Serum- and glucocorticoid-induced kinase 3 in recycling endosomes mediates acute activation of Na\(^{+}/H^{+}\) exchanger NHE3 by glucocorticoids

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**ABSTRACT** Na\(^{+}/H^{+}\) exchanger 3 (NHE3) is the major Na\(^{+}\) transporter in the intestine. Serum- and glucocorticoid-induced kinase (SGK) 1 interacts with NHE regulatory factor 2 (NHERF2) and mediates activation of NHE3 by dexamethasone (Dex) in cultured epithelial cells. In this study, we compared short-term regulation of NHE3 by Dex in SGK1-null and NHERF2-null mice. In comparison to wild-type mice, loss of SGK1 or NHERF2 significantly attenuated regulation of NHE3 by Dex but did not completely obliterate the effect. We show that transfection of SGK2 or SGK3 in PS120 cells resulted in robust activation of NHE3 by Dex. However, unlike SGK1 or SGK2, SGK3 rapidly activated NHE3 within 15 min of Dex treatment in both PS120 and Caco-2bbe cells. Immunofluorescence analysis showed that SGK3 colocalized with NHE3 in recycling endosomes, whereas SGK1 and SGK2 were diffusely distributed. Mutation of Arg-90 of SGK3 disrupted the endosomal localization of SGK3 and delayed NHE3 activation. Activation of SGK3 and NHE3 by Dex was dependent on phosphoinositide 3-kinase (PI3K) and phosphoinositide-dependent kinase 1 (PDK1), and Dex induced translocation of PDK1 to endosomes. Our study identifies SGK3 as a novel endosomal kinase that acutely regulates NHE3 in a PI3K-dependent mechanism.

**INTRODUCTION**

Glucocorticoids (GCs) are widely used to treat inflammatory bowel disease (Campieri et al., 1997). In addition to antiinflammatory effects, GCs show a direct effect on NaCl and water absorption in the intestine by targeting Na\(^+/H^{+}\) exchanger 3 (NHE3; Charney et al., 1975), which is the major Na\(^{+}\) transporter located at the brush border membrane of the intestinal epithelium (He and Yun, 2010). Early studies showed that GCs stimulate NHE3 primarily through a genomic effect that increases NHE3 transcripts (Yun et al., 1993). More recently, GCs were implicated in nongenomic regulation of NHE3, with GCs stimulating NHE3 activity without changing NHE3 expression levels, and serum- and glucocorticoid-induced kinase (SGK) 1 was subsequently shown to mediate nongenomic regulation of NHE3 by GCs (Baum et al., 1994; Yun et al., 2002a; Wang et al., 2005, 2007).

SGK1 was initially identified by a differential screen of a rat mammary tumor cell line in search of GC-inducible genes (Webster et al., 1993), and a subsequent protein sequence homology search identified two additional SGK isoforms, SGK2 and SGK3 (also known as cytokine-independent survival kinase; Kobayashi et al., 1999). A body of evidence shows that SGKs share similar functions in regulation of a broad range of ion transporters and channels, including glutamate transporters (Boehmer et al., 2003), Na\(^{+}\) channels (Alvarez et al., 2003), and mediates activation of NHE3 by dexmethasone (Dex) in cultured epithelial cells. In this study, we compared short-term regulation of NHE3 by Dex in SGK1-null and NHERF2-null mice. In comparison to wild-type mice, loss of SGK1 or NHERF2 significantly attenuated regulation of NHE3 by Dex but did not completely obliterate the effect. We show that transfection of SGK2 or SGK3 in PS120 cells resulted in robust activation of NHE3 by Dex. However, unlike SGK1 or SGK2, SGK3 rapidly activated NHE3 within 15 min of Dex treatment in both PS120 and Caco-2bbe cells. Immunofluorescence analysis showed that SGK3 colocalized with NHE3 in recycling endosomes, whereas SGK1 and SGK2 were diffusely distributed. Mutation of Arg-90 of SGK3 disrupted the endosomal localization of SGK3 and delayed NHE3 activation. Activation of SGK3 and NHE3 by Dex was dependent on phosphoinositide 3-kinase (PI3K) and phosphoinositide-dependent kinase 1 (PDK1), and Dex induced translocation of PDK1 to endosomes. Our study identifies SGK3 as a novel endosomal kinase that acutely regulates NHE3 in a PI3K-dependent mechanism.

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Because our initial proposal for the role of SGK1 in NHE3 regulation pertains to short-term rather than chronic regulation (Grahammer et al., 2006), we investigated the effects of Dex treatment for 4 h as a short-term treatment and 24 h as a long-term treatment. The basal rate of Na+/H+ exchange activity in NHE3 was measured in the presence of NHE1 and NHE2 inhibitor Hoe-694 in isolated villi of Sgk1floxflox and Sgk1floxflox;Villin-Cre mice was not changed by the loss of SGK1 (Figure 1). Dex treatment of Sgk1floxflox mice for 4 and 24 h increased NHE3 activity in the ileum by ~120% (Figure 1, A and B). In contrast, the loss of SGK1 in Sgk1floxflox;Villin-Cre mice markedly decreased the effect of Dex resulting in only ~25–30% increase in NHE3 activity (Figure 1, C and D). These findings reveal that SGK1 plays a significant role in, but is not obligatory for, Dex-mediated stimulation of NHE3. To ensure that the remaining increase in Na+/H+ exchange activity in NHE3 in the absence of SGK1 or NHERF2 suggests that additional means for NHE3 activation must be present (Yun et al., 2002a). One potential mechanism involves increased NHE3 expression, but our earlier study showed that NHE3 protein abundance does not change at 4 h (Wang et al., 2007), requiring an alternative explanation.

