Glycogen Synthase Kinase-3β Is a Prosurvival Signal for the Maintenance of Human Mast Cell Homeostasis

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Homeostasis of mature tissue-resident mast cells is dependent on the relative activation of pro- and antiapoptotic regulators. In this study, we investigated the role of glycogen synthase kinase 3β (GSK3β) in the survival of neoplastic and nonneoplastic human mast cells. GSK3β was observed to be phosphorylated at the Y216 activating residue under resting conditions in both the neoplastic HMC1.2 cell line and in peripheral blood-derived primary human mast cells (HuMCs), suggesting constitutive activation of GSK3β in these cells. Lentiviral-transduced short hairpin RNA knockdown of GSK3β in both the HMC1.2 cells and HuMCs resulted in a significant reduction in cell survival as determined with the MTT assay. The decrease in stem cell factor (SCF)-mediated survival in the GSK3β knockdown HuMCs was reflected by enhancement of SCF withdrawal-induced apoptosis, as determined by Annexin V staining and caspase cleavage, and this was associated with a pronounced reduction in SCF-mediated phosphorylation of Src homology 2 domain-containing phosphatase 2 and ERK1/2 and reduced expression of the antiapoptotic proteins Bcl-xl and Bcl-2. These data show that GSK3β is an essential antiapoptotic factor in both neoplastic and nontransformed primary human mast cells through the regulation of SCF-mediated Src homology 2 domain-containing phosphatase 2 and ERK activation. Our data suggest that targeting of GSK3β with small m.w. inhibitors such as CHIR 99021 may thus provide a mechanism for limiting mast cell survival and subsequently decreasing the intensity of the allergic inflammatory response. The Journal of Immunology, 2011, 187: 000–000.

Chronic allergic inflammation is characterized by increased mast cell infiltration and population in the affected tissues. Mast cell burden in these tissues is dependent not only on migration of mast cells/mast cell precursors into these sites but also on prosurvival and antiapoptotic signaling pathways. Processes, including the migration and survival of tissue mast cells, are tightly regulated by stem cell factor (SCF), the ligand for the growth factor receptor KIT (1–3). KIT, which is a member of the growth factor receptors with inherent tyrosine kinase family activity (4, 5), undergoes dimerization and autophosphorylation following SCF-induced ligation. Constitutive activation of KIT activity, through a point mutation (D816V) in the KIT catalytic domain is considered a hallmark of the myeloproliferative disorder mastocytosis (6–8). This disease is characterized by dysregulated growth of mast cells and elevated mast cell numbers in associated skin lesions and tissues such as bone marrow (7). In addition to the documentation of more mast cells harboring the D816V mutation within tissues of mastocytosis patients, the rapidly dividing HMC1.2 human mast cell line also expresses this mutation (9).

Autophosphorylation of KIT at specific tyrosine residues within the cytosolic domain induces recruitment of Src homology 2 domain-containing signaling proteins, which leads to activation of these and other molecules required for transducing KIT-mediated responses, including mast cell growth and survival (10, 11). We have recently provided evidence to support a role for PI3K-regulated mammalian target of rapamycin in cell survival in both human and mouse mast cells (12). Furthermore, the transcription factor FOXO3a has been shown to regulate SCF-mediated survival in mouse mast cells by repressing the expression of proapoptotic proteins via a PI3K-dependent pathway (13). It is clear, however, that other antiapoptotic pathways exist that may contribute to mast cell homeostasis. For example, in the LAD2 human mast cell line, cell survival can be maintained in the absence of SCF despite the lack of an activating mutation in KIT (14). Furthermore, mouse mast cells can proliferate and survive in the presence of IL-3 but in the absence of SCF.

We have previously determined that glycogen synthase kinase 3β (GSK3β), a ubiquitously expressed, conserved serine/threonine protein kinase, contributes to various FcRI/KIT-mediated mast cell functions including cytokine production and KIT-mediated chemotaxis (15). As GSK3β has also been implicated in tumor growth (16), we thus investigated whether, as for PI3K and mammalian target of rapamycin, GSK3β also contributes to the homeostasis of normal and neoplastic mast cells. In this study, we report that gene knockdown of GSK3β by short hairpin RNA (shRNA) and blocking of its activity by the small molecule GSK3β inhibitor CHIR 99021 substantially reduces the ability of the HMC1.2 cell line to expand and survive in culture. Furthermore, we show that the ability of SCF to maintain the survival of terminally differentiated primary cultures of human mast cells (HuMCs) is markedly reduced as a consequence of induction of apoptosis following GSK3β knockdown. Thus, GSK3β appears to represent an important mast cell prosurvival signal. Targeting GSK3β accordingly may provide a mechanism for modulating...
mast cell survival and apoptotic pathways in neoplastic mast cells and potentially mast cells participating in the allergic inflammatory response.

