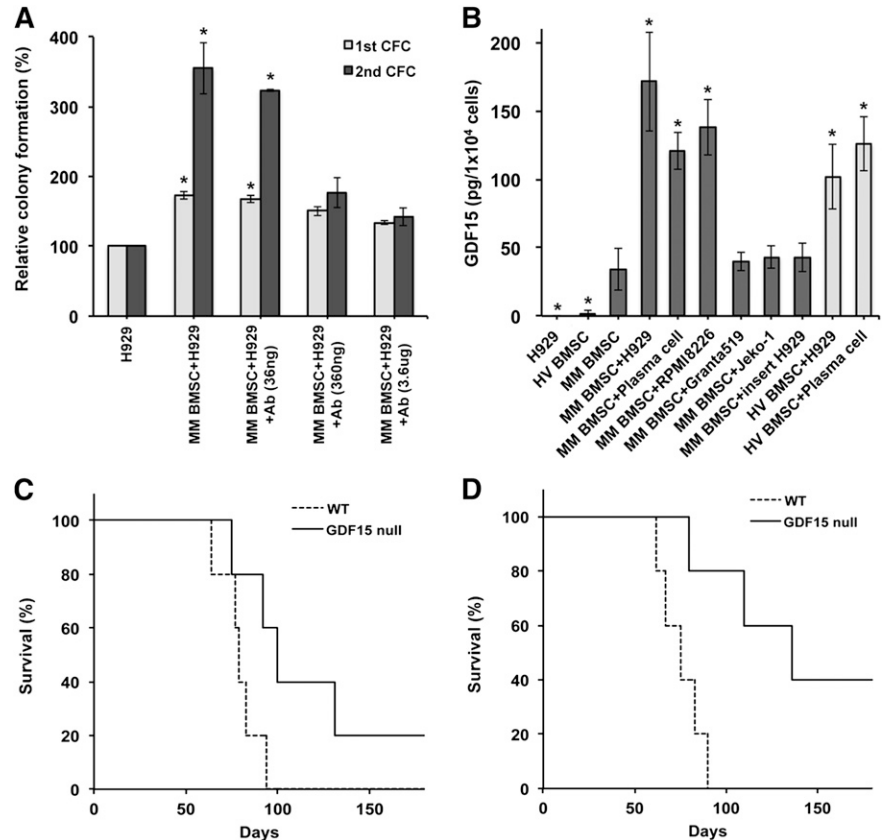
BMSCs promote drug resistance and tumor cell survival in MM, but their impact on tumor initiation and self-renewal is not well understood. We cocultured NCI-H929 cells with BMSCs isolated from MM patients and found that both primary and secondary colony formation were significantly increased compared with MM cells cultured alone (Figure 3A). To determine whether GDF15 mediates this interaction, we treated cocultures with a neutralizing anti-GDF15 antibody and found that it modestly affected initial colony formation compared with an isotypic control antibody. However, the anti-GDF15 antibody significantly inhibited secondary colony

formation at the 2 highest doses ($P = .004$), suggesting that BMSC-derived GDF15 primarily enhances the self-renewal of MM cells.

We also investigated the production of GDF15 by BMSCs and found that those isolated from MM patients secreted more GDF15 than BMSCs from healthy volunteers (HVs), as previously reported (Figure 3B),⁴ but these levels were significantly increased during cocultures with NCI-H929 cells ($P < .03$). BMSCs, rather than MM cells, were responsible for increased GDF15 production because they exclusively expressed *GDF15* by qRT-PCR after coculture (supplemental Figure 3). Moreover, direct cell-cell contact was required because cell culture inserts separating the 2 cell types prevented increased GDF15 production (Figure 3B). Interestingly, normal plasma cells isolated from HVs also induced GDF15, whereas cocultures with Granta 519 and Jeko-1 mantle cell lymphoma cells that phenotypically resemble B cells did not induce GDF15 production.

To determine the effects of BMSC-derived GDF15 on in vivo tumor-initiating potential, we examined the engraftment of immunocompetent *Gdf15*^{-/-} (null) or wild-type C57/BL6 mice with tumor cells from *Vk**Myc transgenic mice that develop plasma cell dyscrasias closely resembling human MM.^{24,25} The engraftment of

Figure 3. BMSC-derived GDF15 enhances clonogenic MM growth. (A) Serial colony formation by NCI-H929 cells cocultured with MM BMSC in the presence or absence of anti-GDF15 antibody (Ab) for 4 days. **P* < .05 compared with NCI-H929 cell only. (B) GDF15 secretion by primary BMSCs derived from HVs (light gray) and MM patients (dark gray) cultured alone or with NCI-H929 (H929), Granta519, Jeko-1, or normal CD138⁺ plasma cells. Cell culture inserts inhibited cell-cell contacts between NCI-H929 cells and MM BMSCs. **P* < .05 compared with MM BMSC only. (C) Survival of *Gdf15*^{null} or wild-type (WT) mice after injection of Vk*Myk MM cells (*n* = 5 for each group). (D) Secondary transplantation of BM derived from mice initially engrafted with Vk*Myk MM cells.



