

Molecular Cloning, Sequence Identification and Tissue Expression Profile of Three Novel Sheep (*Ovis aries*) Genes-*RHOA*, *RCHY1* and *RSU1*

¹Liu Yonggang and ²Xia Xueshan

¹College of Animal Science and Technology,
Yunnan Agricultural University, 650201 Kunming, China

²Faculty of Life Science and Technology,
Kunming University of Science and Technology, 650224 Kunming, China

Abstract: The complete coding sequences of three sheep genes-*RHOA*, *RCHY1* and *RSU1* were amplified using the Reverse Transcriptase-Polymerase Chain Reaction (RT-PCR). The sheep *RHOA* gene encodes a protein of 193 amino acids that shares high homology with the ras homolog gene family, member A (RHOA) proteins of fifteen species- cattle (100%), rat (99%), human (100%), mouse (99%), chicken (99%), dog (99%), turkey (99%), sumatran orangutan (99%), poephila guttata (99%), Atlantic salmon (97%), green anole (96%), rainbow trout (96%), African clawed frog (96%), zebrafish (96%) and Western clawed frog (95%). The sheep *RCHY1* gene encodes a protein of 261 amino acids that shares high homology with the ring finger and CHY zinc finger domain containing 1 (RCHY1) proteins of nine species-cattle (94%), giant panda (92%), pig (91%), human (90%), white-tufted-ear marmoset (90%), chimpanzee (90%), rabbit (89%), mouse (86%) and rat (84%). The sheep *RSU1* gene encodes a protein of 277 amino acids that shares high homology with the ras suppressor protein 1 (RSU1) proteins of twelve species-cattle (99%), chimpanzee (98%), dog (99%), human (98%), giant panda (98%), horse (98%), rabbit (98%), chicken (96%), mouse (97%), African clawed frog (94%), Western-clawed-frog (94%) and zebrafish (91%). Finally, these three novel sheep genes were assigned to GeneIDs: 100302078, 100302546 and 100302548. The phylogenetic analysis revealed that the sheep *RHOA*, *RCHY1* and *RSU1* genes all have a closer genetic relationship with the *RHOA*, *RCHY1* and *RSU1* genes of cattle. Tissue expression profile analysis demonstrated that sheep *RHOA*, *RCHY1* and *RSU1* genes were all generally but differentially expressed in detected tissues.

Key words: Sheep, *RHOA*, *RCHY1*, *RSU1*, tissue expression, China

INTRODUCTION

Ras homolog gene family, member A (RHOA) belongs to the RhoA-like sub-family. The RhoA-like sub-family consists of RHOA, RHOB and RHOC. Latest researches demonstrated that RHOA promotes the formation of stress fibers and focal adhesions, regulating cell shape, attachment and motility (Su *et al.*, 2006). RHOA is vital for muscle contraction in vascular smooth muscle cells, RHOA plays a key role in cell contraction, differentiation, migration and proliferation (Peyton *et al.*, 2008). RHOA activities appear to be elaborately regulated in a time- and space-dependent manner to control cytoskeletal changes (Maekawa *et al.*, 1999; Ohta *et al.*, 1999; Saeki *et al.*, 2005). Ring finger and CHY zinc finger domain containing 1 (RCHY1), a member of proteins containing the ring finger and CHY zinc finger domain has ubiquitin ligase activity. It mediates E3-dependent

ubiquitination and proteasomal degradation of target proteins including TP53, HDAC1 and CDKN1B thus, regulating their levels and cell cycle progression (Wu *et al.*, 2010; Yan *et al.*, 2010; Shloush *et al.*, 2011).

Ras Suppressor protein 1 (*RSU1*) gene encodes a protein that is involved in the ras signal transduction pathway, growth inhibition and nerve-growth factor induced differentiation processes as determined in mouse and human cell line studies. In mouse, the encoded protein was initially isolated based on its ability to inhibit v-Ras transformation. Multiple alternatively spliced transcript variants for this gene have been reported; one of these variants was found only in glioma tumors (Chunduru *et al.*, 2002; Dougherty *et al.*, 2005, 2008).