**RESULTS**

**SGK1 and NHERF2 play important roles in Dex-mediated stimulation of NHE3 in ileum**

It was shown previously that treatment of SGK1-null mice with Dex for 96 h resulted in a ~40% increase in NHE3 activity compared with ~110% in wild-type (WT) mice (Grahammer et al., 2006). For the current study, we mated Sgk1floxflox mice with Villin-Cre mice to generate Sgk1floxflox;Villin-Cre mice, in which the Villin promoter directs the deletion of Sgk1 gene outside the epithelial cells in the intestine (Pinto et al., 1999). The expression of Sgk1 was examined by reverse transcription PCR (RT-PCR) and Western blotting to confirm the absence of Sgk1 in the intestinal epithelial cells of Sgk1floxflox;Villin-Cre mice (Supplementary Figure S1, A and B).

Because our initial proposal for the role of SGK1 in NHE3 regulation pertains to short-term rather than chronic regulation (Grahammer et al., 2006), we investigated the effects of Dex treatment for 4 h as a short-term treatment and 24 h as a long-term treatment. The basal rate of Na+/H+ exchange activity in NHE3 was measured in the presence of NHE1 and NHE2 inhibitor Hoe-694 in isolated villi of Sgk1floxflox and Sgk1floxflox;Villin-Cre mice was not changed by the loss of SGK1 (Figure 1). Dex treatment of Sgk1floxflox mice for 4 and 24 h increased NHE3 activity in the ileum by ~120% (Figure 1, A and B). In contrast, the loss of SGK1 in Sgk1floxflox;Villin-Cre mice markedly decreased the effect of Dex resulting in only ~25–30% increase in NHE3 activity (Figure 1, C and D). These findings reveal that SGK1 plays a significant role in, but is not obligatory for, Dex-mediated stimulation of NHE3. To ensure that the remaining increase in Na+/H+ exchange activity in NHE3 in the absence of SGK1 or NHERF2 suggests that additional means for NHE3 activation must be present (Yun et al., 2002a). One potential mechanism involves increased NHE3 expression, but our earlier study showed that NHE3 protein abundance does not change at 4 h (Wang et al., 2007), requiring an alternative explanation.

**DIFFERENTIAL REGULATION OF NHE3 BY SGK1, SGK2, AND SGK3**

Because other SGK isoforms, SGK2 and SGK3, can mediate similar effects, we sought to determine whether SGK2 or SGK3 regulates NHE3 activity in response to Dex treatment. The presence of SGK2 and SGK3 in small intestine was confirmed by RT-PCR. The expression levels of SGK2 and SGK3 mRNA in intestinal epithelial cells were not changed by the loss of SGK1 in Sgk1floxflox;Villin-Cre mice (Supplementary Figure S1A). PS120 cells stably expressing rabbit NHE3 with a vesicular stomatitis virus glycoprotein (VSVG) epitope fused at the carboxy-terminal, PS120/NHE3V, were transfected with hemagglutinin (HA)-SGK1, -SGK2, or -SGK3, and the resulting cells were treated with 1 μM Dex (Figure 2A). Overexpression of SGK1 did not mediate an apparent activation of NHE3 by Dex, as PS120 cells lacking NHERF2 expression (Figure 2B), which is required for SGK1-mediated NHE3 regulation (Yun et al., 2002a). In contrast, both SGK2 and SGK3 showed significant increases in NHE3 activity at both 4 and 24 h of Dex treatment (Figure 2, C and D), suggesting that the remaining NHE3 activation in Sgk1floxflox;Villin-Cre mice might have been due to the activities of SGK2 and SGK3. Surprisingly, Dex treatment showed a marked increase in NHE3 activity in PS120/NHE3V/HA-SGK3 cells at 15 min (Figure 2D), but not in cells expressing SGK1 or SGK2. To our knowledge, this is the first observation of such a rapid effect by Dex on NHE3. However, we did not observe any effect at 5 min, and we considered 15 min as the earliest time needed for Dex to regulate NHE3.
Intriguingly, NHE3 activity was significantly increased (~25%) in Sgk1flox/flox mice (Figure 3A), validating the acute regulation of NHE3 by Dex. Moreover, Dex stimulated NHE3 activity in both Sgk1flox/flox;Villin-Cre and Nherf2−/− villi to the same extent as WT villi (Figure 3, A and B), further demonstrating that Dex-induced acute (15 min) stimulation of NHE3 in the intestine is independent of SGK1 and NHERF2. We confirmed that the acute pH recovery was due to NHE3 by using S-3226 (Supplementary Figure S2C).

