Osmotic Stress-induced Phosphorylation of H2AX by Polo-like Kinase 3 Affects Cell Cycle Progression in Human Corneal Epithelial Cells*

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Background: To study Plk3-activated γ H2AX affecting hyperosmotic stress-induced cell fate. **Results:** Plk3 directly catalyzed γ H2AX in hyperosmotic stress-induced cells, resulting in an accumulation of G₂/M phase, altered population in the G₁ and S phases, and increased apoptosis.

Conclusion: Hyperosmotic stress-activated Plk3 elicited γ H2AX, which regulates cell cycle progression and cell fate. **Significance:** Plk3-mediated γ H2AX in stress-induced cells plays important roles to determine cell growth and death.

Increased concentrations of extracellular solutes affect cell function and fate by stimulating cellular responses, such as evoking MAPK cascades, altering cell cycle progression, and causing apoptosis. Our study results here demonstrate that hyperosmotic stress induced H2AX phosphorylation (γ H2AX) by an unrevealed kinase cascade involving polo-like kinase 3 (Plk3) in human corneal epithelial (HCE) cells. We found that hyperosmotic stress induced DNA-double strand breaks and increased γ H2AX in HCE cells. Phosphorylation of H2AX at serine 139 was catalyzed by hyperosmotic stress-induced activation of Plk3. Plk3 directly interacted with H2AX and was colocalized with γ H2AX in the nuclei of hyperosmotic stress-induced cells. Suppression of Plk3 activity by overexpression of a kinase-silencing mutant or by knocking down Plk3 mRNA effectively reduced γ H2AX in hyperosmotic stress-induced cells. This was consistent with results that show yH2AX was markedly suppressed in the Plk3^{-/-} knock-out mouse corneal epithelial layer in response to hyperosmotic stimulation. The effect of hyperosmotic stress-activated Plk3 and increased yH2AX in cell cycle progression showed an accumulation of G₂/M phase, altered population in G₁ and S phases, and increased apoptosis. Our results for the first time reveal that hyperosmotic stressactivated Plk3 elicited yH2AX. This Plk3-mediated activation of yH2AX subsequently regulates the cell cycle progression and cell fate.

Changes of the extracellular osmotic pressure are often found in some pathologic conditions, such as diabetes mellitus, uremia, dehydration after exercise, heat shock, fatal burns, and inflammation sites in various tissues including the cornea (1-4). It has been shown that persistent hyperosmotic stress can induce DNA damage, cell cycle arrest, and apoptosis (5, 6). Hyperosmotic stress extracts water out of the cell to induce cell



shrinkage. To restore the volume, cells undergo a regulated volume increase (RVI) process within several minutes by uptake of inorganic ions and water (7–9). Cell shrinkage and increased ionic strength alternate cell architecture compartments, denature proteins, and disturb cell function (5, 10, 11). Hyperosmotic stress is one of the important stimuli for cells on the surface of the body. For example, the corneal epithelial layer in the front of the eye forms the first defense line to protect injuries of eye structures behind the cornea from insults of environmental hazards including hyperosmotic stresses (12). Hyperosmotic stresses from environmental and pathological conditions strike the corneal epithelial cells and alter the fluid balance resulting in cell shrinkage, which may attribute to delayed wound healing and dry eye diseases (13).

Hyperosmotic stress as one of the extracellular stimuli induces a series of changes in intracellular kinase cascades that include activation of JNK, p38, and Polo-like kinase 3 (Plk3)² signaling pathways to activate the important AP-1 (activating protein-1) transcription factor complex (11, 14-18). Particularly, Plk3, one of the four members in the Polo like kinase family, plays a significant role in hyperosmotic stress-induced signaling cascades in addition to the JNK signaling pathway to transmit the stress signals in corneal epithelial cells. As a protein kinase, it catalyzes downstream phospho-proteins at serine/threonine sites. There are two functional domains in the Plk3 protein including a kinase domain (KD) at the N terminus and a Polo-box domain (PBD) at the C terminus that is able to interact with interactive proteins (19, 20). Following cell cycle progression, Plk3 undergoes substantial alterations in abundance, kinase activity, and subcellular distribution. As a multifunctional protein, active Plk3 induced by hyperosmotic stress plays an important function in the formation of c-Jun and ATF-2 heterodimers and homodimers in the AP-1 complex (21-25). Plk3 also involves the pathways of IR/UV irradiation-

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² The abbreviations used are: Plk, polo-like kinase; ROS, reactive oxygen species; MMS, methylmethane sulfonate; ATM, mutated in ataxia telangiectasia; ATR, ATM-related; DSB, double strand break; SCGE, single cell gel electrophoresis assay; HCE, human corneal epithelial; MEF, mouse embryonic fibroblast.

induced DNA damage by regulating p53 activity and cell cycle progress in G₂/M phase (26, 27). Plk3 activity is rapidly increased upon stress stimulation by ionizing radiation (IR), reactive oxygen species (ROS), methylmethane sulfonate (MMS), UV irradiation, and hypoxia (17, 18, 28–31). Plk3 plays roles in the pathway of DNA damage-dependent activation of p53 downstream from both ATM (mutated in ataxia telangiectasia)/ATR (ATM-related), in the family of phosphatidylinositol 3-kinase-related kinases, and Chk2 (32, 33).

