

New consensus nomenclature for mammalian keratins

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Keratins are intermediate filament-forming proteins that provide mechanical support and fulfill a variety of additional functions in epithelial cells. In 1982, a nomenclature was devised to name the keratin proteins that were known at that point. The systematic sequencing of the human genome in recent years uncovered the existence of several novel keratin genes and their encoded proteins. Their naming could not be adequately handled in the context of the original system. We propose a new consensus nomenclature for keratin genes and proteins that relies upon and extends the 1982 system and adheres to the guidelines issued by the Human and Mouse Genome Nomenclature Committees. This revised nomenclature accommodates functional genes and pseudogenes, and although designed specifically for the full complement of human keratins, it offers the flexibility needed to incorporate additional keratins from other mammalian species.

What's in a name? That which we call a rose, by any other name would smell as sweet (*Romeo and Juliet*, Act II, Sc. ii).

Keratins (previously also called cytokeratins) are filament-forming proteins of epithelial cells and are essential for normal tissue structure and function. Keratin genes account for most of the intermediate filament genes in the human genome, making up the two largest sequence homology groups, type I and II, of this large multigene family. They are highly differentiation-specific in their expression patterns, implying functional differences. Mutations in most of them are now associated with specific tissue-

fragility disorders, and antibodies to keratins are important markers of tissue differentiation and, therefore, tools in diagnostic pathology. Since the first keratins were sequenced and identified as type I and II intermediate filament proteins, the increasing numbers of keratins has provided an ongoing challenge for their clear identification and logical classification across species.

The first attempt at providing a comprehensive keratin nomenclature dates back to 1982. Moll et al. (1982) used 2D isoelectric focusing and SDS-PAGE to map the keratin profiles of a large number of normal human epithelia, tumors, and cultured cells. They grouped the basic-to-neutral type II keratins as K1–K8 and the acidic type I keratins as K9–K19 (Moll et al., 1982). Although not open-ended for type II keratins, this system has so far proven manageable, as the incorporation of a few novel type II keratins could be accomplished by the addition of discriminatory suffix letters to keratins exhibiting similar gel-electrophoretic properties (Collin et al., 1992a,b; Winter et al., 1998). Moreover, the Moll nomenclature has not been further challenged by the “hard” α -keratins of hair and nail (hair keratins), as these keratins were named Ha (acidic, type I) or Hb (basic to neutral, type II) followed by a number, with H standing for hair (Heid and Franke, 1986; Rogers et al., 1998, 2000). Overall, however, the present naming of keratins has not been systematic, and a reorganized and durable scheme is long overdue.

Genome analyses have recently demonstrated that humans possess a total of 54 functional keratin genes, i.e., 28 type I and 26 type II keratins, forming two clusters of 27 genes each on chromosomes 17q21.2 and 12q13.13 (the gene for the type I keratin K18 being located in the type II keratin gene domain; Hesse et al., 2001, 2004; Rogers et al., 2004, 2005; Table I). Recognition of the extent of this large mammalian gene family led to a suggested revised nomenclature (Hesse et al., 2004) based on an extended Moll system, K1–K8, and K9–K24 (Moll et al., 1982, 1990; Chandler et al., 1991; Zhang et al., 2001; Sprecher et al., 2002), and conceptually close to an earlier proposal (Rogers and Powell, 1993). In this nomenclature, all

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Abbreviation used in this paper: HGNC, Human Genome Nomenclature Committee.

Table I. Human keratin genes

Human keratin genes	Type I genes	Type II genes
Total genes	33	34
Functional genes	28	26
Pseudogenes	5	8
Epithelial keratin genes	17	20
Hair keratin genes	11	6

human type I keratins were named Ka9 to KaX and all type II keratins were named Kb1 to KbY, thus, enabling type I and II keratins of other mammalian species to be added consecutively into this open-ended system. At the 2004 Gordon Conference on Intermediate Filaments in Oxford, an initiative to achieve international consensus led to the formation of a broad-based Keratin Nomenclature Committee that included active investigators in the keratin field and members of the Human Genome Nomenclature Committee (HGNC) and the Mouse Genome Nomenclature Committee. This committee evaluated several potential nomenclature schemes and, after extensive deliberation and consultation with other colleagues in the intermediate filament field, arrived at the consensus nomenclature system that is detailed in the following sections.

To structure the new nomenclature system, the 54 human keratins and their genes are divided into three categories: (1) epithelial keratins/genes, (2) hair keratins/genes, and (3) keratin pseudogenes. The nomenclature is also structured to allow for the inclusion of a fourth category of nonhuman epithelial and hair keratins of other mammalian species, whose genes are either absent or occur as pseudogenes in the human genome.

