

ARPP ‐16, protein kinase A and MAST3 kinase: a newly identified pathway for regulation of protein phosphatase PP2A in striatal neurons.

MUSANTE V. 1 , ANDRADE E. 2 , CANIO J. 4 GREENGARD P. 3 , NAIRN A.C. 1,3

1Dept. of Psychiatry, Yale Univ. Sch. of Med., New Haven, CT; 2Lab. of Mol. Biol., 3Lab. of Mol. and Cell. Neurosci., Rockefeller Univ., New York, NY; 4W.M. Keck Foundation, Keck MS and Proteomics, Yale Univ., New Haven, CT

INTRODUCTION

tion is the most common post‐translational modification in neuronal signaling and result from the balanced action of kinases and phosphatases. We previously identified ^a family of striatal‐enriched phospho‐proteins substrates for PKA: DARPP‐32, RCS and

ARPP‐16 (Walaas et al.,1983, Horiuchi et al.,1990) While the role of DARPP-32 and RCS in the dopamine (DA)-mediated regulation of serine/threonine phosphatases

in striatal medium spiny neurons (MSNs) has been well characterized (Walaas et al. 2011) little has been known about the role of ARPP-16. ARPP‐16 is related to two other members of ARPP family, ARPP‐19 and ENSA. ARPP‐19 and ENSA are ubiquitously

distributed and have been identified as phosphatase PP2A inhibitors in mitotic cells. In Xenopus oocytes, phosphorylated by Greatwall kinase (GWT), ARPP‐19/ENSA inhibit PP2A during the G2/M phase (Lorca & Castro 2013).

Recently, we found ARPP‐16 directly interacts with PP2A in striatum and it is phosphorylated at Ser46 by sociated serine/threonine kinase 3 (MAST3), a mammalian analogous of GWT, enriched in s

and still poorly characterized.
Phosphorylation of ARPP-16 at Ser46 by MAST3 com werts the protein into an inhibitor of PP2A towards selection substrates including DARPP‐32. Moreover, Ser46 of ARPP‐16 is phosphorylated to ^a high basal stoichiometry in

striatum, while activation of PKA by cAMP leads to marked dephosphorylation of Ser46 in striatal slices. In the current study we further investigate the role of the phosphorylation mechanisms in ARPP‐16 regulation. We demonstrated, both in vitro and in intact cells, that PKA plays a fundamental role in the MAST-mediated
phosphorylation of ARPP-16 and regulation of PP2A. PKA phosphorylation of ARPP-16 at Ser88 negatively acts in
an intra

phosphorylation of Ser88 does not have any effect on the phosphatse innibilion. We also find that PACA
phosphorylation of Ser88 does not have assets in the state of the state and the state of the forestion in
transfected H understand the role of this kinase in the regulation of the synaptic activity in striatum.

METHODS:

Cell line culture: HEK 293‐T cells were grown on un‐coated plates in DMEM (Invitrogen), supplemented with 10% FBS **GST pull-downs:** His-ARPP-16 was immebilized onto 80 µl/sample [of 50/50 slurry] Talon metal affinity resin (Clomech Laboratory Inc, Mountain View;
CA). Increasing amounts of purlined PP2A-A (0, 20 ng, 40 org. 00 ng/ben a

CONCLUSIONS

- **ARPP‐16 interacts with phosphatase PP2A by direct binding with the A subunit (Fig. 1),**
- **ARPP‐16 is phosphorylated at S46 by MAST3 kinase (Fig. 2),**
- **Three heterotrimers of PP2A are inhibited by p‐Ser46‐ ARPP16, but to varying extents (Fig. 3), phosphorylation on Ser88‐ARPP‐16 does not affect PP2A activity,**
- **Phosphorylation of ARPP‐16 by PKA (on S88) or MAST3 (on S46) mutually attenuate each other's ability to phosphorylate ARPP‐16** *in vitro* **(Fig. 4),**
- **PKA phosphorylates MAST3** *in vivo* **and** *in vitr***^o and decreases its activity** *in vitro* **(Fig. 5),**
- **ARPP‐16 phosphorylation on Ser88 is able***, in vivo***, to regulate in an intra‐molecular fashion the phosphorylation on Ser46, at the same time, phosphorylation on Ser46 is just partially able to influence the phosphorylation on Ser88 by PKA (Fig.6),**
- **cAMP plays ^a key role toward MAST3 in the regulation of phosphorylation on S46‐ARPP16 (Fig. 7).**

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Figure 2: ARPP‐16 is phosphorylated at Ser46 in intact cells and *in vitro* **by MAST3 Kinase**

c.

b.

a.

