

Nomenclature Paper

A comprehensive nomenclature for serine proteases with homology to tissue kallikreins

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Abstract

The human kallikrein locus on chromosome 19q13.3–13.4 contains kallikrein 1 – the tissue kallikrein – and 14 related serine proteases. Recent investigations into their function and evolution have indicated that the present nomenclature for these proteins is inadequate or insufficient. Here we present a new nomenclature in which proteins without proven kininogenase activity are denoted kallikrein-related peptidase. Names are also given to the unique rodent proteins that are closely related to kallikrein 1.

Keywords: cancer; evolution; hormone; inflammation; kininogen.

Introduction

The term kallikrein (derived from Greek, *kallikreas*, for pancreas) was coined by Kraut and colleagues in 1930, when they demonstrated that an earlier described hypotensive substance in urine is present at high concentration in the pancreas (Frey and Kraut, 1926; Kraut et al., 1930). Today, the substance is known as kallikrein 1 or tissue kallikrein (EC 3.4.21.35), an enzyme that generates Lys-bradykinin by specific proteolysis of kininogen 1. There is also a proteolytic enzyme in blood plasma that gives rise to bradykinin that is known as plasma kallikrein (EC 3.4.21.34). Several other proteases also exhibit kallikrein activity, albeit usually less efficiently than the tissue and plasma kallikreins. A recent review of the kallikrein-kinin system is provided by Moreau et al. (2005).

Some 25 years ago, it was shown that mouse and rat salivary glands secrete proteins with homology to tissue kallikrein – at that time known as glandular kallikrein (Bothwell et al., 1979). Owing to their close relationship, glandular kallikrein and its homologs were assigned to a subfamily of serine proteinases, which was named the glandular kallikrein family. Although some of the novel glandular kallikreins displayed potent kallikrein – i.e., kininogenase – activity, they were primarily considered to be involved in prohormone processing, as some of them formed complexes and cleaved precursor proteins of epidermal and nerve growth factors (Thomas et al., 1981; Blaber et al., 1987). A comprehensive analysis showed that there were 24 or 25 glandular kallikrein genes in the mouse genome, designated *mGK-1* to *mGK-25* (Evans et al., 1987). Similar analysis of the rat genome identified 10 or 11 glandular kallikrein genes, denoted *rGK-1* to *rGK-10* (Wines et al., 1989). In a revision of the kallikrein nomenclature, the gene family was renamed the tissue kallikrein gene family and the symbol *GK* was replaced by *KLK*, e.g., the new designation of *rGK-4* was *rKLK4*

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(Berg et al., 1992). In contrast to the large number of murine genes, the human tissue kallikrein family seemed to consist of only three genes, which coded for tissue kallikrein, prostate-specific antigen (PSA) and human glandular kallikrein 1 (hGK-1) – later renamed human kallikrein 2 (hK2) (Fukushima et al., 1985; Lundwall and Lilja, 1987; Schedlich et al., 1987).

The discrepancy in the number of genes was recently explained by comparative studies on the kallikrein locus in mammals (Olsson and Lundwall, 2002; Olsson et al., 2004a,b). These investigations showed that several duplications of the tissue kallikrein gene (*KLK1*) occurred very late in phylogeny and created 23 *KLK1* paralogs that seem to be unique to the mouse and nine *KLK1* paralogs that seem to be unique to the rat. Late duplication of *KLK1* was also observed in the horse, but not in artiodactyls, carnivores, cavian rodents and primates. Another, presumably primate-specific, duplication yielded the hK2 (*KLK2*) and PSA (*KLK3*) genes. A functional gene related to the progenitor of this duplication is present in the dog, whereas in the mouse and rat there is a non-functional pseudogene.

Around the turn of the millennium, investigators identified several genes of simple serine proteases adjacent to the human tissue kallikrein locus on chromosome 19q13.3–13.4 (Gan et al., 2000; Harvey et al., 2000; Yousef et al., 2000). The mutual sequence agreement of the tissue kallikrein family members was higher than the similarity between any of the novel serine protease genes. However, the chromosomal location in combination with overlapping expression in hormone-dependent tissues

suggested both a common ancestry and overlapping functionality. Therefore, the old tissue kallikrein family was expanded with the adjacently located serine proteinases into what has become known as the extended kallikrein family (Yousef and Diamandis, 2001; Borgoño and Diamandis, 2004; Borgoño et al., 2004).