As mentioned before, *RHOA*, *RCHY1* and *RSU1* genes are three genes which have important functions. Until today, *RHOA*, *RCHY1* and *RSU1* genes had been reported in human and other animals but the sheep *RHOA*,

RCHY1 and *RSU1* genes have not been reported yet. In present experiment, the researchers will isolate the coding sequences of sheep *RHOA*, *RCHY1* and *RSU1* genes based on the coding sequence information of *RHOA*, *RCHY1* and *RSU1* genes from human or other mammals and their highly homologous sheep ESTs sequence information, subsequently perform sequence and tissue expression profile analysis for these genes. These will establish the primary foundation of understanding these three sheep genes.

MATERIALS AND METHODS

Animals and sample preparation: Five adult Yunnan local sheep were slaughtered. Spleen, skin, lung, fat, muscle, heart, liver, kidney and ovary samples were collected, frozen in liquid nitrogen and then stored at -80°C . The total RNA was extracted using the total RNA extraction kit (Gibco, USA). First-strand cDNA synthesis was performed as that described by Liu *et al.* (2004). These first-strand cDNA samples were used to perform RT-PCR for the isolation of sheep *RHOA*, *RCHY1* and *RSU1* genes and for the tissue expression profile analysis.

Isolation of the sheep *RHOA*, *RCHY1* and *RSU1* genes:

The primers for sheep *RHOA* gene isolation were designed based on the coding sequence information of human *RHOA* gene and its highly homologous sheep EST sequences: DY503081 and EE758341. Similarly, the primers for sheep *RCHY1* gene isolation were designed based on the coding sequence information from human *RCHY1* gene and its highly homologous sheep EST sequences; DY519128 and EE754481. The primers for sheep *RSU1* gene isolation were designed based on the coding sequence information from human and mouse *RSU1* genes and their highly homologous sheep EST sequences; DY495018 and FE027712. These primer sequences and their annealing temperature for RT-PCR reaction were shown in Table 1. The RT-PCR was performed to isolate these three sheep genes using the pooled cDNAs from different tissues above. The 25 μL reaction system was; 2.0 μL cDNA, 2.5 μL 2 mM mixed dNTPs, 2.5 μL 10 \times Taq DNA polymerase buffer, 2.5 μL 25 mM MgCl_2 , 2.0 μL 10 μM forward primer, 2.0 μL 10 μM reverse primer, 2.0 units of Taq DNA polymerase (1 U/1 μL) and 9.5 μL sterile water. The PCR program initially started with a 94°C denaturation for 4 min followed by 35 cycles of $94^{\circ}\text{C}/50$ sec, $\text{Ta}^{\circ}\text{C}/50$ sec, $72^{\circ}\text{C}/50$ sec then 72°C extension for 10 min, finally 4°C to terminate the reaction (Table 1).

These PCR products for sheep *RHOA*, *RCHY1* and *RSU1* genes were then cloned into PMD18-T vector and sequenced bidirectionally with the commercial fluorometric method. At least five independent clones were sequenced for every gene.

Table 1: Primers for sheep *RHOA*, *RCHY1*, *RSU1* and *beta-actin* genes and their annealing Temperature ($\text{Ta}^{\circ}\text{C}$)

Genes	Primer sequence	$\text{Ta}^{\circ}\text{C}$
<i>RHOA</i>	Forward 5'- ATGGCTGCCATCCGGAAG-3'	61
	Reverse: 5'-TCACAAGACAAGGCACCCAG-3'	
<i>RCHY1</i>	Forward 5'-ATGGCTTCTCGACGCTG-3'	61
	Reverse: 5'-TCATTGCTGATCTAATGAAATTT-3'	
<i>RSU1</i>	Forward: 5'-ATGTCCAAGTCTCTGAAA-3'	52
	Reverse: 5'-TTATTTATTCTTGGCTGCC-3'	
<i>Beta-actin</i>	Forward: 5'-CTTGATGTCACGGACGATT-3'	56
	Reverse: 5'-CACGGCATTGTCCAACT-3'	