For both type I and II keratins, these four categories are numerically arranged in the following order (Table II): (1) human epithelial keratins, (2) human hair keratins, (3) nonhuman epithelial/hair keratins, and (4) human keratin pseudogenes.

Because of historical reasons and the extensive number of existing publications, the Moll designation for the epithelial keratins K1–K8 and K9–K24 (Moll et al., 1982, 1990; Chandler et al., 1991; Zhang et al., 2001; Sprecher et al., 2002) is retained, as is the existing HGNC gene designation scheme (i.e., *KRT#*).

New nomenclature for type I keratins

Human epithelial keratins

After the classical type I epithelial keratins K9–K24, the numbers 25–28 were consecutively assigned to the four recently identified epithelial keratins K25irs1–4, which are differentially expressed in the inner root sheath of the human hair follicle (Bawden et al., 2001; Rogers et al., 2004). Thus, the numbers 9–28 represent the 17 human type I epithelial keratins and their genes (Tables I, II, and III [columns 4 and 5]). Numbers 11, 21, and 22 are unused for historical reasons. K11, for example, was originally thought to be a unique gene product (Moll et al., 1982), but was later confirmed as a polymorphic variant of K10 (Korge et al., 1992; Mischke, 1998). K21 (Chandler et al., 1991) has now been shown to be a rat orthologue for human and mouse K20 (Moll et al., 1993; Zhou et al., 2003), and it has therefore been removed from the category of human type I epithelial keratins. Position number 22 was previously reserved by the HGNC, but has never been used.

Table II. Numbering scheme of keratin categories

Category	Number range
Human type I epithelial keratins	9–28
Human type I hair keratins	31–40
Nonhuman type I epithelial and hair keratins	41–70
Human type II epithelial keratins	1–8 and 71–80
Human type II hair keratins	81–86
Nonhuman type II epithelial and hair keratins	87–120
Type II keratin pseudogene	121–220
Type I keratin pseudogenes	221 →

Human hair keratins

For the 11 human type I hair keratins (Table I), positions 29 and 30 were skipped and the keratins were numbered from 31–40 (Table II) to achieve a last digit matching between the current system (Table III, column 1) and the numbering system proposed herein (Table III, column 4; i.e., Ha1 → K31, etc). Note that K33a and K33b (the former Ha3-I and -II) are isoforms.

Nonhuman epithelial and hair keratins

We projected that 30 positions, 41–70 (Table II), i.e., considerably more than those comprising the human type I epithelial keratins, should be sufficient to cover this category of keratins. Pending their characterization at the gene/protein and tissue expression level, any new members of this type shall be added chronologically, independent of their nature as epithelial or hair keratins. Presently, there are only two keratins that fulfill these criteria (Table III). K41 represents a hair keratin that is differentially expressed in chimpanzee and gorilla hairs, whereas the human orthologue is a pseudogene (Winter et al., 2001). Similarly, K42 is the new designation for the recently described mouse and rat epithelial keratin K17n that is expressed mainly in nail tissue (Tong and Coulombe, 2004). The orthologous gene in humans is also a pseudogene (Trojanovsky et al. 1992; Hesse et al., 2004; Rogers et al., 2004; Tong and Coulombe, 2004). Another candidate for this group is the keratin currently called Ka11, which is a rat keratin whose gene is nonfunctional in mice and absent from the human genome (Hesse et al., 2004).

New nomenclature for type II keratins

Human epithelial keratins

Compared with the classical type I epithelial keratins, significantly more adjustments had to be made within the family of type II epithelial keratins K1–K8 (Table IV). The numbering of additional type II epithelial keratins begins with K71, after the nonhuman mammalian type I keratins (Table II). Thus, K71–K74 were assigned to the four type II inner root sheath keratins K6irs1–4 (Langbein et al., 2003; Table IV).