GDF15 null mice was modestly decreased compared with wild-type recipients (Figure 3C; 80% vs 100%) but a significantly prolonged median survival was observed (79 vs 96 days, *P* < .05). We also carried out secondary transplants by injecting equal numbers of tumor cells isolated from GDF15 null or wild-type primary recipients into wild-type secondary recipients. The engraftment of tumor cells from GDF15 null primary recipients was significantly impaired compared with those from wild-type mice (Figure 3D; 60% vs 100%) and was associated with a significant prolongation of survival (136 vs 75 days, *P* < .002). Therefore GDF15 produced by BMSC in response to contact with plasma cells enhances the clonogenic growth and self-renewal of MM cells both in vitro and in vivo.

GDF15 enhances the clonogenic growth and self-renewal of phenotypically defined MM TICs

In many diseases, self-renewal is enriched within subpopulations of TICs that are phenotypically distinct from bulk tumor cells. Several methods have been used to identify tumorigenic MM cells, but the precise TIC phenotype has remained unclear.^{11,12,26,27} Normal and MM plasma cells uniformly display high levels of CD138 (Syndecan-1), but human MM cell lines and clinical specimens contain relatively small subpopulations that lack or express low levels of this surface antigen.^{11,12,20,26} We previously found that CD138^{neg} MM cells give rise to CD138⁺ plasma cells and are enhanced for long-term tumor colony formation in vitro and engraftment in vivo.^{11,12} Similarly, increased ALDH expression may also identify highly tumorigenic and self-renewing MM cells similar to normal hematopoietic stem cells and TICs in breast and pancreatic carcinoma.^{12,28-30} To better characterize MM TICs, we compared the expression of CD138 and ALDH in NCI-H929 and

RPMI 8226 cells. In each cell line, CD138^{neg} and ALDH⁺ cells accounted for 1.43% to 1.77% and 2.17% to 2.53% of all cells, but these populations were not completely overlapping because 43% to 63% of the ALDH⁺ cells were also CD138^{neg} (supplemental Table 1). We examined the engraftment of NSG mice with NCI-H929 and RPMI 8226 cells isolated based on CD138 or ALDH expression and found that the TIC frequency of CD138^{neg} cells was significantly increased compared with CD138⁺ cells (Table 2 and supplemental Table 2; *P* < .000000093 and .0000069, respectively). Similarly, the TIC frequency of ALDH⁺ NCI-H929 and RPMI 8226 cells was significantly greater than ALDH^{neg} cells (*P* < .00000021 and .0000015, respectively) but similar to CD138^{neg} cells. Therefore human MM TICs can be enriched based on CD138 and ALDH expression.

To determine the impact of GDF15 on MM TICs, we isolated CD138^{neg} cells from the NCI-H929 cell line and found that GDF15 significantly increased total cell numbers compared with vehicle-control-treated cells (Figure 4A; *P* < .014). To determine whether GDF15 induces the expansion of CD138^{neg} NCI-H929 cells or their

Table 2. Tumor-initiating frequency of NCI-H929 cells in NSG mice

| | Tumor formation (cells) | | | | | TIC frequency ⁻¹ (95% CI) | P value |
|----------------------|-------------------------|-----------------|-----------------|-----------------|-----------------|--------------------------------------|------------|
| | 10 ⁶ | 10 ⁵ | 10 ⁴ | 10 ³ | 10 ² | | |
| CD138 ⁺ | 2/2 | 4/5 | 1/5 | 0/5 | 0/5 | 59 000 (23 000-160 000) | .000000093 |
| CD138 ^{neg} | — | — | 3/3 | 3/5 | 1/5 | 920 (320-2 600) | |
| ALDH ^{neg} | 3/3 | 2/3 | 1/3 | 0/3 | 0/3 | 68 000 (19 000-240 000) | .00000021 |
| ALDH ⁺ | — | — | 3/3 | 2/3 | 0/3 | 1 100 (270-4 300) | |

—, missing data; 95% CI, 95% confidence interval.

