

## Development of a Single-Molecule Identification Method for Hydroxymethyl Cytosine in Tumor Marker Molecules

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Cancer is the most common cause of death in Japan, and it is said that one in two people will develop cancer at some point in their lifetime <sup>1)</sup>. Early diagnosis can reduce the burden of treatment and lower the mortality rate <sup>2)</sup>. Therefore, there is a demand for fast and accurate diagnostic methods. Hydroxymethyl cytidine (5-hmC: See Figure) is gaining attention as an epigenetic marker for cancer. 5-hmC can be detected before symptoms appear, enabling early detection. Conventional detection methods include mass spectrometry and conversion of modified nucleic acids (5-hmC) to different nucleobases followed by sequencing.

In this study, we focused on hydroxymethyl cytidine (5-hmC), an oxidized modification of cytosine in DNA and RNA, and employed single-molecule quantum measurement for the detection of this modified molecule (5-hmC). This method has the advantage of directly detecting and identifying molecules in an aqueous solution on a single-molecule basis. So far, we have successfully identified single molecules of nucleobases and amino

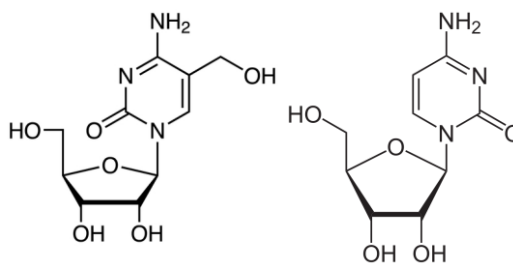


Figure:(Left) Hydroxymethyl Cytidine (5-hmC), (Right): Cytidine

acids using highly sensitive and rapid electrical measurement technology with a nano-gap electrode device <sup>3)-6)</sup>. For the measurement solution, we adjusted the concentration to 1μM using 5-hmC and cytidine (C), a control molecule. We conducted rapid current measurements with the nano-gap electrode. As a result, we successfully obtained current signals for each molecule. From these signals, we extracted characteristic features and attempted to identify single molecules through machine learning. As a result, we successfully created a classifier with an F-measure value of over 0.8 for distinguishing between 5-hmC and C. This technology is expected to be applied to early diagnosis and prognostic evaluation of cancer.

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