Histone H2AX is one of the nucleosome core histone H2A variants and has been considered as a marker that indicates the occurrence of DNA double strand breaks (DSBs) in mammalian cells (34, 35). One of the early events in IR/UV irradiationinduced DNA DSBs is ATM/ATR-dependent phosphorylation of the histone variant H2AX at the C terminus serine 139 subsequently termed yH2AX (36, 37). In DNA DSBs cells, DNA damage response (DDR)-induced and ATM/ATR-catalyzed phosphorylation of H2AX interacts with checkpoint- and DNA repair-associated proteins such as Rad50, Rad51, and Brca1 as well as the p53 binding protein 1 (53BP1), resulting in cell cycle arrest (38-41). The questions remaining in hyperosmotic stress-induced cells are whether there is a phosphorylation of H2AX and whether extracellular stress-sensitive Plk3 interacts with H2AX in the ATM-Chk2 pathway to regulate cell fate. In the present study, we demonstrate important findings to show that hyperosmotic stress-induced DNA DSBs in corneal epithelial cells increased phosphorylation of H2AX, dependent on activation of Plk3. As an alternative signaling pathway, Plk3 served as a transducer to amplify the hyperosmotic stress-induced cellular response by directly interacting with H2AX at the protein level and to catalyze phosphorylation of H2AX at the position of serine 139, resulting in alteration of cell fate.

EXPERIMENTAL PROCEDURES

Cell Culture and Treatments

Primary human corneal epithelial (PHCE) cells and human telomerase immortalized corneal epithelial (HTCE) cell were cultured in a serum-free keratinocyte medium (Defined Keratinocyte-SFM; Invitrogen, Carlsbad, CA). The human corneal epithelial (HCE) cell line was grown in Dulbecco's modified Eagle's medium/F-12 (1:1) containing 10% fetal bovine serum and 5 μ g/ml insulin. Mouse embryonic fibroblasts (MEF) were isolated for wild-type and Plk3^{-/-} mouse embryos. Primary MEFs were derived from embryonic day 14.5 embryos with the respective genotype of both wild-type and PLK3^{-/-} mice (42). The MEF cells were cultured in high glucose DMEM supplemented with 10% FBS and antibiotics (100 μ g of penicillin and 50 μ g of streptomycin sulfate per ml). In all of the MEF experiments, primary MEFs were restricted only in third to fifth passages. The culture conditions for corneal epithelial cells and MEFs were in an incubator supplied with 95% air and 5% CO₂ at 37 °C. The medium was replaced every 2 days, and cells were subcultured by treatment with 0.05% trypsin-EDTA. For experiments of hyperosmotic stress stimulation, various concentrations of sorbitol, sucrose, NaCl, and glucose were added to the culture medium following a time course. Osmotic pressures of hyperosmotic media were measured using a Vapor Pressure Osmometer (VAPRO5520).

Cell Cycle Analysis

Cell cycle analysis was performed using a flow cytometry (BD LSRII, BD Biosciences, San Jose, CA). HCE cells and MEFs were treated with the control condition and hyperosmotic stresses following a time course, and then the attached cells were trypsinized and fixed with 70% ethanol and 50 mM glycine. The cells were resuspended in PBS containing RNase A (100 ng/ml) and propidium iodide (PI, 25 ng/ml). Cell populations in different phases were mapped with BD FAC Diva Software V6.11, and cell cycle progression was analyzed with MedFit LTTM V3.1 (Verity Software House). Statistical analysis was performed as described in the statistical section below.

Comet Assay

Hyperosmotic stress-induced DNA damage in the individual HCE cell was detected by the comet assay or the single cell gel electrophoresis assay (SCGE). The classic "comet tail" shapes resulting from damaged cellular DNA (containing fragments and strand breaks) were separated from intact DNA in the electrophoretic field. The assay was performed according to the manufacturer's instructions (Cell Biolabs, San Diego, CA). Briefly, cells were trypsinized to detach from culture dishes and washed once with PBS. Cells were loaded onto a low meltingpoint agarose gel and spread on the OxiSelect[™] Comet Slide. Embedded cells were treated for 30 min at 4 °C in the dark with a lysis buffer and then alkaline solution, which relaxes and denatures the DNA. Electrophoresis was performed for 15-30 min in an electric field with 1 volt/cm. After rinsing the slide three times in distilled water and once in 70% ethanol, DNA was stained with Vista Green DNA Dye. Preparations were photographed using a Nikon T*i* fluorescent microscope.