Differential regulation of SGK1, SGK2, and SGK3 kinase activity by Dex

Despite the similarity in SGK isoforms, SGK1 is the only isoform for which expression is regulated by GCs. However, there is the possibility that Dex treatment regulates the kinase activity without a change in the expression level. Therefore we examined whether the kinase activity of these SGKs is regulated by Dex. We again used PS120/NHE3V cells expressing HA-SGK1, HA-SGK2 or HA-SGK3 to determine the effect of Dex on SGK kinase activity. These cells have an advantage in that the expression of SGKs is under the control of cytomegalovirus (CMV) promoter and is not altered by Dex. Cells were treated with Dex for 0–24 h, and HA-SGKs were immunoprecipitated with anti-HA antibodies. The amounts of SGKs were normalized by Western immunoblot analysis of an aliquot of immunoprecipitated proteins to ensure that the identical amount of SGK protein was used in the kinase assay. We determined SGK kinase activity by two approaches: by using [γ-32P]ATP and by a nonradioisotope time-resolved fluorescence–fluorescence resonance energy transfer (TR-FRET) method. Figure 4 shows that the kinase activities of SGK1, SGK2, and SGK3 were similarly increased at 4 and 24 h of Dex treatment. However, unlike that of SGK1 or SGK2 (Figure 4, A and B), SGK3 activity was markedly increased at 15 min (Figure 4C). The nonradioactive TR-FRET assay yielded similar results (Supplementary Figure S3). Although the extent of increases was smaller in the TR-FRET assay, it is important to note that the results were qualitatively identical to [γ-32P]-labeling of Crosstide. We therefore used the TR-FRET method to determine kinase activity in the subsequent experiments.

In light of our unexpected findings, we examined whether Dex acutely regulates NHE3 in vivo. Because 15-min treatment of animals in vivo is not feasible, isolated villi were treated with 1 μM Dex ex vivo for 15 min prior to measurement of NHE3 activity.

SGK2 and SGK3 mediate exocytic trafficking and phosphorylation of NHE3

We showed previously that SGK1 stimulates NHE3 activity by increasing NHE3 expression at the cell surface (Wang et al., 2005). To
In contrast, the presence of SGK3 resulted in a significant increase in surface NHE3 at both 15 min and 4 h of Dex treatment (Figure 5A).

We have previously shown that SGK1-mediated activation of NHE3 is dependent on phosphorylation of NHE3 at S663 (Wang et al., 2005). Because SGK2 and SGK3 recognize the same phosphorylation motif (Kobayashi et al., 1999), we sought to determine the importance of S663 in NHE3 regulation by SGK2 or SGK3. PS120/ opposum NHE3 (OKNHE3) and PS120/OKNHE3-S663A (OKNHE3 with a mutation of serine 663 to alanine) cells transfected with HA-SGK2 or HA-SGK3 were treated with Dex. Dex treatment for 4 h stimulated the activity of NHE3 as we have shown earlier, but the mutation of S663A ablated the effect (Figure 5B). For SGK3, we treated the cells for 15 min, but as for SGK2, NHE3-S663A did not respond to Dex treatment (Figure 5C).

Although direct phosphorylation of NHE3 by SGK2 or SGK3 was not shown here, the results demonstrate that the presence of S663 is required for NHE3 regulation by SGK2 and SGK3.

SGK3 and NHE3 colocalize in recycling endosomes

We then asked whether the distinct response of NHE3 to SGK3 could be related to their spatial localizations. Immunofluorescence analysis of PS120/NHE3V cells (Figure 6A, top and middle) showed that both SGK1 (green) and SGK2 (green) were diffusely expressed throughout the cells. SGK1 and SGK2 partially colocalized with NHE3 (red) based on the overlap of fluorescence labels but did not show any distinct compartmentalization with NHE3. In contrast, SGK3 labeling showed localization in endosome-like compartments (Figure 6A, bottom) and, importantly, SGK3 (green) colocalized with NHE3 (red) in these vesicles. The endosomal localization of SGK3 was confirmed by costaining with Rab5 (Figure 6B, bottom, red), an early endosome marker. In contrast, neither SGK1 nor SGK2 colocalized with Rab5 in early endosomes (Figure 6B, top and middle). Furthermore, we examined whether the SGKs are expressed in recycling endosomes. PS120/NHE3V cells were transfected with green fluorescent protein (GFP)-Rab11 (a marker for recycling endosome) together with FLAG-tagged SGK1, SGK2, or SGK3. Figure 6C shows that SGK1 and SGK2 (red) were not found in recycling endosomes where Rab11 (green) is expressed. In contrast, SGK3 (red) and NHE3 (blue) colocalized with Rab11 (green) in recycling endosomes (Figure 6D).

We next examined whether the endosomal localization of SGK3 is important for acute NHE3 regulation. It was shown that...
the PX domain of SGK3 is critical for its endosomal localization, and mutation of R90 located in the PX domain disrupts the endosomal localization of SGK3 (Tessier and Woodgett, 2006). Therefore we expressed SGK3 or SGK3-R90A (SGK3 containing a mutation of Arg90 to Ala) in PS120/NHE3V cells (Figure 7A). Immunofluorescence staining confirmed that the R90A mutation perturbed the cellular location of SGK3 and, unlike WT SGK3, the SGK3-R90A mutant diffusely expressed throughout the cytoplasm, which appears similar to the expression patterns of SGK1 and SGK2; consequently, SGK3-R90A did not colocalize with NHE3 (Figure 7B). Intriguingly, the effect of the R90A mutation on endosomal localization of SGK3 was also extended to regulation of its activity in response to Dex. The kinase activity of SGK3-R90A was elevated at 4 and 24 h, but not at 15 min, of Dex treatment (Figure 7C), suggesting that the acute regulation of SGK3 by Dex is related to the endosomal localization of SGK3. Moreover, SGK3-R90A failed to activate NHE3 activity at 15 min, but its effect was delayed until 4 h of Dex treatment (Figure 7D). These findings indicate that endosomal localization enables SGK3 to acutely regulate NHE3.

