- 1 Novel inhibitory activity for serine protease inhibitor Kazal type-3 (Spink3) on human recombinant
- 2 kallikreins
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Kallikrein-related peptidases (KLKs) are trypsin-like and chymotrypsin-like serine proteases which are expressed in several tissues. Their activity is tightly controlled by inhibitors including members of the serine protease Kazal-type (SPINK) family. These enzymes are promising targets for the treatment of skin desquamation, inflammation and cancer.

Spink3 or caltrin I is expressed in mouse pancreas and males accessory glands and the resulting mature protein has been associated with different activities such as an inhibitor of trypsin and acrosin activity, calcium transport inhibitor in sperm and inhibitor of cell proliferation during embryogenesis. In this study we produced a soluble recombinant Spink3 from mouse seminal vesicle (rmSpink3) that inhibited the activity of human KLKs. Using FRET substrates, rmSpink3 exhibited a potent inhibitory activity against human KLK2, KLK3, KLK5 (KLKn) (Ki ranging from 260 to 1500 nM), and to a lesser extent against KLK6, KLK1 and KLK7 (Ki around 3000 nM). As shown by mass spectrometry analysis of rmSpink3 incubated with trypsin, the inhibitor was not truncated by the target enzyme. Based on the analysis in

silico of the expression of Spink3/SPINK1 and KLKs it is speculated that some KLKs may be natural targets of Spink3/SPINK1, however experimental confirmation using both proteins from mouse or either from human origin is needed.

This work shows rmSpink3 is a potent inhibitor of various human KLKs members suggesting the potential of this molecule in the diagnosis/prevention of several human diseases.

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1- Introduction

Kazal-type protease inhibitors play important roles in many biological and physiological processes, such as blood clotting, the immune response and reproduction [1]. Serine protease inhibitor Kazal type 3 (Spink3) is a mouse 6-kDa protein, also called P12 [2, 3], which was initially described as an inhibitor of trypsin and acrosin [4]. The orthologue human protein, designated SPINK1 or PSTI, was originally isolated from pancreas where its expression is constitutive [5-7] and essential for the control of trypsin activity. SPINK1 was reported to differ from the other pancreatic inhibitor BPTI because the former one lacks of inhibitory effect on chymotrypsin and pancreatic kallikreins (KLKs) [7], however the effect was evaluated over porcine pancreatic kallikrein [8]. It has been reported that mutations in Spink1 are associated with hereditary pancreatitis [9]. Recent evidence suggests that Spink3 is a multifunctional protein that functions as a growth factor for the regeneration of acinar cells in pancreas as well as for the proliferation of various types of cells during early embryogenesis [3, 7, 10, 11]. These are novel functions of this molecule apart from that already known as a protease inhibitor. Spink3 has also been detected in the male accessory sexual glands and female uterus under androgen and progesterone control, respectively. In this context, Spink3 secreted from the mouse seminal vesicles and prostate, contributes to the seminal fluid [2]. MatureSpink3 attaches to the sperm head [12], inhibits calcium uptake by spermatozoa [12, 13] and is involved in the sperm acrosome

reaction and zona binding [2, 14], independently of its trypsin inhibitory activity [12, 15]. However, the target proteinase of Spink3 (if there is one) in organs other than the pancreas, remains unknown. Kazal domain, which comprises most of the mature form of Spink3, is composed of 40-60 amino acid residues that include six well-conserved cysteine residues capable of forming three intra-domain disulfide bridges (cys 1–5, 2–4, 3–6). This assembly results in the characteristic three-dimensional structure consisting of one α -helix surrounded by an adjacent β -sheet and three loops of peptide segments [16]. The inhibitory specificity of a Kazal domain varies with a different reactive P1 amino acid, which is the second amino acid after the second Cys [1]. The reactive loop harbors the specificity determinant P1 amino acid and provides easy access to the active site of the cognate proteinase. The Kazal domain acts as a substrate analogue that stoichiometrically and competitively binds to the active site of the target proteinase forming a proteinase–proteinase inhibitor complex, much more stable than the Michaelis enzyme-substrate complex [16]. KLKs are secretory serine proteases of small size (approximately 30 kDa). The human glandular KLK gene locus on chromosome 19q13.4 is composed of 15 genes encoding serine proteases. Many members of the human KLK family, in addition to KLK2 and KLK3 (prostate specific antigen, PSA), are promising biomarkers for ovarian, prostate, and breast cancers since KLK proteinase expression, which in many cases is under the control of steroid hormones, changes dramatically in a variety of malignant diseases [17, 18]. KLK genes are widespread along mammalian [19] and non-mammalian [20] genomes indicating their importance in the physiology. In mouse, the total number of glandular Klk genes is 37 of which 11 are pseudogenes. Interestingly, there are 12 genes that represent the orthologues of human KLK genes (KLK1, KLK4-KLK15) being KLK2 and KLK3 the only ones that do not have mouse orthologues [21, 22]. Given the pleiotropic roles of KLKs, both activators and inhibitors of KLK activities either from synthetic or natural origin are of therapeutic interest [17, 23].

