## **REVIEW**



# A systematic nomenclature for mammalian tropomyosin isoforms

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**Abstract** Tropomyosin, a ubiquitous protein in animals and fungi, is associated with the actin cytoskeleton and is involved with stabilising actin filaments and regulating the interaction of the filament with other actin binding proteins. The protein is best known for its role in regulating the interaction between actin and myosin in muscle contraction but in recent years its role as a major player in the organisation and dynamics of the cytoskeleton has been increasingly recognised. In mammals Tpm is expressed from four distinct genes and alternate splicing of each gene can produce a total of up to 40 different mRNA variants most of which are expressed as proteins. We are expecting a renaissance in the study of tropomyosins as the roles of these different isoforms are beginning to be deciphered. However, it is our belief that such a renaissance is being limited by confusion over the naming systems for the tropomyosin isoforms. These result in even experienced workers struggling to reconcile work done in different laboratories and at different times. We propose here a systematic nomenclature for tropomyosin based on the best current practice. We recommend the adoption of these

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names and a cross-reference to the table of alternate names and accession numbers for protein sequences is included here. The National Center for Biotechnology Information (NCBI) website has been amended to include the nomenclature for the human, mouse and rat genes.

**Keywords** Cytoskeleton · Actin binding protein · Muscle thin filament

Tropomyosin (Tpm) is a protein family associated with the stabilisation and regulation of the actin cytoskeleton (Gunning et al. 2008). It is expressed in opisthokonts including animals and fungi, but is not documented in plants, protists or procaryotes (Barua et al. 2011; Cranz-Mileva et al. 2013). Tpm polymerises head to tail as a twochained coiled coil along both sides of a helical actin filament stabilising the actin structure and interacting with a range of other actin binding proteins to regulate the actin cytoskeleton. The protein was discovered in the 1940s as a component of the actin filaments of striated muscle (Bailey 1946) and this remains the best studied example of Tpm function. In striated muscle, together with troponin, it enables calcium regulation of striated muscle contraction by blocking  $(-Ca^{2+})$  or allowing  $(+Ca^{2+})$  myosin access to its binding site on actin (Gordon et al. 2000; Geeves 2012). Since the 1940s the different possible roles of Tpm and the number of interacting partners has multiplied (see special issue of J Muscle Res. Cell Motil., Marston & Gautel 2013). We now know that in mammalian cells Tpm is expressed from four separate genes and alternate promoter selection and splicing of each gene can produce up to 40 different mRNA variants most of which have been shown to be expressed as proteins in different tissues (Pittenger et al. 1994; Vindin and Gunning 2013).



The steady discovery of different splice variants in specific tissues has, over the years, led to a proliferation of different names for the isoforms, some identified with specific tissues but later found with a broader distribution. This has resulted in a great deal of confusion in the literature and it is often difficult, even for those of us in the field, to be sure which isoforms we are each referring to. It must be an even bigger nightmare for those trying to enter the field. The lack of a coherent nomenclature is an impediment to progress. We firmly believe that Tpm research is due for a renaissance from both the structurefunction perspective in muscle research and the cytoplasmic perspective in the localisation-function of distinct isoforms and their binding partners. This is evidenced by the collection of articles published in a special issue of J Muscle Res Cell Motil devoted to Tpm Marston & Gautel (2013). This renaissance will be enhanced if we can establish a widely agreed-upon nomenclature that the majority of active researchers will use. It will be much better if we can establish this now as the field begins to expand and other pressures develop on the nomenclature.

After consulting colleagues working with Tpm we now make a specific proposal for how the nomenclature could be simplified and rationalised and to make it compatible with current best practice for gene and protein names. The proposal keeps as close as possible to some well-established historical precedents.

Our suggestion is that any publication dealing with the tropomyosin protein should designate, at the first use, the formal name of the protein isoform as proposed here, and then define the short form or abbreviation that will be used thereafter in the article text. Reference to Tables 1, 2, 3 and 4 here will allow cross-reference to any earlier versions of the names used e.g.,  $\alpha$ -fast skeletal tropomyosin is Tpm1.1.