Gene Transfection and Recombinant Proteins

Human corneal epithelial cells were transfected with Plk3 wild type and kinase-defective Plk3^{K52R} mutant (a full-length Plk3 that contains a mutation to substitute the lysine 52 with an arginine) using Lipofectamine reagents (Invitrogen). Transfected cells were subjected to Western analysis and immunocomplex kinase assays. Transfections of Plk3-specific siRNA (Qiagen, SI02223473 and SI02223466) were done by adding Plk3-specific siRNAs with a final concentration of 25 nm mixed with 12 μ l of HiPerFect in 100 μ l of serum-free culture medium. The mixtures were incubated for 20 min at room temperature. The mixture was evenly added into culture cells. Transfected cells were cultured under normal growth conditions for 48-84 h before performing experiments. Non-silencing siRNA-transfected cells were used as the controls with the same transfection method. In addition, human H2AX fulllength cDNA in a pCR2.1-TOPO plasmid was subcloned into the EcoRI cloning sites in vector pFlag-CMV-4 (Sigma). H2AX^{S139A} mutant was generated by site-directed mutagenesis using the QuikChange Lightning Site-directed Mutagenesis Kit (Agilent Technologies, Inc.), and the mutant sequence was confirmed by DNA sequencing. The fusion protein of GST-H2AX^{wt} and GST-H2AX^{S139A} was produced by cloning the



wild type H2AX and H2AX^{S139A} mutant into EcoRI sites within the bacterial expression vector pGEX-4T-3. Purification of GST-H2AX^{wt} and GST-H2AX^{S139A} was performed under standard conditions. Briefly, *Sf*9 cells (ATCC) infected with H2AX baculovirus were cultured in Grace's insect cell culture medium. Infected cells were harvested on day 3 and lysed in a lysis buffer (50 mM NaH₂PO₄, 300 mM NaCl, 1% Nonidet P-40 20 mM imidazole, 1 mM PMSF, 2 μ M pepstatin A, 10 units/ml aprotinin). Cell lysates were incubated with Ni-NTA agarose resins for 3 h at 4 °C. Fusion proteins were eluted from Ni-NTA resins using the lysis buffer containing 200 mM imidazole after extensive wash of the resins with lysis buffer. Fusion proteins were purified by dialyzing in a storage buffer (25 mM Tris, pH 7.4, 5 mM EGTA, 2 mM DTT, 0.1% Triton X-100, and 50% glycerol) and stored at 80 °C for subsequent uses.

Immunocytochemistry

Immunostaining Experiments—corneal epithelial cells were grown on glass slides and hyperosmotic sorbitol solutions were used to treat HCE cells. Mouse corneal sections and HCE cells were fixed for 15 min in 4% paraformaldehyde, and then permeabilized with PBS-0.2% Triton X-100 (PBS-T) for 30 min at room temperature. The cells were blocked by incubation with 10% normal horse serum (NHS) or 10% normal goat serum in PBS-T for 1 h at room temperature, followed by double immunostaining with the corresponding antibodies. Corneal tissue and HCE cell slices were washed with PBS and stained with DAPI. A Nikon fluorescent Ti microscope was used to capture stained tissue imaging. Imaging data were analyzed using a Nikon NIS Element Software program.

Immunoprecipitation and Immunocomplex Kinase Assay-Corneal epithelial cells (5 \times 10⁷) were rinsed with PBS and incubated in 1 ml of lysis buffer (20 mM Tris, pH 7.5, 137 mM NaCl, 1.5 mM MgCl₂, 2 mM EDTA, 10 mM sodium pyrophosphate, 25 mM glycerophosphate, 10% glycerol, 1% Triton X-100, 1 mM sodium vanadate, 1 mM phenylmethylsulfonyl fluoride, 250 μ M 4NPP, 10 μ g/ml leupeptin, and 10 μ g/ml aprotinin) on ice for 30 min. The cell lysates were spun at 13,000 \times g for 10 min at 4 °C and incubated at 4 °C overnight with antibodies against Plk3 and yH2AX, respectively. The immunocomplexes were recovered by incubation with 50 μ l of 10% protein A/G-Sepharose (Santa Cruz Biotechnology). The immunocomplex beads were rinsed twice with lysis buffer and once with kinase buffer, and then subject to immunoblotting and kinase assay. The effect of active Plk3 on catalyzing H2AX phosphorylation was measured using immunocomplex kinase assays by incubation of immunoprecipitated Plk3 with H2AX fusion protein in 30 μ l of kinase buffer (20 mm HEPES, pH 7.6, 5 mm MgCl₂, 10 µM MnCl₂, 25 mM glycerophosphate, 1 mM sodium orthovanadate, 2 mM dithiothreitol, 20 μ M ATP, and, 10 μ Ci of $[\gamma^{-32}P]ATP$ for 30 min at 37 °C. Kinase reactions were terminated by adding an equal volume of $2 \times$ Laemmli buffer and boiling for 5 min. Equal volumes of the samples were displayed on 10-15% SDS-PAGE and visualized by exposure on x-ray films. For cold kinase assay, $[\gamma^{-32}P]ATP$ was omitted from the kinase buffer. Western blot assay was performed by harvesting 5×10^5 cells in 0.5 ml of lysis buffer. Samples were loaded onto 10-15% SDS-PAGE gels and fractionated by electrophoresis.

