

## Product datasheet for **SC108675**

### ZNF592 (NM\_014630) Human Untagged Clone

#### Product data:

|                           |                                                                                                   |
|---------------------------|---------------------------------------------------------------------------------------------------|
| Product Type:             | Expression Plasmids                                                                               |
| Product Name:             | ZNF592 (NM_014630) Human Untagged Clone                                                           |
| Tag:                      | Tag Free                                                                                          |
| Symbol:                   | ZNF592                                                                                            |
| Synonyms:                 | CAMOS; SCAR5                                                                                      |
| Mammalian Cell Selection: | None                                                                                              |
| Vector:                   | <u>pCMV6-XL4</u>                                                                                  |
| E. coli Selection:        | Ampicillin (100 ug/mL)                                                                            |
| Fully Sequenced ORF:      | >OriGene ORF within SC108675 sequence for NM_014630 edited (data generated by NextGen Sequencing) |

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ATGGGGGATATGAAAACCCAGATTTTGTATGACCTTCTGGCTGCCTTTGACATCCCAGAC
CCCACCAGCCTTGATGCCAAGGAGGCCATCCAGACACCCAGTGAGGAGAATGAGAGTCCC
CTCAAACCTCCAGGCATATGTATGGATGAAAGTGTGTCTTGTCTCACTCAGGATCAGCC
CCCGATGTGCCGGCCGTGAGTGTCAATTGTCAAGAACACCAGCCGCCAGGAGTCATTTGAA
GCGGAGAAAGACCACATTACTCCAGTCTCCTACACAATGGATTCCGGGGCTCAGATCTG
CCTCCAGATCCCCACAACCTGTGGGAAATTTGATTCTACTTTTATGAATGGAGACAGTGCC
AGGAGTTTCCCTGGCAAACCTGGAGCCTCCCAAGTCAGAGCCATTACCCACCTTCAACCAG
TTCAGTCCAATCTCCAGCCAGAACCTGAGGATCCCATCAAAGATAACGGATTTGGGATA
AAGCCCAAACACTCTGACAGTTATTTCCACCCCTCTTGGGTGCGGGGCTGTGGGAGGC
CCAGTCCCTGGAGGCTCTGGCTAAGTTTCCGGTTCCAGAGCTGCATATGTTTGATCATTTT
TGTAAGAAAGAACCCAAGCCAGAACCCCTGCCCTTGGGGAGCCAGCAGGAACACGAGCAA
AGTGGGCAGAACACAGTGAACCTCACAAAGGATCCGGATGCCACTCGATTCTTCGGGGAA
GCTTTGGAGTTCAACAGCCATCCTAGCAACAGTATTGGAGAGTCCAAGGGGCTTGCCCGG
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AAGAAGCAACAGAGCACAGCACTGCAGGCATCCACCCTGGCCCCCTGCCAACCTCTGCCC

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AAAGCCGTGCACTTGGCCAACCTGAACCTCGTCCCCACAGTGTTGCTGCATCAGTGACA  
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 GGCCGCTGGGGTAGGCCTGAAGCCACCCGAGGTGGAAGCCAGGCCGCGGCTGAGGAAC  
 ACTGGCTGGACCTGCCAGGAGTGCCAGGAGTGGGTTCCAGATCGGGAGAGCTACGTGTCC  
 CACATGAAAAAGAGCCACGGTCCGACATTGAAGCGGTACCCATGCCGGCAGTGTGAACAG  
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 AAGTCCCTTTTTAGTGCAGCAATGTAGTTTTGCCACAGACTCGGGGCTCGAGTTTCAG  
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 CCAGAGGAGGGCTCCGGGGAGGAGGTGCCATGGAGACTAGAGAGAATGGACTGGAAGAA  
 TGTGCCGGTGAGCCTTTGTCAGCTGACCAGAGGCGAGGAGATTGCTGGGCCCGGCCCT  
 GAGGACGATGGTGGCCACAATGATCAGTCAACCACAGGCCTCTCAGGACCAGGACAGC  
 CACACACTGTCCCCTCAGGTGTGA