# **Detailed proposals**

- 1. The human tropomyosin genes should be known as *TPM1* through *TPM4* (*Tpm1* through *Tpm4* for mouse and rat tropomyosin) to be consistent with other gene nomenclatures.
- 2. The protein short name Tm is historically well established but Tpm is consistent with standard protein nomenclature and is therefore preferred in the formal name.
- 3. Tpm1 or Tpm2 specifies the protein is from gene *TPM1* or *TPM2* etc.
- 4. Alternate isoforms are numbered systematically from one (e.g., Tpm1.1 etc.), keeping as close as possible to

- any precedent. Thus Tpm2.3 is the tropomyosin isoform from gene 2 isoform 3.
- In the formal name a subscript designates the tissue most closely associated, historically, with the protein isoform. Many such isoforms have been since found in multiple tissues and such designations are not always useful.
  - St striated muscle, cardiac/skeletal tissue
  - Sm smooth muscle,
  - Br brain
  - Cy other cytoplasmic
- 6. The splicing of the four exons that vary in vertebrates can be designated by a four letter code; a.a.b.d indicates exon 1 is splice form a, exon 2 is splice form a, exon 6 is splice form b and exon 9 is splice form d.
  - A dash as in b.-.b.d indicates that exon 2 is missing and therefore this is a short form of Tpm.
  - Previous publications of Tpm isoforms have indicated that exon 9 can be a combination of two splice forms at the mRNA level (e.g., Vindin and Gunning 2013) but these reflect splice variants in non-coding regions at the 3' end of the mRNA transcripts. We have therefore omitted such isoforms from our list of expressed proteins.
- 7. The formal names of the two tropomyosins from the *TPM1* gene, found in the contractile filaments of smooth muscle, are then

$$Tpm1.3_{Sm(a,a,b,d)}$$
 and  $Tpm1.4_{Sm(a,a,a,d)}$ 

The short form of the name would be Tpm1.3 and Tpm1.4 Tables 1, 2, 3 and 4 list the current, known Tpm protein isoforms and the proposed formal names. As far as possible the alternate names for the same protein that have been used in the past are included in columns two and three. We also include, in Table 5, the full set of human Tpm amino acid sequences in the order that the exons appear in the gene. This will allow any ambiguity in the literature to be checked against the names and exon sequences used here. A fuller list of animal Tpm sequences can be found in Barua et al. 2011 and Schevzov et al. 2011.

It is almost certain that additional isoforms of tropomyosin will be identified in these species. Consideration of all possible splicing combinations already established in at least one isoform leads to a calculation of 48 possible distinct isoforms in humans without consideration of currently unknown splicing alternatives. For example, Cooley and Bergtrom (2001) identified an increased number of



**Table 1** List of tropomyosin isoforms derived from the human TPM1 gene (geneID:7168), the mouse *Tpm*1 gene(geneID:22003) or the rat