Materials and Methods

Mast cell Culture

HuMCs were developed from CD34+ peripheral blood progenitor cells in StemPro-34 culture medium containing StemPro supplement (Invitrogen Life Technologies), l-glutamine (2 mM), penicillin (100 U/ml), streptomycin (100 μg/ml), recombinant human IL-3 (30 ng/ml; first week only), recombinant human IL-6 (100 ng/ml), and recombinant human SCF (100 ng/ml) (PeproTech, Rocky Hill, NJ). Experiments were conducted 7–9 wk after the initiation of HuMC cultures. The CD34+ cells were obtained from normal volunteers under a protocol (98-L-0027; principal investigator, Dr. A. Kirshenbaum) approved by the National Institute of Allergy and Infectious Diseases Institutional Review Board and with appropriate informed consents. The growth factor-independent human mast cell line HMC1.2 was cultured in IMDM medium supplemented with FBS (10%), l-glutamine (2 mM), penicillin (100 U/ml), and streptomycin (100 μg/ml).

Lentivirus shRNA transfection of cell lines and transduction of HuMCs

The following GS3Kβ-targeted shRNA were purchased from Sigma-Aldrich (St. Louis, MO): 5′-CCGGGTGGTAGGCGGATCCACAAACTGATCCAGAATGTTGGATAGAACCTCAGGTTCCTACTCTCACTGATCCACACTTCTTTTT3′ (TRCN0000010352) (construct A); 5′-CCGGGGAAGCTTGTGAAAGCACCATGAGTTGGGTTTACGCAGGTCTATAATGCTGTAGTCTTTTT3′ (TRCN0000010352) (construct B); and 5′-CCGGCACAAGATGAGAAAGAGACACCACACTGAGTTGGTCCTCATCTCTTTGTTTTTT3′ (SHC002) (control nontarget vector construct). Packaging vector (MissionLentiviral packaging mix; Sigma-Aldrich), pLKO1 transfer vectors with shRNA (Sigma-Aldrich), or scrambled control shRNA were cotransfected into 293T packaging cells. The transfected 293T cells were grown in DMEM containing FBS (10%), l-glutamine (2 mM), penicillin (100 U/ml), and streptomycin (100 μg/ml) with FuGENE6 transfection reagent (Roche, Indianapolis, IN) as described (15). The transfected 293T cells were grown in DMEM containing FBS (10%), l-glutamine (2 mM), penicillin (100 U/ml), and streptomycin (100 μg/ml). Following 16–18 h of transfection, medium was removed and replaced with fresh DMEM. After 62–65 h of transfection, virus was collected by centrifugation, and the virus pellet was resuspended in 3 ml prewarmed StemPro medium (Invitrogen Life Technologies). Transduction of HuMCs was conducted by transferring the 3 ml resuspended virus to a T75 culture flask containing 3 to 4 × 10^6 HuMCs in 15 ml supplemented StemPro cell culture medium (Invitrogen Life Technologies) or 2 to 3 × 10^6 HMC1.2 cells in 10 ml IMDM medium. Two days postinfection, the medium was changed to virus-free medium, and antibiotic selection was initiated (0.2 μg/ml puromycin [Sigma-Aldrich] for HuMCs and 2 μg/ml puromycin for HMC1.2 cells). Experiments were conducted on days 7 to 8 posttransduction. We have previously demonstrated that this construct does not affect the expression of other signaling proteins including phospholipase Cγ (PLCγ)1, Lyn, and NF-kB in HuMCs (15).

Cell activation

HuMCs were starved in cytokine-free media overnight following rinsing with HEPES buffer containing 0.04% BSA and activated by addition of SCF. Cells were pretreated with the Src family tyrosine kinase inhibitor PP2 (3 μM), or the PI3K inhibitor wortmannin (100 nM; Calbiochem) 20 min prior to the activation with SCF. Cell activation was measured by cellular Annexin V binding assay (Annexin V, FITC apoptosis detection kit I; BD Biosciences, San Jose, CA). In brief, cells were stained with Annexin V and propidium iodide (PI) according to the manufacturer’s instructions and then analyzed using FACScan flow cytometry (BD Biosciences). All PI-positive cells were considered dead. PI-negative and Annexin V-positive cells were considered early apoptotic cells, and the remaining double-negative cells were considered viable.

Apoptosis assay

Apoptosis was evaluated by flow cytometry using cellular Annexin V binding assay (Annexin V, FITC apoptosis detection kit I; BD Biosciences, San Jose, CA). In brief, cells were stained with Annexin V and propidium iodide (PI) according to the manufacturer’s instructions and then analyzed using FACScan flow cytometry (BD Biosciences). All PI-positive cells were considered dead. PI-negative and Annexin V-positive cells were considered early apoptotic cells, and the remaining double-negative cells were considered viable.