RT-PCR for tissue expression profile analysis: RT-PCR for tissue expression profile analysis was performed as previously described elsewhere (Liu and Gao, 2009; Yonggang and Shizheng, 2009; Liu, 2009). The researchers selected the housekeeping gene *beta-actin* (Accession No.: NM_001009784) as a positive control. The primers of sheep *RHOA*, *RCHY1* and *RSU1* genes which were used to perform the RT-PCR for tissue expression profile analysis were same as the primers for isolation RT-PCR above. The PCR reactions were optimized for a number of cycles to ensure product intensity within the linear phase of amplification. The 25 μL reaction system was: 1 μL cDNA ($100 \text{ ng } \mu\text{L}^{-1}$), 5 pmoles each oligonucleotide primer, 2.5 μL , 2 mmol L^{-1} mixed dNTPs, 2.5 μL 10 \times Taq DNA polymerase buffer, 2.5 μL 25 mmol L^{-1} MgCl_2 , 1.0 unit of Taq DNA polymerase and finally add sterile water to volume 25 μL . The PCR program initially started with a 94°C denaturation for 4 min followed by 25 cycles of $94^{\circ}\text{C}/50$ sec, $\text{Ta}^{\circ}\text{C}/50$ sec, $72^{\circ}\text{C}/50$ sec then 72°C extension for 10 min, finally 4°C to terminate the reaction (Table 1).

Sequence analysis: The cDNA sequence prediction was conducted using GenScan software (<http://genes.mit.edu/GENSCAN.html>). The protein prediction and analysis were performed using BLAST tool at the National Center for Biotechnology Information (NCBI) server (<http://www.ncbi.nlm.nih.gov/BLAST>) and the ClustalW software (<http://www.ebi.ac.uk/clustalw>).

RESULTS AND DISCUSSION

RT-PCR results for sheep *RHOA*, *RCHY1* and *RSU1* genes:

Through RT-PCR with pooled tissue cDNAs, for sheep *RHOA*, *RCHY1* and *RSU1* genes, the resulting PCR products were 582, 786 and 834 bp (Fig. 1).

Sequence analysis: These cDNA nucleotide sequence analysis using the BLAST software at NCBI server (<http://www.ncbi.nlm.nih.gov/BLAST>) revealed that these three genes were not homologous to any of the known sheep genes and they were then deposited into the GenBank database (Accession No.: FJ943984, FJ937958 and FJ937963). The sequence prediction was carried out

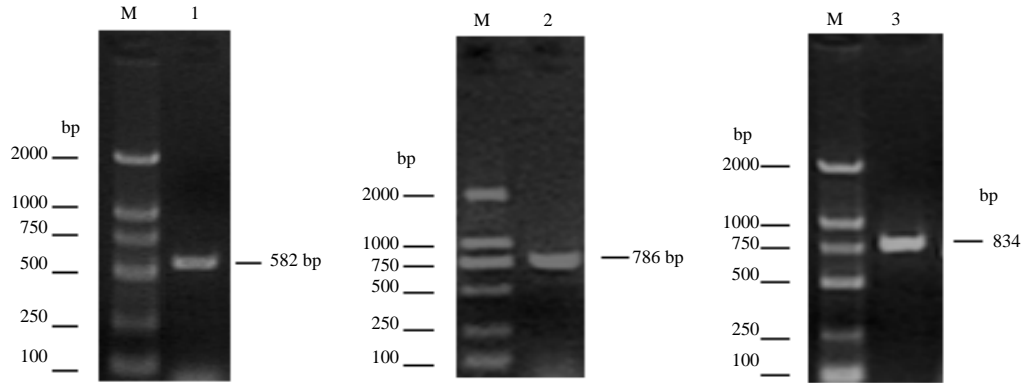


Fig. 1: RT-PCR results for sheep *RHOA*, *RCHY1* and *RSU1* genes. M: DL 2000 DNA markers; 1: PCR product for sheep *RHOA* gene; 2: PCR product for sheep *RCHY1* gene; 3: PCR product for sheep *RSU1* gene

using the GenScan software and results showed that the 582, 786 and 834 bp cDNA sequences represent three single genes which encoded 193, 261 and 277 amino acids, respectively. Finally, these three novel sheep genes were assigned to GeneIDs; 100302078, 100302546 and 100302548.

Further BLAST analysis of these proteins revealed that the sheep *RHOA* protein has high homology with the ras homolog gene family, member A (*RHOA*) proteins of fifteen species-cattle (Accession No.: NP_788818; 100%), rat (Accession No.: NP_476473; 99%), human (Accession No.: NP_001655; 100%), mouse (Accession No.: NP_058082; 99%), chicken (Accession No.: NP_990035; 99%), dog (Accession No.: NP_001003273; 99%), Turkey (Accession No.: ADX97247; 99%) sumatran orangutan (Accession No.: NP_001124755; 99%), poephila guttata (Accession No.: ACH45076; 99%), Atlantic salmon (Accession No.: ACI33725; 97%), green anole (Accession No.: XP_003216698; 96%), rainbow trout (Accession No.: NP_001158532; 96%), African clawed frog (Accession No.: AD40671; 96%), zebrafish (Accession No.: NP_998515; 96%) and Western clawed frog (Accession No.: CAJ81715; 95%) (Fig. 2).