The number of distinct variants of the *K6* gene has been a matter for some discussion, but sequencing of the human genome has revealed that there are, in fact, only three K6 variants, each encoded by their own gene (Rogers et al., 2005). These are now designated K6a (*KRT6A*), K6b (*KRT6B*), and K6c (*KRT6C*). Because of its lack of conformity to the rules of the HGNC, the hair follicle-specific keratin K6hf (Winter et al., 1998; Wang et al., 2003) has been repositioned after the last inner root sheath

Table III. Type I keratins

Current protein designation	Current gene designation	Protein designation according to Hesse et al., 2004 ^a	New protein designation	New gene designation
Human epithelial keratins				
K9	<i>KRT9</i>	Ka9	K9	<i>KRT9</i>
K10	<i>KRT10</i>	Ka10	K10	<i>KRT10</i>
K12	<i>KRT12</i>	Ka12	K12	<i>KRT12</i>
K13	<i>KRT13</i>	Ka13	K13	<i>KRT13</i>
K14	<i>KRT14</i>	Ka14	K14	<i>KRT14</i>
K15	<i>KRT15</i>	Ka15	K15	<i>KRT15</i>
K16	<i>KRT16</i>	Ka16	K16	<i>KRT16</i>
K17	<i>KRT17</i>	Ka17	K17	<i>KRT17</i>
K18	<i>KRT18</i>	Ka18	K18	<i>KRT18</i>
K19	<i>KRT19</i>	Ka19	K19	<i>KRT19</i>
K20	<i>KRT20</i>	Ka20	K20	<i>KRT20</i>
K21 rat	no designation	—	—	—
K23	<i>KRT23</i>	Ka23	K23	<i>KRT23</i>
K24	<i>KRT24</i>	Ka24	K24	<i>KRT24</i>
K25irs1 ^b , K10C ^c , hIRSa1 ^d	<i>KRT25A</i>	Ka38	K25	<i>KRT25</i>
K25irs2 ^b , K10D ^c	<i>KRT25B</i>	Ka39	K26	<i>KRT26</i>
K25irs3 ^b , K10B ^c , hIRSa3.1 ^d	<i>KRT25C</i>	Ka40	K27	<i>KRT27</i>
K25irs4 ^b , hIRSa2 ^d	<i>KRT25D</i>	Ka41	K28	<i>KRT28</i>
Human hair keratins				
Ha1	<i>KRTHA1</i>	Ka25	K31	<i>KRT31</i>
Ha2	<i>KRTHA2</i>	Ka26	K32	<i>KRT32</i>
Ha3-I	<i>KRTHA3A</i>	Ka27	K33a	<i>KRT33A</i>
Ha3-II	<i>KRTHA3B</i>	Ka28	K33b	<i>KRT33B</i>
Ha4	<i>KRTHA4</i>	Ka29	K34	<i>KRT34</i>
Ha5	<i>KRTHA5</i>	Ka30	K35	<i>KRT35</i>
Ha6	<i>KRTHA6</i>	Ka31	K36	<i>KRT36</i>
Ha7	<i>KRTHA7</i>	Ka32	K37	<i>KRT37</i>
Ha8	<i>KRTHA8</i>	Ka33	K38	<i>KRT38</i>
		Ka35	K39	<i>KRT39</i>
		Ka36	K40	<i>KRT40</i>
Nonhuman hair/epithelial keratins				
	φ hHaA ^b , Ka34P ^a man		K41 chimp, gorilla	<i>KRT41</i> ^a chimp, gorilla <i>KRT41P</i> man
K17n mouse, rat ^f	no designation	Ka22 mouse, rat	K42 mouse, rat	<i>KRT42</i> mouse, rat <i>KRT42P</i> man
Human keratin pseudogenes				
	φ hHaB ^b			<i>KRT221P</i>
	φ KRT1 ^b , Ka21P ^a			<i>KRT222P</i>
	φ KRT1 ^b , Ka37P ^a			<i>KRT223P</i>

^aHesse et al., 2004; ^bRogers et al., 2004; ^cHesse et al., 2001; ^dBawden et al., 2001; ^eWinter et al., 2001; ^fTong and Coulombe, 2004.

keratin K74 and so renamed K75 (Table IV). For similar reasons of identity and nomenclature conformity, the epidermal keratin K2e has been redesignated K2, and the palatal keratin K2p (Collin et al., 1992a,b) has been renamed K76 (Table IV, column 4).

Although the completion of sequencing and analysis of the human type I keratin gene domain did not reveal any new keratin genes, that of the type II keratin gene domain led to the detection of four hitherto unknown genes whose encoded proteins had previously been designated K1b, K5b, K6l, and Kb20 (Hesse et al., 2001, 2004; Rogers et al. 2005; Table IV, column 1). Although there is only limited expression data available for K5b, K6l, and Kb20 (Rogers et al., 2005), keratin K1b has recently been demonstrated to be specifically expressed in eccrine

sweat glands (Langbein et al., 2005). In the new nomenclature, these keratins are designated K77–K80 (Table IV, column 4). Collectively, K1–K8 and K71–K80 cover the twenty human type II epithelial keratins (Tables I and II).