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Many endogenous inhibitors are known to regulate the activity of KLKs forming complexes with plasma inhibitors, including various serpins, such as $\alpha 1$ -antitrypsin ($\alpha 1$ -protease inhibitor), $\alpha 1$ -antichymotrypsin (ACT), protein C inhibitor (a.k.a. plasminogen activator inhibitor-3), plasminogen activator inhibitor-1, antithrombin III, and $\alpha 2$ -antiplasmin [24]. Furthermore, it has been proposed that inhibitory peptides derived from Kazal type serine protease inhibitors may regulate KLK activity due to co-localization; that is the case of KLKs and SPINK5 in the skin [25-27]. Another supporting evidence, is that both the tandem localized *KLK* genes [28] and *SPINK1/Spink3* genes [29] are under transcriptional regulation by steroid hormones. We have been working with Spink3 in the elucidation of its role on sperm physiology [12]. Our experience with Spink3 and the above mentioned evidences prompted us to speculate that Spink3 might in vitro regulate the activity of KLKs. , We have previously obtained a soluble recombinant GST-Spink3 fusion protein which is devoided of trypsin inhibitory activity unless released from the GST portion [12]. Although this recombinant Spink3 retained protease inhibitory activity, protein yield was low which precluded scaling up. In this study, we developed a new construct that allowed the expression of a version of recombinant mouse Spink3 (rmSpink3) with protease inhibitory activity and good yield. This protein was an effective inhibitor on various humanKLKs of pharmacological interest.

2- Materials and methods

- 88 2.1- Heterologous expression of recombinant Spink3 (rmSpink3-His₆)
- 89 A DNA fragment corresponding to mature Spink3 (nucleotides 158 to 331 which encoded a polypeptide
- 90 of 6.1 kDa starting from residue 24 to the stop codon) was previously cloned into pGEX-4T-3 expression
- 91 vector (GE) [12]. To facilitate subcloning of Spink3 cDNA into the expression vector pET-24b(+)
- 92 (Novagen), the restriction sites Ndel and Xhol were added to the 5' end nucleotide sequences of Spink3-
- 93 Fw (5'-GG GAA TTC CAT ATG GCT AAG GTG ACT GG-3') and Spink3-Rev (5'-GGC CTC GAG GCA AGG CCC
- 94 ACC-TTT TC 3') primers, respectively. To obtain a fusion protein with a His-tag at the C-terminus (8,06

kDa), the stop codon of *mSpink3* was excluded in the Rev primer. The PCR reaction was carried out in a mixture (20 μl) containing 1 x buffer, 1.5 mM MgCl₂, 0.2 mM dNTPs mix, 1 μM of primers, 15% DMSO, 300 ng pGEX-*Spink3* template and 2.5 U Taq DNA polymerase. The PCR amplification protocol was as follows: 35 cycles of denaturation at 94°C for 1 min, primer annealing at 59° C for 1 min, and primer extension at 72°C for 1 min. The PCR product of the expected size (204 bp) was agarose gel-purified according to the MinElute purification kit (Qiagen).

The cDNA encoding mature Spink3 and pET -24b(+) expression vector were digested with *Ndel* and *Xhol* (Promega) following the manufacturer's instructions and ligated with T4 DNA Ligase (Promega) according to standard protocols. The generated construct, pET-24b (+) -*Spink3* which encoded recombinant mouse Spink3 followed by a six-histidine His-tag at the C-terminus, (rmSpink3-His₆) was then amplified in *E. coli* DH5α competent cells and the potential positive clones were confirmed by DNA

sequencing.