The sheep *RCHY1* protein has high homology with the ring finger and CHY zinc finger domain containing 1 (*RCHY1*) proteins of nine species-cattle (Accession No.: NP_001077223; 94%), giant panda (Accession No.: XP_002919200; 92%), pig (Accession No.: NP_001090959; 91%), human (Accession No.: NP_056251; 90%), white-tufted-ear marmoset (Accession No.: XP_002745761; 90%), chimpanzee (Accession No.: XP_517222; 90%), rabbit (Accession No.: XP_002717057; 89%), mouse (Accession No.: NP_0808337; 86%) and rat (Accession No.: NP_001007619; 84%) (Fig. 3).

The sheep *RSU1* protein has high homology with the ras suppressor protein 1 (*RSU1*) proteins of twelve species-cattle (Accession No.: NP_001035691; 99%), chimpanzee (Accession No.: XP_001151460; 98%), dog (Accession No.: XP_535177; 99%), human (Accession No.: NP_036557; 98%), giant panda (Accession No.: XP_002918673; 98%), horse (Accession No.: XP_001498103; 98%), rabbit (Accession No.: XP_002717469; 98%), chicken (Accession No.: NP_001186520; 96%), mouse (Accession No.: NP_033131; 97%), African clawed frog (Accession No.: NP_001085943; 94%), Western-clawed-frog (Accession No.: NP_001015695; 94%) and zebrafish (Accession No.: XP_002666722; 91%) (Fig. 4).

Based on the results of the alignment of *RHOA*, *RCHY1* and *RSU1* proteins, three phylogenetic trees were constructed using the Dendrogram procedure of ClustalW software as shown in Fig. 5-7. The phylogenetic analysis revealed that the sheep *RHOA*, *RCHY1* and *RSU1* genes all have a closer genetic relationship with the *RHOA*, *RCHY1* and *RSU1* genes of cattle.

Tissue expression profile: Tissue expression profile analysis was carried out and results revealed that the sheep *RHOA*, *RCHY1* and *RSU1* genes are all generally but differentially expressed in tissues including spleen, lung, muscle, kidney, ovary, skin, liver, heart and fat (Fig. 8).

In the current study, the researchers firstly get the coding sequences of sheep *RHOA*, *RCHY1* and *RSU1* genes by RT-PCR. With the development of modern bioinformatics, establishment of specific sheep NCBI EST database and different convenient analysis tools, researchers can easily find the useful ESTs which were highly homologous to the coding sequences of human