Plk3 Mediates Hyperosmotic Stress-elicited γH2AX

Proteins in the gel were electro-transferred to a PVDF membrane. The membrane was exposed to blocking buffer (5% non-fat milk in TBS-0.1% Tween 20 (TBS-T) for 1 h at room temperature, and then incubated with respective antibodies overnight at 4 °C. After three washes with TBS-T buffer, the membrane was incubated with horseradish peroxidase (HRP)-linked secondary antibody for 1 h at room temperature. Expression of proteins was detected with a Western blot detection kit (Santa Cruz Biotechnology).

RESULTS

Effects of Hyperosmotic Stresses on DSBs and H2AX Phosphorylation/Activation-In previous studies, we found that Plk3 activity is one of the major regulators to mediate cellular responses to various stresses including UV irradiation, hypoxic stimulation, and hyperosmotic pressure. The effect of hyperosmotic stress-induced Plk3 activation on downstream events involves increases in phosphorylation and formation of c-Jun and ATF-2 hetero/homodimers in the AP-1 complex. In corneal epithelial cells, hyperosmotic stress-induced H2AX phosphorylation was studied by treating cells with increased extracellular osmotic pressures that were created by adding high concentrations of sorbitol of up to 1,000 mM. Phosphorylation levels of γ H2AX were markedly increased at a sorbitol concentration of 250 mM sorbitol in the culture medium in three different origins of human corneal epithelial HCE, PHCE, and HTCE cells (Fig. 1A). In addition, four different solutes of sorbitol, glucose, sucrose, and NaCl at concentrations of 300 mM were applied to cell cultures to verify the hyperosmotic pressure effects on the phosphorylation levels of H2AX in HCE cells (Fig. 1B). Phosphorylation of H2AX was increased by 250 mM sorbitol within 15 min and reached the peak level at 60 min (Fig. 1C). Hyperosmotic stress-induced increases in H2AX phosphorylation were also demonstrated in the nuclei by immunostaining in sorbitol-treated cells (Fig. 1D). It is known that H2AX has been considered as a hallmark for DNA DSBs in mammalian cells. To verify the effect of hyperosmotic stress on DNA damage response in corneal epithelial cells, cells were treated with sorbitol and the antitumor drug etoposide that forms a DNA and topoisomerase II enzyme complex resulting in DNA DSBs (43). DNA DSBs were detected using the Comet assay to compare the effects of hyperosmotic sorbitol (250 mm) and etoposide (20 μ g/ml for 4 h) (Fig. 1*E*). The results of these experiments suggest that hyperosmotic stresses significantly increased H2AX phosphorylation and DNA DSBs in corneal epithelial cells.

Effect of Hyperosmotic Stress-induced Active Plk3 on H2AX Phosphorylation—One of the important questions in the present study is to determine whether hyperosmotic stress-induced Plk3 activity is associated with H2AX phosphorylation. To detect interactions between Plk3 and H2AX, H2AX fusion proteins were purified and used as substrates in immunoprecipitation kinase assays to test the effect of hyperosmotic stressinduced Plk3 kinase activity on H2AX phosphorylation. Hyperosmotic stress-induced active Plk3 was immunoprecipitated and used to catalyze phosphorylation of the H2AX fusion proteins. Stress stimuli were applied by treating cells with UV irradiation, 250 mM sorbitol, and adding 300 mM NaCl to corneal epithelial



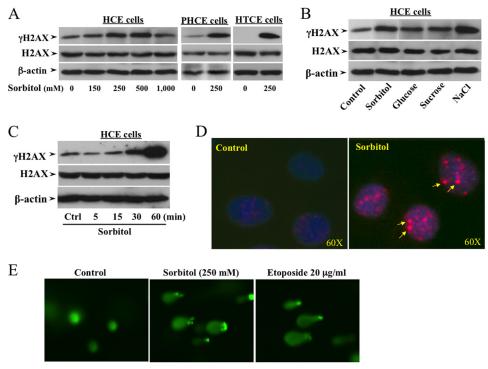


FIGURE 1. Effects of altered osmotic stresses on DNA DSBs and H2AX phosphorylation. *A*, increases in γ H2AX following increased sorbitol concentrations in various human corneal epithelial cells. *B*, effects of high concentrations of extracellular solutes: sorbitol (300 mM), glucose (300 mM), sucrose (300 mM), and NaCl (300 mM) on increases in γ H2AX in HCE cells. *C*, high concentration sorbitol-induced γ H2AX following a time course in HCE cells. *D*, localization of hyperosmotic stress-induced γ H2AX in nuclei of HCE cells. *E*, hyperosmotic stress-induced DNA DSBs in HCE cells. DNA DSBs were detected with Comet assays in sorbitol (250 mM)-treated cells, and the anti-tumor drug etoposide (20 μ g/ml)-treated cells served as positive controls.

cells. Plk3 that was immunoprecipitated from stress-induced cells phosphorylated the wild-type H2AX fusion protein at Ser-139 recognized by a γ H2AX Ser-139 phosphorylation-specific antibody (Fig. 2A). Hyperosmotic stress-induced Plk3 activation and H2AX phosphorylation occurred within 15 min and reached the peak effect at 60 min following a time course (Fig. 2B). A mutant of Flag-H2AX^{S139A} fusion proteins made by replacing serine139 with an alanine residue was subject to immunoprecipitation kinase assay for detecting hyperosmotic stress (sorbitol 250 and 500 mM)-induced difference in Plk3 activity to affect the phosphorylation level of serine 139 (Fig. 2C). Plk3 was also detected as a loading control. In addition, a constitutively active Plk3 was found to be able to catalyze phosphorylation of the H2AX^{wt} fusion protein, but it failed to induce phosphorylation of the Flag-H2AX^{S139A} mutant (Fig. 2D). These results indicate that active Plk3 in hyperosmotic stress-induced corneal epithelial cells is able to phosphorylate H2AX at serine 139 (S139).

