Improved Single-molecule Imaging Based On Photon Counting With An EMCCD Camera

Abbas Padeganeh¹, Etienne Lareau², Olivier Daigle², Anne-Marie Ladouceur¹, Paul Maddox^{1,3}. ¹Institute for Research in Immunology and Cancer (IRIC), University of Montreal, Montreal, QC, Canada, ²Nüvü Camēras Inc., Montreal, QC, Canada, ³Department of Pathology and Cell Biology, Faculty of Medicine, Université de Montréal, Montreal, QC, Canada.

Introduction

New breakthroughs in digital imaging have historically opened opportunities to scientists in bio-medical research. One important landmark of the last decade has been the development of the electron-multiplying charge coupled device (EMCCD), an imaging detector that uses an avalanche phenomenon to amplify the weak photo-electron signals. EMCCDs are well suited for ultralow light applications in that a single photo-electron can be amplified sufficiently to allow the possibility of photon counting. However, as with all electronic detectors, the noise increases greatly with an increase in gain, diminishing the overall sensitivity and dynamic range and thus compromising images.

Recently, a new camera has been developed by Nüvü Cameras that significantly reduces the noise generated during the read-out process (up to 10 times) and allows more efficient photon counting. This major reduction of the noise threshold represents an opportunity for various low-light imaging applications including single-molecule studies.

Methods







and a high EM gain are necessary.





It is used as a reference to compare the blue curves.

tions.

We present a demonstration of its applicability in a total internal reflection fluorescence microscopy (TIRFM) assay to visualize single biomolecules. In molecular biology, it is well established that most proteins exert their biological roles in complex with other proteins. Thus, detecting individual components and the structure of these complexes is essential to understand the mechanism of biological events. To this aim, we employ TIRFM imaged in photon counting mode for visualization of single centromeric nucleosomes, containing a histone variant known as CENP-A.

We show that this technique is well suited to TIRFM and that it is capable of detecting single fluorescently labelled CENP-A in the nucleosome with an enhanced signal to noise ratio. The desired and improved properties of the assay such as high sensitivity and photon counting shall pave the way for comprehensive studies of single molecules in this and other biological contexts.

Proposed models for nucleosomal composition at centromeres



H2A







GFP



Raw data before correction

Manual automated analysis confirmed the presence of a mixed population of one and two CENP-A containing nucleosomes in Hela cells. Cell cycle analysis also revealed a dynamic change in the proportion of each population.

Conclusions

- Our TIRFM assay is capable of detecting single fluorescently-labeled biomolecules

- Using an EMCCD camera in photon-counting mode enables visualization of single molecules and photobleaching behaviour at a much lower laser power compared with analog mode and thus may provide a better time resolution

- There is a mixture of two different CENP-A nucleosomes in the mammalian system

- Cell-cycle dependent changes appear to occur in the structure of CENP-A nucleosomes

Acknowledgements

We would like to thank all members of the Maddox labs for their support and useful contributions to this work. Special thanks to Joel Ryan for his assistance. This work was supported by NSERC, CIHR, the Canadian Cancer Research Society and Nüvü Cameras.





INSTITUTE FOR RESEARCH IN IMMUNOLOGY AND CANCER