Figure 3: P‐S46‐ARPP‐16 inhibits PP2A heterotrimer ability to dephosphorylate P‐T75‐DARPP‐32 *in vitro***.**

RESULTS

a. (left panel) Re a. (left panel) Recombinant purified ARPP-16 or P-588-ARPP-16 (1 uM) were incubated with ATP++³⁹P and
immunoprecipitated MAST3 Kinase overexpressed in HEK293 cells, for various times (as indicated); the proteins
were sep immunoprecipitated MA5T3 Kinase overespressed in HEK293 cells, for 2 min; proteins were separated by SDS
PAGE and phosphorylation of Ser46 was measured by autoradiography. Phosphorylation by MAST is modulated by
b. (left commercial purified ARPP-16 (1 µM) or P-S46-ARPP-16 were incubated with ATP+p-²¹P and purified PKA, for various times (as indicated); the proteins were separated by SDS-PAGE and purified PKA, for various times (as indica phosphorylation of Ser46-ARPP-16 was measured by autoradiography. (**right panel**) Increasing concentrations (10-
500 nM) of ARPP-16 or P-S46-ARPP16 were incubated with ATP-y-¹²P and commercial purified PKA, for 2 min; th ins were separated by SDS‐PAGE and phosphorylation of Ser88 was measured by aut
phorylation by PKA is modulated by previous MAST3 phosphorylation of ARPP‐16 with decrease Phosphorylation by PKA is modulated by previous MAST3 phosphorylation of ARPP-16 with decrease of
Vmax (from 6999 to 1027 ³²Pincorporation/min).

(**a.** and **b.** left panels) The resulting values for phosphorylation are expressed in arbitrary densitometric units $(a.u.)$ as mean \pm SEM of five independent experiments.

a. Schematic model of MAST3 kingse phosphonolation sites identified by MS analysyi, MAST3-HA was over a. Schematic model of MAST3 kinase phosphorylation sites identified by MS analysyl. MAST3-HA was overexpressed
in HEK293T cells and after 24 hours cells were treated with forskolin (10 uM for 30 min). MAST3-HA was then
iso staining. The phosphorylated sites on MAST3 sequence were identified by Orbitrap MS/MS analysis. **b.** Immunoprecipitated MAST3 kinsse, overexpressed in HEX293T cells, was incubated with ATP+-^{11P} and commercial
purified PKA for various times (as indicated). The proteins were separated by SOS-PAGE and phosphorylation of
MA indicated). The proteins were separated by SDS‐PAGE and phosphorylation of Ser46‐ARPP‐16 was measured by autoradiography. **b.** and **c.** Results shown the average from three experiments (error bars show SEM).

ents (error bars show SEM)

d. To confirm the PKA phosphorylation effect on MAST3 kinase activity we used mutant construct sfor T389‐MAST3. The unphosphorylatable T389A‐ MAST3‐HA and the phosphomimetic T389D‐MAST3‐ HA were overexpressed in HEK293T cells. Immunoprecipitated MAST3 mutants were incubated with ARPP‐16 (100 nM) in the presence of ATP‐γ‐32P for different times. T389A‐MAST3 was previously incubated for 20 min with PKA before the assay with ARPP‐16. The proteins were resolved by SDS‐PAGE and phosphorylation of Ser46‐ARPP‐16 was measured by autoradiography. Results represent the average from

Time (min)

d.

a. Hek293T cells were transfected with ARPP16‐HA (WT) or the phosphomutants S46D‐ARPP‐16 or S88D‐ARPP‐16 alone or in presence of MAST3‐HA kinase. After 24 hours cells were treated with 10 ^M Forskolin (FSK) for 30 minutes. Levels of phosphorylation for S46 and S88 ARPP-16 were measured by immunoblotting with phospho-specific antibodies on SDS-
PAGE-recolved cell bisates, Phospho-site signals were normalized for total APPP-HA expression with anti HA an PAGE-resolved cell lysates. Phospho-site signals were normalized for total ARPP-HA expression with anti HA antibody.
b. Graph of summary data shows phosphorylation on the different sites expressed in arbitrary densitomet **b.** Graph of s mean ± SEM of six independent experiments run in triplicate. *, ^p < 0.05; *** ^p < 0.001, on way ANOVA, multiple comparison test.

Figure 7: cAMP‐dependent signaling regulates ARPP‐16 phosphorylation via MAST3/PKA in ^a cell line model

a. The effect of cyclic AMP (cAMP) signaling as ^a mutual regulator of ARPP16 phosphorylation (S46 and S88) in intact cells was investigated. Hek2931 cells were transfected with ARPP16-FM (AT6-FM) alone or plus MAST3+1-Mor MAST3+T380-HA
After 24 hours: cells were treated with 10 juM Forskolin (FSK). Levels of S46 and S88 A16+MA were measured by phospho-specific A16 antibodies on cell lysates (upper panel). Phospho-site signals were normalized for total ARPP‐HA
expression with anti‐HA antibody Graph of summary data (lower panel) shows phosphorylation on the differ expression with anti-HA antibody. Graph of summary data (lower panel) shows phosphorylation on the different sites
expressed in arbitrary densitometric units (a.u.) as mean ± SEM of four independent experiments run in trip 0.01; *** p < 0.001, one way ANOVA, multiple comparison test. ##, p<0.01