SGK3-mediated activation of NHE3 is dependent on GR, phosphoinositide 3-kinase, and phosphoinositide-dependent kinase 1

Recent evidence has emphasized the short-term effects of GCs, but the mechanism of how GCs exert acute effects is incompletely understood. The presence of a G protein–coupled GC receptor (GR) that is insensitive to the GR antagonist, RU486 (RU), has been proposed (Maier et al., 2005). Hence we first examined whether RU-insensitive GR is involved in the acute regulation of SGK3 by Dex. The preincubation of PS120/NHE3V/HA-SGK3 cells with RU (1 μM) did not affect the basal SGK activity but completely abrogated the Dex-induced rapid effect on SGK3 activity, suggesting the conventional GR and not the membrane-bound GR regulates SGK3 (Figure 8A). SGKs are related to Akt and are regulated by phosphoinositide 3-kinase (PI3K) and phosphoinositide-dependent kinase 1 (PDK1; Kobayashi et al.,...
induced PI3K-PDK1 pathway is not specific for acute (15 min) NHE3 activation by SGK3. We did not examine the role of PDK1 for SGK1 in PS120 cells, since SGK1 showed no effect on NHE3 (Figure 2B) unless NHERF2 was coexpressed, but we showed previously that regulation of NHE3 by SGK1 is PI3K-dependent (Yun et al., 2002a). If PDK1 is important for both SGK2- and SGK3-dependent regulation of NHE3, what allows the acute regulation of SGK3? To delineate the role of PDK1 in the rapid activation of SGK3 and NHE3, we examined the temporospatial location of phosphorylation of PDK1 by using the PI3K inhibitor, LY294002 (LY). Pretreatment with LY (20 μM) completely blocked the activation of SGK3 activity by 15 min of Dex treatment (Figure 8A). Similarly, RNA interference of PDK1 by short-hairpin RNA (shRNA), which reduced the PDK1 expression by ∼70% (Figure 8B), ablated the stimulatory effect of Dex on SGK3 activity (Figure 8C). Consistent with the inhibition of SGK3 kinase activity, the acute regulation of NHE3 transport activity by Dex in SGK3-expressing cells was completely blocked when cells were pretreated with RU or LY (Figure 8D), or with knockdown of PDK1 (Figure 8E). Taken together, these data show that acute activation of SGK3 and NHE3 by Dex is through the GR-PI3K-PDK1 pathway. Because the effects of SGK2 and SGK3 on NHE3 at a later time point (4 h) are similar, we determined whether the delayed regulation of NHE3 by SGK2 and SGK3 is also dependent on PDK1. Knockdown of PDK1 ablated activation of NHE3 activity in cells expressing SGK2 or SGK3 (Supplementary Figure S4), suggesting that the Dex-induced PI3K-PDK1 pathway is not specific for acute (15 min) NHE3 activation by SGK3. We did not examine the role of PDK1 for SGK1 in PS120 cells, since SGK1 showed no effect on NHE3 (Figure 2B) unless NHERF2 was coexpressed, but we showed previously that regulation of NHE3 by SGK1 is PI3K-dependent (Yun et al., 2002a).

If PDK1 is important for both SGK2- and SGK3-dependent regulation of NHE3, what allows the acute regulation of SGK3? To delineate the role of PDK1 in the rapid activation of SGK3 and NHE3, we examined the temporospatial location of phosphorylation of PDK1 (p-PDK1: an active form of PDK1 kinase) at Ser241 relative to SGK3 in response to Dex treatment. Fluorescence labeling of SGK3 (Figure 8F, green) partially colocalized with p-PDK1 (Figure 8F, red) under basal conditions, whereas treatment with Dex for 15 min resulted in increased overlap of their fluorescence labels (Figure 8F). A quantitative determination of colocalization of SGK3 and p-PDK1 showed a significant increase in colocalization coefficient induced by Dex treatment (Figure 8G). An x,z scan shows that Dex also increased colocalization of p-PDK1 and SGK3.
in the perimembrane areas (Supplementary Figure S5). To determine whether Dex mobilizes p-PDK1 to endosomal compartments, we determined the expression patterns of p-PDK1 in PS120/NHE3V cells upon 15 min of Dex treatment (Figure 8, H and I). The results show that the expression of p-PDK1 in recycling endosomes increased following Dex treatment. We also examined whether Dex-induced mobilization of p-PDK1 occurs in intestinal epithelial cells. Consistent with the findings in PS120 cells, Dex increased p-PDK1 expression in recycling endosomes in human intestinal epithelial Caco-2bbe cells (Supplementary Figure S6). These findings demonstrate that Dex acutely activates NHE3 by mobilizing PDK1 to endosomal compartments in the cytoplasm and on the membrane, where it activates SGK3 that in turn phosphorylates NHE3.