For overexpression of rmSpink3-His₆, *E. coli* Rosetta cells (Novagen) were transformed with the pET-24b (+)-*Spink3* and the cells harboring the plasmid were grown overnight at 37 °C in Luria Bertani (2YT-LB) medium (1% yeast extract, 2% tryptone, 0.5% NaCl) supplemented with kanamycin (50 μ g/ml) and chloramphenicol (35 μ g/ml). This starting culture was used to inoculate (1/100) the same medium (50 ml) and the culture was grown at 37 °C under continuous shaking until the optical density at 600 nm (OD₆₀₀) reached 0.4, then shifted to 18 °C and further incubated to an OD₆₀₀ ~0.6. Synthesis of rmSpink3 was induced by addition of 0.1 mM IPTG for 16 h at 18° C. Bacteria were harvested by centrifugation (1700 x g for 8 min at 4° C), suspended in PBS (ICN tablet 2810305; 10 mM phosphate buffer pH 7.4, 137 mM NaCl, 2.7 mM KCl) containing 1 mg/ml lysozyme, 0.2% (v/v) Triton X-100 and 0.5 mM PMSF and lysed by sonication (6 x 30 s 40 W, Cole Parmer 4710). The soluble (S) and insoluble (I) cell lysate

fractions were obtained by centrifugation (12000 xg for 25 min at 4 °C) and analyzed by SDS-PAGE (15% w/v polyacrylamide gel) [30].

HiTrap IMAC HP affinity chromatography (28-4046-23 AA, GE) was applied for purification of rSpink3-His $_6$ from the soluble fraction of induced cells. The column was loaded with 0.1 M NiSO $_4$, equilibrated and washed with binding buffer (20 mM sodium phosphate, 0.5 M NaCl, 30 mM imidazole, pH 7.4). Then, the soluble extract was loaded, washed and the recombinant protein was eluted with 20 mM sodium phosphate, 0.5 M NaCl and 500 mM imidazole, pH 7.4. Purified rmSpink3 was concentrated by ultrafiltration in YM-3000 membranes (Amicon) and the purity of the protein was evaluated by SDS-PAGE (w/v 15% polyacrylamide gel). The identity of rmSpink3 was validated by Western blot.

2.2- Western blot

Proteins were fractionated on 15% (w/v) polyacrylamide gels and transferred to PVDF membranes for 2 h 80 mA in towbin buffer (25 mM Tris, 192 mM glycine and 20% methanol) using a TE 70 PWR transfer equipment (Amersham). Nonspecific binding sites on the membrane were blocked with 3% skimmed milk in TBS-N (10 mM Tris-HCl pH 7.5, 0.15 M NaCl, 0.1% ipegal CA630). Incubation with primary antibody was performed for 1 h at 37 °C with either anti-SPINK1 (1:4500, Sigma HPA027498) or anti-His (1:3000, GE 27-4710-01) antibodies in blocking solution. After washing with TBS-N, the membrane was incubated either with alkaline phosphatase-conjugated anti-rabbit IgG (1:10000, A8025 Sigma) or alkaline phosphatase-conjugated anti-mouse IgG (1:5000, A3562 Sigma) and developed with 0.33 mg/ml nitro-blue tetrazolium (NBT) and 0.165 mg/ml 5-bromo-4-chloro-3-indolyl phosphate (BCIP) in alkaline phosphatase buffer (0.1M Tris pH 9.5; 0.1M NaCl and 5 mM MgCl2). The ability of the commercial anti-SPINK1 (raised against an immunogenic sequence of 51 amino acids from SPINK1) to recognize Spink3was previously confirmed [12].

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2.3- Determination of the kinetic parameters of KLK-Spink3 interaction