Soon after its discovery, a rational nomenclature was adopted for members of the extended kallikrein family (Diamandis et al., 2000). The human kallikrein family, as we now know it, consists of 15 genes, designated kallikrein 1–15 and denote by the gene symbols *KLK1*–*KLK15*. The nomenclature has served its purpose and is widely accepted by scientists from many different disciplines. However, some shortcomings of the nomenclature have been recognized, as follows:

- The nomenclature was developed for human genes and does not provide names for unique animal genes;
- The term kallikrein was introduced and has been used for decades to identify enzymes with kininogenase activity. Most of the enzymes in the extended kallikrein family are presumed to not display kininogenase activity and thus use of the term kallikrein might be misleading and confusing;
- At present, the nomenclature does not fully comply with the guidelines provided by the human (<http://www.gene.ucl.ac.uk/nomenclature/guidelines.html>) and mouse (<http://www.informatics.jax.org/mgihome/nomen/gene.shtml>) gene nomenclature committees (HGNC and MGNC), e.g., there should not be two separate symbols, such as hK1 and *KLK1*, to depict the protein and the gene.

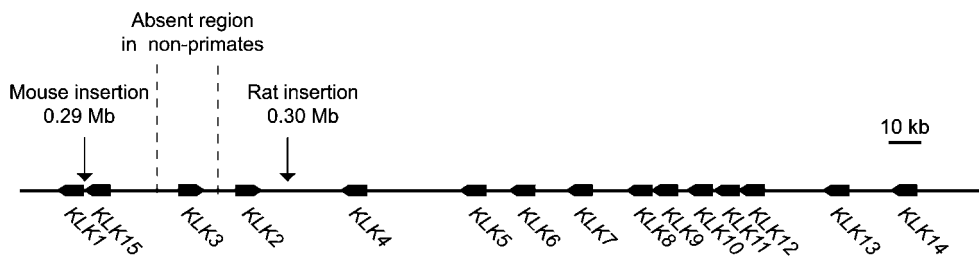


Figure 1 Schematic drawing of the human kallikrein locus on chromosome 19q, 56.0–56.3 Mb. Approximate locations of genes are indicated by their symbols and arrowheads to mark the direction of transcription. The dashed lines surrounding *KLK3* depict the approximate location of the duplicated region that so far has only been detected in primate species. The arrows show the location of expanded regions in murine species that contain closely related *Klk1* paralogs.

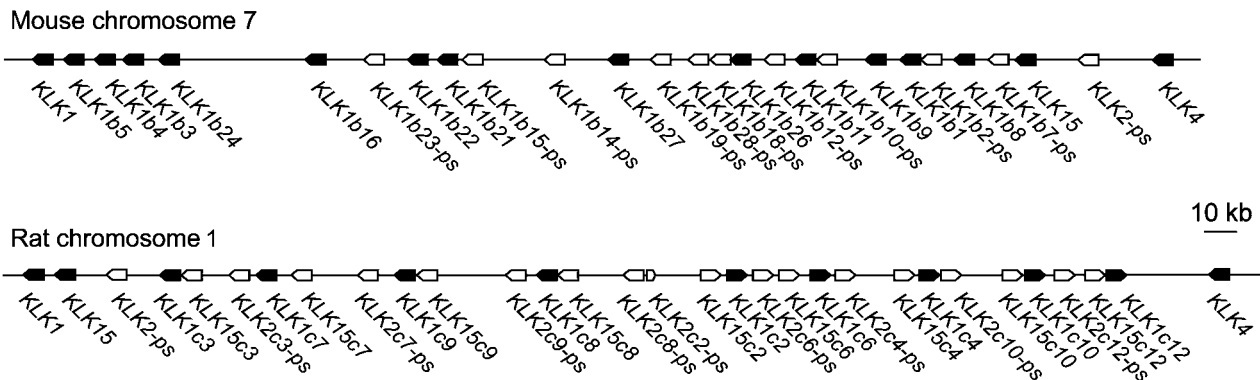


Figure 2 Illustration of the genomic region between *Klk1* and *Klk4*, encompassing taxon-specific genes in the mouse and the rat. Expressed genes are indicated by filled and pseudogenes by empty arrowheads.

Table 1 Proposed new nomenclature for mouse kallikrein 1-related peptidases – family b.

