ProteoGenix SAS
Espace Européen de l'Entreprise
15 rue de La Haye
67300 Schiltigheim, France
Tel. +33 (0)3 90 20 54 70 - Fax +33 (0)3 88 56 16 88
contact@proteogenix.fr
www.ProteoGenix.science

SGCA-sarcoglycan, alpha (50kDa dystrophin-associated glycoprotein) Gene

Reference: PTXBC025702

Product type:	DNA	Storage Condition:	-20°C
Size/Quantity:	2 ug	Delivery Condition:	+4°C, Blue Ice
Synonyms	50DAG; ADL; DAG2; DMDA2; LGMD2D; SCARMD1; adhalin; alpha-sarcoglycan; 50 kDa dystrophin-associated glycoprotein; 50kD DAG; alpha-SG; dystroglycan-2; limb girdle muscular dystrophy 2D; sarcoglycan, alpha (50kDa dystrophin-associated glycoprotein); sarcogly		

For research use only.

GENE DESCRIPTION	sarcoglycan, alpha (50kDa dystrophin-associated glycoprotein)
GENE ID	6442
ORIGIN SPECIES	Human
ACCESSIONS	BC025702
VECTOR	pDONR223
NCBI SYMBOL	SGCA
LENGTH	ProteoGenix bp
SEQUENCE PRIMERS	Forward primer M13R (5'→3'): GTAAAACGACGGCCAGT
	Reverse primer T7P (5'→3'): TAATACGACTCACTATAGG

ORF (Only for reference, product provided without STOP codon)

atggctgagacactcttctggactcctctcctcgtggttctcctggcagggctgggggacaccgaggcccagcagaccacgctacacccacttgtgggccgtgtctttgtgcacaccttggaccatgagacgtttctgagccttcctgagcatgtcgctgtcccacccgctgtccacatcacct
accacgcccacctccagggacacccagacctgccccggtggctccgctacacccagcgcagccccacccacccctggcttcctctacggctctgccacccccagaagatcgtgggctccaggtcattgaggtcacagcctacaatcgggacagctttgataccactcggcagaggctggt
gctggagattggggacccagaaggccccctgctgccataccaagccgagttcctggtgcgcagccacgatgcggaggaggtgctgccctcaacacctgccagccgcttcctctcagccttgggggactctgggagcccggagagcttcagctgctcaacgtcacctctgccttggac
cgtgggggccgtgtccccttcccattgagggccgaaaagaagggtatacattaaggtgggttctgcctcacctttttctacttgcctgaagatggtggcatcccccgatagccacgcccgctgtgcccagggccagcctccacttctgtcttgctacgacaccttggcaccccacttccgc
gttgactggtgcaatgtgaccctggtggataagtcagtgccggagcctgcagatgaggtgcccaccccaggtgatgggatcctggagcatgacccgttcttctgcccacccactgaggccccagaccgtgacttcttggtggatgctctggtcaccctcctggtgcccctgctggtggccct
gcttctcaccttgctgctggcctatgtcatgtgctgcccggcgggagggaaggctgaagagagacctggctacctccgacatccagatggtccaccactgcaccatccacgggaacacagaggagctgcggcagatggcggccagccgcgaggtgccccggccactctccaccctgcc
catgttcaatgtgcacacaggtgagcggctgcctccccgcgtggacagcgcccaggtgcccctcattctggaccagcactga