

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

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INTRODUCTION

Protein phosphorylation is the most common post-translational modification in neuronal signaling 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 phosphatase

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 ubiquitor

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

ntly, we found ARPP-16 directly interacts with PP2A in striatum and it is phosphorylated at Ser46 by otubule-associated serine/threonine kinase 3 (MAST3), a mammalian analogous of GWT, enriched in striatum logous of GWT, enriched in

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stratum, wine activation of YAA by CAMM keaks the marked appropriority/inten of series in training as memorarized, being the series of the se

phosphorylation of SerBB does not have any effect on the phosphatase inhibition. We also that that PAA hopophorylate Natifiant's in writer resulting in knase inhibition, and that activation of CAMP by forsionin in transfetched HEX cells significantly decreases MATS1 activity. Overall these data sugget that PS-aetAPP-16 acts to basaly control PPZA in MSNs, but that DA, acting via PKA, regulates this pathway by inactivating ARPP-16 and/or MATS1 leading to selective potentiation of PPZA signaling. Turther oppenments are in progress to beliently other specific MATS1 substrates and better and the selective that the selective and better understand the role of this kinase in the regulation of the synaptic activity in striatum.

METHODS:

wes: His-ARPP-16 was immobilized onto 80 µl/cample (of 50/50 slurry) Talon metal affinity resin (Clontech Laboratory Inc, Mountain View, og amounts of purified PP2A-4 (0, 20 ng, 40 ng, 200 ng, or 400 ng) were added to the beads and samples were incubated for 1 hr at 4° C A sa control, 40 ng PP2A - Awas added to beads with on the ARPP-16 immobilized.

Plasmids and Transfection: MAST3-HA, MAST3T389A/L HEK293 cells with Lipofectamine 2000 (Invitrogen) in Opt	0-HA, ARPP-16-HA, Balfa-FLAG, B568-FLAG and PR72-FLAG constructs were transfected in -MEM media (Invitrogen) an. Protein expression was assayed at least 24 hr later.
APPP 126 in vitro phosphorylation: Recombinant APP- Agrance (Gapes), hurring divisit-ApPP-16 () µully user a presence of 200 µul ATP (Sigma) or 1 mM tile-vATP (Bo CAMPP-32 in vitro phosphorylation and PP2A activity or in buffer containing 50 mM til-i+CL pH 7.1, 150 mM K for the IP2A accog, different PP2A heterotriners were presence of 73 go (TP4) AdPP-22 for 10 mil at 30 phosphorobins with cold TCA. Call line teachers to a 20 with 250 mM CA. (150 mM K 40, 150 mm) K 40.1 (150 mm) K 40.1 with 250 mm) K 40.1 (150 mm) K 40.0	So fixed to derive tag was expressed in Extended car (IE32) and purified using KMAR approximation buffy of purifying the second
Striatal disc preparation, striatal clics rom 6-3 week ((Kish) et al., 1997). The brains were rapidly removed at NaHCO3, 1.5 mM CaC2, 1.25 mM KH2PO4, 1.5 mM KH3 microtone (VT10005, kicia Meror) (Kisha were et HCO3 buffer and allowed to recover for 30 mis with co- with advected edaminase added to fresh KHbLHCO phosphatase and protease inhibitor cocktails and quan parformed	di nice or 90-110 p zet were prepared za describel min tight procedural modifications of pactor in iso-colo, paysmatel in Rote-Ho 20 battle (21 am Mix 2), and Mix 3, and Mix 4, and Mix

- 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 vitro 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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b



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



RESULTS



a. (left panel) Recombinant purified ARPP-16 or P-S88-ARPP-16 (1 uM) were incubated with ATP-y-32P and panely recommanic pumper and provide the processing of the provided with AIP+P-16 or processing the commentance pumper and a processing the relation of the processing of t Increasing conce Increasing concentrations [L0-soo inn) or AmP+10 or P-soo-AMP+10 were includated with AmP+2P+ immunoprecipited MASTI Kinase overexpressed in HRX93 cells for 2 min; proteins were separated by SDS-PAGE and phosphorylation of Ser46 was measured by autoradiography. Phosphorylation by MAST is modulated by previous PKA phosphorylation of Ser46 was received in the Carse in the Masti (from 1398 to 857-Phi Incorporation/min). periods FAC phosphorylation of purper 10 with a declease in Mina (1001 1536 to dot) + incorporation (1001 1536 to dot) 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-32P and commercial purified PKA, for 2 min: the proteins were separated by SDS-PAGE and phosphorylation of Ser88 was mea ured by autoradiography. vlation by PKA is modulated by previou m 6999 to 1027 ³²Pincorporation/min is MAST3 phosphorylation of ARPP-16 with dec

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



a Schematic model of MAST3 kinase phosphonylation sites identified by MS analysyi MAST3-HA was on a. Schematic model or invols skinase prospirorylauton sites identified by waranays), invols s-ne was overexpressed in HK293T cells and after 24 hours cells were treated with forskolin (Jo UM for 30 min). MAST3-HA was ther isolated by immunoprecipitation, resolved by eletrophoresis and the MAST3 gel band identified by Coomassie staining. The phosphorylated sites on MAST3 sequence were identified by Orbitrap MS/MS analysis. b. Immunoprecipitated MAST3 kinase, overexpressed in HEK293T cells, was incubated with ATP-y-32P and comr purified PKA for various times (as indicated). The proteins were separated by SDS-PAGE and phosphorylation of MAST3 was measured by autoradiography. c. The activity of PKA-phosphorylated MAST3 kinase (P-MAST3) was measured as the ability to phosphorylate ARPP-16 in comparison with unphosphorylated MAST3. Recom purified ARPP-16 (100 nM) was incubated with P-MAST3 or MAST3 in the presence of ATP-γ-³²P, for various times (as indicated). The proteins were separated by SDS-PAGE and phosoborylation of Ser46-ARPP-16 was measured by autoradiography. b. and c. Results shown the average from three experiments (error bars show SEM).

tated MAST3 mutants were

from three





a. Hek293T cells were transfected with ARPP16-HA (WT) or the phosphomutants S46D-ARPP-16 or S88D-ARPP-16 alone o in presence of MAST3-HA kinase. After 24 hours cells were treated with 10 μM Forskolin (FSK) for 30 minutes. Levels of phosphorylation for S46 and S88 ARPP-16 were measured by immunoblotting with phospho-specific antibodies on SDSblotting with phospho-specific antibodies on SDS-ARPP-HA expression with anti HA antibody. PAGE-resolved cell lysates. Phospho-site signals were normalized for total ARPP-HA expr b. Graph of summary data shows phosphorylation on the different sites expressed in arbitrary densitometric units (a.u.) as mean ± SEM of six independent experiments run in triplicate. *, p < 0.05; *** p < 0.001, on way ANOVA, multiple</p> 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 a. Interest, ot typical waver (such as a structure) and the structure interest of the structu phospho-specific A16 antibodies on cell lysates (upper panel). Phospho-site signals were normalized for total ARPP-HA sion with anti-HA antibody. Graph of summary data (lower panel) shows phosphorylation on the different site: d in arbitrary densitometric units (a.u.) as mean ± SEM of four independent experiments run in triplicate. **, p p < 0.001, one way ANOVA, multiple comparison test. ##, p<0.01.