1a 2a 2b 1b 3 4 5 6a 6b 7 8 9a 9b 9c 9d

Short name	Common names	Alternate names in use	Formal protein name	Accession numbers Full length sequences only <sup>a</sup>	Exon usage <sup>b</sup>
Tpm1.1	Tm skα	αTm, fast skTm , cardiac Tm, striated Tm	Tpm1.1st (a.b.b.a)	Human: NP_001018005.1 Mouse: NP_001157720.1 Rat: NP_001288265.1	1a.2b.3.4.5.6b.7.8.9a 1a.2b.3.4.5.6b.7.8.9a
Tpm1.2	Tm skα1	κTm, Tmskα1-1	Tpm1.2st (a.a.b.a)	Human: NP_001288173.1 Mouse: none Rat: NP_001288271.1	1a.2a.3.4.5.6b.7.8.9a 1a.2a.3.4.5.6b.7.8.9a
Tpm1.3	Tm smα-1	$\alpha$ smooth Tm	Tpm1.3sm (a.a.a.d)	Human: NP_001018020.1 Mouse: none Rat: NP_001029243.1	la.2a.3.4.5.6a.7.8.9d la.2a.3.4.5.6a.7.8.9d
Tpm1.4	Tm smα	Tm6, $\alpha$ smooth Tm Tm $\alpha 1.2$	Tpm1.4sm (a.a.b.d)	Human: NP_001018007.1 Mouse: NP_001157721.1 Rat: AAA21804.1	1a.2a.3.4.5.6b.7.8.9d 1a.2a.3.4.5.6b.7.8.9d
Tpm1.5	Tm3-1	_	Tpm1.5cy (a.b.a.a)	Human: NP_000357.3 Mouse: NP_001157723.1 Rat: none	1a.2b.3.4.5.6a.7.8.9a 1a.2b.3.4.5.6a.7.8.9a
Tpm1.6	Tm2	_	Tpm1.6cy (a.b.b.d)	Human: NP_001018004.1 Mouse: NP_077745.2 Rat: NP_001029241.1	1a.2b.3.4.5.6b.7.8.9d 1a.2b.3.4.5.6b.7.8.9d
Tpm1.7	Tm3	Tm 1.4	Tpm1.7cy (a.b.a.d)	Human: NP_001018006.1 Mouse: NP_001157722.1 Rat: NP_001029242.1	1a.2b.3.4.5.6a.7.8.9d 1a.2b.3.4.5.6a.7.8.9d
Tpm1.8	Tm5a	-	Tpm1.8cy (bb.d)	Human: NP_001288218.1 Mouse: NP_001157724.1 Rat: NP_001029245.1	1b3.4.5.6b.7.8.9d 1b3.4.5.6b.7.8.9d
Tpm1.9	Tm5b	_	Tpm1.9cy (ba.d)	Human: none Mouse: NP_001157725.1 Rat: NP_001029246.1	1b3.4.5.6a.7.8.9d 1b3.4.5.6a.7.8.9d
Tpm1.10	TmBr1	_	Tpm1.10br (a.b.b.c)	Human: none Mouse: NP_001157727.1 Rat: NP_001029244.1	1a.2b.3.4.5.6b.7.8.9c 1a.2b.3.4.5.6b.7.8.9c
Tpm1.11	TmBr2	_	Tpm1.11br (bb.b)	Human: none Mouse: none Rat: NP_062004.1	1b3.4.5.6b.7.8.9b 1b3.4.5.6b.7.8.9b
Tpm1.12	TmBr3	Tmbrα, Tmbrα-1 Tm α-1.6	Tpm1.12br (bb.c)	Human: NP_001018008.1 Mouse: NP_001157728.1 Rat: NP_001288665.1	1b3.4.5.6b.7.8.9c 1b3.4.5.6b.7.8.9c
Tpm1.13	_	_	Tpm1.13 (bb.a)	Human: none Mouse: NP_001157726.1 Rat: NP_001029247.1	1b3.4.5.6b.7.8.9a 1b3.4.5.6b.7.8.9a

<sup>&</sup>lt;sup>a</sup> The accession numbers are from September, 2014 and for full-length human, mouse and rat protein sequences that have been documented at the nucleic acid level. RefSeq numbers are listed when available. The N-terminal Met is removed post translation in all exon 1b-encoded sequences. As far as is currently known the N-terminal amino acid is N-acetylated post translation in all isoforms. NCBI website has been updated to include the above terminology

<sup>&</sup>lt;sup>b</sup> The exon notation includes only the protein-coding exons and does not reflect splice variants in non-coding regions at the 3' end of the mRNA transcripts



**Table 2** List of tropomyosin isoforms derived from the human TPM3 gene (geneID:7170), the mouse *Tpm3* gene (geneID:59069) or the rat *Tpm3* gene (geneID:117557). Gene structure:

