# Scienze della Vita



The informational content of cellular imaging in studies of dynamic biological processes

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of fixed samples and living cells.

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	INTRODUCTION	
High content imaging of dynamic biological processes	THE IBPM-CNR MICROSCOPY PLATFORM - NIKON REFERENCE CENTER	
Image-based assays in the biomedical field are rapidly evolving from qualitative to more	N The elettrice heater with a sector of a bight marketing interview.	

quantilative approaches ( high content ) that provide high resolution spatio-temporal information

Time-lapse video recording of living cells depicts dynamic information that would remain otherwise unnoticed with analyses of fixed samples and that are fundamental to understand complex biological processes (e.g., cell division, cell differentiation, senescence, intra- and intercellular signalling, cellular migration, infection, host/pathogen interactions, response to drug treatments, induction of cell death).



(modified from Neumann et al., 2010)



http://bbcd.bio.uniroma1.it/bbcd/archivionotizie/cnr-microscopy-platform-nikon-reference-center-ibpm



### **CELLULAR IMAGING: FIELDS OF APPLICATION**

The IBPM microscopy platform supports versatile applications, in which dynamic studies (time-lapse recording) are coupled with high resolution qualitative and quantitative image analysis of cells and cellular structures

#### **Applications:**

#### Real-time visualization of dynamic processes (cellular signalling, intracellular transport, organization of organelles and subcellular structures)

Single-cell analysis, to visualise cell heterogeneity and rare behaviours within a cell population

Recording of cellular morphological changes or cell death in response to particular stimuli (physical / chemical damage

### **Biological processes**

40 50

30

Studies of cell division and checkpoints

Tumour cell growth and inhibitory drugs / molecules /modulating genes

Differentiation of stem and progenitor cells

Assay / design of innovative therapeutic

## **CREATION OF CELL MODELS AND SET-UP OF AD-HOC WORKFLOWS FOR HIGH** THROUGHPUT (HT) AND HIGH CONTENT (HC) AUTOMATED IMAGE ANALYSIS

#### Creation and validation of informative cell models useful for automated analysis

Cell lines can be engineered to visualize cellular components using fluorescent proteins or fluorescent dyes suitable for live imaging

Set-up of specific protocols for live cell imaging (will take into account: phototoxicity assessment, definition of exposure time and intervals; concentration of staining dye etc.)

#### Automation of image acquisition and data analysis

- 1. Creation of *ad-hoc* acquisition workflow to maximize the amount and the quality of imaging data (JOBS module of Nikon proprietary Nis Elements software)
- 2. Classification of phenotypes of interest using the machine learning-based Nis Elements classifier
  - 3. Automated analysis of phenotypes of interest



Measurements of cell migration

High definition analysis of subcellular structures (5 fluorescence excitation channels, simultaneous visualization of 4 stainings, image deconvolution, 3D reconstruction)

Proximity ligation assays for *in situ* protein interactions and *in* situ post-translational modifications.

strategies Cell response to parasites, bacteria and viral infectious agents

Novel biocompatible matrices to support cell growth in tissue regeneration

The uptake of nanoparticles and functionalised nanomaterials within cells



### **EXEMPLIFYING RESULTS**

**1. Single-cell recording depicts stochastic phenomena in cell** biology: heterogeneous responses to novel mitotic inhibitors, including rare yet biologically significant behaviours



**TAX** (n=101) **ATI 33** (n=193)



is modulated by different genetic backgrounds (precision medicine) MycN -

**2.** Time-lapse recording shows that the outcome of therapeutic treatments





interphase catastrophe

Time lapse video-recording of neuroblastoma cells: shows that treatment with PARP inhibitors induces differential fates, depending of the status of MYCN amplification (a driver of aggressiveness in neuroblastoma); the fraction of cells that undergo a specific cell fate can be quantified.

(Colicchia et al., 2017)

Amplified

MycN



### 3. Automated detection and analysis of protein-protein interactions in fixed cells



probes to reveal interaction



We developed a workflow for the automated detection of PLA signals. This yields rapid and accurate information on i) genuine validation, and ii) subcellular localization of Importin beta interactors selected in proteomewide screening.







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### SHARE SCIENCE WORKSHOP SAPIENZA 2019