Mature humanKLK1, KLK2, KLK3, KLK5, KLK6 and KLK7 were expressed and purified from a baculovirus/insect cell line system as previously described [31]. Briefly, DNA fragments encoding the mature form of KLKs were inserted in frame downstream the enterokinase propeptide nucleotide sequence. To facilitate the purification of the recombinant proteins, a 6x His-tag codon was included at the 3'-end of the enterokinase propeptide sequence. After purification by Ni²⁺ affinity chromatography, KLKs were activated by incubation (24 h) with enterokinase at 1:100 ratio enterokinase:pro-KLKs (w/w). Hydrolysis of the enterokinase propeptide proceeded to apparent completion and mature KLKs were purified by size exclusion chromatography. The final yield of purified mature KLKs was typically 20-25 mg per liter of culture. Purity of mature KLKs was assessed by SDS-PAGE (w/v 15% polyacrylamide gel) and Coomassie Brilliant Blue staining, N-terminal sequencing, and MALDI-TOF mass spectrometry. Protein samples were about 98% homogeneous, with no visible degradation products and had the expected mature N-terminus corresponding to the mature proteases. All FRET peptides were obtained by solid-phase peptide synthesis as previously described [32, 33] and using the Fmoc-procedure in an automated bench-top simultaneous multiple solid-phase peptide synthesizer (PSSM-8 system from Shimadzu-Tokyo, Japan). The peptides were synthesized in TGR-resin (loading 0.2 mmol/g) using HBTU/HOBt as coupling reagent and the cleavage o peptide resins was accomplished with TFA:anisol:EDT:water (85:5:3:7). All peptides obtained were purified by semipreparative HPLC on an Econosil C-18 column. The molecular weight and purity (94% or higher) of synthetized peptides were checked by MALDI-TOF mass spectrometry, using a Microflex LT mass spectrometer (Bruker Daltonics, Billerica, USA). Stock solutions of the peptides were prepared in DMF and the concentrations were measured by colorimetrically determination of the 2,4-dinitrophenyl group (molar extinction coefficient of 17,300 M-1 cm-1 at 365 nm) [34]

To determine kinetic parameters, enzymes were mixed with varying substrate concentrations (0.5-200 μM). The FRET peptides were followed at λex 320 nm and λem 420 nm (excitation and emission wavelengths for Abz) in a spectrofluorometer Shimadzu RF-1501 (Shimadzu Corporation, Kyoto, Japan), previously calibrated with standard solutions of Abz hydrolyzed. KLK1 (5 nM) was assayed in 50 mM Tris-HCl buffer pH 9.0 containing 1 mM of EDTA, using the substrate Abz-KLRSSKQ-EDDnp (Km 1.5 μM); KLK2 and KLK3 (5 nM) were assayed in 50 mM Tris-HCl buffer containing 1 M sodium citrate final pH 7.5 (adjusted with 2N NaOH after citrate addition), using the substrate Abz-KLRSSKQ-EDDnp (Km 5.9 μM and 1.2 µM respectively); KLK5 (5 nM) and KLK6 (5 nM) were assayed in 50 mM Tris-HCl buffer pH 7.5 containing 1 mM of EDTA, using the substrate Abz-KLRSSKQ-EDDnp (Km 0.69 and 4.0, respectively); KLK7 (10 nM) was assayed in 50 mM Tris-HCl buffer containing 1 mM of EDTA, pH 7.5 using the substrate Abz-KLFSSKQ-EDDnp (Km 1.8 uM). As an internal control of our inhibition study chymotrypsin (10 nM) was assayed in 50 mM Tris-HCl buffer pH 8.0 containing 5 mM CaCl2, using the substrate Abz-KLFSSKQ-EDDnp (Km 4.0 μM) and trypsin (10 nM) was assayed in 50 mM Tris-HCl buffer pH 8.0 containing 5 mM CaCl₂, using the substrate Abz-KLRSSKQ-EDDnp (Km 1.9 μM). To determine rmSpink3 inhibition constant, assays were performed at 37°C at substrates concentration of 5 μM. The enzymes were pre-incubated in the assay buffer with different concentrations of rmSpink3-His₆ for 3 min at room temperature before substrate addition. Enzyme concentrations for initial rate determinations were selected such that less than 5% of the added substrate would be hydrolyzed during the time course of data collection. Enzymatic velocities in the presence or absence of rmSpink3-His₆ were calculated from the increments of fluorescence in the time-course of the reactions and the values were expressed as the percentage of the residual activity. The inhibitory domain of Kazal-type serine proteinase inhibitors is a typical cysteine-rich motif, similar to the bovine pancreatic secretory trypsin inhibitor (PSTI), which prevents protease activity in a substrate-like manner and binds tightly to the active site of the enzyme [35]. Thus,

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it was assumed that rmSpink3 was a competitive tightly-binding inhibitor and the inhibition constants (Ki) were obtained according the equation below [36, 37].

Equation 1

$$[I]_t/((1-v_i/v_0)) = K_i(1+S/K_m)v_i/v_0 + [E]_t$$

In the equation, *S* represents substrate concentration, *Km* the Michaelis-Menten constant, *[E]* represents the enzyme concentration, *v0* the velocity of substrate hydrolysis at time zero and *vi* the velocity at each inhibitor concentration. The reactions were performed at different inhibitor concentrations [I] of and the Ki values were obtained by linear regression.