Interactions of Plk3 and γ H2AX in Hyperosmotic Stress-induced Cells—As shown in Fig. 2, Plk3 activated in hyperosmotic stress-stimulated HCE cells was able to directly phosphorylate H2AX fusion proteins. We performed further experiments to substantiate the evidence that Plk3 indeed mediates the phosphorylation process of γ H2AX by demonstrating that there are interactions between Plk3 and H2AX proteins in hyperosmotic stress-induced cells. A bidirectional interaction between Plk3 and H2AX was detected by immunocoprecipitation and Western analysis. An anti-phospho- γ H2AX antibody was used to pull down Plk3 protein, and in turn, the other anti-Plk3 antibody was applied to pull down phospho- γ H2AX in hyperos-

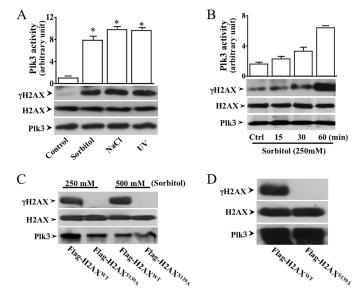


FIGURE 2. **Hyperosmotic stress-activated Plk3 phosphorylated H2AX.** *A*, phosphorylation of H2AX by activated Plk3 from HCE cells stimulated with hyperosmotic stress and UV irradiation in HCE cells. *B*, phosphorylation of H2AX fusion protein by hyperosmotic stress-activated Plk3 following a time course in HCE cells. *C*, hyperosmotic stress-activated Plk3 failed to phosphorylate the H2AX mutant (H2AX^{5139A}) in which Ser-139 was substituted by an alanine residue. The Plk3 level was detected by Western blot and served as the loading control in HCE cells. *D*, effect of constitutively activated Plk3 upon phosphorylation of wild type H2AX and H2AX^{5139A} mutant fusion proteins in HCE cells. The asterisk represents a significant difference between untreated control and hyperosmotic stress-induced HCE cells (p < 0.05, n = 3).

motic stress-induced corneal epithelial cells (Fig. 3, A and B). Activations of Plk3 and γ H2AX were found and colocalized in the nuclei of hyperosmotic stress-induced corneal epithelial

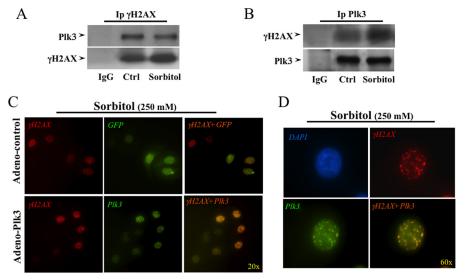


FIGURE 3. **Determining interaction between Plk3 and H2AX in hyperosmotic stress-induced HCE cells.** *A*, detection of Plk3 and H2AX interaction by immunoprecipitation using an anti-phospho-H2AX (anti- γ H2AX) antibody. Plk3 was detected by Western analysis. *B*, detection of Plk3 and H2AX interaction by immunoprecipitation using an anti-Plk3 antibody. γ H2AX was detected by Western analysis using an anti- γ H2AX antibody. *C*, immunocolocalization of Plk3 and γ H2AX in the nucleus of hyperosmotic stress-induced cells. *D*, high amplification view (60×) to show colocalization of Plk3 and γ H2AX in the nucleus of hyperosmotic stress-induced cells. *D*, high amplification view (60×) to show colocalization of Plk3 and γ H2AX in the nucleus of hyperosmotic stress-induced HCE cells. Immunocolocalization experiments were done by immunostaining of hyperosmotic stress-induced to γ H2AX. Photo images were taken using a Nikon fluorescent microscope at 20× and 60×, respectively. Data were analyzed with a Nikon software program.

cells detected by immunostaining and microscopy imaging. Both Plk3 and γ H2AX were activated to form immune active clusters within 45 min upon 250 mM sorbitol stimulation, and both active Plk3 and γ H2AX were colocalized in cell nuclei that were indicated by DAPI nuclear staining under a Nikon fluorescent microscope (20×) (Fig. 3*C*). In addition, the hyperosmotic stress-induced interaction between Plk3 and γ H2AX in the nuclei was more clearly demonstrated by increasing the microscopic power to a higher magnification (60×) in immunostaining experiments (Fig. 3*D*). In control immunostaining experiments using Plk3 knock-down cells, Plk3 expression was not detected (data not shown). Results of immunocoprecipitation and immunocolocalization indicate that there is indeed a protein-protein interaction between Plk3 and γ H2AX upon hyperosmotic stress stimulation in corneal epithelial cells.

