| Revisi           | on 4   |                              |  |
|------------------|--|------------------------------|--|
| Store at<br>-20C | 5-Carboxylcytosine (5-caC) (D7S8U) Rabbit<br>mAb | Ce<br>T E                    | <b>ll Signaling</b><br>сн N о L о G Y*       |
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| Applications: F<br>IF-IC, Dot Blot | Reactivity:<br>All | Sensitivity:<br>Transfected Only  | Source/Isotype:<br>Rabbit IgG                        |                                    |
|------------------------------------|--------------------|---|--|------------------------------------|
| Product Usage<br>Information       |                    | <b>Application</b><br>Immunofluorescence (Imn<br>DNA Dot Blot   | nunocytochemistry)                                   | <b>Dilution</b><br>1:200<br>1:1000 |
| Storage                            |                    | Supplied in 10 mM sodium HEPES (pH 7.5), 150 mM NaCl, 100 μg/ml BSA, 50% glycerol and less than<br>0.02% sodium azide. Store at –20°C. Do not aliquot the antibody.   |  |                                    |
| Specificity/Sensitivi              | ty                 | 5-Carboxylcytosine (5-caC) (D7S8U) Rabbit mAb detects 5-caC by IF in cells over-expressing the TET1 catalytic domain and by dot blot using double-stranded PCR fragments containing 5-caC. Many cells and tissues contain very low endogenous levels of 5-caC that may fall below the detection limits of this antibody. This antibody has been validated using ELISA, dot blot, and synthetic spike-in DNA MeDIP assays and shows high specificity for 5-caC.  |  |                                    |
| Source / Purification              | ı                  | Monoclonal antibody is produced by immunizing animals with 5-carboxylcytidine.  |  |                                    |
| Background                         |                    | Methylation of DNA at cytosine residues is a heritable, epigenetic modification that is critical for proper regulation of gene expression, genomic imprinting, and mammalian development (1,2). 5-<br>methylcytosine is a repressive epigenetic mark established <i>de novo</i> by two enzymes, DNMT3a and DNMT3b, and is maintained by DNMT1 (3, 4). 5-methylcytosine was originally thought to be passively depleted during DNA replication. However, subsequent studies have shown that Ten-Eleven Translocation (TET) proteins TET1, TET2, and TET3 can catalyze the oxidation of methylated cytosine to 5-hydroxymethylcytosine (5-hmC) (5). Additionally, TET proteins can further oxidize 5-hmC to form 5-formylcytosine (5-fC) and 5-carboxylcytosine (5-caC), both of which are excised by thymine-DNA glycosylase (TDG), effectively linking cytosine oxidation to the base excision repair pathway and supporting active cytosine demethylation (6,7). TET protein-mediated cytosine hydroxymethylcation has been discovered in many tissues, with the highest levels found in the brain (9). While 5-fC and 5-caC appear to be short-lived intermediate species, there is mounting evidence showing that 5-hmC is a distinct epigenetic mark with various unique functions (10,11). The modified base itself is stable <i>in vivo</i> and interacts with various readers including MeCP2 (11,12). The global level of 5-hmC increases during brain development, and 5-hmC is enriched at promoter regions and poised enhancers. Furthermore, there is an inverse correlation between levels of 5-hmC and Distone H3K27 trimethylation, suggesting a role for 5-hmC in gene activation (12). Lower amounts of 5-hmC have been reported in various cancers including myeloid leukemia and melanoma (13,14). |  |                                    |
| Background Referer                 | nces               | <ol> <li>Hermann, A. et al. (2004) <i>Cell Mol Life Sci</i> 61, 2571-87.</li> <li>Turek-Plewa, J. and Jagodziński, P.P. (2005) <i>Cell Mol Biol Lett</i> 10, 631-47.</li> <li>Okano, M. et al. (1999) <i>Cell</i> 99, 247-57.</li> <li>Li, E. et al. (1992) <i>Cell</i> 69, 915-26.</li> <li>Tahiliani, M. et al. (2009) <i>Science</i> 324, 930-5.</li> <li>He, Y.F. et al. (2011) <i>Science</i> 333, 1303-7.</li> <li>Ito, S. et al. (2011) <i>Science</i> 333, 1303-3.</li> <li>Kriaucionis, S. and Heintz, N. (2009) <i>Science</i> 24, 929-30.</li> <li>Globisch, D. et al. (2010) <i>PLoS One</i> 5, e15367.</li> <li>Gao, Y. et al. (2012) <i>Cell Stem Cell</i> 12, 453-69.</li> <li>Mellén, M. et al. (2012) <i>Cell</i> 151, 1417-30.</li> <li>Wen, L. et al. (2014) <i>Genome Biol</i> 15, R49.</li> <li>Delhommeau, F. et al. (2009) <i>N Engl J Med</i> 360, 2289-301.</li> <li>Lian, C.G. et al. (2012) <i>Cell</i> 150, 1135-46.</li> </ol>   |  |                                    |
| Species Reactivity                 |                    | Species reactivity is determ  | nined by testing in at least one approved applica    | tion (e.g., western blot).         |
| Applications Key                   |                    | IF-IC: Immunofluorescence   | e (Immunocytochemistry) <b>Dot Blot:</b> DNA Dot Blo | ot                                 |

| Cross-Reactivity Key          | All: All Species Expected  |  |  |
|-------------------------------|--|--|--|
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