Rat	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDGKQVELALWDT
Mouse	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDGKQVELALWDT
Sheep	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDGKQVELALWDT
Cattle	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDGKQVELALWDT
Human	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDGKQVELALWDT
Sumatran orangutan	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDGKQVELALWDT
Dog	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDGKQVELALWDT
Poephila guttata	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDGKQVELALWDT
Chicken	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDGKQVELALWDT
Turkey	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDGKQVELALWDT
Zebrafish	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDSKQVELALWDT
Green anole	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDGKQVELALWDT
Rainbow trout	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDSKQVELALWDT
Atlantic salmon	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDSKQVELALWDT
African clawed frog	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDGKQVELALWDT
Western clawed frog	MAAIRKKLVIVGDGACGKTCLLIVFSKQFPEVYVPTVFENYVADIEVDSKQVELALWDT
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Rat	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
Mouse	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
Sheep	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
Cattle	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
Human	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
Sumatran orangutan	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
Dog	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
Poephila guttata	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLQNIPEKWTPEVKHFCPNVPIILVGNKKD
Chicken	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
Turkey	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
Zebrafish	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
Green anole	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
Rainbow trout	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
Atlantic salmon	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
African clawed frog	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
Western clawed frog	AGQEDYDRLRPLSYPDTDVILMCFIDSIDSLSLENIPEKWTPEVKHFCPNVPIILVGNKKD
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Rat	LRNDEHTRRELAKMKQEPVKPEEGRDMANRIGAFGYMECSAKTKDGVREVFEMATRAALQ
Mouse	LRNDEHTRRELAKMKQEPVKPEEGRDMANRIGAFGYMECSAKTKDGVREVFEMATRAALQ
Sheep	LRNDEHTRRELAKMKQEPVKPEEGRDMANRIGAFGYMECSAKTKDGVREVFEMATRAALQ
Cattle	LRNDEHTRRELAKMKQEPVKPEEGRDMANRIGAFGYMECSAKTKDGVREVFEMATRAALQ
Human	LRNDEHTRRELAKMKQEPVKPEEGRDMANRIGAFGYMECSAKTKDGVREVFEMATRAALQ
Sumatran orangutan	LRNDEHTRRELAKMKQEPVKPEEGRDMANRIGAFGYMECSAKTKDGVREVFEMATRAALQ
Dog	LRNDEHTRRELAKMKQEPVKPEEGRDMANRIGAFGYMECSAKTKDGVREVFEMATRAALQ
Poephila guttata	LRNDEHTRRELAKMKQEPVKPEEGRDMANRIGAFGYMECSAKTKDGVREVFEMATRAALQ
Chicken	LRNDEHTRRELAKMKQEPVKPEEGRDMANRIGAFGYMECSAKTKDGVREVFEMATRAALQ
Turkey	LRNDEHTRRELAKMKQEPVKPEEGRDMANRIGAFGYMECSAKTKDGVREVFEMATRAALQ
Zebrafish	LRNDEHTRRELAKMKQEPVKPEEGRDMANRINAFGYLECSAKTKDGVREVFEMATRAALQ
Green anole	LRNDEHTRRELAKMKQEPVKPEEGRDMANRIGAFGYLECSAKTKDGVREVFEMATRAALQ
Rainbow trout	LRNDEHTRRELAKMKQEPVKPEEGRDMANRISAFGYMECSAKTKDGVREVFEMATRAALQ
Atlantic salmon	LRNDEHTRRELAKMKQEPVKPEEGRDMANRIGAFGYMECSAKSKDGVREVFEMATRAALQ
African clawed frog	LRNDEHTRRELTKMKQEPVKPEEGRDMANRISAYAYMECSAKTKDGVREVFELATRAALQ
Western clawed frog	LRNDEHTRRELTKMKQEPVKPEEGRDMANRISAYGYMECSAKTKDGVREVFELATRAALQ
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Rat	ARRGKKKSGCLIL
Mouse	ARRGKKKSGCLIL
Sheep	ARRGKKKSGCLVL
Cattle	ARRGKKKSGCLVL
Human	ARRGKKKSGCLVL
Sumatran orangutan	ARRGKKKSGRLVL
Dog	ARRGKKKSGCLVL
Poephila guttata	ARRGKKKSGCLFL
Chicken	ARRGKKKSGCLLL
Turkey	ARRGKKKSGCLLL
Zebrafish	ARRGKKKSGCLLL
Green anole	AKRGRKKTSCQLL
Rainbow trout	ARRGKPRNKCLLL
Atlantic salmon	ARRGKKSNNKCLLL
African clawed frog	ARRGKKKPRCLLI
Western clawed frog	ARRGKKKTTCLLI
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Fig. 2: The alignment of the protein encoded by sheep *RHOA* gene and fifteen other kinds *RHOA* proteins