Short name	Common names	Alternate names	Formal protein name	Accession numbers Full length sequences only <sup>a</sup>	Exon usage <sup>b</sup>
Tpm3.12	Tm skα-slow	in use γTm αsTm1,	Tpm3.12st (a.b.b.a)	Human: NP_689476.2	1a.2b.3.4.5.6b.7.8.9a
		slow skTm		Mouse: NP_001280677.2 Rat: NP_001288214.1	1a.2b.3.4.5.6b.7.8.9a
Tpm3.13	_	_	Tpm3.13cy (a.b.a.d)	Human: none	1a.2b.3.4.5.6a.7.8.9d
				Mouse: NP_071709.2	1a.2b.3.4.5.6a.7.8.9d
				Rat: none	
Tpm3.1	Tm5NM1	_	Tpm3.1cy(ba.d)	Human: NP_705935.1	1b3.4.5.6a.7.8.9d
				Mouse: NP_001240667.1	1b3.4.5.6a.7.8.9d
				Rat: NP_775134.1	
Tpm3.2	Tm5NM2	_	Tpm3.2cy(bb.d)	Human: NP_001036816.1	1b3.4.5.6b.7.8.9d
				Mouse: NP_001240669.1	1b3.4.5.6b.7.8.9
				Rat: none	
Tpm3.3	Tm5NM3	-	Tpm3.3cy(bb.a)	Human: EAW53237.1	1b3.4.5.6b.7.8.9a
				Mouse: none	1b3.4.5.6b.7.8.9a
				Rat: NP_001288215.1	
Tpm3.4	Tm5NM4	_	Tpm3.4cy (bb.c)	Human: NP_001036818.1	1b3.4.5.6b.7.8.9c
				Mouse: NP_001280678.1	1b3.4.5.6b.7.8.9c
				Rat: NP_476556.2	
Tpm3.5	Tm5NM5	_	Tpm3.5cy(ba.a)	Human: NP_001265118.1	1b3.4.5.6a.7.8.9a
				Mouse: NP_001258693.1	1b3.4.5.6a.7.8.9a
				Rat: none	
Tpm3.7	Tm5NM7	_	Tpm3.7cy (ba.c)	Human: NP_001036817.1	1b3.4.5.6a.7.8.9c
				Mouse: none	1b3.4.5.6a.7.8.9c
				Rat: none	
Tpm3.8	Tm5NM8	_	Tpm3.8cy (ba.a/c) <sup>c</sup>	Human: none	1b3.4.5.6a.7.8.9a/c
				Mouse: none	1b3.4.5.6a.7.8.9a/c
				Rat: none	
Tpm3.9	Tm5NM9	_	Tpm3.9cy (bb.a/c) <sup>c</sup>	Human: none	1b3.4.5.6b.7.8.9a/c
				Mouse: none	1b3.4.5.6b.7.8.9a/c
				Rat: none	

<sup>&</sup>lt;sup>a</sup> The accession numbers are from September, 2014 and for full-length human, mouse and rat protein sequences that have been documented at the nucleic acid level. RefSeq numbers are listed when available. The N-terminal Met is removed post translation in all exon 1b-encoded sequences. As far as is currently known the N-terminal amino acid is N-acetylated post translation in all isoforms. NCBI website has been updated to include the above terminology

potential isoforms from the *TPM1* gene using RT-PCR. Our ability to detect and confirm additional isoforms will depend on both the extent of expression across, and the levels of expression in, different cell types.

The NCBI accession numbers for each isoform are included in Tables 1, 2, 3 and 4 and the human, mouse and rat genes on the NCBI website have been amended to use the nomenclature as listed here.



b The exon notation includes only the protein-coding exons and does not reflect splice variants in non-coding regions at the 3' end of the mRNA transcripts

<sup>&</sup>lt;sup>c</sup> The designation a/c indicates that the ninth exon includes exon 9a and five amino acids from exon 9c

Table 3 List of tropomyosin isoforms derived from the human TPM2 gene (geneID:7169), the mouse *Tpm*2 gene (geneID:22004) or the rat Tpm2 gene (geneID:500450)<sup>a</sup>. Gene structure