**SGK3 mediates acute activation of NHE3 in Caco-2bbe epithelial cells**

To reproduce the regulation of NHE3 via SGK3 in intestinal epithelial cells, we expressed HA-SGK3 in Caco-2bbe cells stably expressing human NHE3V, Caco-2bbe/hNHE3V (Lin et al., 2010). Dex treatment for 15 min stimulated NHE3 activity in HA-SGK3–expressing cells but not in pcDNA-transfected control cells (Figure 9A). Confocal
immunofluorescence analysis showed that SGK3 (green) colocalized with NHE3 (red) in the subapical compartments (Figure 9B). A cross-sectional view of the cells near the top surface and a stacked view show the apical expression of NHE3, which overlaps with fluorescent wheat germ agglutinin (WGA) that labels glycoprotein at the plasma membrane (Figure 9C, Cont). Treatment of the cells with Dex for 15 min enhanced the overlap of fluorescence labels of NHE3 and WGA, indicating increased surface NHE3 abundance in response to Dex (Figure 9C, Dex). The results from immunofluorescence analysis were confirmed by a surface biotinylation assay, which showed increased NHE3 protein abundance at the apical membrane (Figure 9D).

DISCUSSION

We have previously shown that SGK1 plays a critical role in Dex-induced activation of NHE3 in cultured fibroblast and epithelial cells (Yun et al., 2002a; Wang et al., 2005). However, the critical importance of SGK1 for NHE3 regulation in vivo has remained elusive until the recent report by Grahammer et al. (2006). In that study, the effects of Dex on intestinal NHE3 regulation were compared between WT and SGK1-null mice following a 4-d Dex treatment. Therefore we revisited Dex-induced regulation of NHE3 in the SGK1-null intestine to determine the effects at earlier time points at which NHE3 activity is increased but NHE3 abundance is not changed by Dex treatment. Moreover, we used mice with intestinal epithelia–specific deletion of SGK1. Our data showed that SGK1 plays a major role in short-term (4 h) regulation of NHE3 by Dex, and also confirmed the earlier report by Grahammer and coworkers. In addition, in agreement with our previous in vitro study that NHERF2 is required for SGK1-dependent regulation of NHE3, the NHERF2-null intestine showed decreased responses to Dex. However, the loss of neither SGK1 nor NHERF2 completely ablated the stimulatory effect of Dex. Our previous study showed that NHE3 abundance is not changed until much later than 4 h of Dex treatment and hence it is unlikely that the residual effect on NHE3 is simply due to changes in NHE3 abundance. Because SGKs share similar functions in regulation of various ion channels and transporters (Lang et al., 2003, 2006), we determined whether SGK2 or SGK3 is involved in NHE3 regulation. Our current study using PS120 cells showed that both SGK2 and SGK3 activate NHE3. Our previous study has shown that SGK1, but not SGK2 and SGK3, interacts with NHERF2 (Yun et al., 2002a). Consistently, stimulation of NHE3 activity by SGK2 and SGK3 was independent of NHERF2, occurring in PS120 cells that lack endogenous NHERF2, whereas SGK1 failed to regulate NHE3. A recent study showed that constitutively active SGK2, but not SGK1, stimulates NHE3 activity in OK cells (Pao et al., 2010). This finding is consistent with previous studies that OK cells express NHERF2 at low levels (Yun et al., 2002a). Apart from not needing NHERF2, all SGKs share common mechanisms that involve increased NHE3 abundance at 4 and 24 h of Dex treatment and the inability to regulate NHE3-S663A. In light of the high similarity of the kinase domains of SGK1–3, it is not surprising that they mediate Dex-induced activation of NHE3 via similar mechanisms. However, accumulating evidence shows that SGK3 plays distinct roles from SGK1. SGK3 is important for postnatal hair follicle growth (McCormick et al., 2004), and a recent study showed that expression of SGK3 is estrogen-inducible in breast cancer cells (Wang et al., 2011). In the current study, we showed for the first time that SGK3, but not SGK1 or SGK2, mediates an acute activation of NHE3 at as early as 15 min in response to Dex.

GC-mediated effects generally involve GR-dependent gene regulation. However, recent studies have highlighted short-term effects of GCs (Losel and Wehling, 2003), although the underlying mechanisms remain unclear. It was shown that Dex stimulates PI3K activity by potentiating binding of GR to the p85 subunit of PI3K (Hafezi-Moghadam et al., 2002; Hu et al., 2009). Unlike the classical activation of PI3K by insulin/IGF-1 receptors or tyrosine kinase receptors that occurs as a membrane-based event, GR-dependent activation of PI3K happens most likely in the cytoplasm with PI3K being sequestered away from the membrane (Hu et al., 2009), which in turn activates PDK1. A previous study showed that all SGKs are phosphorylated by PDK1 (Kobayashi et al., 1999), and knockdown of PDK1 abolished Dex-induced regulation of SGK2, SGK3, and NHE3 (Figures 8 and S4). We did not determine whether PDK1 directly regulates SGK1, but it was shown that PDK1 is required for PI3K-mediated activation of SGK1 and that SGK1dependent activation of NHE3 involves PI3K (Park et al., 1999; Yun et al., 2002a). Hence it is highly likely that PDK1 links PI3K to SGK1.

