



Reference: PTXBC014662

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|----------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------|----------------|
| Product type:  | DNA                                                                                                                                                                                                                                      | Storage Condition:  | -20°C          |
| Size/Quantity: | 2 ug                                                                                                                                                                                                                                     | Delivery Condition: | +4°C, Blue Ice |
| Synonyms       | PACE-1; PACE1; protein-associating with the carboxyl-terminal domain of ezrin; SCY1-like 3; SCY1-like protein 3; SCY1-like, kinase-like 3; ezrin-binding partner PACE-1 (PACE-1); ezrin-binding protein PACE-1; SCY1 like pseudokinase 3 |                     |                |

**For research use only.**

|                         |                                                                                                   |
|-------------------------|---------------------------------------------------------------------------------------------------|
| <b>GENE DESCRIPTION</b> | SCY1-like 3 ( <i>S. cerevisiae</i> )                                                              |
| <b>GENE ID</b>          | 57147                                                                                             |
| <b>ORIGIN SPECIES</b>   | Human                                                                                             |
| <b>ACCESSIONS</b>       | BC014662                                                                                          |
| <b>VECTOR</b>           | pDONR223                                                                                          |
| <b>NCBI SYMBOL</b>      | SCYL3                                                                                             |
| <b>LENGTH</b>           | ProteoGenix bp                                                                                    |
| <b>SEQUENCE PRIMERS</b> | Forward primer M13R (5'→3'): GTAAAACGACGGCCAGT<br>Reverse primer T7P (5'→3'): TAATACGACTCACTATAGG |

atggagatcagagaacatgctcttaaaagagctatatactgagagaaacacacattacccttacccttggactgctgtttatcccgcgtactgcgaatgagcaaatgtcttcagttttgtgtataagagagaaaatgagacaaggttaataaagctgccaaagcattggaagacacttgcacactctgtctgaagtatttctatctgtactgcgaagccgagctcatctatctgtcactgagcagatgcacccctggaaagtctgttgaaaccttcttcgcagagactgtgctgcggatcatgatcatatctgcgcctcatctcatcatgacagagacactcaacacaataatgtctgtttatcctgtttgtgtgaggaagatcactgcgaagctagggaagatggaacctgttttaaaattttcacaacacacagactgtctgaggatgagcatcaataaagagaccacatctctctgcgaagatgtctcacaattcaaacactctccagatgtgtcattgcagatgcctgggatgcctttcatttggacaatctgttggaaagtgtgtccacaactcttaaatgacacagctttcagcggatgtcttccagctttcaacagaccttgcactcaactttgtgtaactccatccaaaatgtcggccagcgtctgcacacttactatctatgacttctcagaatgattttctggaagtgttaaaattctgaaagtgttaaacatgagatgtgagaagagaaaacgaaattcttcaatctctgcgcagacagctgcagctctctgcagagaatgatagctcaaaagttgtgcctctctcttcaatcagctgtgttttgcagagcagcagctgttgtaaaagttttcttctatctctgtcccccataaaagatctgcagcgaagaacactctctctctcactcacagcccttcaatctgtctctcactcagcgcactgcaggacactcaacagacagctcagaagaagatctgtccacacagttttgtcggcctgcgtgatactagtattccattgtggcaattactctgcatagcctagcgtctggtctctctctgttggaccagagctgtgttggaggaggagaacgaaccaagatctcaaacgcactccccaaagtttactaaaaatactgactcttctcagaaggtgatactcttccagctttaaatttccataaacggactcagatgtataaaaataactctggagacagatgaaacctccatcaagtgtctaaaaagtctgagaagtggcctcactgcagatgaactgcagagagcctgaaatacaactctacacatagatgttggcctagagacactctgtgatgttcagatctccagctcactactctgttgggaagtactctgtggagactgcgcgcagcagctagatagataccacagccagctgagatcactcagaaaactcttactcagcgcagcagaagccttactctgttcttctactcaatgaagatctatgcctgtgaaatacagcttaccctccaaaagattagcctgtfatacaaaaggggggatgacgcagacaaatcgcagcccaaaagtgtcatcagaanaagcggccttaaggttctatcagaaactgtgtttaggagaggaatcacacaaactgtgtttaggagaggaatcacataaactcaagtaaaaaagagccagtaaaaagcctgagatgatttgcgtatgatacccaaaaatgaacctctctgtctcttattatattactgaactgagacagaanaagtcccaaaaaggaatgatgtctcccagtgatgatttttctcaaaattgtctgcagcagaataacttactggggagagcgtgaaggtctggagaagcgtgaaggtctggagaagaagaaggggaactgaactgttgggaatataactcgtgta