2.4- Mass spectrometry analysis

Trypsin (100 pmol) and rmSpink3-His₆ (1 nmol) were incubated in water (10 μL) for 30 minutes and the released peptides were directly analyzed by MALDI-TOF/MS. Trypsin kept at least 20% its activity on Z-FR-MCA in water. In five times higher concentration of trypsin we got the same result. Aliquots of peptide mix (2μl) incubated with sinapinic acid (10 mg/ml) matrix solution (1:1) were spotted onto a stainless steel MALDI target plate and dried at room temperature before analysis. Mass spectra were obtained with a Bruker Daltonics Microflex LT instrument operating in linear, positive ion mode previously calibrate with serum bovine albumin, proteinase A and trypsinogen. For analysis of protein interactions, mass spectra were acquired utilizing the following instrument parameter: pulsed ion extraction delay of 250 ns, ion source voltage one, 20 kV, ion source voltage two, 18.25 kV, and ion source lens voltage 7.00 kV. For each sample, mass spectra were acquired by accumulating 50 laser shots at 90% laser power in the m/z range of 18-40 kDa.

3- Results

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We have previously reported the construction of a recombinant vector encoding the fusion protein GST-Spink3 [12] that was active as a trypsin inhibitor after removing the GST portion. However, protein yields were not sufficient to allow further characterization of this molecule. In this study, we constructed a new expression vector encoding recombinant C-terminal His₆ tagged Spink3 (rmSpink3-His₆) that was synthesized as a soluble protein in E. coli (Figure 1, lane 2) and purified to apparent homogeneity by Ni⁺² -affinity chromatography (Figure 1, lane 3) with good yield (6 mg/L of culture). Its identity was confirmed by Western blot with anti-hSPINK1 (the human orthologue protein) and by anti-His antibodies (Figure 1, lanes 4 and 5, respectively). The activity of the recombinant protein was first assayed on trypsin showing the typical competitive behavior described for Kazal type inhibitors (data not shown). Thereafter, the effect of rmSpink3-His₆ on the activity of several recombinant KLKs was tested (Figure 2). Calculation of the IC₅₀ from the inhibition plots showed that rmSpink3 was found to be a potent compound against KLK2 and KLK5 (IC₅₀ = 4 and 4.1 μ M, respectively), followed by KLK6 and KLK3 (IC₅₀ = 4.25 and 4.75 μ M, respectively). KLK7 IC₅₀ was not calculated since inhibition assay was performed with a different enzyme concentration. These inhibitory concentrations are lower than those reported for natural isocoumarins [38]. When the reaction was performed in the presence of high affinity substrates, substrate addition reversed the inhibition caused by rmSpink3 or the inhibitor did not affect enzyme activity at all (for example, KLK3 was not inhibited when Abz-KLFSSKQ-EDDnp was used as substrate; Km = $0.04 \mu M$). Although KLK3 is considered a chymotrypsin-like human kallikrein due to its high preference for Phe or Tyr, Andrade et al [39] reported that Abz-KLRSSKQ-EDDnp is cleaved at R-S bond in presence of sodium citrate with a Km value 25 times higher than the hydrolysis of Abz-KLFSSKQ-EDDnp. The evaluation of inhibitors is more convenient using substrates with high Km and high kcat that was the case for the hydrolysis of Abz-KLRSSKQ-EDDnp by KLK3 [39].