So, what makes SGK3 different from other SGKs? We showed that SGK1 and SGK2 are diffusely distributed throughout the cytoplasm, with some in nuclei, but none on endosomes (Figure 6). On the other hand, SGK3 is distinctly localized to recycling and early endosomes, which is consistent with previous reports using HEK293 and COS cells (Jun et al., 2001; Virbasius et al., 2001; Tessier and Woodgett, 2006). On activation with Dex, p-PDK1 is recruited to endosomes, where it activates SGK3, as demonstrated by increased colocalization of p-PDK1 with SGK3 and Rab11 (Figures 8 and S6). In agreement with our findings, a recent study demonstrated that SGK3 facilitates the recruitment of p-PDK1 to endosomes when SGK3 is preactivated at S486 through PI3K signaling (Slagsvold et al., 2006). Hence our current study demonstrates that the core of the acute regulation of SGK3 is its endosomal localization. The importance of endosomal residence of SGK3 for its regulation of NHE3 was further demonstrated by the finding that the R90A mutation in SGK3, which perturbed the endosomal localization of SGK3, lost its acute effect on NHE3 regulation, although delayed activation of NHE3 was not affected. How Dex induces delayed effects on NHE3 is not clear. We propose that SGK1 and SGK2, as well as the SGK3-R90A mutant deficient in the endosome-targeting PX domain, lack the spatial arrangement for a rapid access by PDK1 in response to Dex. Instead, much longer time is required for PDK1 and these SGKs to come in contact for activation of SGKs. In addition, cytosolic SGK1 and SGK2 may need to move through the cytoplasmic matrix to interact with and phosphorylate NHE3 (Figure 10).

Since the seminal studies by Grinstein and his colleagues in the late 1990s that showed NHE3 in recycling endosomes (D’Souza et al., 1998; Kurashima et al., 1998), the trafficking of NHE3 protein from recycling endosomes has been recognized as the pivotal step in NHE3 regulation. The former studies showed that inhibition of PI3K results in a pronounced loss of NHE3 from the cell surface and its accumulation in an intracellular endosomal compartment. Moreover, PI3K regulates NHE3 by several hormones and growth factors, including insulin, lysophosphatidic acid, and basic fibroblast growth factor (Janecki et al., 2000; Lee-Kwon et al., 2003; Fuster et al., 2007). However, the identity of the effector(s) of PI3K signaling targeting NHE3 in recycling endosomes has remained undefined. Our present study is the first to identify SGK3 as a PI3K-signaling effector in recycling endosomes that activates NHE3. We therefore speculate that the endosomal localization of SGK3 has a broader implication on NHE3 regulation. In addition, the localization of SGK3 in recycling endosomes and the broad tissue expression of SGK3 suggest that SGK3, under basal or induced conditions, might be involved in regulating other proteins that traffic between intracellular pools and the plasma membrane.
In summary, our study showed that SGK1 plays a major role in Dex-mediated activation of NHE3 in vivo in the intestine, but some NHE3 activity regulation is independent of SGK1 and NHERF2. We showed that both SGK2 and SGK3 contribute to NHE3 activation, but SGK3 uniquely colocalizes with NHE3 to recycling endosomes, enabling acute regulation of NHE3 via the PI3K-PDK1 pathway.

MATERIALS AND METHODS

Animals

Generation of mice with the sgk1 gene flanked by loxP (Sgk1^floxed/) in C57BL/6 background was reported previously (Fejes-Toth et al., 2008). Intestinal epithelia-specific deletion of SGK1 was achieved by mating Sgk1^floxed/ mice with Villin-Cre transgenic mice. Nherf2-null (Nherf2^−/−) mice in C57BL/6 background have been previously described (Lin et al., 2009). Genotypes were determined by PCR on tail DNA. In all experiments, age- and sex-matched littermates of Sgk1^floxed/ and Sgk1^floxed/Villin-Cre or WT and Nherf2^−/− mice were studied. Animals were maintained and experiments were performed under the institutional guidelines of Emory University.

Antibodies and plasmid constructs

Anti-HA, anti-PDK1, and anti-p–PDK1 (Ser241) antibodies were purchased from Cell Signaling Technology (Danvers, MA). Anti-HA antibody was obtained from Covance (Denver, PA). Rabbit and mouse anti-FLAG antibodies were purchased from Sigma (St. Louis, MO) and anti-Rab5 antibody from Santa Cruz Biotechnology (Santa Cruz, CA). The pB8 plasmid harboring HA-tagged human SGK1 has been previously described (Wang et al., 2005). HA-tagged human SGK2 and SGK3 and FLAG-tagged SGK1 and SGK2 were cloned in pcDNA3.1-hygro. FLAG-SGK3 and pEGFP-Rab11 were obtained from Addgene (Cambridge, MA). The pLKO.1 plasmid harboring sh-PDK1 was purchased from Open Biosystems (Huntsville, AL). OKNHE3-S663A has been previously described (Wang et al., 2005). SGK3-R90A was generated with the QuickChange II XL Site-Directed Mutagenesis Kit according to the recommendation by the manufacturer (Stratagene, Santa Clara, CA). The presence of mutation was confirmed by nucleotide sequencing.

Cell culture and transfection

PS120/NHE3V and PS120 expressing OKNHE3, PS120/OKNHE3, have been previously described (Wang et al., 2005; He et al., 2008). Caco-2bbe/hNHE3V cells were cultured as previously described (Lin et al., 2010). Transfection was performed using the Neon Transfection System according to the manufacturer’s recommendation (Invitrogen, Carlsbad, CA). Stable cell lines expressing SGK plasmids were selected with 800 μg/ml of neomycin or 600 μg/ml of hygromycin. Both PS120 fibroblasts and Caco-2bbe epithelial cells were cultured in DMEM supplemented with 1 mM sodium pyruvate, 50 U/ml penicillin, 50 μg/ml streptomycin, and 10% fetal bovine serum in a 5% CO2 humidified incubator at 37°C.