Effect of Altered Plk3 Activity on Hyperosmotic Stress-induced $\gamma H2AX$ —To study functional interaction of Plk3 and γ H2AX in response to hyperosmotic stress, Plk3 activities in corneal epithelial cells and mouse cornea were altered by different approaches including: 1) transfection of a kinase-defective Plk3^{K52R} mutant; 2) Plk3 mRNA knock-down with siRNA; and 3) Plk3 knock-out (Plk3^{-/-} transgenic mice). In corneal epithelial cells, transfection of the Plk3^{K52R} mutant resulted in a reduced hyperosmotic stress-induced phosphorylation of H2AX compared with cells transfected with the Plk3 wild type (Plk3^{WT}) construct (Fig. 4A). In Plk3-specific siRNAtransfected corneal epithelial cells, partial knock-down of Plk3 effectively minimized hyperosmotic stress-induced phosphorylation of H2AX at 25 and 50 min (Fig. 4B). Further studies were conducted in mouse corneas obtained from both wild type and Plk3 knock-out (Plk $3^{-/-}$) mice. In the cornea of wild type mice, localization of yH2AX in the nuclei of hyperosmotic stressinduced basal epithelial layer was detected using immunostaining and microscopy imaging. Active yH2AX was observed

to form immune active clusters within 45 min upon hyperosmotic sorbitol stimulation, and yH2AX was colocalized in cell nuclei with DAPI nuclear staining (Fig. 4C). However, there was much less yH2AX found in the nuclei of hyperosmotic stressinduced corneas obtained from Plk3^{-/-} mice. Our results clearly indicate that alterations of Plk3 activity in corneal epithelial cells and the epithelial layer of the mouse cornea markedly interrupted the functional interaction between Plk3 and yH2AX in the nuclei in response to hyperosmotic stress stimulation. It was reported that H2AX is phosphorylated by ATM/ ATR that is activated in response to DNA damage, respectively. To study the possible involvement of Plk3 in these pathways, DNA damage-induced activation of ATM/ATR was blocked with 5 to 10 mM caffeine. Blockade of ATM/ATR by caffeine suppressed hyperosmotic stress-induced H2AX phosphorylation and Plk3 activity (Fig. 4, D and E). The relevance of Plk3 in hyperosmotic stress-induced H2AX phosphorylation was further characterized by inhibition of the p38 kinase. Hyperosmotic stress-induced H2AX phosphorylation in the absence or presence of SB202190 (an inhibitor of p38) was also measured in corneal epithelial cells. Inhibition of p38 had no effect on either hyperosmotic stress-induced H2AX phosphorylation or Plk3 activity (Fig. 4, F and G). To measure Plk3 activity without interference by JNK and p38, an immunodepletion (ID) procedure was used to remove JNK and p38 in cell lysates. After the ID procedures, JNK and p38 activities in lysates were markedly diminished to non-detectible levels (data not shown).

Effects of Hyperosmotic Stress-induced Plk3 Activity on Cell Cycle Progression—It has also been shown that Plk3 acts downstream of both ATM and Chk2 in the DNA damage-dependent signaling pathway to regulate progression of the cell cycle. In the present study, the effect of Plk3 activity on mediating hyperosmotic stress-induced changes in cell cycle progression was analyzed by flow cytometry. In hyperosmotic stress-induced