Proliferation assay: BrdU incorporation

Cell proliferation was determined by the BrdU (a thymidine analog) assay that measures its incorporation into DNA. HMC1.2 cells, transfected with shContr or shGS3Kβ, were cultured overnight in IMEM without FBS and resuspended in IMEM containing 10% FBS. Cells were then incubated for 24 h at 1.5 × 10^6 cells/100 μl in 96-well plates. Incorporation of BrdU into the shContr- or shGS3Kβ-transduced HMC1.2 cells was assessed using a BrdU cell proliferation assay kit (Calbiochem, San Diego, CA) as described (19).

Statistical analysis

Data are presented as the mean ± SEM. Significant difference between two groups was analyzed by unpaired Student t test. A p value <0.05 was considered statistically significant.

Results

Knockdown of GS3Kβ decreases survival of the HMC1.2 human leukemia mast cell line

To explore the potential role of GS3Kβ as a mast cell survival/antiapoptotic signal, we adopted a lentivirus-mediated knockdown approach (15), as GS3Kβ deficiency in mice results in an embryonic lethal phenotype (20). We initially examined the consequences of GS3Kβ knockdown in the SCF-independent HMC1.2 mast cell line. As shown in Fig. 1A, GS3Kβ-targeted shRNA substantially decreased the expression of GS3Kβ in the HMC1.2 cells without reducing the expression of the internal controls KIT and Syk. GS3Kβ activity is regulated by the phosphorylation status of Y216 and S9. The Y216 residue is reported to be constitutively phosphorylated in resting cells, thus maintaining GS3Kβ in an active state, whereas the phosphorylation of the S9 site usually follows growth factor stimulation and activation of the PI3K–Akt signaling pathway (21–23). As shown in Fig. 1A, Y216 and S9 are constitutively phosphorylated in the HMC1.2 cells, and, as expected, the phosphorylation of these residues is substantially reduced in cells treated with the GS3Kβ-targeted shRNA.

Having successfully knocked down GS3Kβ in the HMC1.2 cells, we next determined the consequences of GS3Kβ knock-
down on the proliferation and/or survival of HMC1.2 cells. As shown in Fig. 1B, GSK3\(\beta\) knockdown significantly decreased the number of viable HMC1.2 cells as assessed by trypan blue dye exclusion. Cell survival was assessed by means of an MTT assay in which the starting cell number was 1 \times 10^5 cells/well (C), and proliferation was assessed by BrdU incorporation (D). In all cases, \(n = 3–5\). *\(p < 0.05\), **\(p < 0.01\) for comparison with shContr-transduced HMC1.2.

**Expression and phosphorylation of GSK3\(\beta\) in primary human mast cells**

We first confirmed the ability of GSK3\(\beta\)-targeted shRNA to downregulate GSK3\(\beta\) expression and phosphorylation in SCF-challenged and quiescent HuMCs. Cells, starved of cytokines overnight and treated with control or GSK3\(\beta\)-targeted shRNA, were either unchallenged or challenged with SCF for 2 min and cell lysates assessed for expression and phosphorylation of GSK3\(\beta\). Despite overnight starvation of SCF, and as was observed in the HMC1.2 cells, GSK3\(\beta\) was found to be constitutively phosphorylated at position Y216 in resting HuMCs and no consistent increase in the phosphorylation of this residue was observed in cells rechallenged with SCF (Fig. 2A). Regardless, the expression of GSK3\(\beta\) and, as a consequence, GSK3\(\beta\) phosphorylated at Y216 was markedly reduced in HuMCs treated with GSK3\(\beta\)-targeted shRNA. In contrast to the HMC1.2 cells, there was minimal constitutive phosphorylation of the S\(\theta\) residue of GSK3\(\beta\) in resting primary HuMCs. However, this phosphorylation was enhanced in SCF-challenged cells (Fig. 2A). As expected, this phosphorylation was markedly reduced in cells treated with GSK3\(\beta\)-targeted shRNA. Nevertheless, the inability of SCF alone to further enhance the observed constitutive phosphorylation of GSK3\(\beta\) at Y216 again suggests that GSK3\(\beta\) may be constitutively active in the resting state in HuMCs and that this activity cannot be further enhanced through KIT activation. Furthermore, the lack of reduction of the phosphorylation of GS by SCF (Fig. 2B), a response that could be reduced by GSK3\(\beta\)-targeted shRNA (15), suggests that the phosphorylation of GSK3\(\beta\) at S\(\theta\) in the HuMCs minimally impacted GSK3\(\beta\) activity, at least over the time frame examined.

