LH Stimulation of Meiotic Resumption in Ovarian Follicles by Protein Kinase A and a PPP-Family Phosphatase leremy R. Egbert¹, Ivan Silbern², Katie M. Lowther¹, Tracy F. Uliasz¹, Siu-Pok Yee¹, Henning Urlaub², Laurinda A. Jaffe¹ P2872 ¹Department of Cell Biology, UConn Health, Farmington, CT, USA egbert@uchc.edu ²Bioanalytical Mass Spectrometry Group, Max Planck Institute for Biophysical Chemistry, 37077 Göttingen, Germany

Abstract

In mammalian preovulatory ovarian follicles, meiotic arrest of fully grown oocytes is maintaned by cyclic guanosine monophosphate (cGMP) that is produced by the membrane guanylyl cyclase natriuretic peptdide receptor 2 (NPR2) in the follicle's somatic cells and diffuses through gap junctions into the oocyte¹. Phosphorylation of several juxtamembrane serines and threonines is essential for full NPR2 activity. The cyclic surge of luteinizing hormone (LH) acts on receptors in the outermost somatic cell layers to rapidly dephosphorylate and inactivate NPR2, lowering cGMP levels in the follicle and oocyte to trigger meiotic resumption^{2,3}. Our goals are to determine whether protein kinase A (PKA) signaling mediates this process⁴, and which LH-activiated PPPfamily phosphatase dephosphorylates NPR2². Application of the specific PKA inhibitor Rp-8-CPT cAMPS (Rp) to isolated mouse follicles prior to LH treatment (30 min) prevented NPR2 dephosphorylation, showing that the phosphatase activation is mediated by PKA. To identify PPP-family regulatory subunits that undergo rapid a LH-stimulated increase in phosphorylation, lysates of rat follicles treated with or without LH were enriched for phosphorylated peptides and analyzed by liquid chromatography-tandem mass spectrometry (LC-MS/MS) utilizing chemical labeling with tandem mass tags (TMT) for quantification. PPP1R12A and PPP2R5D emerged as two primary candidates. In PPP1R12A, peptides containing phosphorylated S507 residue had ~5-fold higher intensity in LH-treated follicles. PPP2R5D had four sites with elevated intensity with LH: S53 and S566 (10-fold and 4-fold, respectively), and S81/82 (~2-fold). Quantitative western blotting confirmed these results and showed that inhibition of PKA activity with Rp prevented the LH-induced phosphorylation changes. To determine whether phosphorylation of one or both proteins mediates NPR2 dephosphorylation after LH, we have made mice in which S507 of PPP1R12A or S53/S81/S82/S566 of PPP2R5D are replaced with alanines, and are testing their role in this process. Thus, we have identified two PPP-family regulatory subunits that are rapidly phosphorylated in response to LH-PKA signaling as candidates for effecting NPR2 dephosphorylation, an event required for timely oocyte meiotic resumption.

Graphical Background





Results



 16.20 ± 0.03

 16.42 ± 0.06

 19.94 ± 0.32

 16.45 ± 0.13

Amino

acid

53

507

566

Gene

name

PPP2R5D

PPP1R12A

PPP2R5D

PPP2R5A

Phosphopeptide intensity (mass spec)

(mean ± SEM)

Control (PBS) 300 nM oLH change

 19.49 ± 0.27

 18.57 ± 0.59

 1.90 ± 0.35

 7.92 ± 0.07

Fold

10.4

run on a Phos-tag gel and probed with anti-HA. (B) Summary of 3 experiments showing mean \pm SEM.

2a) Identification of candidate PPP-family subunit(s) that are phosphorylated in response to LH/PKA signaling.





and meiotic resumption are currently being tested.