



The role of prolyl oligopeptidase, understanding the puzzle

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Prolyl oligopeptidase or prolyl endopeptidase (POP or PREP) was first described 50 years ago as an oxytocin cleaving enzyme (1) and it was further characterized as a peptidase able to cleave short peptides at the C-side of an internal proline (2). Because of its specificity, and the occurrence of internal proline residues in several neuropeptides, PREP was soon regarded as a peptidase relevant in neuropeptide metabolism, with a great drug target potential for neurological disorders' therapy (3). Accordingly, synthesis and testing of PREP inhibitors became the attention of academia and industry, especially since evidence emerged that PREP inhibition improved cognitive decline and dementia in several animal models [reviewed in (4)]. Some of the very potent and specific inhibitors were taken to preclinical/clinical trials (5). The impetus started to fade away when further research determined that the role of PREP, as neuropeptide regulator, was not that clear. The hypotheses to support its role in learning and memory were not solid enough (6). One of the main concerns was the notion that PREP was considered intracellular and mainly expressed within neurons. This idea seemed to be contradictory with the extracellular environment of bioactive peptide regulation, where the receptor-neuropeptide interaction takes place. Furthermore, additional evidence pointed out that PREP was present in almost all tissues, at significant levels, and not just within the cytoplasm, but within neuron nuclei, attached to membranes, and extracellularly [see for example (7) and papers therein]. This, along with the poor information on the role of PREP in any of those locations,

ceased the interest of pharmaceutical industry to continue the efforts to validate PREP for clinical treatment of dementia.

Only few laboratories had been working to try to unveil the role of PREP in the body (2-4,6-8), and this task has not been easy. During more than 20 years, experimental work has indicated that probably PREP has not one physiological unique distinctive role, but many, which depend on the milieu PREP is located: in or out of the cell, the type of cell or tissue, or the metabolic or pathological conditions in cells, tissues or whole organism. These roles could be also determined by the kind of interaction with physiological peptides (cleavage substrates or inhibitors) or with other protein partners (8).

Information about involvement of PREP in processes non-dependent of its peptidase activity have been substantiated. Genetically or biochemically inactivated versions of this peptidase are able to participate in neuron growth cone, decrease of alpha-synuclein deposits, and other interactions with key proteins as GAP43, tubulins, etc. (8).

As a peptidase, PREP seems to have a clear role on peptide regulation in the renin-angiotensin system, participating in processing angiotensin II (9). On the other hand, there is evidence that PREP is also involved in the digestion of thymosin β 4 to generate the tetrapeptide Ac-SDKP, which promotes angiogenesis, reduces fibrosis and apoptosis, and has anti-inflammatory effects (10).

In brain, the evidence of PREP participating in neuropeptide metabolism has not been conclusive (8).

However, changes of PREP activity have been found in pathologies where neuroinflammation has a central role, as multiple sclerosis, cirrhosis, depression, neurodegeneration and cancer (11-14). Indeed, it has been shown that PREP is secreted by immunoactivity cells (15,16), this being obnoxious to the close parenchymal cell population. However, the molecular mechanisms involved in this activity have not been substantiated. The peptide PGP has been described to correlate with PREP levels in inflammatory lung diseases as chronic obstructive pulmonary disease, and cystic fibrosis (11), by activating neutrophils through CXCR2 receptors.

In previous *in vitro* studies Zhou *et al.*, showed that PREP increases upon hepatocyte steatosis and that PREP inhibitors reduce lipid accumulation (17). In their report, Jiang *et al.* (18), hypothesized that PREP disruption would ameliorate disorders of lipid metabolism and hepatic inflammation to prevent nonalcoholic fatty liver disease (NAFLD) progression to nonalcoholic steatohepatitis (NASH). This work indeed shows that PREP disruption reduces visceral adipose tissue, downregulates free fatty acid transporters, and improves *de novo* lipogenesis in steatohepatitis mouse model. Authors, in part, relate these findings to previous reports on changes in mitochondrial protein turnover due to PREP inhibition (19), but also invoke to a possible gene expression control mediated by this peptidase (20), through a mechanism which involves PPAR- γ expression modulation, a key gene in the regulation of lipid metabolism.

The other important finding of this study, represents the observation that the lack of PREP decreased considerably the inflammatory response produced by high fat diet in wild type animals. Certainly, alteration of inflammatory response has been reported in the brain from mice in PREP-knockout mice in our laboratory (21). Jiang *et al.* (18) find that the hepatic macrophage population, as well as PGP peptide levels, are decreased in high-fat diet (HFD) PREP-deficient mice. As mentioned above, PGP peptide and its N-acetylated derivative are strong neutrophil activators. This peptide is derived from the degradation of the extracellular matrix (ECM) collagen by a concerted action of metalloproteinases and PREP. Thus, the lack of PREP, and the decreased expression of metalloproteins, observed in the PREP deficient mice in this study, are then culprit for the fall in neutrophils activation in the steatosis model. Furthermore, the receptor of this peptide (CXCR2) was found also increased at some extent in the HFD wild type mice, compared with those levels in the liver of low-fat diet (LFD) animals from the same genotype. However,

those levels were found decreased in the liver of both dietary conditions in the PREP-deficient mice, especially on HFD mice. Accordingly, Jiang *et al.* (18) proposed that PREP disruption inhibits the accumulation of hepatic macrophages by regulating the generation of PGP (Ac-PGP) and neutrophil chemotaxis upon HFD stimulus. Finding whether the reduction on inflammatory response is due directly to PREP absence itself, or to the lower hepatic fat accumulation in these mice, would be a further hypothesis for a follow-up research.

ECM remodeling has multiple participation on disease (22). Previous research has showed that ECM digestion by physiological proteases and peptidases, lead to the formation of “matrikines”, which are capable to regulate inflammatory cell phenotypes both *in vitro* and *in vivo* (23). In the commented article (18), and in previous research (17), Jiang *et al.* provide evidence that PREP is indeed having a central part in the relation of ECM remodeling, and thus, in inflammation, paving the road ahead in the research of this interesting protein.

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Footnote

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