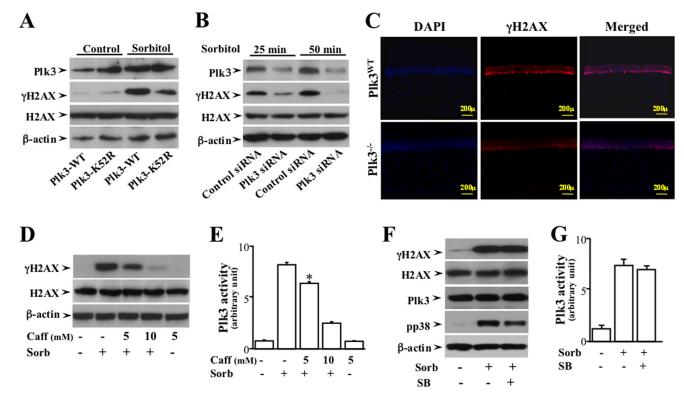


FIGURE 4. **Effects of altered Plk3 activity on hyperosmotic stress-induced** γ **H2AX in HCE cells.** *A*, effect of overexpressing wild type Plk3 and kinasesilencing Plk3^{k52R} mutant on hyperosmotic stress-induced γ H2AX in HCE cells. *B*, effect of knocking down Plk3 mRNAs on hyperosmotic stress-induced Plk3 γ H2AX in HCE cells. *C*, suppression of hyperosmotic stress-induced γ H2AX in the epithelial layer of the Plk3-deficient (Plk3^{-/-}) mouse corneas. Immunostaining experiments were performed to detect hyperosmotic stress-induced activation of γ H2AX using a γ H2AX specific antibody, and cell nuclei were stained by DAPI in corneas of wild type and Plk3^{-/-} knock-out mice. Photo images were taken using a Nikon fluorescent microscope at 10×. *D*, effect of inhibiting ATM/ATR with Caff (caffeine) on hyperosmotic stress-induced H2AX phosphorylation. *E*, effect of inhibiting ATM/ATR with Caff on hyperosmotic stress-induced of ymASB (SB202190, an inhibitor of p38) on hyperosmotic stress-induced ATF-2 phosphorylation. *G*, effect of hyperosmotic stress on Plk3 activity. Plk3 activity was determined by immunocomplex kinase assay, and the H2AX fusion protein was used as the substrate. The asterisk symbol indicates a significant difference (p < 0.05, n = 3).

corneal epithelial cells, significantly accumulated population of the cells in the G_2/M phase, and apoptotic cells were detected following a 12 h time course (Fig. 5, A and B). In contrast, knock-down of Plk3 mRNA effectively blocked hyperosmotic stress-induced changes in the cell cycle distribution (Fig. 5C). Further study was conducted in mouse embryonic fibroblasts (MEFs) obtained from primary cultures of wild type and Plk3^{-/-} knock-out mice. Consistent with HCE cells, hyperosmotic stress-induced γ H2AX activity in Plk3^{-/-} MEFs was significantly suppressed compared with wild type MEFs (Fig. 5D). Cell cycle analysis of wild type MEFs demonstrated that hyperosmotic stress induced significant increases in late G_2/M phase populations in apoptotic cells. However, effects of hyperosmotic stress on the changes of the G_2/M phase and apoptosis were significantly suppressed in Plk3 knock-out Plk3^{-/-} MEFs (Fig. 5, *E* and *F*). The results of hyperosmotic stress-induced cell cycle alterations in Plk3 mRNA knock-down HCE cells and Plk3 knock-out Plk3^{-/-} MEFs were consistent, suggesting that the role of Plk3 in interacting with γ H2AX is associated with DNA replication in response to hyperosmotic stress-induced double-strand breaks in S- and G₂-phase cells.

DISCUSSION

The process of corneal epithelial wound healing can be delayed by various environmental conditions including hyperosmotic stimulation that can result in compromising physiological function of the cornea. It has been shown that cellular responses to hyperosmotic stress include activation of MAP kinase cascades including JNK and p38, and Plk3 signaling pathways to further downstream activation of transcription factors, such as c-Jun and ATF-2 (5, 13, 25, 44, 45). In the present study, we found that extracellular hyperosmotic stresses activated vH2AX phosphorylation as a result of the stress-induced DNA DSBs in corneal epithelial cells, which is consistent with previous reports that hyperosmotic stress can result in DNA damage responses in other types of mammalian cells (34, 46, 47). Interestingly, DNA DSB-induced yH2AX phosphorylation is mediated by Plk3-induced cascade in addition to both ATM and Chk2 in DNA damage-dependent signaling pathways. The effect of hyperosmotic stress on activation of Plk3 subsequently resulting in yH2AX phosphorylation is consistent with previous findings indicating that Plk3 is an important mediator of the cellular response to genotoxic stress in ATM and Chk2 signaling pathways. Furthermore, activity of DNA damage-activated ATM/ATR was inhibited by application of caffeine, an inhibitor of ATM/ATR (25, 48, 49). Inhibition of ATM/ATR by caffeine resulted in suppression of Plk3 activity and H2AX phosphorylation. In addition, inhibition of hyperosmotic stress-induced p38 activation by the SB inhibitor had no effects on H2AX phosphorylation and Plk3 activity (Fig. 4). These results provide further evidence that Plk3 is involved at

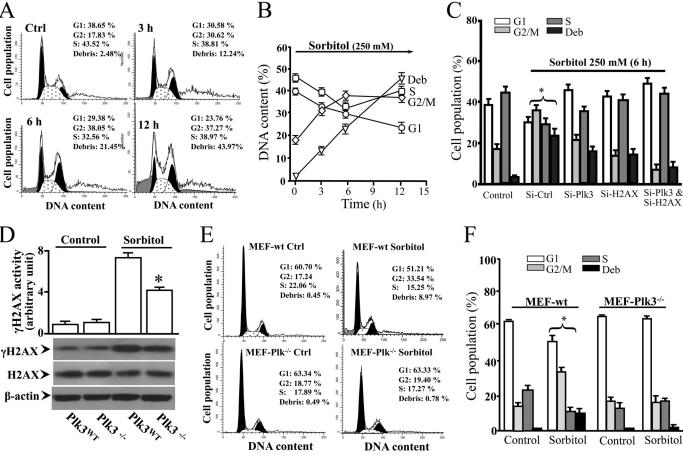


FIGURE 5. Effect of altered Plk3 activity on cell cycle distribution in hyperosmotic stress-induced HCE cells and MEFs. A, effect of hyperosmotic stimulation on cell cycle progression in HCE cells. B, time course of hyperosmotic stress-induced alteration in cell cycle progression in HCE cells. C, effects of knocking down Plk3 mRNA and/or H2AX mRNA on alterations of cell cycle progress in hyperosmotic stress-induced HCE cells. D, effect of hyperosmotic stress on vH2AX levels in MEFs obtained from wild type (Plk3^{wt}) and Plk3^{-/-} knock-out mice. E, effect of hyperosmotic stimulation on cell cycle progression in Plk3^{wt} and MEFs. F, comparisons of cell cycle distributions between wild type and Plk3^{-/-} knock-out MEFs in the absence or presence of hyperosmotic stimula-Plk3^{-/} tion. Asterisk symbols represent a significant difference between hyperosmotic stress-induced control cells and Plk3 and/or H2AX activities suppressed cells (p < 0.05, n = 3).

