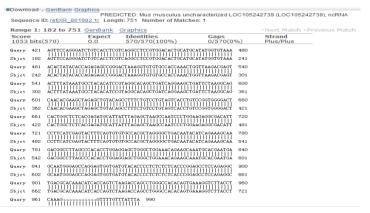


## LncRNA 3'及 5'RACE 实验

合同号: PG1-1601004

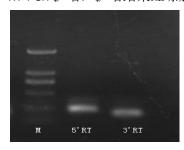
1. 将提供的序列在 NCBI 上进行比对, 比对结果如下:



2. 我们采用能够比对得上的保守区域进行 5′及 3′两端引物的设计, 所用设计引物的序列标注红色如下:

TAGCAGTGCAGGTCATAGGAAAGCGAGTATGTGTCATAGAGGGTTGCCTTCACGTTCAGGCAGCCAATAAAGTCCT
GAGGGTTCAGCTTGTACAGGTCGAAGGTTACCCGGGCCACATCGATCTTCTTAGTTGGCTTCTGGGAAAGGGCTAGCTG
GGCTCTCTTCTTGCGCTGCAAGACAAGCCCGTGTAAGCTCACCTCTGGAACATCTCCCGAGCTCTCGACATTCACCCCTGC
CCCGGGAAGGCCGTCACCTGTTCTGATGGGGGCTTCCACTTCTGCCCCTTCAGCAGGACCATGAACATGGTATCTTTTGC
CAGGGCTTGGAAGTATTCTTCTGTCTCCACGATTGTGCCATCTTCCTCCAGCACCAGGGAGAAGGGCTTGTCTTTAAGTTT
CAAGATGTCCTGGAACGA

2.1 以红色部分作为原始序列进行引物的设计,以提取的 RNA 逆转录后的 cDNA 为模板进行 RT-PCR 扩增,扩增结果经琼脂糖凝胶电泳检测如下:



注: 采用 DL2000 的 Marker

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2.2 RT-PCR 产物送测序,将测序结果进行比对,比对结果如下:

#### 3′比对结果:

# □ Download ∨ GenBank Graphics PREDICTED: Mus musculus uncharacterized LOC105242739 (LOC105242739), ncRNA

 Sequence ID: ref|XR 861982.1| Length: 751 Number of Matches: 1

 Range 1: 512 to 674 GenBank Score
 Graphics
 ▼ Next Match ▲ Previous Match

 Score
 Expect Identities
 Gaps Ji/163(0%)
 Strand Plus/Plus

 294 bits(159)
 3e-76
 162/163(99%)
 1/163(0%)
 Plus/Plus

 Query
 11
 GCTAGGGGCTGAC-ATACATCAGAAAGCAAGACGGCCTTAGCCCACACCTGGAGGAGCTG
 69

 Query
 11
 GCTAGGGGCTGAC-ATACATCAGAAAGCAAGACGGCCTTAGCCCACACCTGGAGGAGGCTG
 69

 Sbjet
 512
 GCTAGGGGCTGACAATACATCAGAAAGCAAGACGGCCTTAGCCCACACCTGGAGGAGCTG
 571

 Query
 70
 GGCTGGAAACAGAAGCAAATGCACGAATGAGCAATGGGAGCCAGGAGTGGTGATGTACAC
 129

 Sbjet
 572
 GGCTGGAAACAGAAGCAAATGCACGAATGAGCAATGGGAGCCAGGAGTGGTGATGTACAC
 631

 Query
 130
 CCCTCTCTCTCACCCGGAGCCTCCAGAGGCTGACGCACAAACA
 172

 Sbjet
 632
 CCCTCTCTCTCACCCGGAGCCTCCAGAGGCTGACGCACAAACA
 674

#### 5'比对结果:

PREDICTED: Mus musculus uncharacterized LOC105242739 (LOC105242739), ncRNA Sequence ID: ref[XR 861982.1] Length: 751 Number of Matches: 1

Range 1: 286 to 509 GenBank Graphics ▼ Next Match ▲							Previous Mato
Score 390 bits(211)		Expect 5e-105	Identities 220/224(98%)	Gaps 1/224(0%)	Strand Plus/Plus		
Query	15	•		AAATGCCTACACATCCGTAGGC		, , , , , , , , , , , , , , , , , , , ,	
Sbjet	286	ACTGGTTAAGA	.CGAGTACTTTAT	AAATGCCTACACATCCGTAGGC		5	
Query	74	GCTGATTCTAA	GGCAGCAACACG	AAGCTAGAGCTGTACAGCCTTT	CTGTCCTGTAGTCAC 13	3	
Sbjet	346	GCTGATTCTAA	GGCAGCAACACG	AAGCTAGAGCTGTACAGCCTTT	CTGTCCTGTAGTCAC 40	5	
Query	134	CTGTCCGGTGG	GGACTCACTGGC	TCTCACGAGATGCATTATTTAG	AGCTAAGCCAATCCC 19	3	
Sbjet	406	CTGTCCGGTGG	GGACTCACTGGC	TCTCACGAGATGCATTATTTAG	AGCTAAGCCAATCCC 46	5	
Query	194	TGGAAGAGGCG	ACATTCCTTCAT	CGAGTACTTTAAATGTGTGCC	237		
Sbjet	466	TGGAAGAGGCG	ACATTCCTTCAT	CGAGTACTTTCAGTGTGTGCC	509		

2.3 RT-PCR 结果比对正确,下一步安排进行 5'及 3'race 实验扩增尝试