# 5-Hydroxymethylcytosine Is a Stable DNA Modification

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### HIGHLIGHTS

5-Hydroxymethylcytosine (hmC) is an oxidation product of 5-methylcytosine (mC) present in DNA of most mammalian cells. Reduction of hmC levels is a hallmark of cancers. Elucidating the dynamics of this oxidation reaction and the lifetime of hmC in DNA is fundamental to understanding hmC function.

#### We have developed:

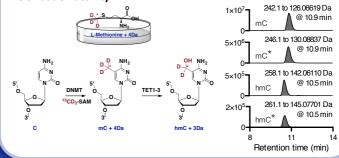
- a sensitive LCMS method for quantification of DNA modifications
- a labelling strategy to track the fate of modified cytosines

#### We have found that:

- mC oxidation does not occur on nascent DNA during replication
- hmC is stable in DNA, and not an intermediate of demethylation
- the rate of proliferation governs the global levels of hmC

# ISOTOPIC LABELLING OF DNA AND LCMS

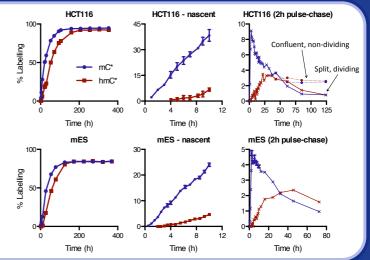
Cells can be grown in custom media containing **L-methionine**( $^{13}$ C,D<sub>3</sub>), a source of methyl group for mC in DNA. Accurate LCMS analysis then provides ratios of labelled to total (labelled + unlabelled) modification (% labelling), and therefore **information about DNA methylation and mC oxidation activity.** 



# TIMING OF mC OXIDATION ON NASCENT DNA

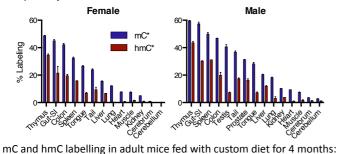
Plots on the right show mC and hmC labelling in 2 different cell lines:

- mC labelling is always higher or equal to hmC, consistent with hmC being a product of mC (left plots)
- A gap between labelling curves reflects a **difference in timing between mC formation and its oxidation to hmC** on a newly synthesized DNA strand (left and middle plots)
- Focus on early time points confirms that the first mC oxidation activity happens several hours after replication (middle plots)
- Pulse-chase experiments with a 2h labelling pulse allow tagging a small proportion of mC and hmC and follow their lifetime in DNA
- Labelled hmC is present >125h after labeling pulse, hmC is therefore a stable modification in cultured cells



# hmC IS STABLE IN DNA In vivo

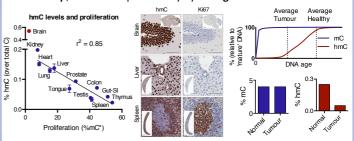
Custom mouse diet containing **L-methionine**( $^{13}$ C,D<sub>3</sub>) is a useful tool to study the **dynamics of DNA modifications** *in vivo*.



- Gap between mC and hmC labelling >> delayed hmC 'maintenance'
- Lack of labelling in slow-/non-dividing tissues >> hmC must be stable

#### GLOBAL hmC LEVELS AND PROLIFERATION

Plotting global levels of hmC against proliferation rate shows a **linear relationship**, confirmed qualitatively by **IHC against hmC and Ki67**.



hmC stability and mC oxidation timing are responsible for:

- Tissue-specific levels of hmC
- Reduced levels of hmC reported in all studied cancers