Human hair keratins

Somewhat by design, the direct continuation of the numbering of the six human hair keratins (Table I), i.e., K81–K86, led to last digit matching between the old (Table IV, column 1) and the new nomenclature (Table IV, column 4), so that this kind of “aide-mémoire” exists for both types of human hair keratins.

Nonhuman epithelial/hair keratins

Unlike the situation for type I keratins, there are currently no type II keratins in this category. According to a recent study,

Table IV. Type II keratins

Current protein designation	Current gene designation	Protein designation according to Hesse et al., 2004 ^a	New protein designation	New gene designation
Human epithelial keratin				
K1	<i>KRT1</i>	Kb1	K1	<i>KRT1</i>
K2e	<i>KRT2A</i>	Kb2	K2	<i>KRT2</i>
K3	<i>KRT3</i>	Kb3	K3	<i>KRT3</i>
K4	<i>KRT4</i>	Kb4	K4	<i>KRT4</i>
K5	<i>KRT5</i>	Kb5	K5	<i>KRT5</i>
K6a	<i>KRT6A</i>	Kb6	K6a	<i>KRT6A</i>
K6b	<i>KRT6B</i>	Kb10	K6b	<i>KRT6B</i>
K6e/h, ^{a,b}	no designation	Kb12	K6c ^c	<i>KRT6C</i>
K7	<i>KRT7</i>	Kb7	K7	<i>KRT7</i>
K8	<i>KRT8</i>	Kb8	K8	<i>KRT8</i>
K6irs1	no designation	Kb34	K71	<i>KRT71</i>
K6irs2	no designation	Kb35	K72	<i>KRT72</i>
K6irs3	no designation	Kb36	K73	<i>KRT73</i>
K6irs4	no designation	Kb37	K74	<i>KRT74</i>
K6hf	no designation	Kb18	K75	<i>KRT75</i>
K2p	<i>KRT2B</i>	Kb9	K76	<i>KRT76</i>
K1b ^d	no designation	Kb39	K77	<i>KRT77</i>
K5b ^d	no designation	Kb40	K78	<i>KRT78</i>
K6l ^d	no designation	Kb38	K79	<i>KRT79</i>
Kb20 ^{a,b}	no designation	Kb20	K80	<i>KRT80</i>
Human hair keratins				
Hb1, K2.9 ^e	<i>KRTHB1</i>	Kb21	K81	<i>KRT81</i>
Hb2	<i>KRTHB2</i>	Kb22	K82	<i>KRT82</i>
Hb3, K2.10 ^e	<i>KRTHB3</i>	Kb23	K83	<i>KRT83</i>
Hb4	<i>KRTHB4</i>	Kb24	K84	<i>KRT84</i>
Hb5, K2.12 ^e	<i>KRTHB5</i>	Kb25	K85	<i>KRT85</i>
Hb6, K2.11 ^e	<i>KRTHB6</i>	Kb26	K86	<i>KRT86</i>
Nonhuman epithelial/ hair keratins				
—	—	—	—	—
Human keratin pseudogenes				
	φ hHbD ^b , Kb31P ^a			<i>KRT121P</i>
	φ hHbC ^b , Kb30P ^a			<i>KRT122P</i>
	φ hHbB ^b , Kb29P ^a			<i>KRT123P</i>
	φ hHbA ^b , Kb28P ^a			<i>KRT124P</i>
	φ KRTH ^b			<i>KRT125P</i>
	φ KRTG ^b , Kb19P ^a			<i>KRT126P</i>
	φ KRTF ^b ,			<i>KRT127P</i>
	φ KRTE ^b ,			<i>KRT128P</i>

^aHesse et al., 2004; ^bRogers et al., 2005; ^cNot identical with the polymorphic variant described previously by Takahashi et al., 1995; see Rogers et al., 2005 for further information; ^dHesse et al., 2001; ^ePowell and Beltrame, 1994.

putative mouse and rat candidate genes might be located in a region syntenic to the *KRT6B*, *KRT6C*, and *KRT6A* subcluster in the human genome (Hesse et al., 2004). The numbering allocated to such potential keratins ranges from 87 to 120 (Table II), thus, covering approximately the same range as that assigned to their type I counterparts.

New nomenclature for human type I and II pseudogene

Human keratin pseudogenes of both types were also included in the new nomenclature system, to give all-inclusive schemes of the type I and II keratin gene chromosomal domains, as

shown in Fig. 1. Whereas the slots reserved for type II pseudogenes extend from 121–220, type I pseudogenes have an open system starting with 221 (Table II), as this is the last category to be named.

