



**THESIS APPROVAL**  
**GRADUATE SCHOOL, KASETSART UNIVERSITY**

Master of Science (Genetic Engineering)

DEGREE

Genetic Engineering

FIELD

Interdisciplinary Graduate Program

PROGRAM

TITLE: *TRPV6* Expression in Mammary Tissues of Pregnant and Lactating Rats

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**THESIS**

***TRPV6* EXPRESSION IN MAMMARY TISSUES OF PREGNANT AND LACTATING RATS**

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**A Thesis Submitted in Partial Fulfilment of  
the Requirements for the Degree of  
Master of Science (Genetic Engineering)  
Graduate School, Kasetsart University  
2006**

**ISBN 974-16-2608-8**

Jirawan Siriwetwivat 2006: *TRPV6* Expression in Mammary Tissues of Pregnant and Lactating Rats. Master of Science (Genetic Engineering), Major Field: Genetic Engineering, Interdisciplinary Graduate Program. Thesis Advisor: Assistant Professor Siriwan Prapong, Ph.D. 69 pages.  
ISBN 974-16-2608-8

Transient receptor potential vanilloid channel 6 (TRPV6) is a highly selective  $\text{Ca}^{2+}$  channel. It plays an important function as gatekeeper of transepithelial  $\text{Ca}^{2+}$  transport in many tissues. The mammary secretory cells during the transition from pregnancy to lactation require essentially large transcellular  $\text{Ca}^{2+}$  fluxes into milk. In this study, the expression profile of TRPV6 mRNA was investigated by using semiquantitative RT-PCR analysis. TRPV6 mRNA expression was found in rat mammary during pregnancy, partum and lactation (-7, -1, 0, +1, +7 and +14). The expression of TRPV6 mRNA increased 5 and 3 folds at +7 and +14 days of lactation periods compared with that at pregnant, partum and early lactating periods ( $p \leq 0.01$ ). Its expression profile was correlated to level of total RNA present in mammary tissue and calcium concentrations in rat milk at various periods.

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## LIST OF ABBREVIATIONS

TRP	=	Transient receptor potential channel
TRPV	=	Transient receptor potential vanilloid channel
CaT1	=	Calcium transporter type 1
ECaC	=	Epithelial calcium channel
PMCA	=	plasma membrane $\text{Ca}^{2+}$ -ATPase
SERCA	=	Sarco(endo)plasmic reticulum $\text{Ca}^{2+}$ -ATPase
SPCA	=	Golgi $\text{Ca}^{2+}$ -ATPase
SOC	=	Store-operated channels
PLC	=	Phospholipase C
$\text{IP}_3$	=	Inositol-1, 4, 5-triphosphate
$\text{IP}_3\text{R}$	=	Inositol-1, 4, 5-triphosphate receptor
DAG	=	Diacylglycerol
GPCR	=	G protein-coupled receptor
$\text{PIP}_2$	=	Phosphatidylinositol 4, 5-bisphosphate
$\mu\text{l}$	=	Micro-litre
$\mu\text{g}$	=	Micro-gram
M	=	Molar
$^{\circ}\text{C}$	=	Degree Celsius



# **TRPV6 EXPRESSION IN MAMMARY TISSUES OF PREGNANT AND LACTATING RATS**

## **INTRODUCTION**

Rat transient receptor potential vanilloid channel 6 (rTRPV6), renamed after CaT1 or ECaC2 (Montell *et al.*, 2002), have been cloned from rat and shown to mediate calcium influx when expressed in *Xenopus* oocytes (Peng *et al.*, 1999). This channel is all or part of the  $\text{Ca}^{2+}$  release-activated  $\text{Ca}^{2+}$  channel (CRAC), which its major characteristic is an activation following store depletion (Petersen and Fedirko, 2001). The TRPV6 is a highly selective calcium channel belonging to the transient receptor potential vanilloid channel (TRPV) family (Benham *et al.*, 2002). Its structure is similar to a  $\text{Ca}^{2+}$  channel TRPV5's structure. TRPV6 shares 75.6% amino acid sequence identity with TRPV5 which is one of the first members of the TRP superfamily of ion channels (Peng *et al.*, 2001a). TRPV6 activity is  $[\text{Ca}^{2+}]_i$ - (Bodding, 2005; Bodding and Flockerzi, 2004) and pH-sensitive (Hirnet *et al.*, 2003). It is activated by low levels of  $[\text{Ca}^{2+}]_i$  (Bodding, 2005; Bodding and Flockerzi, 2004) and it is inactivated by high  $[\text{Ca}^{2+}]_i$  (Bodding, 2005; Bodding and Flockerzi, 2004) and by low pH (Hirnet *et al.*, 2003). The TRPV6 channel is the principal mechanism for  $\text{Ca}^{2+}$  entry into enterocytes as part of the transcellular pathway of calcium absorption in the intestine (Hoenderop *et al.*, 1998). Its apical location on membrane of mouse and human gastrointestinal tract was reported (Zhuang *et al.*, 2002). This supported its physiological function in intestinal calcium absorption (Barley *et al.*, 2001).

Milk is a calcium-rich fluid which the mammary gland during lactation extracts large quantities of calcium from plasma to meet the requirements of the growing neonate. The plasma calcium significantly decreased on one day before parturition which occurred in conjunction with the rat's preparation to give birth and initiate lactation (Reinhardt and Horst, 1999). Calcium is secreted into milk in several chemical forms such as free ionized calcium, casein-bound calcium, and calcium complexed to inorganic anions (Neville *et al.*, 1994). Several calcium transporters have been reported in mammary tissue, for example PMCA1-4 (Reinhardt and Horst, 1999), SPCA (Reinhardt *et al.*, 2000), SERCA (Reinhardt and Horst, 1999). A calcium transport from plasma component to milk compartment required a high capacity mechanism rather than only a passive transport

mechanism. The high capacity mechanism of  $\text{Ca}^{2+}$  uptake across the basolateral membrane has not been precisely identified, however it could be through a  $\text{Ca}^{2+}$  channel mechanism. In addition, a stretch-activated  $\text{Ca}^{2+}$  channel may also provide a pathway for  $\text{Ca}^{2+}$  uptake (Shennan and Peaker, 2000).

Since the TRPV6 is a  $\text{Ca}^{2+}$ -selective channel displaying constitutive activity and saturable kinetics in the submillimolar range (Peng *et al.*, 1999; Lambers *et al.*, 2004), it would be an ideal facilitator of mammary epithelial  $\text{Ca}^{2+}$  transport during lactation. However, this channel has not been widely studied in the mammary gland, though its expression has been reported in mammary tumor (Zhuang *et al.*, 2002). Therefore, this TRPV6  $\text{Ca}^{2+}$ -selective channel is likely to be a candidate  $\text{Ca}^{2+}$  channel for supplying the mammary gland  $\text{Ca}^{2+}$  transport for lactational needs.

This study was designed to examine the expression of TRPV6 mRNA in the normal rat mammary tissue. In addition, this study also examined profile of TRPV6 mRNA expression in rat mammary tissues at various stages of pregnancy, parturition and lactation. Knowledge gained from this study would fulfill a basic physiology of calcium transport in the mammary biology.

## LITERATURE REVIEW

### **The transient receptor potential (TRP) channel family**

Many transient receptor potential (TRP) ion channels have been identified in invertebrates and vertebrates, one in yeast, none in bacteria or plants (Harteneck *et al.*, 2000; Zhou *et al.*, 2003; Montell, 2003). The mammalian transient receptor potential (TRP) ion channels are named after the role of the channels in *Drosophila* phototransduction (Montell *et al.*, 2002; Vennekens *et al.*, 2002). The TRP ion channels are a large family of plasma membrane non-selective and selective cationic channels that are either specifically or ubiquitously expressed in excitable and non-excitable cells (Elliott, 2001). The TRP family of ion channels now comprises more than 30 cation channels (Harteneck, 2003). These proteins have been categorized into seven main subfamilies on the basis of sequence homology; the TRPC (Canonical) subfamily, the TRPV (Vanilloid) subfamily, the TRPM (Melastatin) subfamily, the TRPP (Polycystin) subfamily, the TRPML (Mucolipin) subfamily, the TRPA (Ankyrin) subfamily, and the TRPN (no mechanoreceptor potential C, or NOMPC) subfamily (Birnbaumer *et al.*, 2003). The TRPC (Canonical) and TRPM (Melastatin) subfamilies consist of seven and eight different channels, respectively (i.e. TRPC1-TRPC7 and TRPM1-TRPM8) (Vennekens *et al.*, 2002). The TRPV (Vanilloid) subfamily presently comprises six members (TRPV1-TRPV6) (Gunthorpe *et al.*, 2002). The TRPML (Mucolipin) subfamily comprises three members, and the TRPP (Polycystin) subfamily comprises three members. The TRPA (Ankyrin) has only one mammalian member, TRPA1 (Clapham, 2003). Finally, the TRPN (no mechanoreceptor potential C) subfamily has so far only been detected in *Drosophila*, and zebra fish (Figure 1) (Pederson *et al.*, 2005). The TRP ion channels are gated by responding to temperature, touch, pain, osmolarity, pheromones, taste, and other stimuli (Clapham, 2003; Wissenbach *et al.*, 2004). Although research into TRP ion channels is still in its infancy, they have received particular attention within the field of  $\text{Ca}^{2+}$  signalling because they are the most likely candidates for store-operated  $\text{Ca}^{2+}$  channels (Petersen and Fedirko, 2001). Transient receptor potential vanilloid channel 6 (TRPV6) is an example channel for a highly selective calcium channel of these TRP ion channels (Hirnet *et al.*, 2003).

**Role of the transient receptor potential (TRP) channel family in Phospholipase C-Dependent  $\text{Ca}^{2+}$  Influx (PLC)**

In virtually all eukaryotic cells, the major  $\text{Ca}^{2+}$ -storing, buffering and signalling compartment within cells is endoplasmic/sarcoplasmic reticulum (ER/SR), or derived organelles. In lumen of the ER,  $\text{Ca}^{2+}$  is buffered by proteins with high binding capacity and low affinity, calreticulin, which may have additional signalling functions (Bootman *et al.*, 2002; Zitt *et al.*, 2002; Berridge *et al.*, 2003; Hofmann *et al.*, 2002). Stimulus-induced  $\text{Ca}^{2+}$  released from the ER is thought to be mediated by two main types of  $\text{Ca}^{2+}$  release channels, ryanodine receptor (RyR) and inositol-1, 4, 5-triphosphate receptor ( $\text{IP}_3\text{R}$ ) (Williams, 1998; Petersen *et al.*, 2001; Petersen *et al.*, 2005).

Activation of phospholipase-coupled membrane receptors, G protein-coupled receptor (GPCR) or tyrosine kinase receptor, by hormones, neurotransmitters, growth factors, and other ligands leads to an increase in intracellular  $\text{Ca}^{2+}$  (Mikoshiba, 1997; Hofer and Brown, 2003; Sternfeld *et al.*, 2005). Activation of PLC hydrolyzes phosphatidylinositol 4, 5-bisphosphate ( $\text{PIP}_2$ ) to generate inositol 1, 4, 5-trisphosphate ( $\text{IP}_3$ ) and diacylglycerol (DAG). The increase in intracellular  $\text{Ca}^{2+}$  is initially produced by release of  $\text{Ca}^{2+}$  from intracellular  $\text{IP}_3$ -sensitive stores and is followed by an influx of  $\text{Ca}^{2+}$  from the extracellular space through plasma membrane  $\text{Ca}^{2+}$ -permeable channels (Spassova *et al.*, 2004; Vassilev *et al.*, 2001). The entry of  $\text{Ca}^{2+}$  from the extracellular space is important for refilling intracellular stores. The mechanism for  $\text{Ca}^{2+}$  entry has been studied extensively. In one mechanism, termed “store-operated  $\text{Ca}^{2+}$  entry”, the emptying of intracellular stores activates surface membrane  $\text{Ca}^{2+}$  channels via either diffusible factors or a direct interaction between  $\text{IP}_3$  receptors and surface  $\text{Ca}^{2+}$  channels (Figure 2) (Clapham, 1995; Putney, 2001; Holda *et al.*, 1998; Venkatachalam *et al.*, 2002). These  $\text{Ca}^{2+}$ -permeable channels on the cell surface are called store-operated channels (SOC). Not all  $\text{Ca}^{2+}$  entry after PLC activation is dependent on store depletion.  $\text{Ca}^{2+}$  entry may also be due to activation of surface  $\text{Ca}^{2+}$  channels by other intracellular second messengers, such as DAG activating protein kinase C (PKC) (Clapham, 1995; Hardie, 2003).

However, most TRP channels are non-selectively mediate influx of cations but some members are very selective, for example, TRPM4b and TRPM5 are permeant only to monovalent cations, whereas TRPV5 and TRPV6 are highly selective calcium channels (Montell, 1997).

The  $\text{Ca}^{2+}$  influx channels of the TRP family comprise all TRPCs, all TRPVs, TRPM1, 2, 3, 6, 7, and 8, TRPA1, TRPP2, 3, and 5 and TRPML1, 2, and, 3 (Pedersen *et al.*, 2005). The permeability ratios  $P_{\text{Ca}}/P_{\text{Na}}$  for these channels vary considerably, ranging from 0.3 to  $>100$ . For example, the permeability ratios  $P_{\text{Ca}}/P_{\text{Na}}$  is 0.3 for TRPM2 and  $>100$  for TRPV5 and TRPV6 (Pederson *et al.*, 2005).

### **The transient receptor potential vanilloid (TRPV) channel**

The transient receptor potential vanilloid (TRPV) subfamily, is consisted of 6 members; TRPV1 (VR1), TRPV2 (VRL-1), TRPV3 (VRL-3), TRPV4 (OTRPC4), TRPV5 (ECaC, ECaC1 or CaT2) and TRPV6 (CaT1 or ECaC2) (Gunthorpe *et al.*, 2002; Kedei *et al.*, 2001; Wissenbach *et al.*, 2000; Smith *et al.*, 2002). The TRPV subfamily, on the basis of structure and function, comprises four groups of mammalian TRPVs: TRPV1/TRPV2, TRPV3, TRPV4 and TRPV5/TRPV6 (Peng *et al.*, 2001a). Moreover, TRPV1, TRPV2, TRPV3 and TRPV4 are modestly permeable for  $\text{Ca}^{2+}$  with a permeability ratio  $P_{\text{Ca}}/P_{\text{Na}}$  between  $\sim 3$  and  $\sim 12$  so these channels are non-selective  $\text{Ca}^{2+}$  channels, whereas TRPV5 and TRPV6 are the only highly  $\text{Ca}^{2+}$  selective ion channels ( $P_{\text{Ca}}/P_{\text{Na}} > 100$ ) within the TRPV family (Meir, 2002; Suzuki *et al.*, 2000). The TRPV5 and TRPV6 exhibit a similar ion permeation sequence for divalent cations ( $\text{Ca}^{2+} > \text{Ba}^{2+} \sim \text{Sr}^{2+} > \text{Mn}^{2+}$ ) (Peng *et al.*, 2003b). Like voltage-gate  $\text{Ca}^{2+}$  channels, both of epithelial calcium channels exhibit permeation for monovalent cations in the absence of divalent cations with the permeation sequence of  $\text{Na}^+ \sim \text{Li}^+ > \text{K}^+ > \text{Cs}^+$  (Vennekens *et al.*, 2000). In addition, both channels have recently been shown to have the biophysical characteristics of the  $\text{Ca}^{2+}$ -release activated  $\text{Ca}^{2+}$  channel (CRAC) that refills internal  $\text{Ca}^{2+}$  stores following agonist-elicited release (Yue *et al.*, 2001; Prakriya and Lewis, 2003). The TRPV6 function acts likely as  $\text{Ca}^{2+}$ -sensing  $\text{Ca}^{2+}$  channel to depletion  $\text{Ca}^{2+}$  (Bodding *et al.*, 2002; Schindl *et al.*, 2002). The  $\text{Ca}^{2+}$  channels are inhibited by an increase in the cytosolic  $\text{Ca}^{2+}$  concentration. This negative feedback protects against cytosolic  $\text{Ca}^{2+}$  overloading.  $\text{Ca}^{2+}$  entry therefore only occurs when  $\text{Ca}^{2+}$  can be effectively removed from the inner aspect of the plasma membrane (Prakriya and Lewis, 2003).