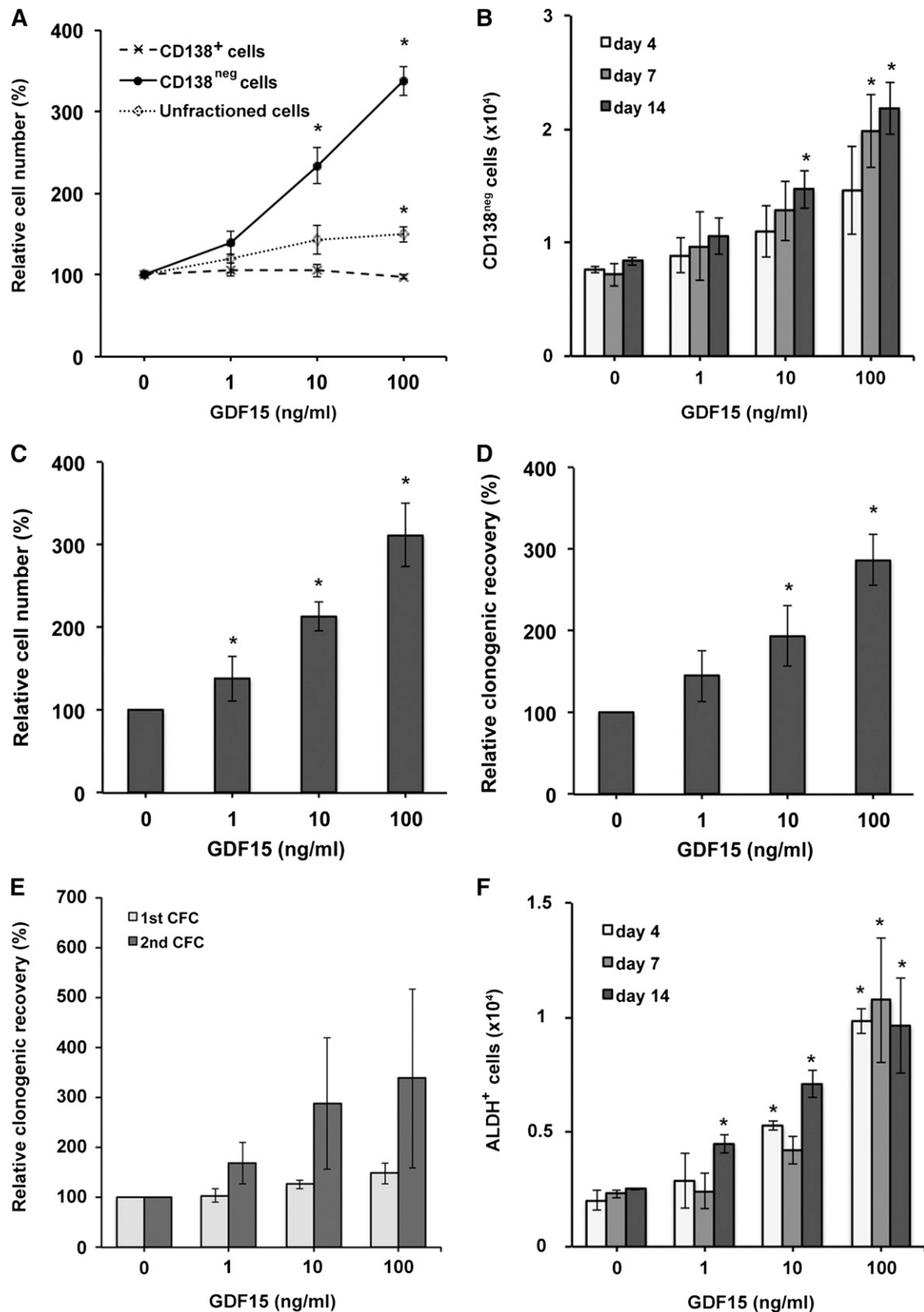


Figure 4. GDF15 expands MM TICs. (A) Relative cell numbers after treatment of unfractionated, CD138^{neg}, and CD138⁺ NCI-H929 cells with GDF15 for 7 days. (B) Absolute number of CD138^{neg} NCI-H929 cells quantified by flow cytometry after treatment with GDF15. (C) Relative expansion of prospectively isolated CD138^{neg} NCI-H929 cells by GDF15. (D) Relative clonogenic recovery of CD138^{neg} NCI-H929 cells after treatment with GDF15 for 7 days. (E) Relative clonogenic recovery of CD138⁺ NCI-H929 cells after treatment with GDF15 for 7 days. (F) Absolute number of ALDH⁺ NCI-H929 cells quantified by flow cytometry after treatment with GDF15. **P* < .05 compared with 0 ng/mL GDF15.