Presently, there are eight keratin pseudogenes designated *KRT121P–KRT128P* located within the type II keratin gene domain on chromosome 12q13.13 (Table IV, column 5). Of these, *KRT121P–KRT124P* are hair keratin pseudogenes and *KRT125P–KRT128P* are epithelial keratin pseudogenes (Fig. 1). Should some of these pseudogenes have active counterparts in other species, the encoded keratins will retain the numbering of the respective pseudogene without the suffix “P,” and,

A chromosome 17q21.2

centromere ...*KRT222P* *KRT24* *KRT223P* *KRT25* *KRT26* *KRT27* *KRT28* *KRT10* *KRT12* *KRT20* *KRT23*
KRT39 *KRT40* *KRT33A* *KRT33B* *KRT34* *KRT31* *KRT41P* *KRT37* *KRT38* *KRT221P* *KRT32* *KRT35*
KRT36 *KRT13* *KRT15* *KRT19* *KRT9* *KRT14* *KRT16* *KRT17* *KRT42P*....telomere

B chromosome 12q13.13

centromere...*KRT80* *KRT7* *KRT121P* *KRT122P* *KRT81* *KRT86* *KRT23* *KRT123P* *KRT85* *KRT84* *KRT82*
KRT124P *KRT75* *KRT6B* *KRT6C* *KRT6A* *KRT5* *KRT71* *KRT74* *KRT72* *KRT73* *KRT125P* *KRT2* *KRT1*
KRT77 *KRT126P* *KRT127P* *KRT128P* *KRT76* *KRT3* *KRT4* *KRT79* *KRT78* *KRT8* *KRT18*¹....telomere

depending on their type, will be included in the respective category of nonhuman epithelial/hair keratins. The positions above 128 can be used to include keratin pseudogenes identified in other mammals.

The type I keratin gene domain contains two hair keratin and three epithelial keratin pseudogenes (Table III and Fig. 1). Two of them, the hair keratin pseudogene *KRT41P* and the epithelial keratin pseudogene *KRT42P*, possess active gene counterparts in other species and have already been named accordingly. The remaining three genes were designated *KRT221P*–*KRT223P* (Table III, column 5).

It should be emphasized that in addition to the aforementioned type I and II keratin pseudogenes in the keratin gene domains on chromosomes 17 and 121, there are at least 61 processed pseudogenes for the type II keratin K8 and 77 for the type I keratin K18, which are dispersed throughout the human genome. Moreover, there are five processed pseudogenes for the type I keratin K19, which are single pseudogenes located on chromosomes 4, 6, and 10, with two pseudogenes located on chromosome 12. None of these contains an intact reading frame. Furthermore, the terminal segment on the human type I keratin gene domain spanning genes *KRT14*, *KRT16*, *KRT17*, and *KRT42P* (Fig. 1) is inserted four times into different regions of chromosome 17. This gives rise to three unprocessed pseudogenes for K14 and K16, and four for K17, as well as four *KRT42P* pseudogenes, which are all assumed to be nonfunctional (Hesse et al., 2001, 2004). The decision as to how these pseudogenes will be included into the respective lists of type I and II keratin pseudogenes will be left to the HGNC.

Concluding remarks

This modified and unifying nomenclature for mammalian keratins preserves the widely used and broadly referenced Moll designation system for the classical human epithelial keratins K1–K8 and K9–K24. The few changes that have been introduced reflect constraints to restrict the protein designation type K#\$ (# = a number; \$ = a letter) to true keratin isoforms. Accordingly, the new nomenclature will have no significant impact on current textbooks and commercial catalogs. Major changes are nearly all restricted to recently identified epithelial keratins. However, laboratories working on hair keratins will have to, under the new system, part with the 20-yr-old designations HaX and HbY and adopt new nomenclature. An effort has therefore been made to preserve the last digit between the old and the new designations, which is hoped will help researchers to adapt the new names. Although the new nomenclature assembles the human type I keratins into an almost uninterrupted series by preserving the original Moll nomenclature where

Figure 1. Human keratin gene domains. (A) Type I. (B) Type II. Red, epithelial keratin genes; black, hair follicle-specific epithelial keratin genes; blue, hair keratin genes; green, pseudogenes.¹ *KRT18* is a type I gene.

possible, there is an unavoidable gap in the numbering of human type II keratins. Sufficient space has been left in the system for keratins occurring in other mammalian species, as well as for keratin pseudogenes. Moreover, we suggest that the term “keratin” rather than “cytokeratin” be used and that mammalian orthologues of human keratins be given the same naming system. As the revised nomenclature should facilitate communication and understanding within the community interested in keratins and their diseases, we advocate that this new system be used in all future studies.

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