Immunoprecipitation and kinase activity assay

PS120/NHE3V cells overexpressing HA-SGK1, HA-SGK2, HA-SGK3, or HA-SGK3-R90A were treated with 1 μM Dex. Cells were then lysed in lysis buffer containing 20 mM Tris-HCl (pH 7.5), 150 mM NaCl, 5 mM β-glycerophosphate, 2.5 mM sodium pyrophosphate, 1 mM Na2EDTA, 1 mM ethylene glycol tetraacetic acid (EGTA), 1 mM Na3VO4, 1 mM sodium fluoride, 10 mM 1 μg/ml leupeptin, 1% Triton X-100, and protease inhibitor cocktail tablets (Roche, Indianapolis, IN). The crude lysate was sonicated twice for 15 s and spun at 14,000 × g for 15 min. Protein concentration was determined by the bicinchoninic acid assay (Sigma). Two milligrams of lysate was incubated overnight with or without anti-HA antibody. The immunocomplex was purified by incubation with 80 μl of protein A-Sepharose beads for 1 h, which was followed by three washes in lysis buffer and two washes in phosphate-buffered saline (PBS). Beads bound with SGK1, SGK2, SGK3, or SGK3-R90A were eluted with 50 μg/ml HA peptide (Covance) in elution buffer (50 mM Tris-HCl, pH 7.5, 150 mM NaCl, 270 mM sucrose, 1 mM Na3VO4, 0.1 mM EGTA, 0.2 mM phenylmethylsulfonyl fluoride, 0.1% β-mercaptoethanol and 10% glycerol). An aliquot of eluted protein was resolved by SDS–PAGE and immunoblotted with anti-HA antibody to ensure an identical amount of protein was used for kinase assay. In vitro kinase assay was determined with two approaches using 1) [γ-32P]ATP and 2) a nonradioisotope TR-FRET method. For the former, the kinase assay was performed for 20 min at 30°C in 50 μl of reaction volume containing 10 μl of immunoprecipitate in reaction buffer (20 mM Tris, 100 μM ATP, 10 mM MgCl2, 1 mM dithiothreitol [DTT], 10 μCi [γ-32P]ATP and 0.3 mM Cross tide (GRPTSSFAEG; Upstate, Lake Placid, NY) as a substrate. The reaction was stopped with stop solution (1 mM ATP, 1% bovine serum albumin, and 0.6% HCl), and the reaction mix was applied to 2.1 cm p81 Whatman filter paper. After drying at room temperature, the filters were washed four times with 25 mM phosphoric acid and once with acetone and were then counted in a scintillation counter. For the latter nonradioisotope TR-FRET method, the assay was performed as previously reported in 384-well plates according to the manufacturer’s recommendation (Zhang et al., 2006). The same amount of eluted protein was incubated with ULight-PLK(Ser137) peptide (PerkinElmer, Waltham, MA) for 60 min at room temperature in kinase assay buffer (50 mM HEPES, pH 7.5, 1 mM MgCl2, 2 mM DTT, and 0.01% Tween-20). The reaction was then stopped by addition of stop solution (detection

FIGURE 8: SGK3-mediated activation of NHE3 is dependent on PI3K and PDK1. (A) PS120/NHE3V/HA-SGK3 cells were pretreated with 1 μM RU or 20 μM LY prior to treatment with 1 μM Dex or carrier for 15 min. SGK3 protein was purified by immunoprecipitation, and the kinase activity was determined by TR-FRET. n = 3. (B) PS120/NHE3V/HA-SGK3 cells were stably transfected with sh-PDK1 or pLKO.1, and the expression level of PDK1 was determined by Western blotting using an anti-PDK1 antibody. (C) PS120/NHE3V/HA-SGK3 cells transfected with sh-PDK1 or pLKO.1 were treated with 1 μM Dex or carrier for 15 min. SGK3 protein was purified, and kinase activity was determined. n = 3. (D) NHE3 activities were determined in PS120/NHE3V/HA-SGK3 cells pretreated with RU or LY prior to treatment with Dex. (E) NHE3 activities in PS120/NHE3V/HA-SGK3 cells transfected with sh-PDK1 or pLKO.1 are shown. (A–E) n.s., not significant. *, p < 0.01 and **, p < 0.05, compared with respective controls. n = 6. (F) PS120/NHE3V cells expressing HA-SGK3 were serum-starved and treated with Dex or carrier for 15 min. Colocalization of SGK3 (green) and phospho-PDK1(p-PDK1, red) was determined using anti-HA and anti-p–PDK1 antibodies, respectively. (G) The graph represents Pearson’s coefficient of SGK3 and p-PDK1 colocalization from 10 independent fields of cells. *, p < 0.01 compared with the control. (H) Colocalization of GFP-Rab11 (green) and p-PDK1 (red) in PS120/NHE3V cells treated with or without Dex for 15 min is shown. Scale bar: 10 μm. Insets show high magnification of merged signals. (I) Pearson’s coefficient of GFP-Rab11 and p-PDK1 colocalization from 10 independent fields of cells. *, p < 0.01 compared with the control.
buffer containing 40 mM EDTA) for 5 min, which was followed by application of detection mix (2 nM Europium-anti-phospho-PLK(Ser137) [PerkinElmer, Winooski, VT] in detection buffer) and incubation for 60 min at room temperature. The signal was determined with BioTek plate reader in a TR-FRET mode (emission at 665 nm, excitation at 320 nm) and recorded for calibration of enzyme activity.