**FIGURE 1.** Antiproliferative effect of GSK3\(\beta\) knockdown in HMC1.2 cell line. A. The HMC1.2 cell line was transduced with shContr or shGSK3\(\beta\). Whole-cell extracts were prepared and immunoblotted with anti-GSK3\(\beta\), anti–phospho-GSK3\(\alpha/\beta\) (p-Y279/Y216), anti–p-GSK3\(\beta\) (S\(\theta\)), anti-Syk, or anti-KIT Abs. B. Cell viability was assessed by trypan blue dye exclusion. Cell survival was assessed by means of an MTT assay in which the starting cell number was 1 \times 10^5 cells/well (C), and proliferation was assessed by BrdU incorporation (D). In all cases, \(n = 3–5\). *\(p < 0.05\), **\(p < 0.01\) for comparison with shContr-transduced HMC1.2.

**FIGURE 2.** Expression and activity of GSK3\(\beta\) inhibition in primary HuMCs. A. HuMCs, transduced with shContr or shGSK3\(\beta\), were starved overnight in SCF-depleted media and then stimulated with SCF for 2 min as described in Materials and Methods. Whole-cell extracts were prepared and immunoblotted with anti–p-GSK3\(\beta\) (Y216), anti–p-GSK3\(\beta\) (S\(\theta\)), or anti-GSK3\(\beta\) Abs. The level of Syk demonstrates equivalent protein loading of samples. B. Kinetics of SCF-mediated phosphorylation of GSK3\(\beta\) in HuMCs. Whole-cell extracts were prepared and immunoblotted with anti–p-GSK3\(\alpha/\beta\) (p-Y279/Y216), anti–p-GS (pGS) (S641), or anti–p-GSK3\(\beta\) (S\(\theta\)) Abs. The level of Syk demonstrates equivalent protein loading of samples. C. HuMCs were preincubated with a Src inhibitor (PP2), a PLC\(\gamma\) inhibitor (U73122), or a PI3K inhibitor (wortmannin [Wortm]) for 20 min prior to stimulation with SCF for 2 min as described in Materials and Methods. Whole-cell extracts were prepared and immunoblotted with anti–p-Akt (S473), anti–p-GSK3\(\beta\) (Y216), anti–p-GSK3\(\beta\) (S\(\theta\)), or anti-pGS (S641) Abs. The level of Syk demonstrates equivalent protein loading of samples.
We have previously determined that phosphorylation of the downstream substrate for GSK3β, GS, is moderately enhanced in HuMCs following a combination of KIT and FcεRI-mediated activation (15). Hence, we examined the phosphorylation status of p-GSK3β (Y216) and GS (S611) in HuMC over a 10-min time period during which maximal SCF-induced KIT phosphorylation is observed (24). As can be seen from Fig. 2B, there was no consistent increase in phosphorylation of either of these respective GSK3β and GS residues. These data overall are thus consistent with the conclusion that the constitutive phosphorylation and activity of GSK3β observed in the HMC1.2 and HuMC cells is regulated at least in part independently of KIT activity.

To provide further support for this conclusion, we investigated whether inhibitors of known signaling processes downstream of KIT block the constitutive phosphorylation of GS and GSK3β, including PP2, an Src family tyrosine kinase inhibitor also capable of inhibiting KIT phosphorylation (24); U71322, a PLCγ inhibitor (25); and wortmannin, a selective PI3K inhibitor (25). As expected, and to serve as a positive control, PP2 inhibited SCF-mediated PI3K activation, as indicated by the attenuation of the expected, and to serve as a positive control, PP2 inhibited SCF-mediated PI3K activation, as indicated by the attenuation of the p-Akt phosphorylation (Fig. 2A). U73122 partially blocked Akt phosphorylation, whereas U73122 partially blocked Akt phosphorylation (Fig. 2C). Although all three agents variably inhibited the increase in phosphorylation of GSK3β at S396 and the observed increase in phosphorylation in GS at S611 in these experiments, they had minimal effect on the constitutive phosphorylation of GSK3β at Y216 and GS at S611. These data suggest that although KIT activates PLCγ1 (25), tyrosine kinases, and PI3K, the constitutive activation pathway is independent of these signaling processes. These data further support the conclusion that the constitutive activation of GSK3β observed in mast cells is independent of KIT activation.