Short name	Common names	Alternate names in use	Formal protein name	Accession numbers Full length sequences only <sup>b</sup>	Exon usage <sup>c</sup>
Tpm2.1	Tm smβ	$\beta Tm\ Tm1$ , $Tm1\beta$ , smooth $Tm$ , smooth $\alpha$ - $Tm$	Tpm2.1sm/cy (a.b.a.d)	Human: NP_998839.1 Mouse: NP_001264805.1 Rat: NP_001019516.1	1a.2b.3.4.5.6a.7.8.9d 1a.2b.3.4.5.6a.7.8.9d
Tpm2.2	Tm skβ	$\beta$ skeletal Tm, $\beta$ cardiac Tm	Tpm2.2st (a.b.b.a)	Human: NP_003280.2 Mouse: NP_033442.2 Rat: NP_001288164.1	1a.2b.3.4.5.6b.7.8.9a 1a.2b.3.4.5.6b.7.8.9a
Tpm2.3		_	Tpm2.3 (a.b.b.d)	Human: NP_001288155.1 Mouse: NP_001264804.1	1a.2b.3.4.5.6b.7.8.9d 1a.2b.3.4.5.6b.7.8.9d
Tpm2.4		-	Tpm2.4 (a.b.a.a)	Rat: none Human: NP_001288156.1 Mouse: none Rat: none	1a.2b.3.4.5.6a.7.8.9a 1a.2b.3.4.5.6a.7.8.9a

<sup>&</sup>lt;sup>a</sup> The TPM2 genes in mouse, rat and human do not contain exon 1b. It is included here as it is found in chicken and zebrafish

**Table 4** List of tropomyosin isoforms derived from the human TPM4 gene (geneID:7171), the mouse *Tpm4* gene (geneID:326618) or the rat gene *Tpm4* (geneID:248512)<sup>a</sup>. Gene structure

1a 2b 1b 3 4 5 6b 7 8 9d

9d

9d

					_
Short name	Common names	Alternate names in use	Formal protein name	Accession numbers Full length sequences only <sup>b</sup>	Exon usage <sup>c</sup>
Tpm4.1	Tm4HMW	$\delta { m Tm}$	Tpm4.1cy (a.b.b.d)	Human: NP_001138632.1 Mouse: none Rat: none	la.2b.3.4.5.6b.7.8.9d la.2b.3.4.5.6b.7.8.9d
Tpm4.2	Tm4	_	Tpm4.2cy (bb.d)	Human: NP_003281.1 Mouse: NP_001001491.1 Rat: NP_036810.1	1b3.4.5.6b.7.8.9d 1b3.4.5.6b.7.8.9d

<sup>&</sup>lt;sup>a</sup> The TPM4 gene of humans (but not rat or mouse) contains exon 9a but there are no reported Tpm4 isoforms expressing exon 9a. Exon 9a is present in other vertebrates e.g., chicken, frog and zebrafish



b The accession numbers are from September, 2014 and for full-length human, mouse and rat protein sequences that have been documented at the nucleic acid level. RefSeq numbers are listed when available. The N-terminal Met is removed post translation in all exon 1b-encoded sequences. As far as is currently known the N-terminal amino acid is N-acetylated post translation in all isoforms. NCBI website has been updated to include the above terminology

<sup>&</sup>lt;sup>c</sup> The exon notation includes only the protein-coding exons and does not reflect splice variants in non-coding regions at the 3' end of the mRNA transcripts

b The accession numbers are from September, 2014 and for full-length human, mouse and rat protein sequences that have been documented at the nucleic acid level. RefSeq numbers are listed when available. The N-terminal Met is removed post translation in all exon 1b-encoded sequences. As far as is currently known the N-terminal amino acid is N-acetylated post translation in all isoforms. NCBI website has been updated to include the above terminology

<sup>&</sup>lt;sup>c</sup> The exon notation includes only the protein-coding exons and does not reflect splice variants in non-coding regions at the 3' end of the mRNA transcripts