All members of the TRPV (Vanilloid) channels family display a membrane topology of six putative transmembrane segments (TM1-TM6), with a re-entrant pore loop (P-loop) between TM5 and TM6, and cytosolic amino and carboxyl termini (Gunthorpe *et al.*, 2002). All TRPV channels form cation channels that are either non-selective or  $\text{Ca}^{2+}$ -selective. TRPV channels have four linked domains of six transmembrane segments so they are thought to form tetramers (Figure 3) (Hellwig *et al.*, 2005).

The TRPV subfamily of calcium-permeable channels is widely distributed in sensory and non-sensory cells from nematodes to mammals (Benham *et al.*, 2002). Members of the TRPV subfamily are activated by a diverse range of stimuli, including heat, proton, lipid, phorbols, phosphorylation, changes in extracellular osmolarity and/or pressure, and depletion of intracellular  $\text{Ca}^{2+}$  stores (Table 1) (Benham *et al.*, 2002; O'Neil and Brown, 2003).

### **Transient receptor potential vanilloid channel 6**

Transient receptor potential vanilloid channel 6 (TRPV6) was first identified from rat duodenum using an expression cloning strategy in *Xenopus laevis* (Peng *et al.*, 1999), and formerly named Calcium transporter type 1 (CaT1) (Peng *et al.*, 1999). It is synonymous with Epithelial calcium channel 2 (ECaC2) (Montell *et al.*, 2002). This protein belongs to the TRPV (vanilloid) subfamily, which is one of the seven subfamilies of the transient receptor potential (TRP) ion channels. TRPV6 is highly selective calcium channel which was supposed to mediate calcium absorption in rat intestine (Peng *et al.*, 1999). In 2000, Peng *et al.* reported that a human orthologue (hTRPV6) of rat TRPV6 (rTRPV6) was cloned from small intestine. Moreover, hTRPV6 expression was studied by using Northern blot analysis. It was reported that hTRPV6 transcripts were most abundantly expressed in duodenum at lower levels, in esophagus, stomach and jejunum, and the signal was absent in ileum (Peng *et al.*, 2000a). It shows that TRPV6 constitutes the rate-limiting influx step of transepithelial  $\text{Ca}^{2+}$  transport in the gastrointestinal tract (Barley *et al.*, 2001). Qiu and Hogstrand (2004) reported an orthologue to the mammalian epithelial calcium channel, TRPV6, was cloned and characterized from pufferfish (*Fugu rubripes*). They demonstrated that this gene predates the evolution of land-living vertebrates. The TRPV6 has been proposed to be responsible for apical calcium entry in the vitamin D-regulated transcellular pathway of  $\text{Ca}^{2+}$  absorption (Hoenderop *et al.*, 1998).

### **Genomic of transient receptor potential vanilloid channel 6**

Genes for the TRP ion channels subunits were first defined in the *Drosophila* visual system (Macpherson *et al.*, 2001). In the *trp* mutation, the light response (receptor potential) decays during prolonged exposure to light. TRP-deficient flies are blinded by intense light because sustained  $\text{Ca}^{2+}$  entry via TRP ion channels and subsequent  $\text{Ca}^{2+}$ -dependent adaptation is disrupted (Clapham *et al.*, 2001; Scott and Zuker, 1998).

The *trpv6* gene encoding TRPV6 protein is juxtaposed on rat chromosome 4 (4q22), mouse chromosome 6 (Peng *et al.*, 2001a), and human chromosome 7 (7q33-34) (Peng *et al.*, 2000a). This gene comprises 15 exons, extends over 15.7 kb, and encodes protein of about 730 amino acid residues. Chromosomal organization analysis revealed that two TRPV members are found as direct repeats; TRPV6 follows TRPV5 (Peng *et al.*, 2001a). This duplication appears to be recent that TRPV5 and TRPV6 are more similar to each other than to other members of TRPV subfamily (Birnbaumer *et al.*, 2003). Therefore, it is quite possible that gene duplication events occurred during evolution that gave rise to these two highly homologous pairs of genes. The various gene products have subsequently acquired their function diversities over the course of evolution.

Human TRPV6 (hTRPV6) shares 30% amino acid sequence identity with the other four proteins (TRPV1-4), whereas hTRPV5 and hTRPV6 share 75.6% overall amino acid identity with one another (Peng *et al.*, 2001a). The main sequence differences are located in the N- and C-terminal tails. However, significant differences exist in the N- and C-termini of TRPV5 and TRPV6, which may account for distinct functional and regulatory features (Nijenhuis *et al.*, 2003). The largest exon of TRPV5 and TRPV6 is the region within the TM domains that encodes part of TM5, the pore region, and TM6 (Peng *et al.*, 2001a). Within the sequence encoded by this exon, TM6 is highly conserved in 20 different proteins belonging to various families of channels of divergent structure and function (Birnbaumer *et al.*, 2003). The conservation of TM6 may suggest a role for it as a key structural element for this diverse group of TM proteins (Peng *et al.*, 2001a). However, the detailed structure of the TM1-TM4 segments of TRPV channels is not available; mutagenesis data provided some clues about their functions (Gunthorpe *et al.*, 2002; Clapham *et al.*, 2001). The TM4 segment in voltage changes across the membrane into movement of the helix, somehow gating the pore by twisting this TM4 helix (Clapham *et al.*, 2001). The TRP channels are very weakly voltage-dependent

and lack the full complement of charged amino acids in the TM4 domain (Clapham *et al.*, 2001). Furthermore, the pore regions of TRPV5 and TRPV6 are quite distinct from those of these other channels. The less-conserved pore region, in contrast, may be one determinant of the characteristic  $\text{Ca}^{2+}$ -selectivity of TRPV5 and TRPV6 (Ellinor *et al.*, 1995; Peng *et al.*, 2001a).

### **Structure of transient receptor potential vanilloid channel 6**

In 1999, Peng *et al.* reported that the 2995 base pair of rat TRPV6 cDNA contains an open reading frame of 2181 base pairs that encodes a protein 727 amino acid residues with a predicted relative molecular mass of 83,245 Da. rTRPV6 is polytopic protein containing six transmembrane domains (TMs). A short hydrophobic stretch between TM5 and TM6 was predicted to be the pore-forming region (Peng *et al.*, 1999). The TM2-TM3 loop is a critical determinant of fast  $\text{Ca}^{2+}$ -dependent inactivation in TRPV6 (Nilius *et al.*, 2002). The amino-terminal hydrophilic segment (326 amino acid residues) of TRPV6 contains three ankyrin repeat domains, which are 33-residue motifs that mediate cytoskeleton anchoring or protein–protein interaction. Thus, TRPV6 may be linked to the cytoskeleton via its ankyrin repeats, enabling anchorage to the plasma membrane and channel regulation via actin filaments (Erler *et al.*, 2004). The carboxyl terminus (150 amino acid residues) contains a highly conserved 25 amino acids, called the TRP domain (Huang, 2004). Putative phosphorylation sites for protein kinase C are present in the cytoplasm domains, suggesting that transport activity could be regulated by phosphorylation. The N- and C-terminal tails of TRPV6 each contain an internal PDZ motif (Hoenderop *et al.*, 2002) which can function as part of a molecular scaffold via interaction with PDZ-domain containing proteins. Via this interaction, transporters, ion channels and receptors are coupled to intracellular signalling molecules, which regulate their activity, and position them to specialized regions in a cell (Figure 4) (Kornau *et al.*, 1997).

Chang and coworkers reported that N-tail (residues 64-77) and C-tail (residues 569-601) of TRPV5 are important for channel subunit assembly, subsequent trafficking of the TRPV5 channel complex to the plasma membrane, and channel activity (Chang *et al.*, 2004). Moreover, Nilius *et al.* (2003b) identified two domains in the carboxyl terminus of TRPV5 that control  $\text{Ca}^{2+}$ -dependent inactivation, whereas hTRPV6 lacks the carboxyl terminal domain (residues 695-725) for  $\text{Ca}^{2+}$ -calmodulin binding site; as a result, it does not form constitutive active channel (Bodding and Flockerzi, 2004).



### **Architecture of transient receptor potential vanilloid channel 5/6**

By using co-immuno precipitations and molecular mass determination of TRPV5/6 complexes using sucrose gradient sedimentation, it was showed that TRPV5 and TRPV6 form homo- and heterotetrameric channel complexes (Hoenderop *et al.*, 2003c). Since TRPV5 and TRPV6 are co-expressed in several tissues (Table 2), that allow oligomerization of these channels could occur *in vivo* (Hoenderop *et al.*, 2003c). The oligomerization of TRPV5 and TRPV6 might influence the functional properties of the  $\text{Ca}^{2+}$  channel. TRPV5 and TRPV6 exhibit different channel kinetics with respect to  $\text{Ca}^{2+}$ -dependent inactivation,  $\text{Ba}^{2+}$  selectivity and sensitivity for inhibition by ruthenium red (Hoenderop *et al.*, 2001a). The influence of the heterotetramer composition on channel properties was investigated.

Concatemers were constructed consisting of four TRPV5 and/or TRPV6 subunits, which configured in a head-to-tail fashion. A different number of TRPV5 and TRPV6 subunits in these concatemers showed that the phenotype resembled the mixed properties of TRPV5 and TRPV6 (Hoenderop *et al.*, 2003c). An increased number of TRPV5 subunits in such a concatemeric displayed more TRPV5-like properties, indicating that the stoichiometry of TRPV5/6 heterotetramers influences the channel properties. The clustering of four subunits is assumed to create an aqueous pore centered at the four-fold symmetry axis (Figure 5) (Hellwig *et al.*, 2005; Hoenderop *et al.*, 2003c).

Nilius and coworkers studied point mutation of the aspartate residue, D542A of TRPV5. They showed that an aspartate-to-alanine mutation at position D<sup>542</sup> of TRPV5, which corresponds to D<sup>541</sup> of TRPV6, abolishes  $\text{Ca}^{2+}$  permeation, and  $\text{Ca}^{2+}$ -dependent current decay, whereas permeation of monovalent cation basically remains intact. It established that D<sup>542</sup>/D<sup>541</sup> is a crucial determinant of the conductive properties of TRPV5/TRPV6 and a key element of the selectivity filter (Nilius *et al.*, 2001).

A characteristic feature of TRPV5/TRPV6 channels is the open block by intracellular  $\text{Mg}^{2+}$  (Nilius *et al.*, 2001). In the absence of extracellular divalent cations, the monovalent currents display characteristic voltage-dependent gating (Voets *et al.*, 2003).  $\text{Mg}^{2+}$  blocks the channel by binding to a single aspartate residue D<sup>541</sup> within TRPV6 (corresponding to D<sup>542</sup> of TRPV5), where it interacts with permeant cations (Voets and Nilius, 2003; Voets *et al.*, 2001; Martinez *et al.*, 2000). The pore helix is followed by the selectivity filter, which has a diameter of approximately 5.4 Å (Voets *et al.*, 2004).

These data indicate that selectivity and permeation properties in TRPV5 and TRPV6 are mainly determined by a ring of four aspartate residues in the channel pore. Sequence similarity is weak between the pore regions of the ring of voltage-dependent gating  $\text{Ca}^{2+}$  channels and TRPV5/TRPV6  $\text{Ca}^{2+}$  channels (Peng *et al.*, 2001a). It was suggested that the highly similarity functional properties are due to the occurrence of a ring of negative charges that structure is a high affinity binding site for cationic molecules at the extracellular entry way (Wood *et al.*, 2001; Owsianik *et al.*, 2006; Vennekens *et al.*, 2000).

### **Tissue distribution of transient receptor potential vanilloid channel 5/6**

The tissue distribution of TRPV5 and TRPV6 has been studied extensively by Northern blot, RT-PCR analysis and immunohistochemistry (Table 2). TRPV5 and TRPV6 are co-expression in the organs that mediate transcellular  $\text{Ca}^{2+}$  transport such as duodenum, jejunum, colon and kidney (Barley *et al.*, 2001; Weber *et al.*, 2001; Hoenderop *et al.*, 2001a; Zhuang *et al.*, 2002). The transport of  $\text{Ca}^{2+}$  across polarized  $\text{Ca}^{2+}$  transporting epithelial within organs such as the intestine, kidney and placenta is a coordinate and active process essential for dietary  $\text{Ca}^{2+}$  absorption, renal reabsorption and maternal-to-fetal transfer (Peng *et al.*, 2003a).

(Re)absorption of  $\text{Ca}^{2+}$  ion in kidney and intestine is essential for homeostasis of  $\text{Ca}^{2+}$  ion in the body. Transepithelial transport of  $\text{Ca}^{2+}$  in the (re)absorptive tissues occurs via paracellular as well as transcellular routes. Paracellular transport is passive, nonsaturable ways of  $\text{Ca}^{2+}$  (re)absorption (Larsson and Nemere, 2002). Transcellular (re)absorption of  $\text{Ca}^{2+}$  is mediated by  $\text{Ca}^{2+}$  entry across the apical membrane through the specialized epithelial  $\text{Ca}^{2+}$  channels, TRPV5 and TRPV6, intracellular buffering and facilitated diffusion bound to  $\text{Ca}^{2+}$ -binding proteins (calbindins) and extrusion across the basolateral membrane by  $\text{Na}^+ / \text{Ca}^{2+}$  exchanger (NCX1) and/or by plasma membrane  $\text{Ca}^{2+}$ -ATPases (Larsson and Nemere, 2002). Immunohistological studied demonstrated the localization of TRPV6 in the brush border membrane of intestinal absorptive cells (Kutuzova and DeLuca, 2004). This finding hints at a role of this protein as the rate apical influx pathway of active  $\text{Ca}^{2+}$  transport (Zhuang *et al.*, 2002; Kutuzova and DeLuca, 2004). TRPV6 was identified as the apical channel responsible for intestinal absorption of  $\text{Ca}^{2+}$  (Forsell *et al.*, 2006; Kutuzova and DeLuca, 2004; Slepchenko and Bronner, 2001; van Abel *et al.*, 2003; Walters *et al.*, 2004), whereas TRPV5 was initially identified as the apical  $\text{Ca}^{2+}$  entry channel in kidney (Hoenderop *et al.*, 2003b).

The TRPV5 was localized to the  $\text{Ca}^{2+}$ -reabsorptive distal convoluted tubules (DCT) and connecting tubules (CNT) (Hoenderop *et al.*, 1998). Recent studies, however, have found that both TRPV5 and TRPV6 are expressed in the DCT and CNT of kidney, where they may form heteromultimers (Peng and Hediger, 2002; Peng *et al.*, 2000b).

The epithelial calcium channels (TRPV5 and TRPV6) present in placenta (Moreau *et al.*, 2002b) might be involved in placental transfer of maternal  $\text{Ca}^{2+}$ . This is a crucial process in fetal development, which is carried out by the placental syncytiotrophoblast layer. The finding that mRNAs of TRPV5 and TRPV6 are expressed in this layer might indicate that these  $\text{Ca}^{2+}$  channels account for the basal  $\text{Ca}^{2+}$  uptake by these cells (Belkacemi *et al.*, 2005; Moreau *et al.*, 2002a; Moreau *et al.*, 2002b; Bernucci *et al.*, 2006).

TRPV6 is not only expressed in (re)absorptive epithelial cells but also in exocrine tissue such as pancreas, prostate, sweat gland, and mammary gland (Weber *et al.*, 2001; Hoenderop *et al.*, 2001a; Zhuang *et al.*, 2002). Moreover, the increasing expression of TRPV6 was found in prostate cancer and correlative with tumor grade. It was also found in carcinoma colon and carcinomas of other tissues such as breast, thyroid, and ovary (Wissenbach *et al.*, 2001; Abeele *et al.*, 2003a; Abeele *et al.*, 2003b; Peng *et al.*, 2001b; Prevarskaya *et al.*, 2004; Zhuang *et al.*, 2002).

### **Regulation of epithelial calcium channel TRPV5/6**

Transcellular  $\text{Ca}^{2+}$  transport is a highly coordinated and regulated process, which can be fine-tuned to the body's specific requirements. TRPV5/6 expression and channel activity have to be conspicuously controlled. This regulation occurs at different levels, including transcriptional and translational regulation, the tetrameric channel stoichiometry, trafficking of channels to and from the plasma membrane and modulation of channel activity at the plasma membrane (Nijenhuis *et al.*, 2005; Hoenderop *et al.*, 2005).