| New gene symbol | Old symbol | New gene name | Alternative and old gene names | GenBank accession no. |
|-------------------|-------------------------------|------------------------------------|---|-----------------------|
| <i>Klk1</i> | <i>mGK-6</i> | Kallikrein 1 | Tissue kallikrein | NM_010639 |
| <i>Klk1b1</i> | <i>mGK-1</i> | Kallikrein 1-related peptidase b1 | | NM_010645 |
| <i>Klk1b2-ps</i> | <i>mGK-2</i> | | | AY152419 |
| <i>Klk1b3</i> | <i>Ngfg, mGK-3</i> | Kallikrein 1-related peptidase b3 | γ subunit of the 7S NGF complex | NM_008693 |
| <i>Klk1b4</i> | <i>Ngfa, mGK-4</i> | Kallikrein 1-related peptidase b4 | α subunit of the 7S NGF complex | NM_010915 |
| <i>Klk1b5</i> | <i>mGK-5</i> | Kallikrein 1-related peptidase b5 | | NM_008456 |
| <i>Klk1b7-ps</i> | <i>mGK-7</i> | | | AY152420 |
| <i>Klk1b8</i> | <i>mGK-8</i> | Kallikrein 1-related peptidase b8 | | NM_008457 |
| <i>Klk1b9</i> | <i>mGK-9, Egfbp3</i> | Kallikrein 1-related peptidase b9 | EGF-BP type C, true EGF-BP | NM_010116 |
| <i>Klk1b10-ps</i> | <i>mGK-10</i> | | | AY152421 |
| <i>Klk1b11</i> | <i>mGK-11</i> | Kallikrein 1-related peptidase b11 | | NM_010640 |
| <i>Klk1b12-ps</i> | <i>mGK-12</i> | | | AY152422 |
| <i>Klk1b14-ps</i> | <i>mGK-14</i> | | | AY152423 |
| <i>Klk1b15-ps</i> | <i>mGK-15</i> | | | AY152424 |
| <i>Klk1b16</i> | <i>mGK-16</i> | Kallikrein 1-related peptidase b16 | γ -renin | NM_008454 |
| <i>Klk1b18-ps</i> | <i>mGK-18</i> | | | AY152426 |
| <i>Klk1b19-ps</i> | <i>mGK-19</i> | | | AY152427 |
| | | | | AY152428 |
| <i>Klk1b21</i> | <i>mGK-21</i> | Kallikrein 1-related peptidase b21 | | NM_010642 |
| <i>Klk1b22</i> | <i>mGk-22, Egfbp1</i> | Kallikrein 1-related peptidase b22 | β -NGF endopeptidase, EGF-BP type A | NM_010114 |
| <i>Klk1b23-ps</i> | <i>mGK-23</i> | | | AY152429 |
| <i>Klk1b24</i> | <i>mGK-24</i> | Kallikrein 1-related peptidase b24 | | NM_010643 |
| <i>Klk1b26</i> | <i>mGK-26, mGK-13, Egfbp2</i> | Kallikrein 1-related peptidase b26 | Prorenin-converting enzyme, EGF-BP type B | NM_010644 |
| <i>Klk1b27</i> | <i>mGK-27</i> | Kallikrein 1-related peptidase b27 | | NM_020268 |
| <i>Klk1b28-ps</i> | <i>mGK-28</i> | | | AY152425 |
| <i>Klk2-ps</i> | <i>mGK-25</i> | | | AY152430 |

Proposed new nomenclature

The organization of the human kallikrein locus is schematically illustrated, with major discrepancies in mouse, rat and dog indicated (Figure 1). The human genes are depicted by the symbols that were introduced in a previous nomenclature paper (Diamandis et al., 2000). The same symbols are used in the new nomenclature, despite the fact that they do not acknowledge the close relationship for *KLK1–KLK3*. However, the gene names are changed for all but *KLK1*, which is still called kallikrein 1. The new names of *KLK2–KLK15* are kallikrein-related peptidase, followed by the number of the gene symbol, e.g., *KLK2* is kallikrein-related peptidase 2. The symbols such as hK1, hK2, etc. previously used to depict the protein should be avoided. To distinguish between the protein and the gene, the former is written in standard font (e.g., KLK2) and the latter in italics (e.g., *KLK2*), as recommended by HGNC. To distinguish a transcript of a gene, the relevant abbreviation is written as a prefix within parentheses, e.g., (mRNA)*KLK2* and (cDNA)*KLK2* to emphasize the message and complementary DNA of the gene for kallikrein-related peptidase 2. If the species needs to be specified, the codes established by SWISS-PROT should be used (<http://www.expasy.ch/cgi-bin/speclist>). The codes are written as a prefix within parentheses, e.g., (HUMAN)*KLK4* and (MOUSE)*Klk4* to distinguish between human and mouse *KLK4*; note that the gene symbols are written in capital letters, with the

exception of the mouse and rat symbols, which are written with an initial capital letter followed by lower case letters. No species-specific prefix is therefore needed in articles only relating to the human and mouse genes.