**GSK3β is required for SCF-mediated HuMCs survival**

To next determine the outcome of GSK3β knockdown on HuMC survival, two different shRNA constructs were selected based on their relative abilities to decrease GSK3β expression in HuMCs as described (15) and as shown in Fig. 3A. Knockdown of GSK3β with these constructs resulted in a significant reduction in viable cells as determined by trypan blue viability count (68.9 ± 8.6% [construct A] and 31.2 ± 10.3% [construct B] versus 91.3 ± 2.7% [control construct] remaining viable cells; n = 3, p < 0.01) (Fig. 3B) and with the MTT assay (54.4 ± 10.4% and 25.7 ± 7.1% versus 95.8 ± 3.3% for survival of cell input; n = 3, p > 0.05 and p < 0.001, respectively) (Fig. 3C). Based on these parameters, the decrease in survival of the cells treated with the two different GSK3β constructs thus correlated to the degree of GSK3β knockdown observed in Fig. 3A. Taken together, these data indicate that, although the requirements for GSK3β in the maintenance of neoplastic mast cells may be partly dependent on effects on cell division, at least in mature primary cultured HuMCs, it principally functions as an antiapoptotic signal for the maintenance of mast cell survival.

**GSK3β knockdown increases SCF withdrawal-induced apoptosis in HuMCs**

To provide further evidence to support the role of GSK3β as an antiapoptotic factor, we next determined whether the degree of knockdown of GSK3β similarly correlated to the respective markers of apoptosis and cell death, Annexin V and PI staining. The GSK3β-targeted shRNA-treated cells were grown in SCF-depleted medium for 20 h to increase the sensitivity of the assay. Under these conditions, the shGSK3β-treated HuMCs displayed a significant increase in apoptosis compared with the scrambled control-treated cells (construct A: 39.9 ± 2.7% and construct B: 54.5 ± 3.9% versus 25.6 ± 3.1% apoptotic cells; n = 4 to 5, p < 0.001) (Fig. 4A). Fig. 4B shows representative scatter plots of HuMCs transfected with shContr or shRNA for two different constructs of GSK3β starved in SCF-depleted media for 20 h and stained for Annexin V and PI. Caspase-3, which is activated upon cleavage, plays a dominant role in the extrinsic apoptotic pathway. To further establish the role of GSK3β in the extrinsic apoptotic pathway, we therefore next examined cleaved caspase-3 by Western blotting. As can be seen in Fig. 4C and 4D, the increase in apoptotic cells was associated with an increase in cleaved caspase-3 in the GSK3β knockdown cells compared with control-treated cells. Taken together, these data support a role for GSK3β as a key regulator of an antiapoptotic signaling pathway required for mast cell homeostasis. These data further support the conclusion that the antiapoptotic signals provided by GSK3β act in conjunction with those initiated by SCF, rather than being directly regulated by SCF-mediated signaling.
Downregulation of GSK3β inhibits SHP2 and ERK1/2 but not PI3K activation

As discussed (26, 27), an intact PI3K signaling axis is required for mast cell homeostasis. We therefore next examined whether the similar requirement for GSK3β for the maintenance of mast cell homeostasis could be explained by potential positive-feedback regulation of PI3K activity. We thus examined if the SCF-stimulated Akt activation in primary HuMCs transduced with scrambled shRNA or GSK3β knockdown cells compared with the scrambled control cells. It is also of note that S9 of GSK3β was not due to positive feedback regulation of PI3K, but requires the regulation of other critical signaling elements required for mast cell homeostasis.

GSK3β knockdown induces apoptosis in HuMC. A, HuMCs transduced with shContr or shRNA for GSK3β (shGSK3β A and B) were starved for 20 h, and apoptosis was measured by Annexin V staining. B, Representative scatter plots of HuMCs transduced with shContr or shRNA for two different constructs of GSK3β starved in SCF-depleted media for 20 h and stained for Annexin V (x-axis) and PI (y-axis). The percent represents total Annexin V+ cells (percent Annexin V+ cells in upper and lower right quadrants). shControl and shGSK3β-transduced cells (construct B) were starved 15–20 h in SCF-depleted medium and then stimulated with SCF for 2 min. C, Whole-cell extracts were prepared and immunoblotted with anti-GSK3β, anti-cleaved caspase-3 Abs. The level of Syk demonstrates equivalent protein loading of samples. Data in D were generated by scanning blots from three to four independent experiments for cleaved caspase-3 and normalized to responses obtained at 2 min with SCF stimulation. In all cases, n = 3–5, *p < 0.05, ***p < 0.001 for comparison with shContr-transduced HuMCs.

In addition to PI3K, studies conducted in mouse bone marrow-derived mast cells suggest that SHP2 is also required for SCF-mediated mast cell proliferation and survival through the regulation of Rac/JNK (28). Furthermore, SHP2/Ras activation has also been shown to be critical for growth factor-induced ERK activation through an undefined mechanism (29). We thus investigated whether the impact of GSK3β deficiency on HuMC survival may be explained by a downregulation of SHP2 and/or ERK activity. As can be seen in Fig. 6A–C, SCF-mediated activation of both SHP2 and ERK was significantly reduced in the GSK3β knockdown cells compared with the control cells. It is also of note that the constitutive phosphorylation of SHP2 but not ERK tended to also be lower in GSK3β knockdown cells. Taken together, these data indicate that the constitutive activation of GSK3β maintains mast cell homeostasis by providing a permissive signal allowing phosphorylated KIT to recruit SHP2 and ERK.