IC₅₀ values can vary with substrate concentration for competitive inhibitors such as Spink3. Moreover, in a tight-binding inhibition model, formation of the EI complex (enzyme-inhibitor) changes the concentration of free inhibitor during the reaction depending on the EI stoichiometry rather than the absolute inhibitor concentration [37, 40]. Therefore, the EI dissociation constant (Ki) calculated using the equation for tight-binding inhibitors as reported for other Kazal-type molecules [41], represents a better way to evaluate the effect of rmSpink3 as protease inhibitor (Table 1). Inhibition assays using FRET substrates indicated the strongest interaction between KLK5 and rmSpink3 (Ki = 0.26±0.03 μM) followed by KLK2 (Ki = $0.63\pm0.03 \,\mu\text{M}$). Slightly weaker inhibition was observed for KLK3 (Ki = $1.53\pm0.12 \,\mu\text{M}$) and to a lesser extent for KLK6 and KLK7 (Ki = 3.02±0.33 and 3.26±0.46 μM). Although KLK1 is a trypsin like enzyme, its isoeletric point is the lowest among all kallikreins evaluated (Table I, inset). This fact might explain the weak inhibition (Ki = $10.71\pm0.76 \mu M$). The rmSPIN3 Kis observed correspond well to those reported for Kazal type inhibitors (10⁻⁷–10⁻¹³ M) [16]. The specificity of rmSpink3 differs from that previously reported for human SPINK6 [41] on different KLKs. However, similarly to SPINK6, the target proteases of rmSpink3 have trypsin-like activity most likely determined by Arg48, which is the predicted P1 residue of the inhibitor, as revealed by sequence comparison with other Kazal-family protease inhibitors (Figure 3). As described for canonical inhibitors, the standard EI mechanism implies that inhibitors are peculiar protease substrates which contain the reactive site P1-P1' peptide bond located in the most exposed region of the protease-binding loop [42]. Kazal-type molecules may or may not be cleaved during inhibition [43]. As shown by the mass spectrometry analysis of rmSpink3 incubated overnight with or without trypsin, the inhibitor molecule was not hydrolyzed by the enzyme (Figure 4). In vivo, the role of mouse Spink3 as a serine protease inhibitor is not clear. Similarly to the human orthologue hSPINK1, Spink3 activity is thought to participate in the prevention of trypsin-catalyzed premature activation of zymogens within the pancreas [9]. Besides of pancreatic trypsin, their target

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enzymes in other tissues are unknown. To explore the potential role of mSpink3 as an inhibitor of KLKs activity, we examined the expression pattern of mSpink3, hSPINK1 and the KLKs used in this study using an in silico approach, combining the information obtained from 4 databases (Gene Expression Atlas EMBL-EBI, Unigene NCBI, Ensembl and MGI) (http://www.ebi.ac.uk; http://www.ncbi.nlm.nih.gov; http://www.Ensembl.org; http://www.informatics.jax.org/). Based on this analysis, Spink3 is expressed in many mouse tissues where KLKs are also expressed (Figure 5). In both mouse and human, mSpink3/hSPINK1 are co-expressed with mKLK1/KLK1 in prostate, pancreas and uterus, suggesting that in vivo KLK1 may be a target protease for Spink3. However, recombinant KLK1 showed the slightest affinity for rmSpink3 *in vitro*. hSPINK1 is also co-expressed with KLK3 and KLK2 in human prostate, and with KLK5 in uterus. As KLK2, KLK3 and KLK5 showed the highest affinities for rmSpink3-His₆ *in vitro* (Table I), based on the structural and functional homology between hSPINK1 and hSpink3, it can be speculated that hSPINK1 may contribute to the regulation of these KLKs in vivo.

4- Discussion

Kazal-type serine protease inhibitors are typically small proteins (40 to 80 amino acids long), with a conserved domain architecture including six cysteine residues that form three intra-domain disulfide bridges in a characteristic pattern (CysI-CysV, CysII-CysIV and CysIII-CysVI). These bridges contribute to a tight three-dimensional conformation which consists of one α -helix surrounded by a three stranded β -sheet and loops of peptide segments [42]. The recombinant expression of these proteins in E. coli usually represents an important challenge due to the presence of disulfide bonds, which may affect protein solubility inside the cell and lead to misfolding and subsequent aggregation. We have been working on the expression of recombinant SPINK3 motivated by the different roles that this protein has been attributed, making it a multifunctional protein [7, 12]. In this context, the possibility that the protein from mouse origin might be a natural molecule capable to inhibit human enzymes with