least partially in the ATM/ATR signaling pathway. The effect of hyperosmotic stress-activated Plk3 on H2AX was also examined by following a time course to show activation processes and the peak time of the Plk3 effect on γ H2AX phosphorylation. It has been shown that the ATM-catalyzed phosphorylation site of H2AX is at serine 139 (Ser-139), termed γ H2AX. The functional role of yH2AX is known to the response of DNA DSBs, and involves the control of cell cycle progression (50). The effect of Plk3 kinase on phosphorylation of H2AX at Ser-139 is supported by crystallographic structure of the nucleosome indicating that residue serine 139 in H2AX is easily accessible to kinases (51). In the present study, effects of hyperosmotic stress- and UV irradiation-induced Plk3 kinase activation on H2AX phosphorylation were observed at the Ser-139 of H2AX using an antibody specifically against Ser-139 in phosphorylated yH2AX. Further verification was done by demonstrating that the phosphorylation of H2AX (becoming γ H2AX) by hyperosmotic stress-induced active Plk3 can be abolished by introducing a mutation to the fusion protein of H2AX^{S139A} substituting Ser-139 with an Ala (alanine) residue (Fig. 2). Thus, we demonstrate for the first time that hyperosmotic stress-activated Plk3 is a functional player in an important event

upstream in the DNA DSB-induced pathway to interact with H2AX, resulting in γ H2AX activation.

Functional characterization of hyperosmotic stress-induced yH2AX phosphorylation through activation of Plk3 in the signal pathway was performed by determination of the interaction of Plk3 and yH2AX at protein levels using immunocoprecipitation and colocalization of Plk3 and yH2AX in stress-induced corneal epithelial cells. Results of immunocoprecipitation experiments show that antibodies against each of the Plk3 and yH2AX bi-directionally pulled down both Plk3 and yH2AX proteins, respectively. In addition, both Plk3 and yH2AX proteins can be colocalized in the nuclei of hyperosmotic stressinduced human corneal epithelial cells using immunostaining and fluorescent microscopy (Fig. 3). To further characterize the hyperosmotic stress-induced yH2AX phosphorylation through activation of Plk3 pathway, Plk3 activities were altered by either transfection of cDNA encoding a kinase-domain silencing Plk3 mutant (Plk3K52R) or knock-down of Plk3 mRNA in corneal epithelial cells, and by comparing γ H2AX activities in corneal epithelia of wild type and Plk3^{-/-} knock-out mice. The results of transfection experiments showed that both introduction of $\text{Plk3}^{\text{K52R}}$ and siRNA specific to Plk3 into corneal epithelial cells

А

Cell population

Cell population

D



markedly suppressed hyperosmotic stress-induced γ H2AX phosphorylation. There were clearly weakened γ H2AX in the epithelial layer of the Plk3^{-/-} mouse cornea when compared with the wild type mouse cornea (Fig. 4). These results provide further evidence that hyperosmotic stress-induced γ H2AX phosphorylation indeed, at least in part, are mediated by the Plk3 pathway.

It has been shown that increased γ H2AX is an important indicator for radiation stress-induced DNA DSBs in the cell cycle, especially critical in late S/G_2 phase. Our studies were conducted to investigate the effect of hyperosmotic stress-induced Plk3 activity on γ H2AX and cell cycle progression in human corneal epithelial cells and MEFs. We found that cell distribution in the cell cycle is very different in control human corneal epithelial and wild type MEFs in G₁ and S phases, which can be explained by different growth patterns of those cells under culture conditions. However, hyperosmotic stimulation resulted in significant alteration of the cell population in G₁ and S phases, and increased cell population in G_2/M phase and apoptotic stage in both cell types. Knocking down mRNAs of Plk3, H2AX, and both Plk3 plus H2AX significantly suppressed the effects of hyperosmotic stimulation on cell cycle redistribution. The most interesting observation is that knock-down and knock-out of Plk3 in human corneal epithelial cells and MEFs presented a consistent effect of protection on cell apoptosis in response to hyperosmotic stimulation (Fig. 5). In addition, hyperosmotic stress-induced alterations of cell cycle distributions in G_1 , G_2/M , and S phases were minimized by suppression of Plk3 and H2AX activities. These results provide important evidence suggesting that regulation of yH2AX by Plk3 in hyperosmotic stress-induced human corneal epithelial cells plays significant and functional roles in cell fate.

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