



London Proteomics Discussion Group presents:
Proteomics: The role of post-translational modification

An LPDG proteomics webinar

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**Data-independent acquisition method
for ubiquitinome analysis reveals
regulation of circadian biology**

**Mapping sites of carboxymethyllysine
modification on proteins reveals
its consequences for proteostasis
and cell proliferation**

Chaired by Prof. Ed Tate

5th February 2021, 14:00 GMT

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