### **Transcriptional and translational regulation**

Parathyroid hormone (PTH) and the biologically active form of vitamin D [ $1,25(\text{OH})_2\text{D}_3$ ] are the main calciotropic hormones controlling  $\text{Ca}^{2+}$  balance (Hoenderop *et al.*, 2005). PTH, which is secreted into the circulation when the extracellular  $\text{Ca}^{2+}$  level decreases (hypocalcaemia), stimulates the activity of 25-hydroxyvitamin- $\text{D}_3$ -1 $\alpha$ -hydroxylase (1 $\alpha$ -OHase), a crucial enzyme in the biosynthesis of  $1,25(\text{OH})_2\text{D}_3$  in kidney (Brown *et al.*, 1999). Studies in several cell models suggested that  $1,25(\text{OH})_2\text{D}_3$  enhances epithelial  $\text{Ca}^{2+}$  channel expression (Hoenderop, 2001b; van Abel *et al.*, 2002; van Abel *et al.*, 2003; van de Graaf *et al.*, 2004; Brown *et al.*, 2005). In vitamin D receptor (VDR) knockout mice, duodenal TRPV6 mRNA levels were significantly down-regulated and associated with decreased intestinal  $\text{Ca}^{2+}$  absorption and hypocalcemia (Weber *et al.*, 2001; Cromphaut *et al.*, 2001; Ferrari *et al.*, 1998). Analysis of putative promoter region of human and murine *TRPV5* and *TRPV6* genes revealed potential vitamin D-response elements (Weber *et al.*, 2001; Nijenhuis *et al.*, 2003) so the expression of TRPV5 and TRPV6 is tightly controlled by  $1,25(\text{OH})_2\text{D}_3$ . The increasing  $1,25(\text{OH})_2\text{D}_3$  affects absorption of  $\text{Ca}^{2+}$  in the small intestine and kidney (Nijenhuis *et al.*, 2003; Wood *et al.*, 2001). However, the transcriptional and translational regulation of TRPV6 by  $1,25(\text{OH})_2\text{D}_3$  was consistently established.

### **Tetrameric channel stoichiometry**

In 2003, Hoenderop *et al.* revealed that TRPV5 and TRPV6 can indeed form homo- and heterotetrameric channel complexes. The concatemeric channels were constructed consisting of four TRPV5 and/or TRPV6 subunits, differences in the ratio of TRPV5 and TRPV6 subunits resulted in mixed properties of TRPV5 and TRPV6 channels because TRPV5 and TRPV6 exhibit different channel kinetics (Hoenderop *et al.*, 2003a). Recent studies showed that, whereas the other TRPV channels preferentially form homotetramers, TRPV5 and TRPV6 seem to be unique in forming heterotetramers (Hellwig *et al.*, 2005). Thus, regulation of the relative expression levels of TRPV5 and TRPV6 may be a mechanism to fine-tune the  $\text{Ca}^{2+}$  transport kinetics in TRPV5/6 co-expressing tissues. Indeed, several studies indicated that certain tissues co-express TRPV5 and TRPV6, including kidney and small intestine, which would allow oligomerization of these channels *in vivo* (Hoenderop *et al.*, 2005).

### **Trafficking of channels to the plasma membrane**

Accurate trafficking of channels and transporters to the plasma membrane is essential for transcellular ion transport. van de Graaf and his coworkers provided the first evidence of a regulatory role for the S100A10-annexin 2 heterotetramer in the trafficking of TRPV5 and TRPV6 (van de Graaf *et al.*, 2003). S100A10, a 97-amino acid protein member of the S100 superfamily, is found tightly associated with annexin 2. S100A10-annexin 2 is a member of the  $\text{Ca}^{2+}$  and phospholipids-binding proteins which is implicated in numerous biological processes including exocytosis, endocytosis and membrane-cytoskeleton interactions (Gerke and Moss, 2002; Gerke *et al.*, 2005). The association of S100A10 with TRPV5 and TRPV6 was restricted to a short conserved peptide sequence located in the C-tail of these channels (van de Graaf *et al.*, 2003). The first threonine of this sequence was identified as a crucial amino acid for binding and channel function. When this particular threonine was mutated, the activity of TRPV5 and TRPV6 was abolished accompanied by a major disturbance in their subcellular localization (Gerke *et al.*, 2005). This indicated that the S100A10-annexin 2 heterotetramer facilitates the translocation of TRPV5 and TRPV6 channels towards the plasma membrane. Interestingly, S100A10 as well as annexin 2 appear to be regulated by  $1,25\text{-(OH)}_2\text{D}_3$ , further supporting the importance of the S100A10-annexin-2 complex in the regulation of  $1,25\text{-(OH)}_2\text{D}_3$ -dependent TRPV5/6-mediated  $\text{Ca}^{2+}$  influx (van de Graaf *et al.*, 2003).

### **Modulation of channel activity at the plasma membrane**

It is well known that metabolic acidosis and alkalosis influence  $\text{Ca}^{2+}$  homeostasis. Therefore, pH might directly influence TRPV5/6 channel activity at the apical plasma membrane. In addition, extracellular pH also affected current kinetics including extracellular  $\text{Mg}^{2+}$  blockade and  $\text{Ca}^{2+}$  affinity. The mean current density decreased at acidic pH and increased at alkaline pH (Peng *et al.*, 1999). Recent studies showed that mutation of the Glu residue at position 522 to Gln near the pore helix decreased the inhibition of TRPV5 by extracellular acidification (Yeh *et al.*, 2003). This Glu-522 in the extracellular loop between TM5 and the pore region, appears to mediate pH sensitivity and, therefore, acts as the pH sensor in TRPV5 but the exact mechanism explaining these effects is unknown (Hoenderop *et al.*, 2005). Alternatively, these data suggest that the luminal pH directly regulates  $\text{Ca}^{2+}$  entry through the epithelial  $\text{Ca}^{2+}$  channels *in vivo*.

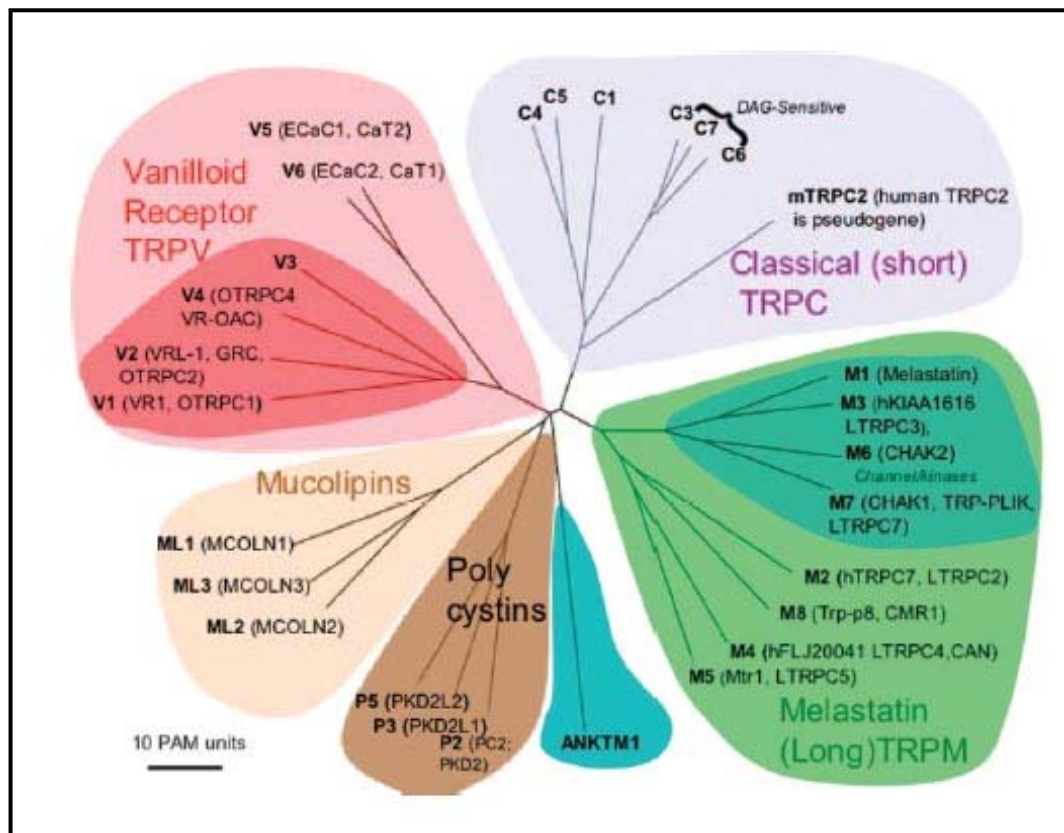
### **Milk calcium and its transportation in mammary epithelial cell**

The mammary gland can extract large quantities of  $\text{Ca}^{2+}$  from the plasma during lactation, to ensure that milk is concentrated with sufficient  $\text{Ca}^{2+}$ .  $\text{Ca}^{2+}$  is secreted into milk as free ionized  $\text{Ca}^{2+}$ ,  $\text{Ca}^{2+}$  bound to citrate or phosphate and as  $\text{Ca}^{2+}$  complexed with milk caseins by exocytotic pathway (McManaman and Neville, 2003; Neville, 2005). Ionized  $\text{Ca}^{2+}$  concentration within human milk is approximately 3 mM, total milk  $\text{Ca}^{2+}$  concentration is around 8 mM, whereas total plasma  $\text{Ca}^{2+}$  concentration, of which half is ionized, is 3 mM. Thus, the mammary gland is required to transport large amounts of  $\text{Ca}^{2+}$ , presumably in an active process due to this concentration gradient. Mammary gland  $\text{Ca}^{2+}$  transport is transcellular and overall unidirectional. The overall direction of transcellular calcium transport is unidirectional but note the presence of two ionized calcium gradients; one across the basolateral membrane and the other across the apical membrane. The  $\text{Ca}^{2+}$  transporters including the plasma membrane  $\text{Ca}^{2+}$ -ATPases (PMCA), sarco(endo)plasmic reticulum  $\text{Ca}^{2+}$  ATPases (SERCAs), Golgi pumps or secretory pathway  $\text{Ca}^{2+}$ -ATPases (SPCAs) as well as  $\text{Na}^+/\text{Ca}^{2+}$  exchangers being the predominant mechanism for  $\text{Ca}^{2+}$  efflux of the plasma membrane are also likely to be important mechanisms of calcium transport during lactation (Reinhardt and Horst, 1999; Reinhardt *et al.*, 2000). These P-type  $\text{Ca}^{2+}$ -transporting ATPases are likely to be the major pathways for lowering  $[\text{Ca}^{2+}]_i$  in mammary gland epithelial cells and due to the high demand for  $\text{Ca}^{2+}$  translocation from plasma to milk during lactation, they may also be important regulators of macro- $\text{Ca}^{2+}$  homeostasis in the mammary gland (Figure 6) (Reviewed by Lee *et al.*, 2006).

Although large transcellular  $\text{Ca}^{2+}$  influxes are required to support the transfer of  $\text{Ca}^{2+}$  from plasma to milk during lactation, uncertainty exists as to the exact mechanisms responsible for the transport of  $\text{Ca}^{2+}$  across the basolateral membranes of secretory mammary gland epithelial cells. Ideally, processes for  $\text{Ca}^{2+}$  transport across the basolateral membrane would have to be of large capacity, unidirectional and inducible. These make  $\text{Ca}^{2+}$  channels likely candidates to perform this role (Reviewed by Lee *et al.*, 2006). TRPV6, a  $\text{Ca}^{2+}$  channels with properties suggesting potential roles in mammary gland epithelial cells  $\text{Ca}^{2+}$  transport, may involve these processes.

**Table 1** Functional properties of members of TRPV subfamily

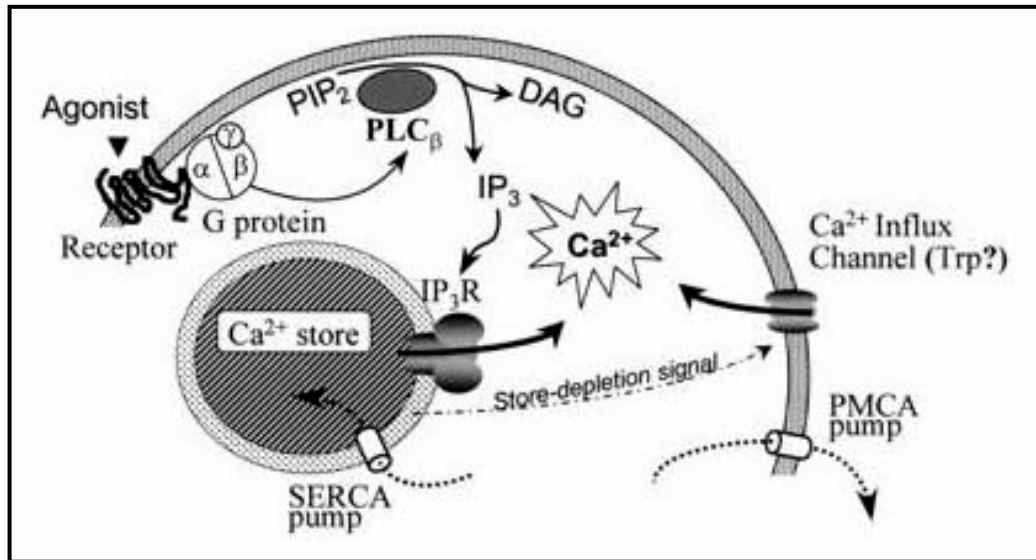
Name	$P_{Ca}/P_{Na}$	Proposed regulation	Function	Tissue distribution	Reference
TRPV1 (VR-1)	3-9	Capsaicin, Heat, Acidic pH	Heat sensor	Dorsal root ganglia	(Kedei <i>et al.</i> , 2001)
TRPV2 (VRL-1)	3	Heat, Growth factors	Heat sensor	Spinal cord, Brain, spleen	(Vennekens <i>et al.</i> , 2002)
TRPV3 (VRL-3)	12	Heat	Heat sensor	Skin, spinal cord, testis	(Smith <i>et al.</i> , 2002)
TRPV4 (OTRPC4)	6	Osmolarity, phorbol derivatives	Osmoreceptor	Kidney, salivary gland	(Nilius <i>et al.</i> , 2003a)
TRPV5 (CaT2 or ECac1)	>100	Low $[Ca^{2+}]_i$	Apical $Ca^{2+}$ transport	kidney, intestine	(Peng <i>et al.</i> , 2003a)
TRPV6 (CaT1 or ECac2)	>100	Store-operated, Low $[Ca^{2+}]_i$	Apical $Ca^{2+}$ transport	Intestine, placenta, kidney	(Hoenderop <i>et al.</i> , 2003a)



**Figure 1** Mammalian TRP family tree. The evolutionary distance between the TRP channels is shown by the total branch lengths in point accepted mutations (PAM) units, which is the mean number of substitutions per 100 residues. The tree was calculated using the neighbor-joining method for human, rat, and mouse sequences.