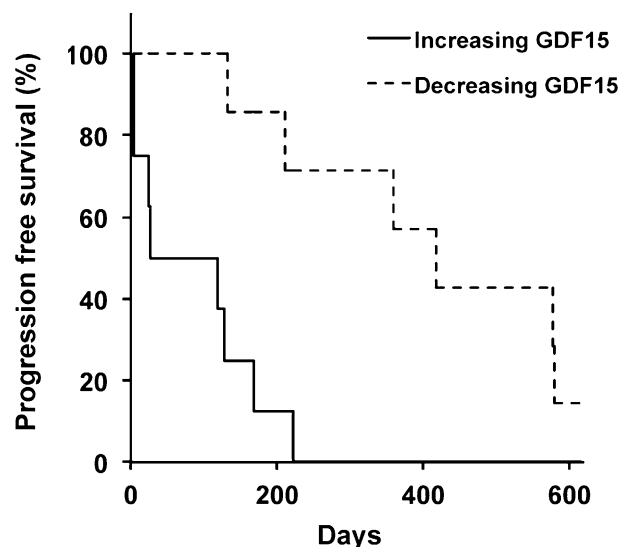


Figure 5. Serum GDF15 levels and clinical outcomes in MM. Progression-free survival of MM patients based on increasing or decreasing serum levels of GDF15.

differentiation into CD138⁺ plasma cells, we examined CD138 expression by flow cytometry and found that GDF15 treatment significantly increased the absolute number of CD138^{neg} cells (Figure 4B; $P < .002$). To more directly examine the impact of GDF15 on specific cell populations, we prospectively isolated CD138^{neg} cells and found that GDF15 significantly expanded both the number of cells (Figure 4C; $P = .03$) and subsequent clonogenic growth (Figure 4D; $P = .02$). In contrast, GDF15 did not significantly alter the total cell number (Figure 4A; 10 ng/mL, $P = .42$; 100 ng/mL, $P = .28$) or colony-forming frequency of isolated CD138⁺ NCI-H929 cells compared with controls (Figure 4E; $P > .06$), consistent with previous findings that GDF15 increases the survival, but not proliferation, of MM plasma cells.⁵ Similarly, treatment of GDF15 did not change the clonogenic potential of CD138⁺ MM cells derived from primary clinical specimens (data not shown). We similarly found that GDF15 expanded CD138^{neg}, but not CD138⁺, RPMI 8226 cells (supplemental Figure 4A-C). We also studied ALDH⁺ MM cells and found that GDF15 treatment significantly increased the absolute number of ALDH⁺ NCI-H929 (Figure 4F; $P < .03$) and RPMI 8226 cells (supplemental Figure 4; $P < .04$). Finally, we also examined the role of AKT and SOX2 on the expansion of MM cancer stem cells (CSCs) and found that the inhibition of AKT or knock-down of SOX2 inhibited the expansion of both CD138^{neg} and ALDH⁺ cells by GDF15 (supplemental Figure 5A-B). Therefore GDF15 primarily expands MM TICs that are CD138^{neg} or ALDH⁺ but has little effect on the proliferation or clonogenic growth of mature CD138⁺ tumor cells.

Serum GDF15 is associated with clonogenic tumor growth and clinical outcomes in MM

The clonogenic potential of TICs suggests they play an essential role in tumor regrowth; therefore changes in the frequency of TICs over time may correlate with clinical outcomes including disease relapse. We previously found that changes in MM colony formation after treatment with high-dose cyclophosphamide and rituximab were associated with progression-free survival.³¹ We quantified serum GDF15 levels in this cohort of patients and found that changes in GDF15 levels correlated with the differences in MM colony formation over the course of treatment in 12 of the 15 patients

studied (80%; Figure 5, supplemental Figure 6, and supplemental Table 3; $P = .026$). We also examined the clinical relevance of changes in GDF15 levels and detected decreasing GDF15 levels in 7 of 15 (47%) patients over the course of treatment that were associated with a significantly longer progression-free survival compared with the remaining 8 patients with increasing GDF15 levels (Table 3; $P = .0088$). Therefore GDF15 levels may serve as a surrogate for the frequency of TICs, as well as predict disease relapse and progression in MM.

Discussion

CSCs have been functionally defined by the capacity to produce differentiated cells that recapitulate the original tumor in the ectopic setting and self-renewal to maintain long-term tumor-initiating potential.³² These properties suggest that the inhibition of CSCs may improve clinical outcomes, and several intrinsic pathways regulating their self-renewal have been identified.³³ Less is known about the extrinsic factors that affect these cells, but we found that GDF15 aberrantly produced by BMSCs within the TME enhances serial tumor cell colony formation as well as primary and secondary MM engraftment in immunodeficient and immunocompetent mice. Therefore GDF15 may play an important role in the pathogenesis of MM through its effects on clonogenic growth and self-renewal.