284 pharmaceutical importance was a challenge. This work shows that mouse Spink3, a member of typical 285 Kazal-type inhibitors, was expressed as a soluble protein in E. coli with good yield, and the recombinant protein rmSpink3-His₆ proved to be a potent inhibitor of human KLK5, KLK2 and KLK3. 286 287 KLKs are interesting targets for the control of many human diseases [17]. They are secreted into various 288 biological fluids and increased levels of expression are unquestionable biomarkers, representing a target 289 point for the pharmaceutical industry. 290 The inhibition of the prostate-specific antigen PSA (KLK3) has been extensively studied. It was initially 291 found that 2-azetidinones inhibits active KLK3 [44]. Thereafter, structural modifications based on the 292 Ser-Ser-Lys-Leu-GIn peptide present in semenogelin II, which is a natural specific substrate for KLK3, 293 served as a template to obtain more potent inhibitors with Ki ranging 65 nm to 6.51 µM. Although they 294 had only a marginal effect in reducing development of cancer, labeled KLK3 inhibitors could find 295 application in diagnosis [17]. It was recently found that natural isocoumarins inhibit KLK5 (the KLK 296 involved in peeling skin syndrome) although their inhibition constants are in the micromolar range [17]. 297 Recombinant mSpink3 inhibits KLK5 and KLK3 with Ki= 0.26 ± 0.03 and 1.53 ± 0.12 μ M, respectively, thus 298 it can be considered as a powerful inhibitor of these KLKs. 299 Besides to synthetic molecules, natural peptides are also interesting candidates to control important 300 KLKs [23]. The activity of Kazal-type inhibitors toward KLKs has already been documented in vivo and in 301 vitro. Recently, SPINK6 was isolated from human skin [45] and has been described as a potent inhibitor 302 of KLK12 and KLK13 [41]. However, there is no evidence of the physiological role of these two enzymes, 303 with the exception that KLK12 expression is diminished in breast cancer [46]. On the other hand, 304 SPINK9/LEKTI-2 inhibits KLK5, considered as one of the most important KLKs in skin desquamation with a 305 Ki value similar to that observed in this work for rmSpink3 with the same enzyme [27]. The murine 306 orthologue of SPINK9, Spink12, inhibits trypsin like activity from primary murine keratinocytes, however 307 the authors of this work do not attribute this activity to KLKs [47]. Additionally, fragments of the

multidomain SPINK5/LEKTI-1 can efficiently and specifically inhibit the epidermal kallikreins KLK5, KLK7, and KLK14 [48]. With an experimental approach similar to ours, human plasma kallikrein (KLKB1) was also inhibited in vitro by a non-typical Kazal-type inhibitor, AdKI (2.9 kDa polypeptide) isolated from the Cuban mollusk Aplysia dactylomela being also identified and characterized [49]. Regarding the inhibitory activity of SPINK1 on kallikreins, it has been reported that PSTI from human gastric mucose was not effective against pancreatic porcine kallikrein [8], however this kallikrein is referred by MEROPS database as KLKB1 [50], which was not assayed by us. Compared to this previous reports for the inhibition of KLKs by Kazal type inhibitors, our results are Kis higher than the ones reported for SPINK6 [41] or SPINK5/LEKTI fragments [51], meaning less efficient inhibition. However, our inhibition constant values correspond well to those reported for Kazal type inhibitors (10⁻⁷–10⁻¹³ M) [16]. The control of excessive activity of KLKs is a result of the balance between these enzymes and their specific inhibitors. Thus, it is hypothesized that in diseases in which the expression of KLKs is increased, a compensatory raise in the expression of inhibitors should occur. In this sense, mSpink3 is believed to have essential roles in the maintenance of integrity and regeneration of acinar cells, since Spink3-/- mice showed enhanced trypsin activity [10] and autophagic degeneration of acinar cells but no signs of apoptosis [3]. The human orthologue SPINK1 is thought to function in the prevention of trypsincatalyzed premature activation of zymogens within the pancreas and the pancreatic duct [9] and mutations in this gene are associated with hereditary pancreatitis [52]. Its expression is augmented in breast cancer [53] and because of that it has been recently proposed as an outlier marker (i.e., a protein with enhanced expression in a subset of cases) in prostate cancer [54, 55]. This increased expression might probably be a mechanism to control the increased activity of KLK3 [56]. Spink3 was also found on the mouse sperm surface bound to an unknown ligand [12, 15] from where it detaches within the female duct and was reported to inhibit a trypsin-like activity from uterine fluid [15].

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Regardless of the evidences cited above, the target proteinases for mSpink3 and hSPINK1 in organs other than pancreas are unknown. Based on the analysis of in silico data one can speculate that mSpink3 might be a potential inhibitor of mKLK1 since both proteins are expressed simultaneously in prostate, pancreas and uterus. This co-expression pattern is consistent with the human orthologues, meaning that there is some functional conservation. Additionally, considering that rmSpink3-His₆ was a potent inhibitor of KLK5 *in vitro* (Table 1) it can be speculated that the human orthologue SPINK1 would inhibit KLK5 in vivo. This is supported by the fact that these proteins are co-expressed in the same tissues (for example, stomach and uterus). Similar extrapolation might be done for SPINK1 and KLK2/KLK3 which are co-expressed in prostate.