Source: Huang (2004)

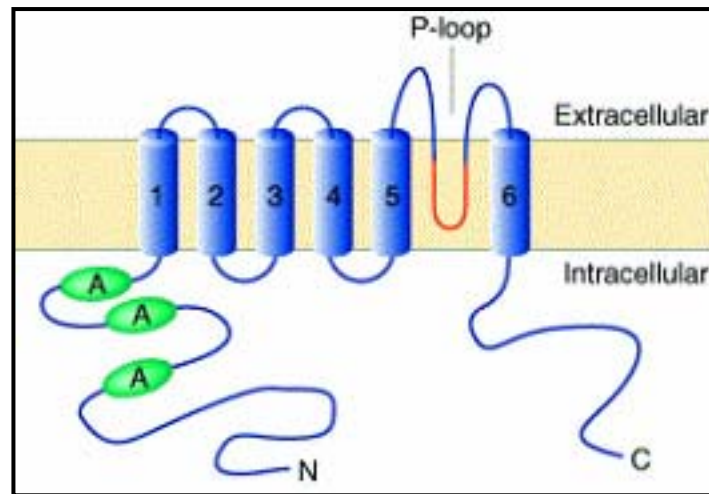




**Figure 2** Diagram of stimulation of phospholipase C (PLC) and activation of Ca<sup>2+</sup> signaling cascade.

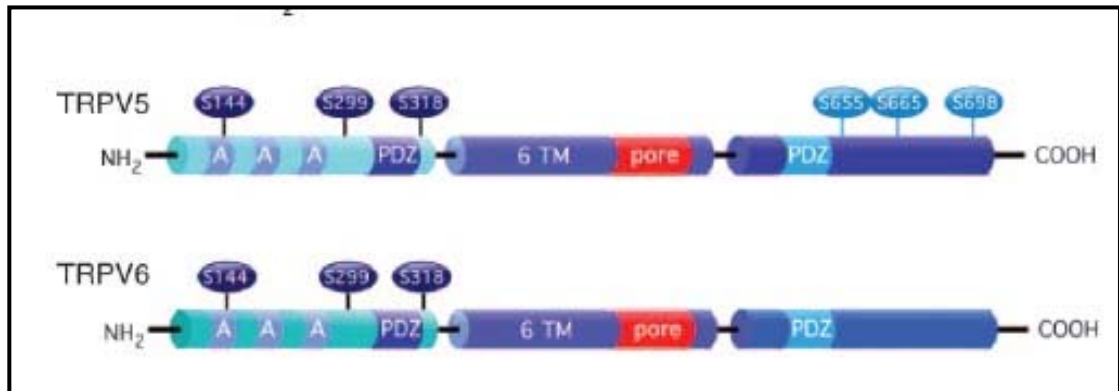
PLC- $\beta$  can be activated by  $\beta\gamma$ -subunits of heterotrimeric G proteins following receptor activation. Two sources contribute to the intracellular Ca<sup>2+</sup> concentration ( $[Ca^{2+}]_i$ ) increase. On inositol 1,4,5-trisphosphate (IP<sub>3</sub>) binding, Ca<sup>2+</sup> in the internal store is released via an intracellular Ca<sup>2+</sup> release channel, the IP<sub>3</sub> receptor (IP<sub>3</sub>R). Ca<sup>2+</sup> also comes in from extracellular space via undefined Ca<sup>2+</sup> influx channels. Influx channel is activated by an undefined store depletion signal. Ca<sup>2+</sup> pumps located on the endoplasmic reticulum (ER) [sarcoplasmic endoplasmic reticulum Ca<sup>2+</sup>-ATPase (SERCA)] and plasma membrane [plasma membrane Ca<sup>2+</sup>-ATPase (PMCA)] actively remove Ca<sup>2+</sup> from cytosol into the ER or external space. PIP<sub>2</sub>, phosphatidylinositol 4, 5-bisphosphate; DAG, 1, 2-diacylglycerol.

Source: Zhu and Birnbaumer (1998)



**Figure 3** The predicted topology of TRPV receptor. TRPV contains six transmembrane domains (6TMs) and a pore loop (P-loop). Three intracellular ankyrin repeats (A) are also present in the N-terminal region.

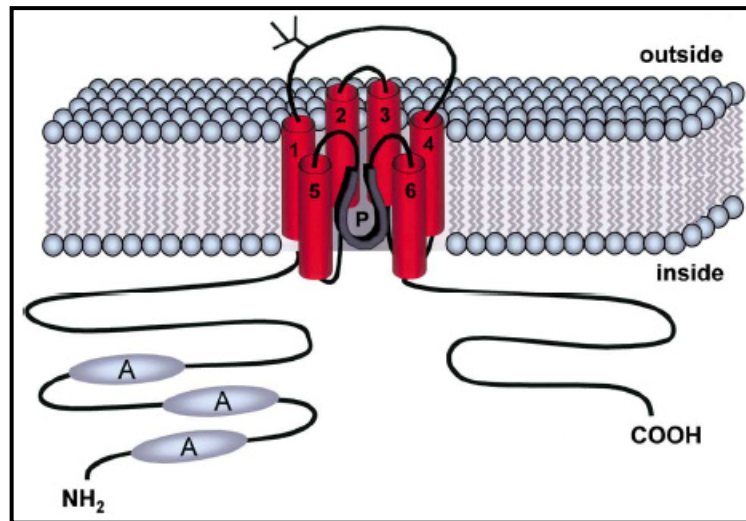
Source: Gunthorpe *et al.* (2002)



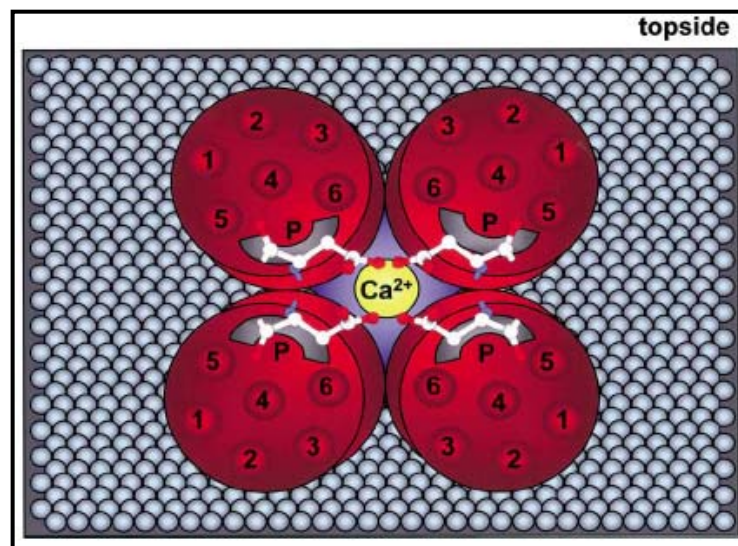
**Figure 4** Structural organization of TRPV5 and TRPV6. Potential regulatory sites in the amino and carboxyl tail of TRPV5 and TRPV6 including ankyrin repeats and PDZ motifs and conserved PKC phosphorylation sites.

Source: Hoenderop *et al.* (2005)

A



B



**Figure 5** The predicted topology of the TRPV5/6 monomer and architecture of (homo/hetero) tetrameric channel complex.

- (A) Schematic representation of TRPV5 and TRPV6 topology. Transmembrane segments are numbered, the pore region is depicted in grey, and ankyrin repeats in the N-terminal tail are indicated with A.
- (B) Top view of the TRPV5/6 (hetero) tetrameric channel shows four monomeric TRPV5/6 subunits forming a tetrameric complex to be the  $\text{Ca}^{2+}$ -binding site within the pore of the channel

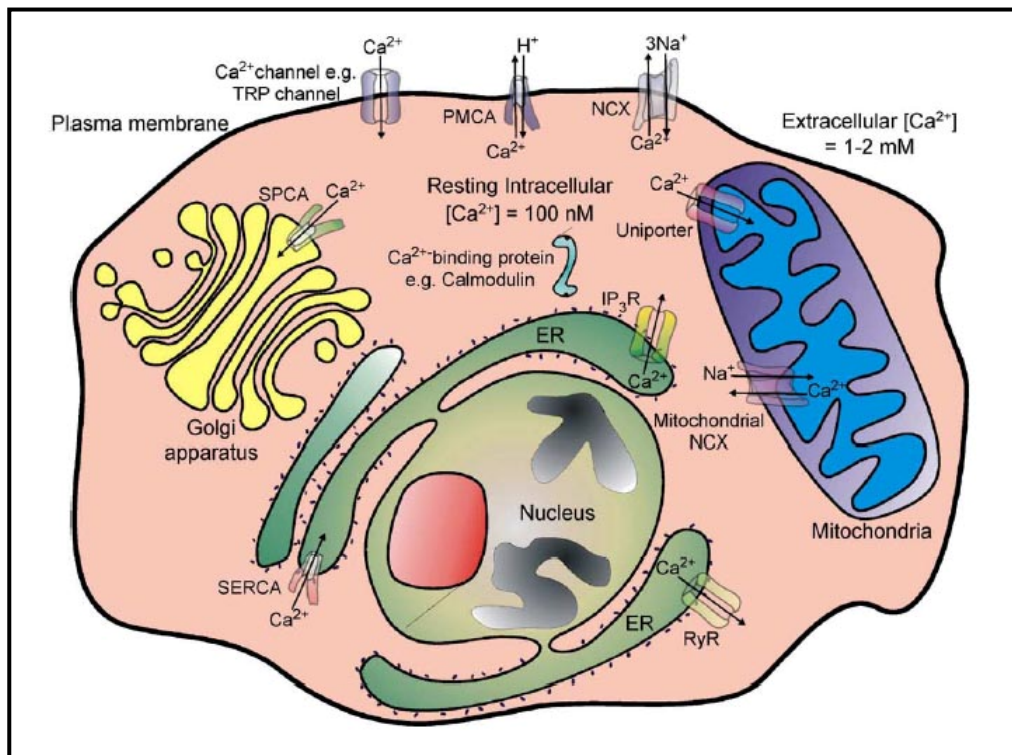
Source: den Dekker *et al.* (2003)

**Table 2** Tissue expression of TRPV5 and TRPV6

Tissue	Mouse		Human		Rat	
	TRPV5	TRPV6	TRPV5	TRPV6	TRPV5	TRPV6
Adrenal gland	-	+	ND	ND	ND	ND
Bone	-	+	ND	ND	ND	ND
Brain	-	+	+	-	-	ND
Caecum	-	+	-	-	-	+
Colon	-	+	+	(-)(+)	ND	ND
Duodenum	-	+	+	+	-	+
Heart	-	-	-	-	ND	ND
Ileum	ND	+	-	(-)(+)	ND	ND
Jejunum	-	+	+	+	ND	ND
Kidney	+	+	+	(-)(+)	+	-
Leukocyte	ND	ND	ND	-	ND	ND
Liver	-	+	-	-	-	ND
Lung	-	+	-	-	-	ND
Mammary gland	ND	ND	ND	+	ND	ND
Oesophagus	ND	+	-	(-)(+)	ND	ND
Osteoblast	-	+	ND	ND	ND	ND
Ovary	-	+	-	-	ND	ND
Pancreas	-	+	+	+	+	ND
Placenta	-	+	+	+	ND	ND
Prostate	-	+	+	+	ND	ND
Rectum	ND	ND	-	-	ND	ND
Salivary gland	ND	ND	ND	+	ND	ND
Skeletal muscle	ND	ND	-	-	ND	ND
Skin	-	+	ND	ND	ND	ND
Small intestine	ND	ND	-	(-)(+)	ND	ND
Spleen	-	+	-	-	-	ND
Stomach	ND	+	-	+	ND	ND
Sweat gland	ND	ND	ND	+	ND	ND
Testis	-	+	+	(+)(-)	-	ND
Thymus	-	+	ND	-	-	ND

TRPV5/6 expression or lack of expression is indicated with + and -, respectively. ND: not determined.

Source: den Dekker *et al.* (2003)



**Figure 6** Calcium regulation in mammary gland epithelial cells. The calcium transporters, channels and binding proteins are likely to contribute to calcium homeostasis and signaling in cells of the mammary gland.

Source: Lee *et al.* (2006)

## **MATERIALS AND METHODS**

### **Animal models and tissue samples**

Twenty-four confirmed pregnant Sprague Dawley rats were purchased from the Animal Facility of National Laboratory Animal Centre (NLAC), Mahidol University. Rats were individually housed in basket cages on sawdust bedding. Throughout pregnancy and lactation the dams were fed standard rat chow. The first day after parturition was assigned as the first day of lactation and the number of pups in each dam was reduced to seven. Rats were randomized block designed into 6 groups assigned as -7, -1, 0, +1, +7, and +14 for totally 4 blocks. Rats in -7 and -1 groups were sacrificed on 7 and 1 day prepartum. Rats in 0 group were sacrificed on parturition day, and rats in +1, +7, and +14 groups were sacrificed on 1, 7, and 14 day of lactation. Rats were sacrificed by 3 times overdose of Sodium Barbiturate (Nembutal<sup>®</sup>) on the date assigned. A 5 ml of blood was collected into sodium citrate anticoagulant tube (BD Vacutainer<sup>®</sup>, USA). Mammary and intestinal tissues were removed from the animal immediately after sacrifice and tissues were flash frozen on ethanol-dry ice bath and stored at -80°C until total RNA was prepared.

### **Total RNA preparation and quantitation**

Total RNA was prepared from tissues by the acid guanidine isothiocyanate-phenol-chloroform extraction using TRIzol reagent (Invitrogen, USA.) according to the manufacturer's protocol with slight modifications of single-step isolation method developed by Chomczynski and Sacchi (Chomczynski and Sacchi, 1987). A 1 gram tissue was homogenized in 15 ml TRIzol reagent at 15 seconds each for 3 times. A 3 ml chloroform was added into the homogenized sample. Let the homogenized sample stand undisturbed for 5 minutes before a 15 seconds of shaking and following with a standing undisturbed for 3 minutes at room temperature. The mixed solution was then centrifuged at 9,500 xg for 25 minutes at 4°C, and an aqueous phase of colorless top layer was collected into a new tube containing 7.5 ml isopropanol, mixed gently 3 times, and let stand undisturbed on ice for 10 minutes before a centrifugation at 11,800 xg for 15 minutes at 4°C. The supernatant was discarded and precipitated RNA pellet was washed with 15 ml 75% iced cold ethanol for 5 minutes at room temperature. The precipitated RNA pellet was then air dry for 10 minutes before being dissolved in diethylpyrocarbonate treated H<sub>2</sub>O (DEPC-ddH<sub>2</sub>O).

Total RNA was further precipitated for 2 hours on ice with 8M lithium chloride (LiCl) at 1:10 dilution. The LiCl precipitated RNA was collected after centrifugation at 14,000 xg for 20 minutes at 4°C. The LiCl precipitated RNA was washed twice in 75% ethanol and resuspended in DEPC-ddH<sub>2</sub>O.

To quantify the extracted total RNA, the RNA concentration was calculated from an UV260 and 280 nm absorbent measurement by spectrophotometer and its concentration was calculated by a following equation:-

$$\mu\text{g}/\mu\text{l of total RNA concentration} = A_{260} \times (0.04 \mu\text{g}/\mu\text{l}) \times \text{DF};$$

DF represents the dilution factor.

A ratio of  $A_{260}$  to  $A_{280}$  was calculated in order to specify a quality of extracted RNA.

### **TRPV6 primer design**

TRPV6 primers were designed by using MacVector 7.2.3 software from conserved regions of rat *TRPV6* gene (Accession number NM\_053686). Primers were always chosen according to following parameters: length between 18 and 25 bases; T<sub>m</sub> in between 55 - 80°C; length of amplification product between 400 and 1000 bp; G+C content between 45-55 %. All primer sequences were shown in Table 3. To determine specificity, all primer sequences were similarly compared against previously reported sequences at the GenBank database located at an URL: <http://www.ncbi.nlm.nih.gov> by using BLAST program version 2.0. Primers with sequences similar to sequence of genes other than TRPV6, they would be not selected. The selected TRPV6 primers were synthesized by KU Vector, Kasetsart University.



**Reverse Transcription Polymerase Chain Reaction (RT-PCR) and Polymerase Chain Reaction (PCR) for primer screening**

The Reverse Transcription Polymerase Chain Reaction (RT-PCR) conditions were as follows in a total volume of 50 µl reaction: 20 mM Tris-HCl (pH 8.3), 50 mM KCl, 1.5 mM MgCl<sub>2</sub>, 0.8 mM each of dNTPs, 50 pmol oligonucleotide primers, 50 U of SuperScript<sup>™</sup> III Reverse Transcriptase (Invitrogen, USA.), 10 U RNaseOut<sup>™</sup> (Invitrogen, USA.), 1.25 U *Taq* polymerase (Invitrogen, USA.), and 1 µg total RNA.

The Polymerase Chain Reaction (PCR) contained similar components as RT-PCR reaction except for SuperScript<sup>™</sup> III Reverse Transcriptase and RNaseOut<sup>™</sup>, which were excluded.

The RT reaction was conducted in a Thermocycler (Biometra-UNO II, Germany) starting at 55°C for 50 minutes, following by 4 minutes at 95°C. The PCR was performed 35 cycles of; 94°C for 15 seconds (denaturing), 55-58°C (primer specific annealing temperature depending on each primer set) for 20 seconds, and 72°C for 40 seconds (extension). The reaction was completed with a 25 minutes extension step at 72°C, and then was cooled to 4°C.