Table 5 Human tropomyosin amino acid sequences in order of the exons in the gene

Exon TPM1 TPM2 TPM3 TPM4	-MDAIKKKMQMLKLDKENALDRAEQAEADKKAAEDRSKQ -MDAIKKKMQMLKLDKENAIDRAEQAEADKKQAEDRCKQ MMEAIKKKMQMLKLDKENALDRAEQAEAEQKQAEERSKQ					
Exon						
TPM1	LEEDIAAKEKLLRVSEDERDRVLEELHKAEDSLLAAEEAAAK					
TPM2						
Exon	2b					
TPM1	LEDELVSLQKKLKGTEDELDKYSEALKDAQEK	LELAEKKATD				
TPM2	~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~	_				
TPM3	~					
TPM4	VEEELTHLQKKLKGTEDELDKYSEDLKDAQEK	LELTEKKASD				
Exon	1b					
TPM1	MAGSSSLEAVRRKIRSLQEQADAAEERAGTLQ	RELDHERKLRET				
TPM2 TPM3	MACTERIA VIVO VI O O O A DDA EED A ED I O	DEVECEDDADEO				
TPM4	MAGITTIEAVKRKIQVLQQQADDAEERAERLQ MAGLNSLEAVKRKIQALQQQADEAEDRAQGLQ					
Exon	3					
TPM1	AEADVASLNRRIQLVEEELDRAQERLATALQK	LEEAEKAADESER				
	AEADVASLNRRIQLVEEELDRAQERLATALQK					
	AEAEVASLNRRIQLVEEELDRAQERLATALQK					
TPM4	AEGDVAALNRRIQLVEEELDRAQERLATALQKLEEAEKAADESER					
Exon	4					
TPM1	~ ~ ~					
TPM2	~ ~					
TPM3	GMKVIENRALKDEEKMELQEIQLKEAKHIAEEADRKYEE					
TPM4	GMKVIENRAMKDEEKMEIQEMQLKEAKHIAEEADRKYEE					
Exon	5					
	VARKLVIIESDLERAEERAELSEG					
	VARKLVILEGELERSEERAEVAES					
	VARKLVIIEGDLERTEERAELAES VARKLVILEGELERAEERAEVSEL					
TPM4	VARKLVILEGELERAEERAEVSEL					
Exon	6a	Exon 6b				
TPM1	QVRQLEEQLRIMDQTLKALMAAEDK	KCAELEEELKTVTNNLKSLEAQAEK				
	RARQLEEELRTMDQALKSLMASEEE	KCGDLEEELKIVTNNLKSLEAQADK				
TPM3 TPM4	RCREMDEQIRLMDQNLKCLSAAEES	KCSELEEELKNVTNNLKSLEAQAEK KCGDLEEELKNVTNNLKSLEAASEK				
Exon	7	Exon 8				
TPM1	YSQKEDRYEEEIKVLSDKLKE	AETRAEFAERSVTKLEKSIDDLE				
TPM2	YSTKEDKYEEEIKLLEEKLKE	AETRAEFAERSVAKLEKTIDDLE				
TPM3	YSQKEDKYEEEIKILTDKLKE	AETRAEFAERSVAKLEKTIDDLE				
TPM4	YSEKEDKYEEEIKLLSDKLKE AETRAEFAERTVAKLEKTIDDLE					



#### Table 5 continued

Exon	9a	Exon 9b
TPM1	DELYAQKLKYKAISEELDHALNDMTSI	DKFLCFTSPKTPSSSWMSHLSELCICLFSS
TPM2	DEVYAQKMKYKAISEELDNALNDITSL	<del>-</del>
TPM3	DELYAQKLKYKAISEELDHALNDMTSI	
TPM4		
Exon	9c	Exon 9d
Exon TPM1	9c DQLYQQLEQNRRLTNELKLALNED	Exon 9d EKVAHAKEENLSMHQMLDQTLLELNNM
TPM1	DQLYQQLEQNRRLTNELKLALNED	EKVAHAKEENLSMHQMLDQTLLELNNM
TPM1 TPM2	DQLYQQLEQNRRLTNELKLALNED	EKVAHAKEENLSMHQMLDQTLLELNNM ETLASAKEENVEIHQTLDQTLLELNNL

The exon sequences are in the order of the coding exons in the genes. Accession numbers used to compile the sequences:

TPM1	NM_001018020.1 NM_001018006.1 NM_000366.5 NM_001018005.1 NM_001018008.1 NM_001018004.1 NM_001018007.1 EL947365, DB542398
TPM2	NM_213674.1 NM_003289.3
TPM3	NM_152263.2 NM_153649.3 NM_001043352.1 NM_001043351.1 NM_001043353.1
TPM4	NM_003290.2 NM_001145160.1

For additional information about tropomyosin sequences and exon organization in animals, refer to Barua et al. 2011 and Schevzov et al. 2011

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