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African clawed frog      MSKSLKKKIVEESREKNVDPIDMCDRGIANMLDVPGLFTLSHITQLILSHNKLTTPVPPNIA
Western clawed frog     MSKSLKKKIVEESREKNVDPIDMCDRGIANMLDVPGLFTLSHITQLILSHNKLTTPVPPNIA
Sheep                    MSKSLKKKIVEESREKNQPEVDMSDRGISNMLDVNGLFTLSHITQLVLSHNKLTTPVPPNIA
Cattle                   MSKSLKKKIVEESREKNQPEVDMSDRGISNMLDINGLFTLSHITQLVLSHNKLTTPVPPNIA
Giant panda              MSKSLKKKIVEESREKNQPEVDMSDRGISNMLDVNGLFSLSHITQLVLSHNKLTTPVPPNIA
Horse                    MSKSLKKKIVEESREKNQPEVDMSDRGISNMLDVNGLFSLSHITQLVLSHNKLTTPVPPNIA
Dog                      MSKSLKKKIVEESREKNQPEVDMSDRGISNMLDVNGLFSLSHITQLVLSHNKLTTPVPPNIA
Mouse                    MSKSLKKKIVEESREKNQPEVDMSDRGISSMLDVNGLFSLAHITQLVLSHNKLTTPVPPNVA
Rabbit                   MSKSLKKKIVEESREKNQPEVDMSDRGISNMLDVNGLFTLSHITQLVLSHNKLTTPVPPNIA
Chimpanzee              MSKSLKKMVEESREKNQPEVDMSDRGISNMLDVNGLFTLSHITQLVLSHNKLTTPVPPNIA
Human                    MSKSLKKKIVEESREKNQPEVDMSDRGISNMLDVNGLFTLSHITQLVLSHNKLTTPVPPNIA
Chicken                  MSKSLKKKIVEESREKNQPEVDMCDRGIANMLDVPGLFTLSHITQLVLSHNKLTTPVPPNIA
Zebrafish                MSKSLKKKIVEESRDKNLPEVDMCDRGIANLLDIPGLFSLSSITQLVLSHNKLSAVPPNIA
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African clawed frog      DLKNLEVLNFFNNQIEELPTQISSLQKCLKHLNLMNRLNGLPRGFGSLPALEVLDTLYNN
Western clawed frog     DLKNLEVLNFFNNQIEELPTQISSLQKCLKHLNLMNRLNSLPRGFGSLPALEVLDTLYNN
Sheep                    ELKNLEVLNFFNNQIEELPTQISSLQKCLKHLNLMNRLNTLPRGFGSLPALEVLDTLYNN
Cattle                   ELKNLEVLNFFNNQIEELPTQISSLQKCLKHLNLMNRLNTLPRGFGSLPALEVLDTLYNN
Giant panda              ELKNLEVLNFFNNQIEELPTQISSLQKCLKHLNLMNRLNTLPRGFGSLPALEVLDTLYNN
Horse                    ELKNLEVLNFFNNQIEELPTQISSLQKCLKHLNLMNRLNTLPRGFGSLPALEVLDTLYNN
Dog                      ELKNLEVLNFFNNQIEELPTQISSLQKCLKHLNLMNRLNTLPRGFGSLPALEVLDTLYNN
Mouse                    ELKNLEVLNFFNNQIEELPTQISSLQKCLKHLNLMNRLNTLPRGFGSLPALEVLDTLYNN
Rabbit                   ELKNLEVLNFFNNQIEELPTQISSLQKCLKHLNLMNRLNTLPRGFGSLPALEVLDTLYNN
Chimpanzee              ELKNLEVLNFFNNQIEELPTQISSLQKCLKHLNLMNRLNTLPRGFGSLPALEVLDTLYNN
Human                    ELKNLEVLNFFNNQIEELPTQISSLQKCLKHLNLMNRLNTLPRGFGSLPALEVLDTLYNN
Chicken                  DLRNIEVLNFFNNQIEELPTQISSLQKCLKHLNLMNRLNTLPRGFGSLPALEVLDTLYNN
Zebrafish                DLKNLEVLNMFNNQIEELPTQISSLQKCLKHLNLMNRLSTLPRGFGSLPALEVLDTLYNN
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African clawed frog      LNENSLPGNFFYLTTLRALYLSDDNDFETLPPDIGKLTQIISLRDNDLISLPKEVGELT
Western clawed frog     MNENSLPGNFFYLTTLRALYLSDDNDFEILPPDIGKLTQIISLRDNDLISLPKEIGDLT
Sheep                    LNENSLPGNFFYLTTLRALYLSDDNDFEILPPDIGKLTQIISLRDNDLISLPKEIGELT
Cattle                   LNENSLPGNFFYLTTLRALYLSDDNDFEILPPDIGKLTQIISLRDNDLISLPKEIGELT
Giant panda              LSENSLPGNFFYLTTLRALYLSDDNDFEILPPDIGKLTQIISLRDNDLISLPKEIGELT
Horse                    LNESSLPGNFFYLTTLRALYLSDDNDFEILPPDIGKLTQIISLRDNDLISLPKEIGELT
Dog                      LNENSLPGNFFYLTTLRALYLSDDNDFEILPPDIGKLTQIISLRDNDLISLPKEIGELT
Mouse                    LNEHSLPGNFFYLTTLRALYLSDDNDFEILPPDIGKLTQIISLRDNDLISLPKEIGELT
Rabbit                   LNENSLPGNFFYLTTLRALYLSDDNDFEILPADIGKLTQIISLRDNDLISLPKEIGELT
Chimpanzee              LSENSLPGNFFYLTTLRALYLSDDNDFEILPPDIGKLTQIISLRDNDLISLPKEIGELT
Human                    LSENSLPGNFFYLTTLRALYLSDDNDFEILPPDIGKLTQIISLRDNDLISLPKEIGELT
Chicken                  LNENSLPGNFFYLTTLRALYLSDDNDFEILPPDIGKLTQIISLRDNDLISLPKEIGELT
Zebrafish                LNESSLPGNFFYLTTLRALYLSDDNDFEILPPDIGKLAQLQIISLRDNDLISLPKEIGDLT
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African clawed frog      QLKELHIQGNRLTVLPPELGNLDTGQKQVFAENNPWVTPADQFQVGVSHVFEYIRSE
Western clawed frog     QLKELHIQGNRLTVLPPELGNLDTGQKQVFAENNPWVTPADQFQVGVSHVFEYIRSE
Sheep                    QLKELHIQGNRLTVLPPELGNLDTGQKQIFKAENNPWVTPADQFQVGVSHVFEYIRSE
Cattle                   QLKELHIQGNRLTVLPPELGNLDTGQKQVFAENNPWVTPADQFQVGVSHVFEYIRSE
Giant panda              QLKELHIQGNRLTVLPPELGNLDTGQKQVFAENNPWVTPADQFQVGVSHVFEYIRSE
Horse                    QLKELHIQGNRLTVLPPELGNLDTGQKQVFAENNPWVTPADQFQVGVSHVFEYIRSE
Dog                      QLKELHIQGNRLTVLPPELGNLDTGQKQVFAENNPWVTPADQFQVGVSHVFEYIRSE
Mouse                    QLKELHIQGNRLTVLPPELGNLDTGQKQVFAENNPWVTPADQFQVGVSHVFEYIRSE
Rabbit                   QLKELHIQGNRLTVLPPELGNLDTGQKQVFAENNPWVTPADQFQVGVSHVFEYIRSE
Chimpanzee              QLKELHIQGNRLTVLPPELGNLDTGQKQVFAENNPWVTPADQFQVGVSHVFEYIRSE
Human                    QLKELHIQGNRLTVLPPELGNLDTGQKQVFAENNPWVTPADQFQVGVSHVFEYIRSE
Chicken                  QLKELHIQGNRLTVLPPELGNLDTGQKQVFAENNPWVTPADQFQVGVSHVFEYIRSE
Zebrafish                QLKELHIQGNRLTVLPPELGNLDTGPKQVFAENNSWVTPADQFQVGVSHVFEYIRSE
*****:*****:*****. *.:*****:*****:*****:*. *.:**