Knockdown of GSK3β downregulates expression of Bcl-2 family member antiapoptotic proteins

Growth factor-induced apoptosis is regulated by the interplay of prosurvival and proapoptotic members of the Bcl-2 family proteins (30, 31). Studies have demonstrated that hyperactivation of phospho-ERK enhances cell survival by inducing expression of the prosurvival molecules Bcl-2 and Bcl-xL and suppression of the proapoptotic BH3-only protein Bim (32, 33). Similarly, gain-of-function SHP2 mutant-expressing cells have elevated levels of Bcl-2 and Bcl-xL and lower levels of Bim compared with wild-type cells (34). Therefore, we next investigated whether the upregulation of Bcl-2 and/or Bcl-xL and downregulation of Bim in SCF-stimulated HuMCs transduced with scrambled shRNA for GSK3β and in GSK3β knockdown cells compared with control cells. These observations indicate that the decrease in survival of HuMCs following knockdown of GSK3β was not due to positive feedback regulation of PI3K by GSK3β, but requires the regulation of other critical signaling elements required for mast cell homeostasis.

FIGURE 4. GSK3β knockdown induces apoptosis in HuMC. A, HuMCs transduced with shContr or shRNA for GSK3β (shGSK3β A and B) were starved for 20 h, and apoptosis was measured by Annexin V staining. B, Representative scatter plots of HuMCs transduced with shContr or shRNA for two different constructs of GSK3β starved in SCF-depleted media for 20 h and stained for Annexin V (x-axis) and PI (y-axis). The percent represents total Annexin V+ cells (percent Annexin V+ cells in upper and lower right quadrants). shControl and shGSK3β-transduced cells (construct B) were starved 15–20 h in SCF-depleted medium and then stimulated with SCF for 2 min. C, Whole-cell extracts were prepared and immunoblotted with anti-GSK3β, anti-cleaved caspase-3 Abs. The level of Syk demonstrates equivalent protein loading of samples. Data in D were generated by scanning blots from three to four independent experiments for cleaved caspase-3 and normalized to responses obtained at 2 min with SCF stimulation. In all cases, n = 3–5. *p < 0.05, ***p < 0.001 for comparison with SCF response in shContr-transduced HuMCs.

FIGURE 5. Effect of GSK3β knockdown on SCF-mediated PI3K activation in primary HuMCs. A, HuMCs, transduced with scrambled shRNA or shGSK3β, were stimulated with SCF for 2 min as described in Materials and Methods. Whole-cell extracts were prepared and immunoblotted with anti-p-Akt (S473), anti-phospho-GSK3β (p-Y279/Y216), or anti-p-GSK3β (S9) Abs. The level of Syk demonstrates equivalent protein loading of samples. A representative blot is shown in figure. B–D, Data were generated by scanning multiple blots as shown in A and normalizing to the response obtained at 2 min with SCF stimulation. In all cases, n = 3 to 4. ***p < 0.001 for comparison with SCF response in shContr-transduced HuMCs.
both Bcl-2 and Bcl-xL and reduced levels of Bim (32), which has previously been shown to be critical for growth factor deprivation-induced mast cell apoptosis (34). As we observed that GSK3β contributed to the regulation of ERK and SHP2 activity, we examined the expression of both antiapoptotic (Mcl-1, Bcl-xL, and Bcl-2) and proapoptotic (Bim, Bid, and Bad) proteins in the SCF-starved and nonstarved HuMCs following treatment with GSK3β-targeted and control shRNA. As shown in Fig. 7A, GSK3β knockdown, if anything, increased the expression of Mcl-1, whereas SCF starvation reduced the expression of this antiapoptotic protein. However, the expression of Bcl-xL and Bcl-2 were reduced by both SCF starvation and GSK3β knockdown, with further reduction observed with the combination of these approaches. In contrast, whereas SCF enhanced the expression of Bim (Fig. 7B) but partially reduced the expression of Bid and Bad, GSK3β knockdown reduced the expression of all three proteins. Nevertheless, the data as a whole suggested that GSK3β acts as a prosurvival signal for human mast cells by regulating the expression of the antiapoptotic proteins following GSK3β knockdown (Bim, Bid, and Bad) and, to a certain extent, by SCF starvation (Bid and Bad) was somewhat counterintuitive but may be a consequence of caspase cleavage (35, 36).