As in most cancers, MM cells display both phenotypic and functional heterogeneity. The vast majority of tumor cells are plasma cells characterized by high surface expression of CD138, but several groups have identified relatively rare subpopulations of MM cells that lack or display low levels of this antigen.^{11,12,20,34-37} Functional heterogeneity also exists in MM, and only a fraction of cancer cells are capable of tumor initiation both in vitro and in vivo.^{38,39} MM cell phenotypes may be linked to their functional properties because CD138^{neg} cells isolated from primarily clinical specimens exhibit enhanced clonogenic growth, give rise to mature CD138⁺ plasma cells, and maintain tumorigenic potential during serial transplantation in immunodeficient NOD/severe combined immunodeficiency (SCID) or NSG mice.^{11,12} The ability to undergo self-renewal and give rise to cells that histologically recapitulate the original tumor suggests that MM TICs are CD138^{neg}, but tumor cells expressing CD138 or high levels of CD38, another marker of mature plasma cells, engraft human fetal bone fragments implanted in SCID (SCID-hu) mice.⁴⁰⁻⁴² Moreover, in the 5T33 murine model of MM, both CD138^{neg} and CD138⁺ cells are capable of clonogenic growth,⁴³ whereas we have found that the engraftment potential of serially passaged tumor cells in the Vk*Myb model is limited to CD138⁺ cells (G.M., unpublished data). Putative human MM TICs have also been identified using the side population and Aldefluor assays, and these cell populations may or may not express CD138.^{12,26} Therefore the functional properties of MM cells may not be confined to a specific phenotypic population but may differ depending on disease stage or specific genetic mutations as described in chronic myeloid leukemia.⁴⁴ We found that GDF15 preferentially

Table 3. Concordance between changes in serum GDF15 levels and MM-CFU in MM patients

| | Increasing GDF15 | Decreasing GDF15 | Total |
|-------------------|------------------|------------------|-------|
| Increasing MM-CFU | 8 | 3 | 11 |
| Decreasing MM-CFU | 0 | 4 | 4 |
| Total | 8 | 7 | 15 |

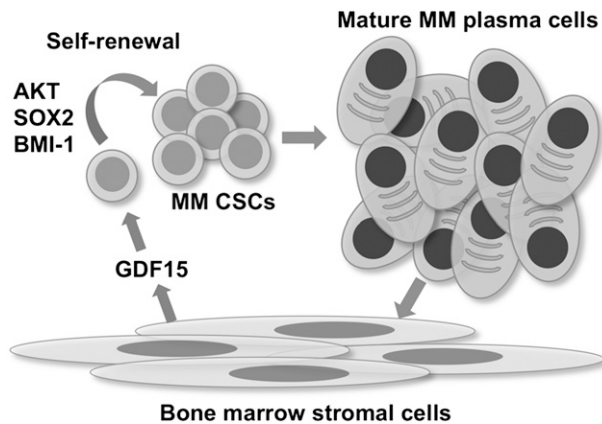


Figure 6. Interactions between MM plasma cells, TICs, and BMSCs. The interaction between BMSCs and mature MM plasma cells induces GDF15 secretion and enhances the self-renewal of MM CSCs by AKT, SOX2, and BMI-1. Subsequent production of mature tumor cells further promotes this process.

induced the proliferation and expansion of CD138^{neg} MM cells, but GDF15 has been found to induce the expression of matrix metalloproteinase 9, which can cleave CD138 from the cell surface.^{45,46} Therefore it is possible that the increased numbers of CD138^{neg} cells after GDF15 treatment reflects the loss of CD138 from the cell surface, but GDF15 also increased the number of ALDH⁺ TICs. Furthermore, GDF15 significantly inhibited serial colony formation in vitro and tumor engraftment in vivo using unfractionated cells, demonstrating that it affects these functional properties in MM regardless of the phenotype of the cell affected. We found that increased tumorigenic potential in response to GDF15 was increased to a greater extent in vivo than in vitro. Therefore, it is possible that GDF15 affects other cellular properties in addition to clonogenic growth, such as homing to the bone marrow or the expansion of tumor cells in vivo.

In contrast to epithelial cancers, in which tumor cells produce GDF15,⁴⁷ MM is characterized by the aberrant secretion of GDF15 by BMSCs.^{4,5} Similar to previous findings,⁴ we found that BMSCs isolated from MM patients secreted higher baseline levels of GDF15 compared with normal donors. GDF15 secretion subsequently decreased with continued passaging, but could be re-induced by co-incubation with CD138⁺ plasma cells. Therefore the initial GDF15 secretion from MM patients is likely a result of interactions between BMSCs and MM plasma cell interactions in situ. We found that BMSCs increased the clonogenic potential of MM TICs, which was dependent on GDF15 because a neutralizing antibody significantly suppressed colony formation. Therefore the TME plays a significant role in regulating MM TICs. Based on these results, we propose a model in which plasma cells induce GDF15 secretion from BMSCs within the TME (Figure 6). GDF15 then activates signaling pathways that enhance the self-renewal and clonogenic potential of MM CSCs. These interactions generate a feed-forward loop that maintains the CSC pool and eventually generates additional MM plasma cells. A previous study found that BMSCs can induce CD138⁺ plasma cells to undergo de-differentiation to CD138^{neg} cells.⁴⁸ We did not observe changes in the expression of CD138 after the incubation of isolated CD138⁺ cells with GDF15, but it is possible that this effect is mediated by factors other than GDF15 or requires longer treatment periods. We found that a blocking antibody against GDF15 partially inhibited the interaction between MM cells