Clone variation with respect to NM\_014630.2  
 2238 a=>g

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|-------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <b>5' Read Nucleotide Sequence:</b> | <p>&gt;OriGene 5' read for NM_014630 unedited</p> <pre> CGGCACGAGCCGCGCCGCCAGGTGGTGTGTTGGACTCTAGACCATGTGCCTAGGTAG AAGTTTTTCTTTCTCCGAGCTCTGCTCCCCTAGCAACGCTCGCCACACCCCTGTTTTG AGATCCTCTCTAAGGAGCGGAGAGTTTAAAGGCAAGAAGGAAGGGAGAGACAGAAGGA AGACGCTCCCCGTACGGAGACAGAGGGAGGGGGGCTCCAAAGCCGAAAGAGGAGGTCC CTACCTGCCACGGATACCACTCAGCCCTTGCCAGCATCCAGCCATGGGGGATATGAAAAC CCCAGATTTTGATGACCTTCTGGCTGCCTTTGACATCCCAGACCCACCCAGCCTTGATGC CAAGGAGGCCATCCAGACACCCAGTGAGGAGAATGAGAGTCCCCTCAAACCTCCAGGCAT ATGTATGGATGAAAGTGTGCTCTGTCTCACTCANGATCAGCCCCGATGTGCCGGCCGT GAGTGTCAATTGTCAAGAACACCGCCAGGAGTCATTTGAAGCGGAGAAAGACCACAT TACTCCAGTCTCTACACAATGGATTCCGGGGCTTCANATCTGCCTTCCAGATCCCCAC AACTGNGGGGNAATTTGATTCTACTTTTATGAATGGAANACAGTGCCAGGAGTTTTCC CTGGCAAACCTGGAGCCTNCCAAGTCAGANCCATTACCCACCTTCACCCA </pre>                                                                                                                                                                                                                                                                                               |
| <b>3' Read Nucleotide Sequence:</b> | <p>&gt;OriGene 3' read for NM_014630 unedited</p> <pre> AACTTTGGACCCGCGCCGCAATCTAGAGTCGAGTTTTTTTTTTTTTTTTTTTGGTGAGA AGATTTTTTATTTGAAATATAGTCTGAGCAACATGAGTGTAGCCAAGAGGAGTAGGGCT GGCCTGCCTGGGCGAGCAGCCTCTCAAATGGCACGAGAGCAGTACAAATATTCCAATGTC TGTGTGCCCAACCCACCTGTCAGGAAAGGCAGAACAGCAGAAACAAAGACCCACCCACA GGCCTTCTCAAGGGTCTGGTTCTTGGTGCCTGCCTTGCCTTACCTCCTGCTCT GCCCAGGTTCAAACAGAGGTGTGCAGCAGAGGTTTCTCCACCCACCCACAGAGACAT GATGAGCCAGTGTGACAATTAGTTCTCAGGACTATTTTGAAGGAAAAGAATCAGAGAAGA CAGGGAGGAGGGCAAGGTGATCAAGGGCCCTCGCCAAACCAGACTAGGCAGAGCTCTCC CAGAGGGACTGTGTCAGCACCCCTCATTTCCCTCCTTCTCAGGATCTAAAGGCAGGAAGG AGAGCTGGTGTCTGCAGGAAAATAGAGCTCTGAGGAGCGCCCTGGCTGAAGGCAGCAAGC GGGAACCTCCCAGAGAGCACAGCCGCTGTGGGGTTCAAGCCTAGTTGACAGCAGTGGGTG TGGGCCACATAAGCAGCTTTGGGTTGGAAGGACACAGACAAGAGATGGGGCACCAAGAAG CAGCGAAAAGCCATANNATACAGAAGCANATTACAGAGCTGTTGAGGGGTCTGCCCGACC CCCACATTTCTGGNGGCTGGNGGGATACCCAGCACTGGNCAGCTCANGGNCAGACACT ACTTCCAGGGTACAACCCGTCACTTTACTGAGGGGGCAGAGCCTTTAATTTTTCCCGG GTCGAGGNCAAGNTGGNAAGACCTTTTGACCAACCCTGACATGCACATGNAAAGTTCCG GTCAACTGAGGGAGCCGTTGTGGTCTCCCTGT </pre> |
| <b>Restriction Sites:</b>           | NotI-NotI                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |
| <b>ACCN:</b>                        | NM_014630                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |
| <b>Insert Size:</b>                 | 4540 bp                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      |
| <b>OTI Disclaimer:</b>              | Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |
| <b>Components:</b>                  | The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         |

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| <b>Reconstitution Method:</b> | <ol style="list-style-type: none"> <li>1. Centrifuge at 5,000xg for 5min.</li> <li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li> <li>3. Close the tube and incubate for 10 minutes at room temperature.</li> <li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li> <li>5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.</li> </ol> |
| <b>RefSeq:</b>                | <u>NM_014630.1, NP_055445.1</u>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         |
| <b>RefSeq Size:</b>           | 5086 bp                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |
| <b>RefSeq ORF:</b>            | 3804 bp                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |
| <b>Locus ID:</b>              | 9640                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |
| <b>UniProt ID:</b>            | <u>Q92610</u>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |
| <b>Cytogenetics:</b>          | 15q25.3                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |
| <b>Domains:</b>               | zf-C2H2                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |
| <b>Protein Families:</b>      | Transcription Factors                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   |
| <b>Gene Summary:</b>          | This gene is thought to play a role in a complex developmental pathway and the regulation of genes involved in cerebellar development. Mutations in this gene have been associated with autosomal recessive spinocerebellar ataxia. [provided by RefSeq, Jan 2011]                                                                                                                                                                                                                                                      |