**A small-molecule GSK3β inhibitor decreases survival of the HMC1.2 human leukemia mast cell line**

Finally, based on our conclusion that GSK3β was an important mast cell prosurvival signal, we investigated whether a small molecule inhibitor of GSK3β could be employed to reduce HMC1.2 survival. To date, CHIR 99021 is the most selective inhibitor of GSK3β reported (37, 38). As can be seen in Fig. 8, this compound dose-dependently reduced the number of viable HMC1.2 cells after 24, 48, and 72 h in culture, thus providing evidence for

![FIGURE 6.](http://www.jimmunol.org/)

**FIGURE 6.** GSK3β knockdown reduces SCF-mediated SHP2 and ERK activation in primary HuMCs. A, HuMCs, transduced with scrambled shRNA or shGSK3β, were stimulated with SCF for 2 min as described in Materials and Methods. Whole-cell extracts were prepared and immunoblotted with anti-GSK3β, anti-p-SHP2 (Y580), or anti-p-ERK (T202/Y204) Abs. The level of Syk demonstrates equivalent protein loading of samples. A representative blot is shown in figure. B–D, Data were generated by scanning the multiple blots as in A and normalizing to the response obtained at 2 min with SCF stimulation in shContr-transduced HuMCs. In all cases, n = 3 to 4. **p < 0.01, ***p < 0.001 for comparison with SCF response.

![FIGURE 7.](http://www.jimmunol.org/)

**FIGURE 7.** Effect of GSK3β knockdown and SCF starvation on pro- and antiapoptotic proteins in primary HuMCs. HuMCs transduced with shContr or shGSK3β A were (SCF+) or were not (SCF−) starved for 20 h. Whole-cell extracts were prepared and immunoblotted with GSK, Mcl-1, Bcl-xL, Bcl-2 (A), or Bim, BID, or Bad-specific (B) Abs. Immunoblotting with Syk-specific Ab was used as sample protein loading control. The relative intensities of the immunoreactive proteins (left panels) were normalized to Syk and evaluated (right panels). In all cases, n = 3 to 4. *p < 0.05, **p < 0.01, ***p < 0.001 for comparison with SCF response.
Discussion
In this paper, we describe studies that support the conclusion that GSK3β represents an essential prosurvival and/or antiapoptotic signaling element in human mast cells and that this may be a consequence of the dependency of SCF-mediated SHP2 and ERK activation, known mast cell survival signals, on GSK3β activity.

Mast cell homeostasis represents a balance between mast cell proliferation/survival signal, and mast cell proapoptotic signals (13, 39). In mature terminally differentiated human mast cells, SCF-mediated KIT activation skews this balance to prosurvival. Similarly, the ability of recognized KIT inhibitors to downregulate the expansion and survival of HMC1.1 and 1.2 cells (40) indicates that the myeloproliferative/survival potential of these cells is, at least in part, a consequence of the presence of activating mutations in KIT in these cell lines. As would be expected for a putative prosurvival/antiapoptotic signal, we observed that GSK3β is constitutively active in the HMC1.2 cells (Fig. 1) and in HuMCs (Fig. 2), as reported (15). This was evidenced by the phosphorylation of the activating Y216 residue in GSK3β and GS at S641 under resting conditions in both the HMC1.2 cells and HuMCs. It is tempting to speculate that the constitutive activation of GSK3β that we observed in the HMC1.2 cell line and the HuMCs may be a consequence of KIT activity. We have observed, for example, that there is a slight enhancement of GS phosphorylation at position S641 in HuMCs coactivated via KIT and the FcεRI. However, the inconsistency of this response in HuMCs challenged with SCF alone (Fig. 2) and the lack of an increase in phosphorylation of GSK3β at the activating Y216 residue in response to SCF would suggest that GSK3β activity in these cells is largely regulated independently of SCF. This conclusion is further supported by the limited ability of inhibitors of SCF-mediated signaling events to attenuate the constitutive GSK3β activity.

The kinase(s) responsible for the phosphorylation of GSK3β at Y216 in the human mast cells remain(s) unclear. However, the studies conducted with PP2 would suggest that Src kinase family members are not responsible. It is evident, however, that the regulation of GSK3β activity at any point in time represents a balance between negative and positive regulatory pathways (21). Although it is apparent that SCF can induce the phosphorylation of GSK3β at the inhibitory site (S9), this was not associated with a decrease in the phosphorylation of GS at S641, which we have demonstrated requires the presence of GSK3β for optimal phosphorylation in mast cells (15). From these data, we conclude that there is net constitutive activation of GSK3β, and, based on the shRNA knockdown studies, it is evident that the constitutive GSK3β activity present in both HMC1.2 cells (Fig. 1) and the HuMCs (Fig. 3) is essential for their survival.