RT-PCR and PCR products were examined by horizontal electrophoresis. Electrophoresis was conducted by prestaining 9 µl of RT-PCR or PCR products with 9 µl of 1:5000 diluted SYBR<sup>®</sup> Gold (Molecular Probes, Eugene, USA) for 20 minutes in the dark. Prestained sample was loaded to a 2 % agarose gel. The gel was electrophoresed in 1X TAE buffer for 1 hour at 110 watts. A 100 bp DNA ladder molecular weight marker (Fermentas) was used as standard maker. The RT-PCR and/or PCR product bands were visualized under UV-transilumainator (Spectroline<sup>®</sup>). Images of the RT-PCR and PCR product were acquired with an EDAS 290 Kodak camera.

Primer set which gave a correct RT-PCR product size with >95% homology to rat TRPV6 gene and gave an undetectable of PCR product would be chosen for further experiment.

### **Sequencing of PCR products**

A DNA fragment, product from the RT-PCR using TRPV6 primer set, was recovered from the gel by using QIAquick Gel Extraction Kit (QIAGEN) as the manufacturer's protocol. The recovered DNA fragment was used as template for PCR reaction using the same primer set to increase a copy number of DNA fragment. The PCR product was confirmed its size by agarose gel electrophoresis, and then it was purified by QIAquick PCR Purification Kit (QIAGEN). The purified PCR product was submitted with its primer set for sequencing at BSU Bio Unit, Thailand. Sequencing was carried out for both forward and reverse primers. Nucleotide sequences of PCR product were analyzed for homology with sequences reported in GenBank database by BLAST program available at the National Center for Biotechnology Information website ([www.ncbi.nlm.nih.gov](http://www.ncbi.nlm.nih.gov)). A multiple sequence alignment was performed to compared with NCBI reported TRPV6 genes.

### **TRPV6 and GAPDH RT-PCR efficiency test**

TRPV6 and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) mRNAs in mammary tissues were detected by RT-PCR technique using TRPV6 primer set 3 and GAPDH primers reported by Gottwald *et al.* (Gottwald *et al.*, 2001), respectively. The GAPDH forward and reverse primer sequences were (5' TGA CCT TGC CCA CAG CCT TG 3') and (5' CAT CAC CAT CTT CCA GGA GCG 3') (Gottwald *et al.*, 2001). The RT-PCR conditions for each gene were analyzed for optimal MgCl<sub>2</sub> concentration, optimal annealing temperature, and optimal number of PCR cycle. The number of PCR cycle was 18-35 and 15-30 cycles for TRPV6 and GAPDH RT-PCR, respectively. The RT-PCR products of TRPV6 and GAPDH mRNAs were electrophoresed and products' band intensity were analyzed by Kodak 1D Image Analysis software<sup>®</sup>.

### **RT-PCR products image analysis**

An image of SYBR<sup>®</sup> Gold prestained DNA band was acquired with an EDAS 290 Kodak camera. The bands' intensity was quantified by KODAK 1D Image Analysis Software 1D. The image file to be analyzed must be saved in TIFF file format.

In order to analyze band intensity, the image file was opened and adjusted by using “Magnification Tool” on Toolbar to increase to 1.50X magnification. An area in image was selected by using “Image Selection Tool”. Lanes in a selected area were defined and bands in each lane were defined and adjusted. The band sensitivity and percentage of the profile with were adjusted to “0” and 80%. The band peak parameter was selected to display band intensity.

### **Semiquantitation of *TRPV6* gene expression**

*TRPV6* gene expression was semiquantitated by comparing amount of RT-PCR products from TRPV6 mRNA RT-PCR reaction to housekeeping gene (GAPDH) mRNA RT-PCR reaction, at the number of PCR cycle at exponential phase (Marone *et al.*, 2001). The 50 µl RT-PCR reaction of 1 µg total RNA was set, as previously described with TRPV6 primer set 3 (as shown in Table 3) and GAPDH forward-backward primers (as reported by Gottwald *et al.*, 2001) for TRPV6 mRNA and GAPDH mRNA, respectively. The PCR reactions were run totally 35 and 19 cycles for TRPV6 and GAPDH reactions, respectively. Their 10 µl RT-PCR products were electrophoresed and products' band intensity was evaluated by Kodak 1D Image Analysis software<sup>®</sup>. A ratio of band intensity of TRPV6 RT-PCR product to GAPDH RT-PCR product was calculated. Duplicate to triplicate reactions were performed for each total RNA sample.

### **Statistical analysis**

The expression of *TRPV6* gene in rat mammary tissues during pregnant, parturition and lactation was shown as mean of TRPV6/GAPDH band intensity ratio  $\pm$  SEM. The statistical analysis was done by using SAS analysis software (SAS Institute, Cary, NC). Differences in TRPV6 mRNA expression in each period were analyzed by One-way ANOVA and it was considered significant when  $p \leq 0.05$ . When expression was significant difference, Duncan's Multiple Range Test was performed and  $p \leq 0.05$  was accepted.

**Table 3** Sequences of TRPV6 primer set designed by using MacVector software

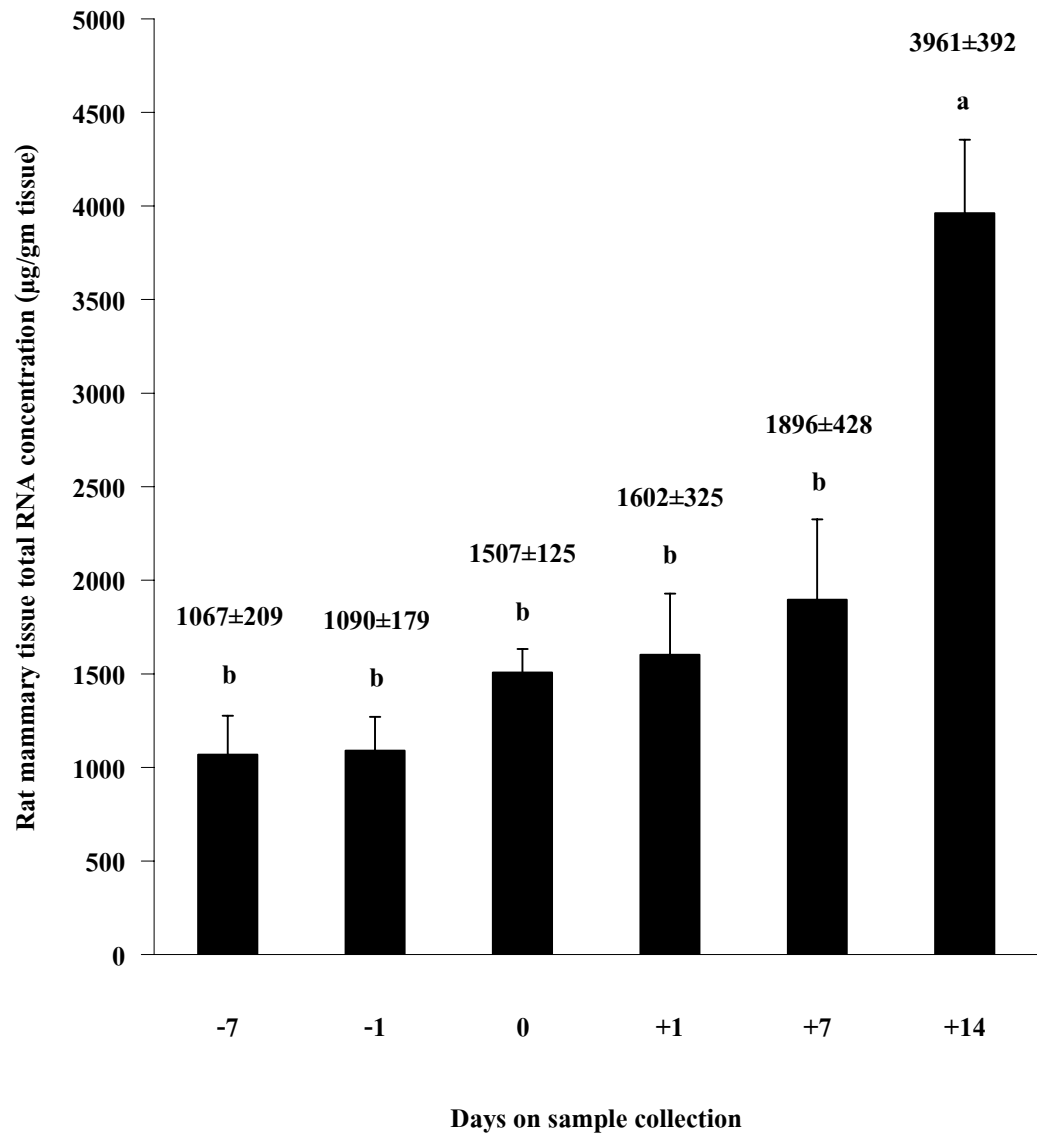
Primer set	PCR product size (bp)	Annealing temperature (°C)	Primer sequences
1	615	57-56	5' AAC AAA GCA GGA GCC CTC TTC G 3' 5' AGC ACG GAC CAA GTT CAC ATT C 3'
2	455	57-56	5' GAG AAC ATC CTT TGT CCT TTG CTG 3' 5' TTC CTT CAC AGG TGT TCT GGT CC 3'
3	570	56.5	5' CCC TTG AAC TTG TGC CCA ATA AC 3' 5' GAA TGG TCT GCC CAA AAA ATC G 3'
4	626	56	5' CTT ATT GTT ACC ACC AAG AAG CGG 3' 5' CCC AAG ATT ACC ACA GCC ATC AG 3'
5	564	58	5' TGG ACC AGA CAC CTG TGA AGG AAC 3' 5' GCC AGC AGA ATC GCA TCA AGT C 3'
6	475	55	5' ATC TTC AGG TTG GGG GTC ACT C 3' 5' TGA TGG CAA AGG CAG CGT AG 3'
7	706	55	5' TCT CAA CAG ACA ACG CAT CCG 3' 5' GAA ATG ACT TTA CTG GAA GGT GGC 3'
8	771	56	5' GAG AAC ATC CTT TGT CCT TTG CTG 3' 5' GAA TGG TCT GCC CAA AAA ATC 3'
9	379	56	5' TGG ACC AGA CAC CTG TGA AGG AAC 3' 5' GAA TGG TCT GCC CAA AAA ATC 3'
10	339	58	5' ATC TTC AGG TTG GGG GTC ACT C 3' 5' GAA TGG TCT GCC CAA AAA ATC 3'

## RESULT

Twenty-four confirmed pregnant Sprague Dawley rats were divided into four blocks. Each block contained six dams in a period of 7 days prepartum (-7), 1 day prepartum (-1), partum (0), 1 day lactation (+1), 7 days lactation (+7), and 14 days lactation (+14). Each block was experimentally designed as described in materials and methods.

### **Quality and quantity of total RNA from mammary tissues**

Total RNA from rat mammary tissues on 7 days prepartum (-7), 1 day prepartum (-1), partum (0), 1 day lactation (+1), 7 days lactation (+7), and 14 days lactation (+14) was expressed in  $\mu\text{g}$  total RNA per 1 gram tissue as shown in Figure 7. The quality of total RNA before LiCl precipitation was poor as  $A_{260}/A_{280}$  ratio being around 1.4-1.6. However, LiCl precipitated total RNA had a better quality as  $A_{260}/A_{280}$  ratio being around 1.6-1.8, and RT-PCR reaction of this RNA was not inhibited (data not shown). As shown in Figure 7, concentration of total RNA, extracted from mammary tissues during prepartum, partum, and early lactation increased gradually during early lactation. It was 2-3 folds total RNA concentration extracted from 14 days lactating samples compared to samples from pregnancy, partum and early lactation periods.

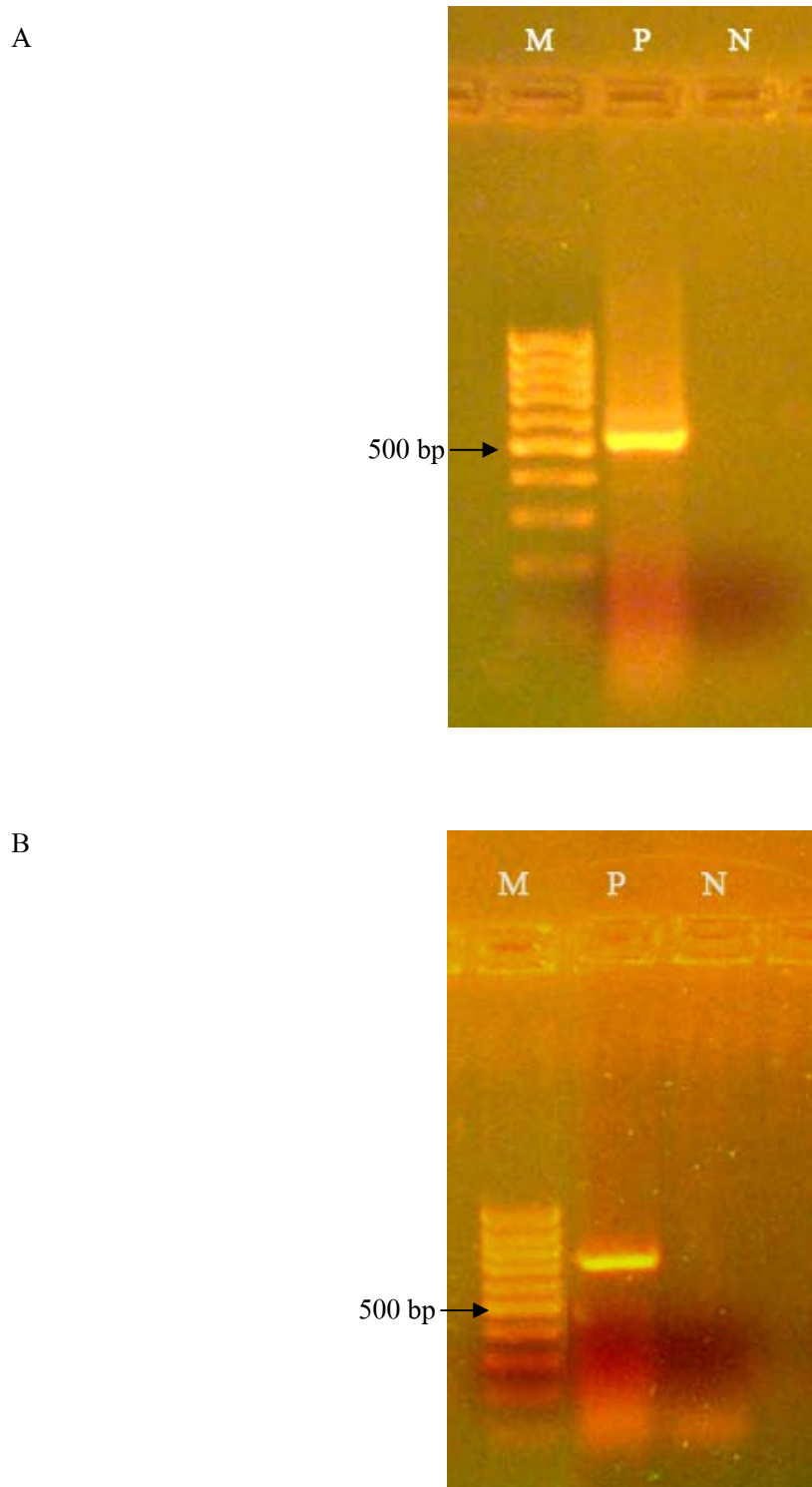


**Figure 7** Total RNA concentration extracted from rat mammary tissues. -7; 7 days prepartum, -1; 1 day prepartum, 0; partum day, +1; 1 day lactation, +7; 7 days lactation, +14; 14 days lactation. (means  $\pm$  SEM, n=4)  
a, b; Significant differences ( $p \leq 0.05$ )