African clawed frog      TYKYLGRHMQANPEPPKKNNDKSKKISRKPLAAKKN
Western clawed frog     TYKYLGRHMQANPEPPKKNNDKSKKISRKPLAAKKN
Sheep                    TYKYLGRHMQANPEPPKKNNDKSKKISRKPLAAKKN
Cattle                   TYKYLGRHMQANPEPPKKNNDKSKKISRKPLAAKKN
Giant panda              TYKYLGRHMQANPEPPKKNNDKSKKISRKPLTAKNK
Horse                    TYKYLGRHMQANPEPPKKNNDKSKKISRKPLTAKNK
Dog                      TYKYLGRHMQANPEPPKKNNDKSKKISRKPLTAKNK
Mouse                    TYKYLGRHMQANPEPPKKNNDKSKKISRKPLAAKKN
Rabbit                   TYKYLGRHMQANPEPPKKNNDKSKKISRKPLTAKNK
Chimpanzee              TYKYLGRHMQANPEPPKKNNDKSKKISRKPLAAKKN
Human                    TYKYLGRHMQANPEPPKKNNDKSKKISRKPLAAKKN
Chicken                  TYKYLGRHMQANPEPPKKNNDKSKKISRKPLAAKKN
Zebrafish                TYKYLGRHMQANPEAPKKNADKSKKISRKPLAAKKN
*****:*****:*. *.:*****:*****:*****:*. *.:

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Fig. 4: The alignment of the protein encoded by sheep *RSU1* gene and twelve other kinds of RSU1 proteins