and BMSCs, but it is likely that other cytokines are also involved. For example, it is well recognized that interleukin-6 is secreted by BMSCs, and we previously demonstrated that IL-6 can enhance MM clonogenic growth in vitro.⁴⁹ Because GDF15 is elevated in a wide variety of inflammatory states that can lead to MGUS,⁵⁰ it is also possible that these interactions play a role in the generation of MGUS and the subsequent transformation to MM.

Our findings may also have significant clinical implications. We found that relative changes in serum GDF15 levels over the course of treatment are significantly associated with progression-free survival. These data support previous findings that GDF15 levels in MM patients are associated with clinical outcomes, although the measurement of GDF15 was limited to a single time point in this study.⁵ We also found that serum GDF15 levels acutely increased after treatment with high-dose cyclophosphamide and then subsequently returned to lower levels. Because GDF15 is a p53 target gene,^{51,52} the acute elevation of serum levels may likely reflect cellular responses to cytotoxic treatment. We also found that GDF15 levels over the course of treatment paralleled changes in the frequency of MM colony formation. Therefore GDF15 may serve as a novel biomarker indicative of the frequency of MM TICs, although larger clinical studies are needed to confirm these results. Because the disruption of growth factor and cytokine signaling by neutralizing antibodies or small molecules is used in the treatment of many tumor types, targeting GDF15 activity may inhibit both the clonogenic potential of MM CSCs as well as the survival of mature plasma cells.

Acknowledgments

We thank Suzanne Sebal and Se-Jin Lee for providing GDF15^{null} mice. W.M. is a Scholar in Clinical Research of the Leukemia and Lymphoma Society.

This study was supported by grants from the National Institutes of Health, National Cancer Institute (R01CA127574 and R21CA155733 [W.M.]) and National Heart, Lung, and Blood Institute (T32HL7525 [T.T.]).

Authorship

Contribution: T.T., Y.L., Q.W., and W.M. carried out the in vitro and in vivo studies; N.G., I.B., and C.A.H. were involved in specimen and clinical data collection; Q.W., M.C., P.L.B., G.M., and R.W.J. were involved in establishment of in vivo models of MM; and T.T. and W.M. wrote the manuscript with the assistance and final approval of all authors.

Conflict-of-interest disclosure: The authors declare no competing financial interests.

The current affiliation for T.T. is Center for Cancer and Immunology Research, Children's National Hospital, Washington, DC.

Correspondence: William Matsui, The Sidney Kimmel Comprehensive Cancer Center, Johns Hopkins University School of Medicine, The Bunting Blaustein Cancer Research Building, Room 245, 1650 Orleans St, Baltimore, MD 21287; e-mail: matsui@jhmi.edu.