### **Primer sequences and specific TRPV6 primer selection by RT-PCR**

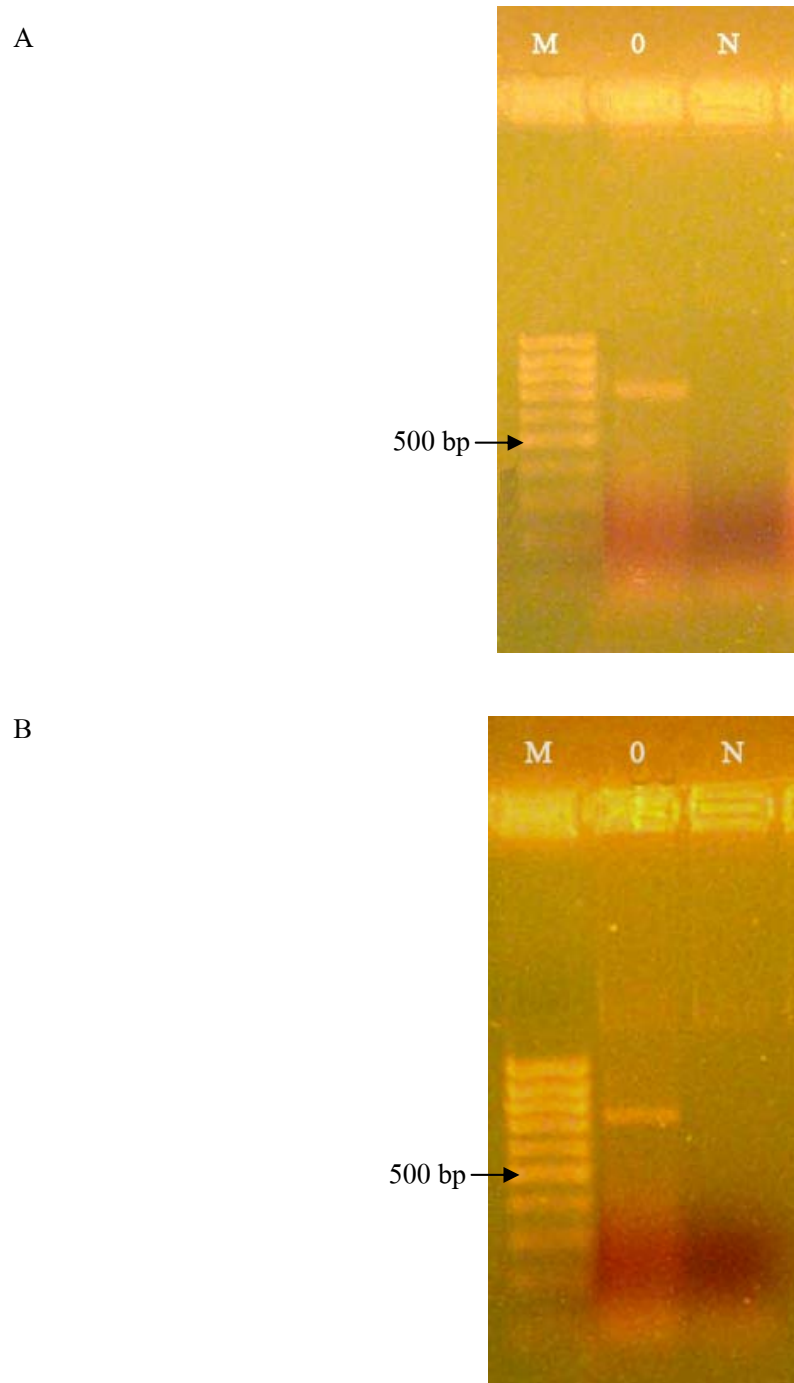
The TRPV6 primers were designed from conserved regions of rat *TRPV6* gene (Accession number NM\_053686) by MacVector 7.2.3 software. All primer sequences were shown in Table 3. Their specificity to TRPV6 mRNA was tested by RT-PCR with rat intestinal tissue total RNA. There were only primer set 3 and set 7 (Figure 8) which their RT-PCR products were corresponded to the expected size as indicated in Table 3. Therefore, primer set 3 and 7 were further tested by RT-PCR and PCR with mammary tissue total RNA.

TRPV6 RT-PCR product sizes were 570 bp and 706 bp when RT-PCR were performed with rat mammary total RNA by primer set 3 and 7, respectively (data not shown). However, primer set 7 gave positive PCR products when PCR reaction was performed without reverse transcriptase enzyme (Figure 9). Primer set 3 was selected for further test by RT-PCR and PCR with total RNA extracted from rat mammary tissues for the whole period of experiment (-7, -1, 0, +1, +7, and +14 days). Figure 10 shown results from RT-PCR and PCR products of primer set 3. As shown in Figure 10, RT-PCR products amplified by primer set 3 were 570 bp and there was no PCR product shown off in PCR reactions of total RNA extracted from mammary tissue of all periods. Therefore, RT-PCR product of primer set 3 was selected for further study by sequencing technique.

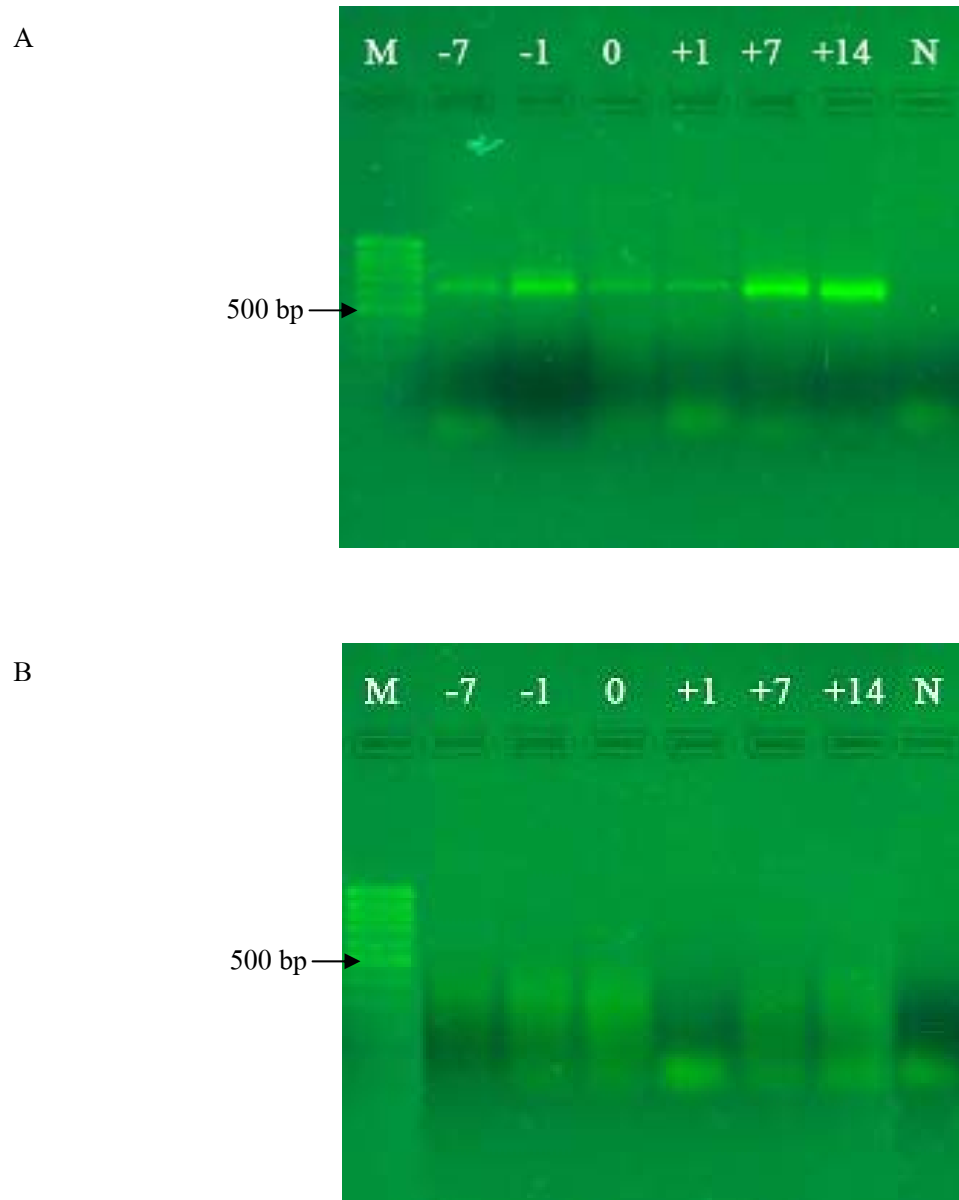


**Figure 8** RT-PCR product of TRPV6, as amplified by primer set 3 (A) and primer set 7 (B) with rat intestinal tissue total RNA. M; 100 bp DNA ladder molecular weight maker (Fermentas), P; rat intestinal tissue, and N; negative control.





**Figure 9** RT-PCR product (A) and PCR product (B) of TRPV6, as amplified by primer set 7, in rat mammary tissue on partum day. M; 100 bp DNA ladder molecular weight maker (Fermentas), 0; partum day, and N; negative control.

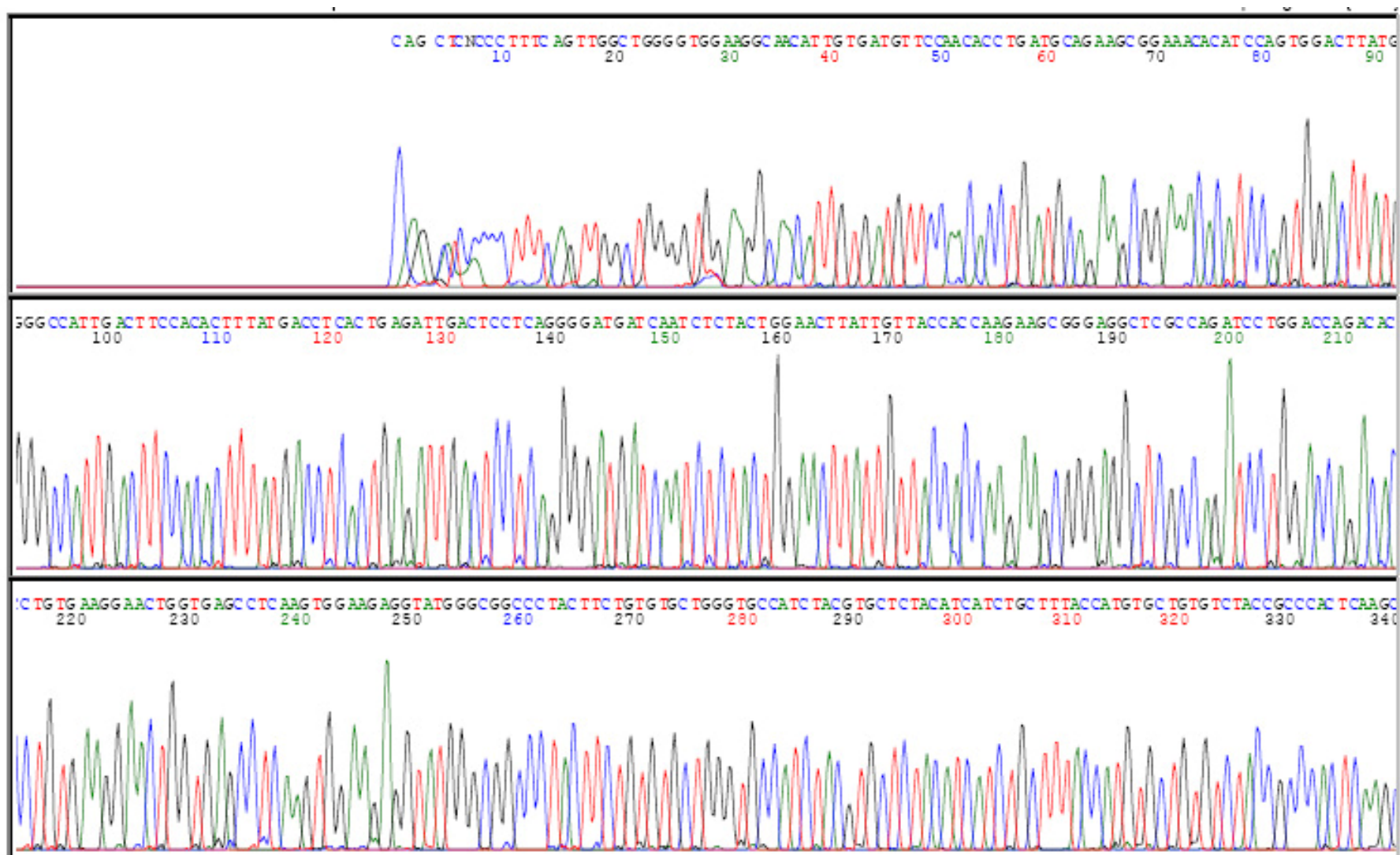


**Figure 10** RT-PCR product (A) and PCR product (B) of TRPV6, as amplified by primer set 3, in rat mammary tissue during pregnant, partum, and lactation. M; 100 bp DNA ladder molecular weight maker (Fermentas), -7; 7 days prepartum, -1; 1 day prepartum, 0; partum, +1; 1 day lactation, +7; 7 days lactation, +14; 14 days lactation, and N; negative control.

### **Sequencing and analysis of primer set 3 RT-PCR product**

A 570 bp RT-PCR product, amplified by primer set 3, of total RNA from mammary tissue on +14 days lactation was recovered and reamplified by PCR with the same primer set. The PCR product was then purified and confirmed its correct size by agarose gel electrophoresis. The purified PCR product was submitted for sequencing. Figure 11 shown sequencing result of RT-PCR product of primer set 3.

A 532 base sequences, shown in Figure 12, starting from base no. 1 to no. 532 was analyzed by multiple sequence alignment with bovine, human, mouse, and rat TRPV6 sequences by using Clustal W Alignment MacVector 7.2.3 software. The multiple sequence alignment result was shown in Figure 12. As shown in Figure 12, there was 98%, 93%, 85%, and 53% homology between RT-PCR product sequence to sequences of TRPV6 from rat, mouse, human, and bovine, respectively.



