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In 10 μ L Volume Using 50 Ng Linearized PXST, 50 Ng Puri-fied Fragment, 5 U T4 DNA Ligase, 1 μ L 10 \times T4 Buffer And 6 μ LddH 20. The Mixture Was Incubated At 22 $^{\circ}$ C For 1 H Andthenaddedinto100 μ L E. Coli DH5 α Chemically Com-petent Cells, Followed By Incubation On Ice For 30 Min And Heat 4th, 2024

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Other SAMS Using Neighbour-Joining Method (Fig. 2). Based On 1000 Replicates, The Phylogenetic Tree Was Divided Into Fig 2: Unrooted Phylogenic Tree Of Protein Sequences Of CcSAMS And Other SAMSs Constructed By The Neighbor-joining Method. Bootstrap Majority Consensus Values O 4th, 2024

2. J. Sambrook And D. W. Russell, Molecular Cloning: A ...

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Protein From L. Acidophilus NCFM Was Extracted And Purified. en The Secondary Structure And Adhesion Ability Ont 1th, 2024

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3, Respectively. The PCR Conditions, With TaKaRa LA Taq Polymerase And GC Buffer I, Were Essentially The Same As Those Described Previ-ously (10). The Nucleotide Sequence Of The Amplified DNA Fragment (507 Bp From The 5-end Of Pld1) Was 1th, 2024

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And Cloned Into The Same Sites Of PcDNA3.1/Hygro (Invitrogen, Carls-bad, CA),

Resulting In PcDNA3.1-C45T-1 (the Name Of C45T-1 Was Given After The Determination Of Acceptor Specificity). The Second Gene Was Initially Identified In AA233362 And AA777237 Derived F 3th, 2024

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Gibson Cloning Procedure • Design Four PCR Primers With Overlap Between Insert/insert Or Insert/vector • Adjacent Segments Should Have Identical Sequences (A And B) • PCR Primers Should Contain 5' End That Is The Same As Adj 2th, 2024

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Proc. Natl. Acad. Sci. USA92 (1995) Withphenol, DNAwasprecipitatedwithethanol Andspooled Onto A Glass Rod. Construction And Screening Ofa AExpression Library. S. Solfataricus Genomic DNAwaspartially Digested With Sau3A And Fragments From 0.5 To 2.5 Kb Were Ligated To BamHI- Digested And Dephosphorylated AZAPExpress Arms (Strat- Agene); Packagingwasperformed According To The ... 4th, 2024

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And 1 Kb Plus DNA Ladder, Invitrogen (catalogue No. 10787-018) Was Used As Marker To Determine The Size Of DNA Fragment. DNA Recovery The Amplified DNA Fragment Verified With Gel Electrophoresis Was Extracted Using TM Purelink Quick Gel Extraction Kit From Invitrogen (catalogue No. K2100-12). Four Tubes Of PCR Reaction (25 1th, 2024

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Bled Using Vector NTI Express Software (Life Technolo-gies, Carlsbad, CA) Based On The Obtained Partial Frag-ment And The 3'-/5'-full RACE Products. Amino Acid ... (Table 1) Were Designed Using Primer Express 3.0 To Develop A Real-time PCR Assay For Detecting Duck MAPK1 Expression. The Target 213-bp Amplicon Was 1th, 2024

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Gao Et Al. 3149 Table 1. Primers Used In This Study. Primers Name Sequence(5' To

3') SMART IVTM Oligonucleotide

AAGCAGTGGTATCAACGCAGAGTGGCCATTACGGCCGGG CDS III/3 ... 2th, 2024

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ization Of Corkscrew (csw), Which Is Known To Be Maternally Required For Normal Determination Of Cell Fates At The Termini Of Drosophila Embryos [10], Showed It To Be A Member Of The Cytoplasmic Family Of PTPs [11]. Mutational Analysis Has Shown That Csw Works In Concert With D-raf To Positively Transduce The Torso Receptor Protein-tyrosine ... 2th, 2024

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4Boyer, R F (1986) Modern Experimental Biochemistry, Addison- Wesley Publishing Co, Reading, MA, USA -5Maniatis, T, Fritsch, E F, And Sambrook, J (1982) Molecular Cloning. A Laboratory Manual, Cold Spring Harbor Laboratory, New York, NY, USA 6Clark, John M, Switzer, R L (1977) Experimental Biochemistry, W H Cited By: 20 Publish Year: 1992 Author: J R L Walker 1th, 2024

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