References

- Palumbo A, Anderson K. Multiple myeloma. *N Engl J Med*. 2011;364(11):1046-1060.
- Scadden DT. The stem-cell niche as an entity of action. *Nature*. 2006;441(7097):1075-1079.
- Wang LD, Wagers AJ. Dynamic niches in the origination and differentiation of haematopoietic stem cells. *Nat Rev Mol Cell Biol*. 2011;12(10):643-655.
- Corre J, Mahtouk K, Attal M, et al. Bone marrow mesenchymal stem cells are abnormal in multiple myeloma. *Leukemia*. 2007;21(5):1079-1088.
- Corre J, Labat E, Espagnolle N, et al. Bioactivity and prognostic significance of growth differentiation factor GDF15 secreted by bone marrow mesenchymal stem cells in multiple myeloma. *Cancer Res*. 2012;72(6):1395-1406.
- Bootcov MR, Bauskin AR, Valenzuela SM, et al. MIC-1, a novel macrophage inhibitory cytokine, is a divergent member of the TGF-beta superfamily. *Proc Natl Acad Sci USA*. 1997;94(21):11514-11519.
- Staff AC, Trovik J, Eriksson AG, et al. Elevated plasma growth differentiation factor-15 correlates with lymph node metastases and poor survival in endometrial cancer. *Clin Cancer Res*. 2011;17(14):4825-4833.
- Wallin U, Glimelius B, Jirstrom K, et al. Growth differentiation factor 15: a prognostic marker for recurrence in colorectal cancer. *Br J Cancer*. 2011;104(10):1619-1627.
- Brown DA, Lindmark F, Stattin P, et al. Macrophage inhibitory cytokine 1: a new prognostic marker in prostate cancer. *Clin Cancer Res*. 2009;15(21):6658-6664.
- Koopmann J, Rosenzweig CNW, Zhang Z, et al. Serum markers in patients with resectable pancreatic adenocarcinoma: macrophage inhibitory cytokine 1 versus CA19-9. *Clin Cancer Res*. 2006;12(2):442-446.
- Matsui W, Huff CA, Wang Q, et al. Characterization of clonogenic multiple myeloma cells. *Blood*. 2004;103(6):2332-2336.
- Matsui W, Wang Q, Barber JP, et al. Clonogenic multiple myeloma progenitors, stem cell properties, and drug resistance. *Cancer Res*. 2008;68(1):190-197.
- Hu Y, Smyth GK. ELDA: extreme limiting dilution analysis for comparing depleted and enriched populations in stem cell and other assays. *J Immunol Methods*. 2009;347(1-2):70-78.
- Kempf T, Eden M, Strelau J, et al. The transforming growth factor-beta superfamily member growth-differentiation factor-15 protects the heart from ischemia/reperfusion injury. *Circ Res*. 2006;98(3):351-360.
- Naka K, Hoshii T, Muraguchi T, et al. TGF-beta-FOXO signalling maintains leukaemia-initiating cells in chronic myeloid leukaemia. *Nature*. 2010;463(7281):676-680.
- Eyler CE, Foo W-C, LaFiura KM, McLendon RE, Hjelmeland AB, Rich JN. Brain cancer stem cells display preferential sensitivity to Akt inhibition. *Stem Cells*. 2008;26(12):3027-3036.
- Peltier J, Conway A, Keung AJ, Schaffer DV. Akt increases sox2 expression in adult hippocampal neural progenitor cells, but increased sox2 does not promote proliferation. *Stem Cells Dev*. 2011;20(7):1153-1161.
- Avilion AA, Nicolis SK, Pevny LH, Perez L, Vivian N, Lovell-Badge R. Multipotent cell lineages in early mouse development depend on SOX2 function. *Genes Dev*. 2003;17(1):126-140.
- Takahashi K, Yamanaka S. Induction of pluripotent stem cells from mouse embryonic and adult fibroblast cultures by defined factors. *Cell*. 2006;126(4):663-676.
- Spisek R, Kukreja A, Chen L-C, et al. Frequent and specific immunity to the embryonal stem cell-associated antigen SOX2 in patients with monoclonal gammopathy. *J Exp Med*. 2007;204(4):831-840.
- Park IK, Qian D, Kiel M, et al. Bmi-1 is required for maintenance of adult self-renewing haematopoietic stem cells. *Nature*. 2003;423(6937):302-305.
- Molofsky AV, Pardoll R, Iwashita T, Park I-K, Clarke MF, Morrison SJ. Bmi-1 dependence distinguishes neural stem cell self-renewal from progenitor proliferation. *Nature*. 2003;425(6961):962-967.
- Jagani Z, Wiederschain D, Loo A, et al. The Polycomb group protein Bmi-1 is essential for the growth of multiple myeloma cells. *Cancer Res*. 2010;70(13):5528-5538.
- Hsiao EC, Koniari LG, Zimmers-Koniari T, Sebald SM, Huynh TV, Lee SJ. Characterization of growth-differentiation factor 15, a transforming growth factor beta superfamily member induced following liver injury. *Mol Cell Biol*. 2000;20(10):3742-3751.
- Chesi M, Robbiani DF, Sebag M, et al. AID-dependent activation of a MYC transgene induces multiple myeloma in a conditional mouse model of post-germinal center malignancies. *Cancer Cell*. 2008;13(2):167-180.
- Jakubikova J, Adamia S, Kost-Alimova M, et al. Lenalidomide targets clonogenic side population in multiple myeloma: pathophysiologic and clinical implications. *Blood*. 2011;117(17):4409-4419.