**Figure 11** Nucleotide sequences of primer set 3 RT-PCR product as sequencing by primer set 3 forward primer.



```

bTRPV6 1301 CCAGGGCCTCACCCCTTTCAAGCTGGCCGGAGTGGAGGGCAACACTGTGA 1350
hTRPV6 930 CCAGGGTCTCACCCCTTTCAAGCTGGCTGGAGTGGAGGGTAACACTGTGA 979
mTRPV6 936 CCAGGGACTCACCCCTTTCAAGTTGGCTGGGGTGGAAAGGCAACATTGTGA 985
rTRPV6 1003 CCAAGGACTCACCCCTTTCAAGTTGGCTGGGGTGGAAAGGCAACATTGTGA 1052
532 bp 1 CAGCTCNCCCTTTTCAGTTGGCTGGGGTGGAAAGGCAACATTGTGA 44
          * * * * * * * * * * * * * * * * * * * * * *
bTRPV6 1351 TGTTCCAGAACCTGGTGCAGAAAGCAGAAGTACATCCAGTGGACATGTGGA 1400
hTRPV6 980 TGTTTCAGCACCTGATGCAGAAAGCGGAAGCACACCCAGTGGACGTATGGA 1029
mTRPV6 986 TGTTCCAACACCTGATGCAGAAAGCGGAAACACATCCAGTGGACATATGGG 1035
rTRPV6 1053 TGTTCCAACACCTGATGCAGAAAGCGGAAACACATCCAGTGGACTTATGGG 1102
532 bp 45 TGTTCCAACACCTGATGCAGAAAGCGGAAACACATCCAGTGGACTTATGGG 94
          *****
bTRPV6 1401 CCGCTGACCTCTACTCTCTATGACCTCACGGAAATCGACTCCTCAGGGGA 1450
hTRPV6 1030 CCACTGACCTCGACTCTCTATGACCTCACAGAGATCGACTCCTCAGGGGA 1079
mTRPV6 1036 CCATTGACCTCCACACTGTATGACCTCACTGAGATTGACTCCTCAGGAGA 1085
rTRPV6 1103 CCATTGACTTCCACACTTTATGACCTCACTGAGATTGACTCCTCAGGGGA 1152
532 bp 95 CCATTGACTTCCACACTTTATGACCTCACTGAGATTGACTCCTCAGGGGA 144
          ** *****
bTRPV6 1451 GGAGCTATCCCTGCTGGAACCTTATCGTCACCTCCAAGAAGCGGGAGGTAC 1500
hTRPV6 1080 TGAGCAGTCCCTGCTGGAACCTTATCATCACCACCAAGAAGCGGGAGGCTC 1129
mTRPV6 1086 TGATCAATCCCTACTGGAACCTTATTGTTACTACCAAGAAACGGGAGGCTC 1135
rTRPV6 1153 TGATCAATCTCTACTGGAACCTTATTGTTACCACCAAGAAGCGGGAGGCTC 1202
532 bp 145 TGATCAATCTCTACTGGAACCTTATTGTTACCACCAAGAAGCGGGAGGCTC 194
          ** * * * * * * * * * * * * * * * * *
bTRPV6 1501 GCGAGGAGATTGTGAGGCTGCTCATTGAGCACGGTGCTGACATCCGGGGCC 1550
hTRPV6 1130 GCCAGATCCTGGACCAGACGCCGGTGAAGGA-GCTGGTGAGCCTCAAGTG 1178
mTRPV6 1136 GCCAGATCCTGGACCAGACACCTGTAAAGGA-ACTGGTGAGCCTCAAGTG 1184
rTRPV6 1203 GCCAGATCCTGGACCAGACACCTGTGAAGGA-ACTGGTGAGCCTCAAGTG 1251
532 bp 195 GCCAGATCCTGGACCAGACACCTGTGAAGGA-ACTGGTGAGCCTCAAGTG 243
          ** ** * * * * * * * * * * * * *
bTRPV6 1551 CAGGACTCCCTGGGAAACACAGTGTTCACATCCT-CGTCCCTCCA-GCCC 1598
hTRPV6 1179 GAAGCGGTACGGGCGGCGGTACTTCTGCTGCTGGGTGCCATATATCTGC 1228
mTRPV6 1185 GAAGAGGTATGGGCGGCCCTACTTCTGCTGCTGGGTGCCATCTATGTGC 1234
rTRPV6 1252 GAAGAGGTATGGGCGGCCCTACTTCTGTGTGCTGGGTGCCATCTACGTGC 1301
532 bp 244 GAAGAGGTATGGGCGGCCCTACTTCTGTGTGCTGGGTGCCATCTACGTGC 293
          * * * * * * * * * * * * * * * *
bTRPV6 1599 AACAAAACCTTTGCCT-GCCAGATGTACAACCTGCTGCTGTCTACGAC- 1646
hTRPV6 1229 TGTACATCATCTGCTTACCATGTGCTGCATCTACCGCCCCCTCAAGCCC 1278
mTRPV6 1235 TGTACATCATCTGCTTTACCATGTGCTGTGTCTACCGCCCGCTCAAGCCC 1284
rTRPV6 1302 TCTACATCATCTGCTTTACCATGTGCTGTGTCTACCGCCCACTCAAGCCC 1351
532 bp 294 TCTACATCATCTGCTTTACCATGTGCTGTGTCTACCGCCCACTCAAGCCC 343
          * * * * * * * * * * * * * * * *
bTRPV6 1647 AGGCGAGGGGACCACCTGCAGTCCCTGGACCTCATGCTCAATCACCAGGG 1696
hTRPV6 1279 AGGACCAATAACCGCACGAGCCCCCGGGACAACACCTCTTTACAGCAGAA 1328
mTRPV6 1285 AGGATCACTAACCGCACCAACCCCGGGACAACACTCTGATGCAACAGAA 1334
rTRPV6 1352 AGGATCACTAACCGCACCAACCCCGAGGACAATAACCTCCTGCAGCAGAA 1401
532 bp 344 AGGATCACTAACCGCACCAACCCCGAGGACAATAACCTCCTGCAGCAGAA 393
          *** * * * * * * * * * * *

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**Figure 12** Multiple alignment of the 532 bases RT-PCR product nucleotide sequences with bTRPV6 (NM\_018646), hTRPV6 (NM\_595413), mTRPV6 (NM\_022413), and rTRPV6 (NM\_053686) by using Clustal W Alignment MacVector 7.2.3 software.

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bTRPV6 1697 CCTCACCCCTTCAAGCTGGCCGGAGTGGAGGGCAACACTGTGATGTTCC 1746
hTRPV6 1329 GCTACTTCAGGAAGCCTACATGACCCCTAAGGACGATATCCGGCTGGTCG 1378
mTRPV6 1335 GCTCCTTCAGGAGGCCTATGTGACCCCTAAGGATGACCTCCGGCTGGTGG 1384
rTRPV6 1402 GCTCCTTCAGGAGGCCTATGTGACCCCTAAGGATGATCTCCGGCTGGTGG 1451
532 bp 394 GCTCCTTCAGGAGGCCTATGTGACCCCTAAGGATGATCTCCGGCTGGTGG 443
          **      *                ***      *      * * * *
bTRPV6 1747 AGCACCTGATGCAGAAGCGGAAGTACATCCAGTGGACGTGCGGGCCGCTG 1796
hTRPV6 1379 GGGAGCTGGTG-ACTGTCATTGGGGC-TATCATCATCCTGCTGGTAGAGG 1426
mTRPV6 1385 GGGAACTCGTG-AGCATCGTTGGGGC-TGTGATCATCCTGCTGGTGGAGA 1432
rTRPV6 1452 GGGAGCTGGTG-AGCATCGTTGGGGC-TGTGATCATCCTGCTGGTGGAGA 1499
532 bp 444 GGGAGCTGGTG-AGCATCGTTGGGGC-TGTGATCATCCTGCTGGTGGAGA 491
          * * * * * *      *      * * *      *      * * * * *
bTRPV6 1797 ACCTCCACTCTCTACGACCTCACAGAGATTGACTCCTGGGGAGAAGACGT 1846
hTRPV6 1427 TT-CCAGACATCTTCAGAATGGGGGTCACTCGCTTCTTTGGACAGACCAT 1475
mTRPV6 1433 TT-CCAGACATCTTCAGGTTGGGGGTCACTCGATTCTTTGGACAGACCAT 1481
rTRPV6 1500 TT-CCAGACATCTTCAGGTTGGGGGTCACTCGATTTTTTGGGCAGACCAT 1548
532 bp 492 TTTCCAGACATCTTCAGGTTGGGGGTCACTCGATTTTTTGGG 532
          *      *** *      *      * * *      *      * *

```

**Figure 12** (Continued)

### **TRPV6 and GAPDH RT-PCR efficiency test**

Semiquantitative RT-PCR analysis was chosen to assess expression levels of TRPV6 transcript from rat mammary tissue during pregnant, partum, and lactation. This method is based on the use of an internal control, which is a housekeeping gene. A highly level expression housekeeping gene, glyceraldehyde-3-phosphate dehydrogenase (GAPDH), was used as an internal control in this experiment. A number of parameters could affect RT-PCR results and were extremely important for the reaction to be quantitative.

An optimal concentration of  $\text{MgCl}_2$  and optimal primer annealing temperature were tested for both TRPV6 and GAPDH RT-PCR. There were 1.5 mM  $\text{MgCl}_2$  concentration and 56.5°C optimal annealing temperature that gave the best results for both TRPV6 and GAPDH (data not shown).

The optimal PCR cycle numbers was evaluated by choosing cycle numbers that provided product band intensity in exponential phase. As shown in Figure 13, exponential phases for TRPV6 RT-PCR and GAPDH RT-PCR were during 28-31 cycle and 16-19 cycle numbers, respectively. However, 31 PCR cycle numbers for TRPV6 RT-PCR of total RNA of samples at prepartum and partum days gave a very poor to undetectable result, hence, a 35 cycle numbers was selected for TRPV6 RT-PCR rather than 31 cycle numbers. A 19 and 35 PCR cycle numbers were used in GAPDH RT-PCR and TRPV6 RT-PCR reactions in next experiments of semiquantitation.

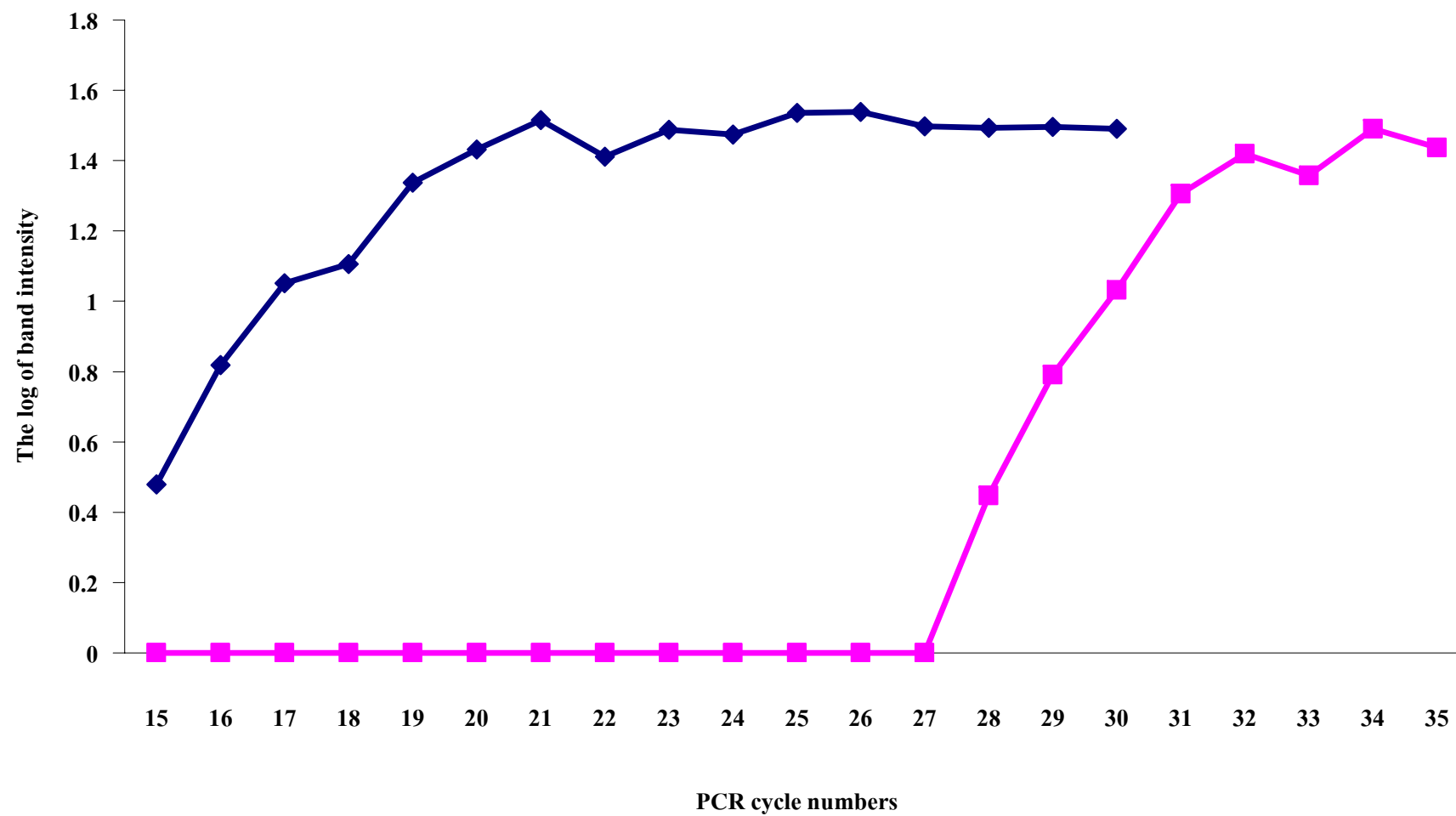


**Figure 13** Determination of a amplification power for TRPV6 and GAPDH. The intensity of TRPV6 and GAPDH RT-PCR products were shown in log band intensity.



GAPDH RT-PCR

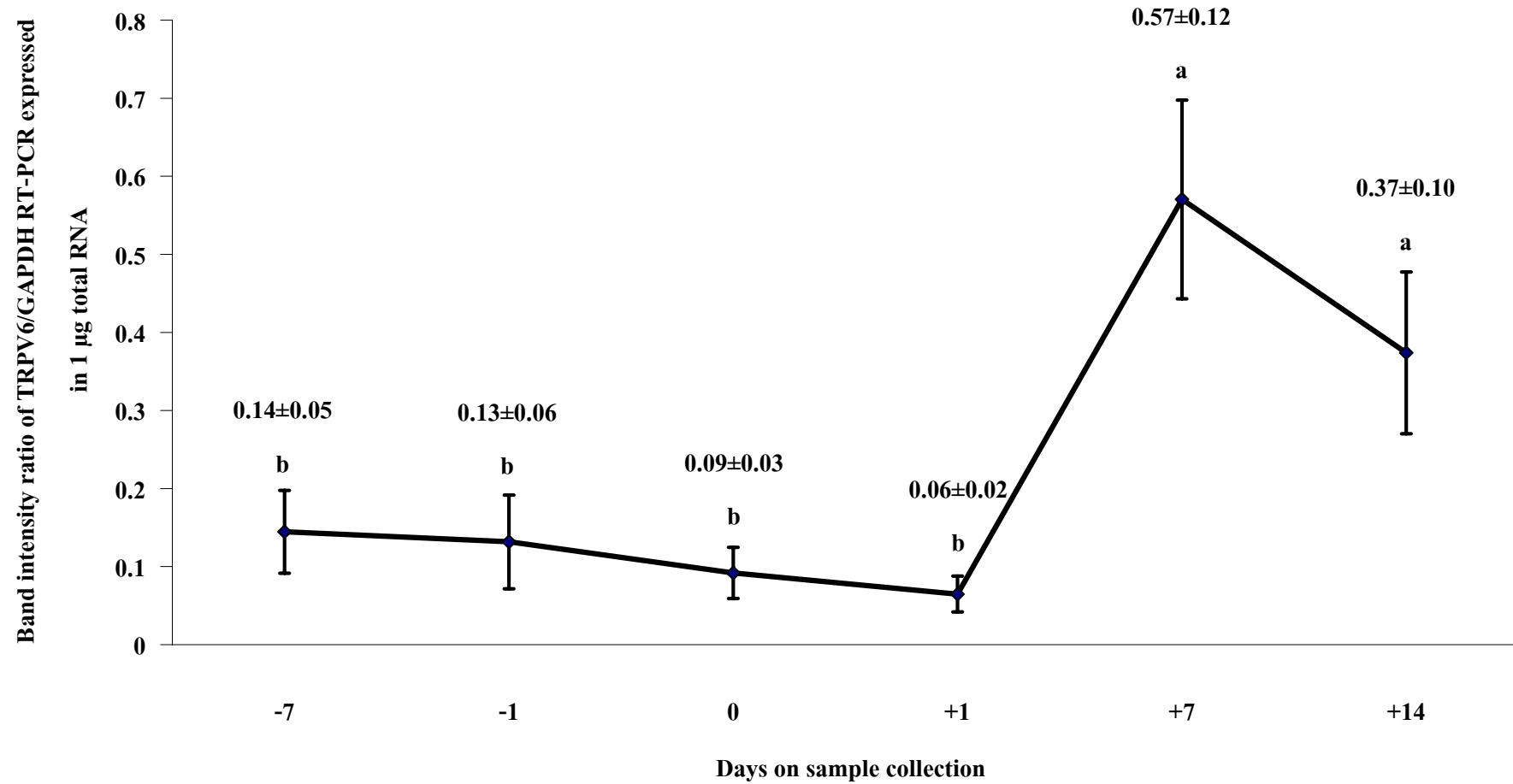
TRPV6 RT-PCR



**Semiquantitative of TRPV6 gene expression in rat mammary tissues during pregnancy, partum, and lactation**

An expression of TRPV6 mRNA in mammary tissues was evaluated during periods of pregnancy, parturition, and lactation. Figure 14 shown band intensity ratio of TRPV6/GAPDH RT-PCR products over a period of -7, -1, 0, +1, +7, and +14 of prepartum, partum and lactation days. During -7, -1, 0, and +1 days, TRPV6 mRNA was expressed at low basal level. The expression of TRPV6 transcript increased to 5 and 3 folds at +7 and +14 days lactation periods ( $p \leq 0.01$ ), Figure 14. This expression profile was evaluated in 1  $\mu$ g total RNA. If one calculated the expression profile of TRPV6 transcripts in mammary tissue per gram tissue, expressions of mammary TRPV6 would be 9 and 11 folds in mammary tissue during +7 and +14 days of lactation compared to TRPV6 level in mammary tissue during prepartum, partum and early lactation (data not shown).