In conclusion, our efforts to find novel KLK inhibitors resulted in the efficient production of recombinant Spink3 from mouse seminal vesicle which showed high inhibitory activity on several tissue human kallikreins. New target KLKs for mSpink3 and hSPINK1 are proposed based on in silico analysis of expression; however, this observation needs experimental validation using the right partners in further

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studies.

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6- Glossary

KLK: Kallikrein

355 SPINK: Serine protease inhibitor Kazal-type

356	FRET: Fluorescence resonance energy transfer
357	GST: glutathione S-transferase
358	PSA: Prostate specific antigen
359	KLKB1: Human plasma kallikrein
360	kDa: kilo Dalton
361	His: Histidine
362	Ki: Inhibition constant
363	Km: Michaelis constant
364	PSTI: Pancreatic secretory trypsin inhibitor
365	MALDI: Matrix-assisted laser desorption/ionization
366	EI: Enzyme-inhibitor complex

Figure 1: Recombinant expression and purification of rmSpink3. Electrophoretic profile by SDS-PAGE of the soluble fractions of *E. coli* cells harboring pET-24b (+)-*spink3-His*₆ after 16 h in absence (-IPTG, lane 1) or presence (+ IPTG, lane 2) of 0.1 mM IPTG. rmSpink3-His₆ was purified by HiTrap affinity chromatography (Ni²⁺, lane 3). Protein samples were stained with Coomasie brilliant blue G250 (lanes 1-3). Purified rmSpink3-His₆ was identified by western blotting with anti-SPINK1 (α-SPINK1, lane 4) and anti-His (α-HIS, lane 5) antibodies. The positions of molecular mass markers (kDa) are indicated on the left. A representative gel of three independent experiments is shown.

Figure 2: Inhibition of KLKs by rmSpink3. Recombinant human kallikrein related peptidases (KLKs) were synthesized from an insect cell/baculovirus expression system as described in M&M. The enzymes (5 nM except of KLK7, tryspin and chymotrypsin that were 10 nM) were pre-incubated with increasing concentrations of rmSpink3 for 3 min at room temperature. After incubation, the fluorescence resonance energy transfer (FRET) peptides Abz-KLRSSKQ-EDDnp or Abz-KLFSSKQ-EDDnp were added at 5 μM concentration as indicated. The reaction progress was monitored in the linear range of the enzyme kinetic curve and the residual activity was plotted against rmSpink3 concentration.

Figure 3: Mouse SPINK family. ClustalW2 sequence alignment of mouse SPINK family members including human SPINK1 that was italicized. Each domain of mSPINK5 has been aligned separately. P1 and P1' sites are framed and Cys residues are shaded. Amino acid sequences of SPINK family inhibitors were obtained from the Uniprot databank (http://www.uniprot.org/). Only the portion corresponding to the mature form of the proteins is indicated as predicted by SignalP software (http://www.cbs.dtu.dk/services/ SignalP-2.0/).

Figure 4: Kazal-type mechanism determination using MALDITOF mass spectrometry. Purified rmSpink3 (1 nmol) was incubated with or without Trypsin (100 pmol) in water for 30 minutes. The prepared

Figure 5: Tissue distribution of Spink3, SPINK1 and KLKs expression in mouse and human. Expression data either at the mRNA or protein level were obtained by combining the information obtained from 4 databases (Gene Expression Atlas EMBL-EBI, Unigene NCBI, Ensembl and MGI) (http://www.ebi.ac.uk; http://www.ncbi.nlm.nih.gov; http://www.Ensembl.org; http://www.informatics.jax.org/). Only the assayed human kallikreins (KLKs) and their mouse orthologues (mKLKs) are included. Bold KLKs indicate KLKs inhibited by rmSpink3.

Table 1: Kinetic parameters of KLKs interaction with rmSpink3-His₆

samples were directly analyzed by MALDI-TOF/MS.

Various recombinant KLKs were incubated with increasing concentrations of rmSpink3-His₆ and the residual activity was determined using the corresponding FRET substrate as indicated in Figure 2. The inactivation constants (Ki) were obtained according the Henderson equation for tight-binding inhibitors [36] where [I]t/(1-vi/v0) = Ki (1+S/Km) vi/v0 + [E]t .Data are presented in μ M±SEM between at least 3 replicates. Inset= Calculated isolectric points for recombinant KLKs.

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