- Hawley TS, Riz I, Yang W, et al. Identification of an ABCB1 (P-glycoprotein)-positive carfilzomib-resistant myeloma subpopulation by the pluripotent stem cell fluorescent dye CDy1. *Am J Hematol*. 2013;88(4):265-272.
- Hess DA, Meyerrose TE, Wirthlin L, et al. Functional characterization of highly purified human hematopoietic repopulating cells isolated according to aldehyde dehydrogenase activity. *Blood*. 2004;104(6):1648-1655.
- Ginestier C, Hur MH, Charafe-Jauffret E, et al. ALDH1 is a marker of normal and malignant human mammary stem cells and a predictor of poor clinical outcome. *Cell Stem Cell*. 2007;1(5):555-567.
- Rasheed ZA, Yang J, Wang Q, et al. Prognostic significance of tumorigenic cells with mesenchymal features in pancreatic adenocarcinoma. *J Natl Cancer Inst*. 2010;102(5):340-351.
- Huff CA, Wang Q, Rogers K, et al. Correlation of clonogenic cancer stem cell (CSC) growth with clinical outcomes in multiple myeloma (MM) patients undergoing treatment with high dose cyclophosphamide (Cy) and rituximab. *AACR Meeting Abstracts*. 2008;2008:LB-87.
- Clarke MF, Dick JE, Dirks PB, et al. Cancer stem cells—perspectives on current status and future directions: AACR Workshop on cancer stem cells. *Cancer Res*. 2006;66(19):9339-9344.
- Lobo NA, Shimono Y, Qian D, Clarke MF. The biology of cancer stem cells. *Annu Rev Cell Dev Biol*. 2007;23:675-699.
- Billadeau D, Ahmann G, Greipp P, Van Ness B. The bone marrow of multiple myeloma patients contains B cell populations at different stages of differentiation that are clonally related to the malignant plasma cell. *J Exp Med*. 1993;178(3):1023-1031.
- Kirshner J, Thulien KJ, Martin LD, et al. A unique three-dimensional model for evaluating the impact of therapy on multiple myeloma. *Blood*. 2008;112(7):2935-2945.
- Chaidos A, Barnes CP, Cowan G, et al. Clinical drug resistance linked to interconvertible phenotypic and functional states of tumor-propagating cells in multiple myeloma. *Blood*. 2013;121(2):318-328.
- Boucher K, Parquet N, Widen R, et al. Stemness of B-cell progenitors in multiple myeloma bone marrow. *Clin Cancer Res*. 2012;18(22):6155-6168.
- Bergsagel DE, Valeriotte FA. Growth characteristics of a mouse plasma cell tumor. *Cancer Res*. 1968;28(11):2187-2196.
- Hamburger AW, Salmon SE. Primary bioassay of human tumor stem cells. *Science*. 1977;197(4302):461-463.
- Yaccoby S, Barlogie B, Epstein J. Primary myeloma cells growing in SCID-hu mice: a model for studying the biology and treatment of myeloma and its manifestations. *Blood*. 1998;92(8):2908-2913.
- Yaccoby S, Epstein J. The proliferative potential of myeloma plasma cells manifest in the SCID-hu host. *Blood*. 1999;94(10):3576-3582.
- Kim D, Park CY, Medeiros BC, Weissman IL. CD19-CD45 low/- CD38 high/CD138+ plasma cells enrich for human tumorigenic myeloma cells. *Leukemia*. 2012;26(12):2530-2537.
- Van Valckenborgh E, Matsui W, Agarwal P, et al. Tumor-initiating capacity of CD138- and CD138+ tumor cells in the 5T33 multiple myeloma model. *Leukemia*. 2012;26(6):1436-1439.
- Jamieson CHM, Ailles LE, Dylla SJ, et al. Granulocyte-macrophage progenitors as candidate leukemic stem cells in blast-crisis CML. *N Engl J Med*. 2004;351(7):657-667.
- Griner SE, Joshi JP, Nahta R. Growth differentiation factor 15 stimulates rapamycin-sensitive ovarian cancer cell growth and invasion. *Biochem Pharmacol*. 2013;85(1):46-58.
- Ramani VC, Purushothaman A, Stewart MD, et al. The heparanase/syndecan-1 axis in cancer: mechanisms and therapies. *FEBS J*. 2013;280(10):2294-2306.
- Bauskin AR, Brown DA, Kuffner T, et al. Role of macrophage inhibitory cytokine-1 in tumorigenesis and diagnosis of cancer. *Cancer Res*. 2006;66(10):4983-4986.
- Yaccoby S. The phenotypic plasticity of myeloma plasma cells as expressed by dedifferentiation into an immature, resilient, and apoptosis-resistant phenotype. *Clin Cancer Res*. 2005;11(21):7599-7606.
- Matsui W, Huff CA, Vala M, Barber J, Smith BD, Jones RJ. Anti-tumour activity of interferon-alpha in multiple myeloma: role of interleukin 6 and tumor cell differentiation. *Br J Haematol*. 2003;121(2):251-258.
- Mimeault M, Batra SK. Divergent molecular mechanisms underlying the pleiotropic functions of macrophage inhibitory cytokine-1 in cancer. *J Cell Physiol*. 2010;224(3):626-635.
- Yang H, Filipovic Z, Brown D, Breit SN, Vassilev LT. Macrophage inhibitory cytokine-1: a novel biomarker for p53 pathway activation. *Mol Cancer Ther*. 2003;2(10):1023-1029.
- Osada M, Park HL, Nagakawa Y, et al. Differential recognition of response elements determines target gene specificity for p53 and p63. *Mol Cell Biol*. 2005;25(14):6077-6089.