

# Product datasheet for PH302455

### CD44 (NM\_001001389) Human Mass Spec Standard

### **Product data:**

#### **Product Type:** Mass Spec Standards CD44 MS Standard C13 and N15-labeled recombinant protein (NP\_001001389) **Description:** Species: Human **HEK293 Expression Host: Expression cDNA Clone** RC202455 or AA Sequence: Predicted MW: 76.6 kDa >RC202455 protein sequence **Protein Sequence:** Red=Cloning site Green=Tags(s) MDKFWWHAAWGLCLVPLSLAQIDLNITCRFAGVFHVEKNGRYSISRTEAADLCKAFNSTLPTMAQMEKAL SIGFETCRYGFIEGHVVIPRIHPNSICAANNTGVYILTSNTSQYDTYCFNASAPPEEDCTSVTDLPNAFD GPITITIVNRDGTRYVQKGEYRTNPEDIYPSNPTDDDVSSGSSSERSSTSGGYIFYTFSTVHPIPDEDSP WITDSTDRIPATSTSSNTISAGWEPNEENEDERDRHLSFSGSGIDDDEDFISSTISTTPRAFDHTKQNQD WTOWNPSHSNPEVLLOTTTRMTDVDRNGTTAYEGNWNPEAHPPLIHHEHHEEEETPHSTSTIQATPSSTT EETATQKEQWFGNRWHEGYRQTPREDSHSTTGTAAASAHTSHPMQGRTTPSPEDSSWTDFFNPISHPMGR GHQAGRRMDMDSSHSTTLQPTANPNTGLVEDLDRTGPLSMTTQQSNSQSFSTSHEGLEEDKDHPTTSTLT SSNRNDVTGGRRDPNHSEGSTTLLEGYTSHYPHTKESRTFIPVTSAKTGSFGVTAVTVGDSNSNVNRSLS GDQDTFHPSGGSHTTHGSESDGHSHGSQEGGANTTSGPIRTPQIPEWLIILASLLALALILAVCIAVNSR RRCGQKKKLVINSGNGAVEDRKPSGLNGEASKSQEMVHLVNKESSETPDQFMTADETRNLQNVDMKIGV TRTRPLEQKLISEEDLAANDILDYKDDDDKV C-Myc/DDK Tag: **Purity:** > 80% as determined by SDS-PAGE and Coomassie blue staining **Concentration:** >0.05 µg/µL as determined by microplate BCA method Labeling Method: Labeled with [U-13C6, 15N4]-L-Arginine and [U-13C6, 15N2]-L-Lysine **Buffer:** 25 mM Tris-HCl, 100 mM glycine, pH 7.3 Storage: Store at -80°C. Avoid repeated freeze-thaw cycles. Stability: Stable for 3 months from receipt of products under proper storage and handling conditions. NP 001001389 **RefSeq: RefSeq Size:** 5619



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### OriGene Technologies, Inc.

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|                  | CD44 (NM_001001389) Human Mass Spec Standard – PH302455                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |  |
|------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--|
| RefSeq ORF:      | 2097                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |  |
| Synonyms:        | CDW44; CSPG8; ECMR-III; HCELL; HUTCH-I; IN; LHR; MC56; MDU2; MDU3; MIC4; Pgp1                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     |  |
| Locus ID:        | 960                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |  |
| UniProt ID:      | <u>P16070</u>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     |  |
| Cytogenetics:    | 11p13                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             |  |
| Summary:         | The protein encoded by this gene is a cell-surface glycoprotein involved in cell-cell<br>interactions, cell adhesion and migration. It is a receptor for hyaluronic acid (HA) and can also<br>interact with other ligands, such as osteopontin, collagens, and matrix metalloproteinases<br>(MMPs). This protein participates in a wide variety of cellular functions including lymphocyte<br>activation, recirculation and homing, hematopoiesis, and tumor metastasis. Transcripts for<br>this gene undergo complex alternative splicing that results in many functionally distinct<br>isoforms, however, the full length nature of some of these variants has not been determined.<br>Alternative splicing is the basis for the structural and functional diversity of this protein, and<br>may be related to tumor metastasis. [provided by RefSeq, Jul 2008] |  |
| Protein Families | : Adult stem cells, Cancer stem cells, Druggable Genome, Embryonic stem cells, ES Cell Differentiation/IPS, Stem cell relevant signaling - DSL/Notch pathway, Transmembrane                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       |  |
| Protein Pathway  | vs: ECM-receptor interaction, Hematopoietic cell lineage                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |  |

## **Product images:**

| 116 - | -  | - |
|-------|----|---|
| 66 -  | -  |   |
| 45 -  | -  |   |
| 35 -  | -1 |   |
| 25 -  | -1 |   |
| 18 -  | -1 |   |
| 14 -  | -  |   |

Coomassie blue staining of purified CD44 protein (Cat# [TP302455]). The protein was produced from HEK293T cells transfected with CD44 cDNA clone (Cat# [RC202455]) using MegaTran 2.0 (Cat# [TT210002]).

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