The proposed new nomenclature for unique kallikrein 1-related peptidases in rodents is according to the proposal by Olsson et al. (2004a). They should be named kallikrein 1-related peptidase followed by a letter depicting the subfamily and the number from the old *GK* nomenclature, e.g., the gene cloned with the designation *mGK-5* has the new gene symbol *Klk1b5* and is called kallikrein 1-related peptidase b5. The proposed new nomenclature for murine kallikrein 1-related peptidases is displayed in Tables 1 and 2, with their location on the chromosome illustrated in Figure 2. The gene subfamilies seem to overlap with single or very closely related animal species, so that the b-family might be confined to *Mus musculus* and the c-family to *Rattus norvegicus* and perhaps also *Rattus rattus*. In non-rodent species, *KLK1* expansion is only known to occur in the horse, where the subfamily is designated by the letter d.

The canine gene with homology to the progenitor of *KLK2* and *KLK3* gives rise to the dog prostate arginine esterase. The proteolytic specificity of this enzyme is similar to that of *KLK2*, but not to that of PSA, which displays an expanded chymotrypsin-like activity (Chapdelaine et al., 1984; Lazure et al., 1984; Malm et al., 2000). Thus, it is proper to assign the symbol *KLK2* to the gene for dog arginine esterase and, as a conse-

Table 2 Proposed new nomenclature of rat kallikrein 1-related peptidases – family c.

| New gene symbol | Old symbols | New gene name | Alternative and old gene names | Associated pseudogenes | Accession no. |
|-----------------|----------------------------------|------------------------------------|---|---|---|
| <i>Klk1</i> | <i>rGK-1, PS</i> | Kallikrein 1 | Tissue kallikrein | | M11563 |
| <i>Klk1c2</i> | <i>rGK-2, RSKG-5, S2, rKLK2</i> | Kallikrein 1-related peptidase c2 | Tonin | <i>Klk15c2-ps</i> <i>Klk2c2-ps</i> | M11565 BK001365 ^a |
| <i>Klk1c3</i> | <i>rGK-3, RSKG-50, S1, rKLK3</i> | Kallikrein 1-related peptidase c3 | | <i>Klk15c3-ps</i> <i>Klk2c3-ps</i> | M11564 BK001366 ^a BK001376 ^a |
| <i>Klk1c4</i> | <i>rGK-4, rKLK4</i> | Kallikrein 1-related peptidase c4 | | <i>Klk15c4-ps</i> <i>Klk2c4-ps</i> | L33839 BK001367 ^a BK001377 ^a |
| <i>Klk1c6</i> | <i>rGK-6, rKLK6</i> | Kallikrein 1-related peptidase c6 | | <i>Klk15c6-p</i> <i>Klk2c6-ps</i> | BK001361 ^a BK001368 ^a BK001378 ^a |
| <i>Klk1c7</i> | <i>RSKG-7, K1, rK7, rKLK7</i> | Kallikrein 1-related peptidase c7 | Esterase B | <i>Klk15c7-ps</i> <i>Klk2c7-ps</i> | M19647 BK001369 ^a BK001379 ^a |
| <i>Klk1c8</i> | <i>rGK-8, P1, rK8, rKLK8</i> | Kallikrein 1-related peptidase c8 | | <i>Klk15c8-ps</i> <i>Klk2c8-ps</i> | M27215 BK001370 ^a BK001375 ^a |
| <i>Klk1c9</i> | <i>S3, rK9, SEV, rKLK9</i> | Kallikrein 1-related peptidase c9 | | <i>Klk15c9-ps</i> <i>Klk2c9-ps</i> | M11566 BK001371 ^a BK001380 ^a |
| <i>Klk1c10</i> | <i>rK10, rKLK10</i> | Kallikrein 1-related peptidase c10 | Endopeptidase k, T-kininogenase, proteinase B, antigen D3b region | <i>Klk15c10-ps</i> <i>Klk2c10-ps</i> | S48142 BK001372 ^a BK001381 ^a |
| <i>Klk1c12</i> | <i>RSKG-3, rKLK12</i> | Kallikrein 1-related peptidase c12 | | <i>Klk15c12-ps</i> <i>Klk2c12-ps</i> | M19648 BK001373 ^a BK001382 ^a |
| <i>Klk2-ps</i> | | | | | BK001374 ^a |

Each functional *Klk1* paralog on the chromosome is followed by associated pseudogenes that are paralogous with *Klk15* and *Klk2*, as illustrated in Figure 2.

^aThird party annotation that has been removed from GenBank, but is still retrievable.

quence, also to the homologous gene in other species, such as the rodent pseudogenes that show equally strong similarity to *KLK2* and *KLK3*.

The nomenclature suggested here is based on our current understanding of genes at the kallikrein locus and may need to be updated in the future as our knowledge widens. If novel genes are discovered, they should have the same stem symbol, but with a novel number (e.g., *KLK16*). Genes created by duplication after the divergence of murine rodents from the lineage leading to primates are exemptions to the rule and should have a name based on the founder gene.

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