Surface biotinylation

Surface biotinylation of NHE3 was performed as previously described (He et al., 2010). Briefly, PS120/NHE3V cells were serum-starved for 4 h and Caco-2bbe/hNHE3V cells for 24 h prior to treatment with 1 μM Dex for indicated time periods (Figure 5A). Cells were rinsed twice in PBS and incubated for 10 min in borate buffer (154 mM NaCl, 7.2 mM KCl, 1.8 mM CaCl₂, and 10 mM H₃BO₃, pH 9.0). Cells were then incubated for 40 min with 0.5 mg/ml NHS-SS-biotin (Pierce) in borate buffer. Unbound NHS-SS-biotin was quenched with Tris buffer (20 mM Tris, 120 mM NaCl, pH 7.4). Cells were then rinsed with PBS, scraped, lysed in the lysis buffer described above, and sonicated twice for 15 s. The lysate was agitated for 30 min and spun at 14,000 × g for 30 min to remove the insoluble cell debris. An aliquot was retained as the total fraction representing the total cellular NHE3. Protein concentration was determined, and 1 mg of lysate was then incubated with streptavidin-agarose beads (Pierce) for 2 h. The streptavidin-agarose beads were washed three times in lysis buffer and twice in PBS. All the above procedures were performed at
inverted microscope and were superfused with NH₄⁺ buffer (40 mM NH₄Cl, 90 mM NaCl, 20 mM HEPES, 5 mM KCl, 1 mM TMA-PO₄, 2 mM CaCl₂, 1 mM MgSO₄, and 25 mM glucose) and subsequently with TMA⁺ buffer (130 mM TMA-Cl, 20 mM HEPES, 5 mM KCl, 1 mM TMA-PO₄, 2 mM CaCl₂, 1 mM MgSO₄, and 25 mM glucose). Na⁺ supplement provided with 50 μM Hoe-694 (NHE1 and NHE2 inhibitor) or 10 μM S-2226 (NHE3 inhibitor) was then reintroduced to drive Na⁺-dependent pH recovery. NHE3 activity in cultured cells was likewise determined. PS120 fibroblasts and Caco-2bbe cells seeded on coverslips were grown to 80–90% confluence and 5–7 d postconfluence, respectively. PS120 and Caco-2bbe cells were serum-starved for 4 h and overnight, respectively, prior to Dex treatment and NHE3 activity measurement. The rate of Na⁺-dependent pH recovery was calculated by determining slopes along the early stage of pH recovery by linear least-squares analysis over a minimum of 9 s. Calibration of the fluorescence signal was performed using the K⁺/H⁺ ionophore nigerin as previously described (Lamprecht et al., 1998). Comparisons of Na⁺/H⁺ exchange were made between measurements made on the same day. The microfluorometry was performed on a Nikon (Melville, NY) TE200 inverted microscope with a Nikon CFI Super Fluor 40× objective, coupled to a Lambda 10-2 filter wheel controller equipped with a multiwavelength filter set designed for BCECF-AM. Photometric data were acquired using MetaFluor software (Molecular Devices, Sunnyvale, CA).

**Confocal immunofluorescence**

PS120 cells grown on coverslips and Caco-2bbe cells grown on Transwell filters were fixed and permeabilized as described previously (He et al., 2008). For labeling of glycoprotein as a marker of apical membrane, Caco-2bbe cells were subjected to 10-min incubation with Alexa Fluor 647-conjugated WGA (Invitrogen) prior to fixation. Cells were then stained with a rabbit monoclonal anti-HA antibody, a rabbit or mouse anti-FLAG antibody, a mouse monoclonal anti-VSVG antibody, a rabbit monoclonal anti–p-PDK1 antibody, or a mouse monoclonal anti-Rab5 antibody for 1 h at room temperature. Following three washes, 10 min each, with PBS, the cells were incubated with Alexa Fluor 488-conjugated goat anti–rabbit immunoglobulin G (IgG), Alexa Fluor 546-conjugated goat anti–rabbit IgG, Alexa Fluor 555-conjugated goat anti–mouse IgG, or Alexa Fluor 647–conjugated donkey anti–mouse IgG (Invitrogen) for 1 h at room temperature. After three 10-min washes with PBS, the specimens were mounted with ProLong Gold Antifade Reagent (Invitrogen) and observed under a Zeiss LSM510 laser confocal microscope (Zeiss Microimaging, Thornwood, NY) coupled to a Zeiss Axioplan2e with 60× Plan-Apochromat oil lenses. For quantitative evaluation of the colocalization of green (SGK3 or GFP-Rab11) and red (p-PDK1) fluorescence signals, 10 random fields of cells were selected. Pearson’s correlation coefficient was determined for each cell using MetaMorph (Molecular Devices).

**Statistical analysis**

Results are presented as the mean ± SE. Statistical significance was assessed by Student’s t test using the SPSS statistical program (SPSS, IBM, New York, NY). p < 0.05 is considered significant.

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