**Figure 14** The band intensity ratio of TRPV6/GAPDH on days during pregnancy, partum, and lactation of rat mammary tissue. -7; 7 days prepartum, -1; 1 day prepartum, 0; partum day, +1; 1 day lactation, +7; 7 days lactation, +14; 14 days lactation. (means  $\pm$  SEM, n=4)  
a, b; Significant differences ( $p \leq 0.01$ )



## DISCUSSION

TRPV6 protein in many organs, such as intestine, kidney, placenta, and mammary gland, has been reported by several investigators (Peng *et al.*, 1999; Nijenhuis *et al.*, 2003; Bernucci *et al.*, 2006; Zhuang *et al.*, 2002). However, there have been no evidences on the changes of TRPV6 mRNA expression in rat mammary gland under different physiological condition. The purpose of this study was to investigate TRPV6 mRNA expression profile in rat mammary tissues during pregnancy, parturition and lactation (-7, -1, 0, +1, +7, and +14) by using semiquantitative RT-PCR analysis.

### **Quality and quantity of total RNA from mammary tissues**

The total RNA of rat intestine and mammary tissues were isolated by TRIzol reagent (Invitrogen, USA). The concentration of total RNA extracted from these mammary tissues increased gradually from pregnancy to early lactation but significantly increased on 14 days of lactation ( $p \leq 0.05$ ) as shown in Figure 7. The increasing total RNA concentration during lactation correlated with an increasing of total protein concentrations in rat milk throughout lactation (Yamamoto *et al.*, 1992). In this experiment, the total RNA extracted from rat mammary tissues must be precipitated with 8 M lithium chloride (LiCl) since non LiCl precipitated total RNA from mammary tissue gave a negative result in RT-PCR reaction (data not shown). LiCl precipitated total RNA had a better quality than the quality of total RNA before LiCl precipitation because RT-PCR inhibitors were removed from RNA extracted from mammary tissue (Mathy *et al.*, 1996).

### **The specific TRPV6 primer selection by RT-PCR and sequencing**

To select specific TRPV6 primer, the LiCl precipitated total RNA of rat mammary tissue was used as template in RT-PCR reaction with primer sets as shown in Table 3. The RT-PCR product from primer set 1, 2, 4, 5, 6, 8, and 10 showed that their RT-PCR product sizes were not corresponded to the expected size as shown in Table 3, when compare with RT-PCR product (positive control) amplified from total RNA of rat intestine as template (data not shown). It might be alternatively RNA splicing forms of this gene in mammary tissue, evidences in mammalian plasma membrane  $\text{Ca}^{2+}$ -ATPases, PMCA1-4, encoded by a multigene family. Additional diversity of PMCA1-4 is generated by alternative RNA splicing (Filoteo *et al.*, 1997; Reinhardt and Horst, 1999).

Further studies to investigate some spliced forms of this *TRPV6* gene in mammary tissues, we need to clone full length of *TRPV6* gene in rat mammary tissue.

There are two primer sets, set 3 and set 7, that could generate positive correct size RT-PCR products, but primer set 7 gave positive PCR products when PCR reaction was performed without reverse transcriptase enzyme (Figure 9). From this result, it was suggested that it might be DNA contaminated in total RNA sample. In addition, the forward and reverse primers of primer set 7 are on the same exon 15 (Peng *et al.*, 1999; Peng *et al.*, 2001a; Hirnet *et al.*, 2003), so PCR gave positive result with correct size of PCR product. Therefore, RT-PCR product primer set 3 was selected to be confirmed by sequencing as shown in Figure 11. The obtained sequences were then analyzed by multiple sequence alignment with bovine, human, mouse, and rat TRPV6 sequences (Figure 12). It had 98%, 93%, 85%, and 53% homology between RT-PCR product sequence to sequences of TRPV6 from rat, mouse, human, and bovine, respectively, so this primer set 3 was selected to assess expression levels of TRPV6 transcript in further study.

#### **TRPV6 gene expression in rat mammary tissue during pregnancy, partum, and lactation**

Results from experiment showed that TRPV6 mRNA expressed in rat mammary tissues during pregnancy, parturition and lactation. The expressions of TRPV6 increased 5 and 3 folds at +7 and +14 days lactation periods compared with that at prepartum, and partum periods ( $p \leq 0.01$ ). The expressions of TRPV6 transcripts in mammary tissue per gram tissue were 9 and 11 folds during +7 and +14 days of lactation compared to TRPV6 level in mammary tissue during prepartum, partum and early lactation. Its expression profile was correlated to level of total RNA present in mammary tissue at various periods (Figure 7 and Figure 14). Calcium concentrations in rat milk increased only during early stages of lactation and remained stable after 7 days lactation (Yamamoto *et al.*, 1992), so this TRPV6 expression profile increase paralleled to milk calcium secretion. An increasing expression of TRPV6 mRNA in lactating mammary tissues suggested that TRPV6 plays important role(s) in calcium transport during lactation and milk production.

Rat milk has calcium concentration in excess of 60 mM. In human and cows, changes in milk calcium appear to be regulated by the amount of casein and citrate rather than ionized calcium [ $\text{Ca}^{2+}$ ] (Neville *et al.*, 1994). In most milk, calcium is associated with casein, citrate, phosphate,

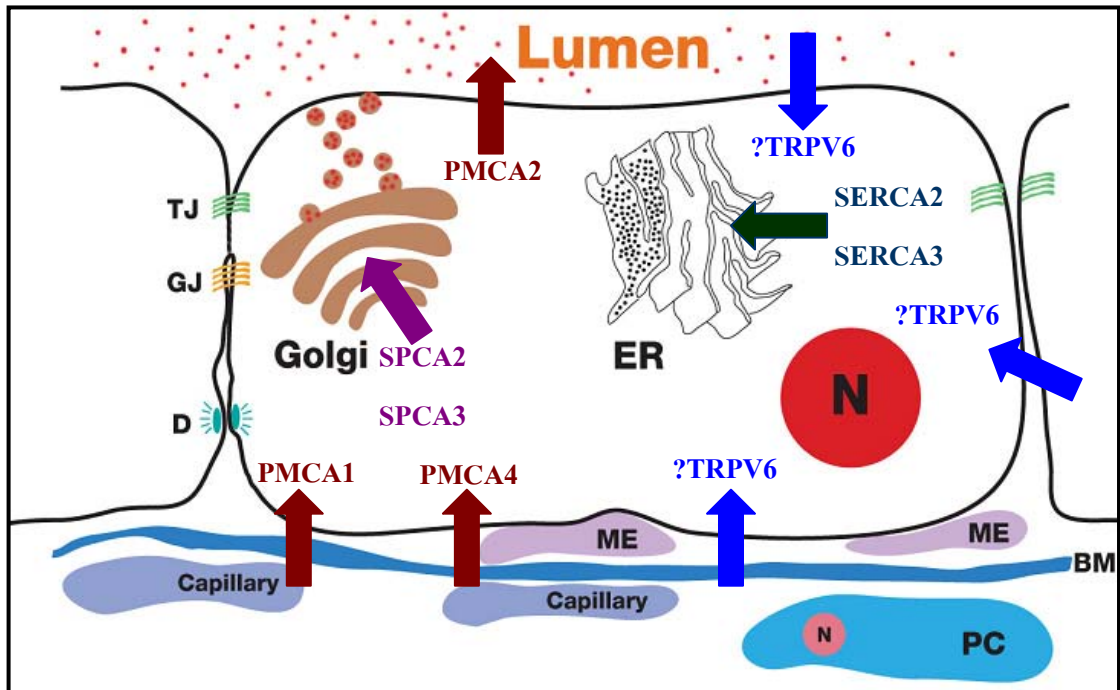
bicarbonate, and carbonate in the sacculars of the terminal Golgi apparatus and is secreted via the Golgi-secretory vesicle system from the mammary alveolar cell (Neville and Peaker, 1979). It seems that the mammary gland does not transport calcium directly from the blood-to-milk compartments direction since calcium cannot cross the apical membrane or tight junctions (Neville and Peaker, 1979). It can be inferred that calcium transport into milk is transcellular (McManaman and Neville, 2003; Neville, 2005). The mechanism by which calcium is secreted into milk should be complex but, in fact, they are simple (Figure 15). During the transition from pregnancy to lactation, calcium transport mediates from blood across the basolateral membrane of mammary epithelial cell to cytoplasm by passive transport. This mechanism is not enough for large transcellular  $\text{Ca}^{2+}$  fluxes which is required to support the transfer of  $\text{Ca}^{2+}$  from plasma to milk during lactation. This profiling expression of TRPV6 in mammary tissue during pregnancy and lactation supported that TRPV6  $\text{Ca}^{2+}$  channel involves in macro calcium in this active mammary tissue. However, the exact mechanism(s) responsible for the transport of  $\text{Ca}^{2+}$  across the basolateral membranes of mammary secretory cells has not been determined (Shennan and Peaker, 2000).

Intracellular calcium is pumped from the cytoplasm into Golgi compartment by the Golgi  $\text{Ca}^{2+}$ -ATPase, SPCA (Taylor *et al.*, 1997). Reinhardt *et al.*, 2000 reported SPCA protein expression increased 1 week before parturition and increased further as lactation proceeded. During lactation, the Golgi compartment requires large amount of calcium concentration for casein synthesis and micelle formation (Neville *et al.*, 1994). The expression of  $\text{Ca}^{2+}$ -ATPases (PMCA1b, PMCA4b, SERCA2, and SERCA3) mRNA and their protein targeted at basolateral membrane and endoplasmic reticulum (ER), respectively, showed the lowest levels of expression during pregnancy and early lactation (Reinhardt and Horst, 1999; Reinhardt *et al.*, 2000). Their levels became significantly elevated on days 14 of lactation (Reinhardt and Horst, 1999). These  $\text{Ca}^{2+}$ -ATPases play an important role in maintaining low  $\text{Ca}^{2+}$  concentration in cytoplasm while facilitating a large transcellular calcium flux. Another  $\text{Ca}^{2+}$ -ATPase, PMCA2b, is located at apical membrane of mammary secretory cell (Reinhardt *et al.*, 2000). The expression of PMCA2b mRNA and protein increased continuously from a starting of lactation and they increased on 21 days of lactation while a macro amount of  $\text{Ca}^{2+}$  was transported into milk. PMCA2b is important for macro-  $\text{Ca}^{2+}$  homeostasis in lactating tissue. (Reinhardt and Horst, 1999; Reinhardt *et al.*, 2000). The large amount of calcium required during lactation correlated with TRPV6 mRNA expression that its mRNA significantly increased ( $p \leq 0.01$ ) at +7 and +14 days lactation periods. This evidence implied that TRPV6 channel might be the



channel mediate calcium transport from blood into mammary epithelial cell during a requirement of large amount of calcium.

However, to evaluate role(s) of TRPV6  $\text{Ca}^{2+}$ -selective channel as a channel to mediate blood calcium into mammary secretory cell, further studies will be required to identify the localization of TRPV6 protein in the mammary gland. Whether apical or basolateral locations are required for physiological explanations.



**Figure 15** Diagram of calcium transport in mammary epithelial cell. PMCA; plasma membrane  $\text{Ca}^{2+}$ -ATPase, SERCA; Sarco(endo)plasmic reticulum  $\text{Ca}^{2+}$ -ATPase, and SPCA; Golgi  $\text{Ca}^{2+}$ -ATPase.

## CONCLUSION

By using semiquantitative RT-PCR analysis, the expression of TRPV6 mRNA was observed in pregnant, parturition and lactating rat mammary tissues (-7, -1, 0, +1, +7, and +14). Measurement of TRPV6 mRNA expression in rat mammary tissues during pregnancy, parturition and lactation indicated that TRPV6 mRNA levels did not change appreciably between pregnancy, parturition, and early lactation, but increased significantly ( $p \leq 0.01$ ) on 7 days lactation and 14 days lactation. This result suggested that the increasing TRPV6 mRNA expression during 7 days lactation until 14 days lactation might respond to calcium requirement of mammary secretory cells during lactation.

Lactating mammary tissue is an organ in which large amounts of calcium are transported. Therefore, it is tempting to hypothesize that TRPV6 channel in the mammary gland contributes to the massive transport of calcium from blood to milk.

However, in order to understand role(s) of TRPV6 protein in rat mammary tissue during pregnancy, parturition and lactation, an immunohistochemistry technique was required to investigate localization of TRPV6 protein in rat mammary tissue.

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## **APPENDIX**

## **Appendix A**

### **Preparation of chemical reagents**

#### **1 litre 0.1% diethylpyrocarbonate treated water (DEPC-ddH<sub>2</sub>O)**

1 ml of diethylpyrocarbonate (BIO BASIC INC, )

1 litre deionized distilled water

Add 1 ml DEPC into 1 litre deionized distilled water and then mixed by using stirrer and incubate overnight at 42°C. Then autoclave 45 mins, 121°C at 15lb/sq liquid cycle before use.

#### **20 ml 8 M Lithium chloride (8 M LiCl)**

6.78 g LiCl

20 ml DEPC-ddH<sub>2</sub>O DEPC

Add 6.78 g. LiCl into 20 ml DEPC-ddH<sub>2</sub>O and shaken well before use

#### **1 litre 50X TAE buffer**

242 g Tris base

57.1 ml glacial acetic acid

100 ml 0.5 M EDTA (pH 8)

1 litre distilled water

Add 242 g. Tris Base, 57.1 ml glacial acetic acid, and 100 ml 0.5 M EDTA (pH 8) into 500 ml distilled water. Then mixed by using stirrer, and add distilled water until 1 liter.

## Appendix B

### The band intensity ratios of TRPV6/GAPDH on days during pregnancy, partum, and lactation of rat mammary tissues

Blocks	Days	Replication I			Replication II		
		TRPV6	GAPDH	Ratio I	TRPV6	GAPDH	Ratio II
1 <sup>st</sup>	-7	18.0700	44.5800	0.40534	4.3451	42.2800	0.10277
	-1	0.0001	52.0200	0.00000	8.4275	48.9400	0.17220
	0	7.7370	45.1900	0.17121	5.7659	42.1600	0.13676
	+1	3.0648	49.8900	0.06143	4.9020	43.5000	0.11269
	+7	7.1171	41.2500	0.17254	30.9300	45.9200	0.67356
	+14	28.4000	43.8100	0.64825	16.5600	37.8400	0.43763
2 <sup>nd</sup>	-7	9.8803	33.3200	0.29653	0.0001	23.6900	0.00000
	-1	0.0001	28.9200	0.00000	0.0001	4.2014	0.00002
	0	0.0001	28.3500	0.00000	0.0001	19.9000	0.00001
	+1	4.5787	33.9400	0.13491	0.0001	32.3500	0.00000
	+7	22.9000	33.4400	0.68481	19.2000	19.9200	0.96386
	+14	5.4804	37.4000	0.14653	0.0001	16.8300	0.00001
3 <sup>rd</sup>	-7	15.4000	72.6600	0.21195	6.9032	49.3300	0.13994
	-1	12.4700	65.2100	0.19123	19.9400	52.7100	0.37830
	0	4.3039	46.9300	0.09171	6.1608	44.7300	0.13773
	+1	0.0001	51.4700	0.00000	0.0001	45.7800	0.00000
	+7	10.9800	43.7500	0.25097	17.2800	52.3100	0.33034
	+14	2.9940	31.8500	0.09400	26.3500	36.0800	0.73032
4 <sup>th</sup>	-7	0.0001	69.2200	0.00000	0.0001	62.7500	0.00000
	-1	5.7130	44.0400	0.12972	8.7078	47.9600	0.18156
	0	11.5000	67.2300	0.17105	1.8192	67.6200	0.02690
	+1	0.0001	65.0000	0.00000	15.3000	72.9800	0.20965
	+7	26.8400	32.9400	0.81481	23.5400	35.0500	0.67161
	+14	29.8200	58.9300	0.50602	24.9700	58.3400	0.42801

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