The requirement for GSK3β in mast cell homeostasis may be dependent on both its regulation of cell division and of antiapoptotic pathways. Although the BrdU assay conducted in the HMC1.2 cells indicates that the dependency of the survival of these cells on GSK3β could be partly explained by prevention of cell division, the studies conducted on the nonproliferating HuMCs would indicate that GSK3β also plays a major role in the prevention of mast cell apoptosis. This conclusion is supported by the close correlation between GSK3β expression and HuMC survival as determined by trypan blue exclusion and the MTT assay (Fig. 3) and by the close correlation between the degree of GSK3β knockdown and the indices of apoptosis, Annexin V staining, and caspase cleavage in the SCF-starved HuMCs (Fig. 4).

GSK3β has been described to regulate multiple cellular events including cell growth, cell survival, metabolism, gene expression, and apoptosis (21, 41–43). However, in certain cases, including the regulation of growth factor- and mitogen-mediated responses, the described roles appear paradoxical in that GSK3β may both positively and negatively regulate cellular processes through tightly coupled activation and/or inactivation. For example, in a variety of cell types including eosinophils (41), GSK3β has been suggested to support cell survival downstream of the PI3K/Akt pathway. However, in studies conducted in both hematopoietic cells and in neuronal cells, it has been proposed that apoptosis induced by growth factor withdrawal or PI3K inhibition is mediated by GSK3β (44–46). Furthermore, GSK3β inhibition has been suggested to modulate radiation resistance in certain cancers as well as promoting tumor growth through stabilization of B-catenin (16, 47). Thus, as with the regulation of GSK3β activity, GSK3β-regulated responses may also represent a fine balance between negatively regulated and positively regulated signaling responses.

Our previous studies in which we explored the regulation of mast cell chemotaxis and cytokine production by GSK3β (15) also indicate that the role(s) of GSK3β in mast cell function are complex. Our results suggested, for example, that constitutively activated GSK3β must be considered in the context of regulation of signaling events in addition to the potential roles of receptor-mediated upregulation and/or downregulation of GSK3β activity (15). Certainly, in the current study, the constitutive activity of GSK3β we observed, the inability of SCF to enhance this response, and the requirement for ongoing GSK3β activity for cell survival would suggest that GSK3β is also a prerequisite signal, rather than an inducible signal, for mast cell homeostasis.

Although PI3K and PI3K-regulated prosurvival pathways are recognized to be critical for mast cell homeostasis, our observation that the phosphorylation of AKT is unaffected by GSK3β knockdown (Fig. 5) suggests that GSK3β does not regulate cell proliferation/survival through the feedback regulation of PI3K activity. However, based on our previous results (15), it is possible that PI3K may, in part, contribute to the regulation of GSK3β activity. It thus appears more likely that the ability of GSK3β to function as a prosurvival factor is related to its requirement for the ability of SCF to induce SHP2 and ERK phosphorylation (Fig. 6) and potentially JNK phosphorylation (15). In mouse bone marrow-derived mast cells, SHP2, through the regulation of Rac/JNK, has been shown to be required for SCF-mediated mast cell prolifera-
tion and survival (28) and to be critical for growth factor-induced ERK activation (48). SHP2 is a ubiquitously expressed non-receptor protein tyrosine phosphatase that participates in signaling events downstream of receptors for growth factors, cytokines, hormones, and control cell growth, differentiation, migration, and death (48). Thus, activation of SHP2 and its association with Gab1 is critical for sustained ERK activation downstream of several growth factor receptors and cytokines (48). The necessity for GSK3β in regulation of the MAPKs ERK and JNK in the manner described above may be due to priming of regulatory components of the MAPK pathways by GSK3β (15). Consequently, a deficiency in GSK3β would result in an inability of SCF, or indeed other stimuli, to regulate cellular responses via the MAPKs. However, it is also possible that SHP2 acts as a GSK3β substrate, because GSK3β-knockdown cells showed an impaired SCF-mediated activation of SHP2. As both SHP2 and ERK have been demonstrated to require the expression of antiapoptotic Bcl-2 family proteins (32, 33), our observation that GSK3β may function as a prosurvival signal in human mast cells. Such a mechanism would involve the requirement of SHP2 and ERK for constitutively active GSK3β to promote cell survival through expression of the Bcl-2 family members Bcl-2 and Bcl-xL.

In summary, the data presented in this study provide evidence that GSK3β is a key regulator of mast cell homeostasis in both neoplastic and primary cultured human mast cells through prevention of apoptosis. The data also indicate that the myeloproliferative capacity of the neoplastic HMC1.2 cells at least in intervention of apoptosis. The data also indicate that the myeloproliferative and primary cultured human mast cells through pre-

References