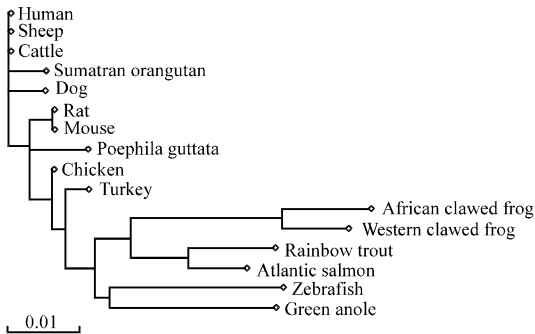


Fig. 5: The phylogenetic analysis for sixteen kinds of *RHOA* genes

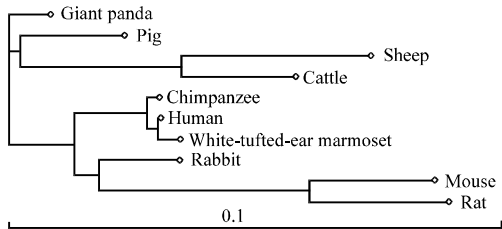


Fig. 6: The phylogenetic analysis for ten kinds of *RCHY1* genes

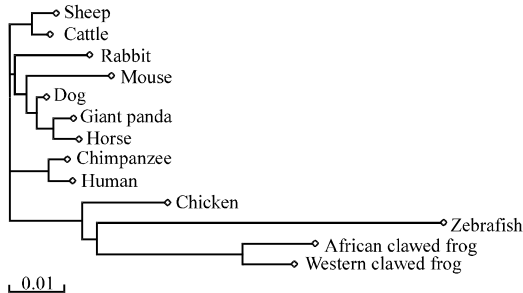


Fig. 7: The phylogenetic analysis for thirteen kinds of *RSU1* genes

genes. Based on these sheep EST sequences, the researchers can obtain the complete coding sequences of some novel sheep genes through the some experimental methods such as RT-PCR. From the clone and sequence analysis of sheep *RHOA*, *RCHY1* and *RSU1* genes, it could be seen that this is an effective method to isolate some novel sheep genes.

Through sequence analysis, the researchers found that the encoding protein of the sheep *RHOA*, *RCHY1* and *RSU1* genes are highly homologous with *RHOA*, *RCHY1* and *RSU1* proteins of human and some other animals. This implied that the *RHOA*, *RCHY1* and *RSU1* genes were highly conserved in some species and the sheep *RHOA*, *RCHY1* and *RSU1* genes will have similar

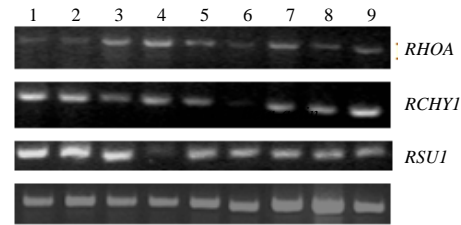


Fig. 8: Tissue expression distribution of sheep *RHOA*, *RCHY1* and *RSU1* genes. The beta-actin expression is the internal control. 1: muscle; 2: heart; 3: lung; 4: spleen; 5: skin; 6: fat; 7: liver; 8: kidney; 9: ovary

functions as the *RHOA*, *RCHY1* and *RSU1* genes of human and other animals. The researchers also found that the sheep *RHOA*, *RCHY1* and *RSU1* proteins do not show complete identity to some animals. This implied that the sheep *RHOA*, *RCHY1* and *RSU1* genes will have some differences in functions to those of other animals.

The phylogenetic analysis revealed that the sheep *RHOA*, *RCHY1* and *RSU1* genes all have a closer genetic relationship with the *RHOA*, *RCHY1* and *RSU1* genes of cattle. This implied that we can use cattle as a model organism to study the sheep *RHOA*, *RCHY1* and *RSU1* genes or use sheep as a model organism to study the cattle *RHOA*, *RCHY1* and *RSU1* genes.

From the tissue distribution analysis in the experiment, it can be seen that the sheep *RHOA*, *RCHY1* and *RSU1* genes were obviously differentially expressed in some tissues. As the researchers did not study functions at protein levels yet there might be many possible reasons for differential expression of sheep *RHOA*, *RCHY1* and *RSU1* genes. The suitable explanation for this under current conditions is that at the same time those biological activities related to the mRNA expression of sheep *RHOA*, *RCHY1* and *RSU1* genes were presented diversely in different tissues.

CONCLUSION

In this study, the researchers first isolated the sheep *RHOA*, *RCHY1* and *RSU1* genes and performed necessary sequence and tissue expression profile analysis. This established the primary foundation for further insight into these novel sheep genes.

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