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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 Jan 8th, 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 Previously (10). The Nucleotide Sequence Of The Amplified DNA Fragment (507 Bp From The 5-end Of Pld1) Was Jan 8th, 2024

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And Cloned Into The Same Sites Of PcDNA3.1/Hygro (Invitrogen, Carlsbad, CA), Resulting In PcDNA3.1-C4ST-1 (the Name Of C4ST-1 Was Given After The Determination Of Acceptor Specificity). The Second Gene Was Initially Identified In AA233362 And AA777237 Derived F Feb 7th, 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 Feb 21th, 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 ... Jan 15th, 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 ... Feb 18th, 2024

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