

## gH2AX/pH3

**Commonly used acronym:** gH2AX/pH3

Created on: 26-06-2022 - Last modified on: 11-07-2022

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## SCOPE OF THE METHOD

<b>The Method relates to</b>	Animal health, Human health
<b>The Method is situated in</b>	Basic Research, Regulatory use - Routine production
<b>Type of method</b>	In vitro - Ex vivo
<b>Specify the type of cells/tissues/organs</b>	Any cell line model or primary cells

## DESCRIPTION

### Method keywords

Histone-PTMs

genotoxicity

mode of action

H2AX

H3

### Scientific area keywords

genotoxicity

aneugen

clastogen

in vitro toxicology

carcinogenicity

Chemical testing

### Method description

A battery of *in vitro* genotoxicity assays is currently used to detect agents with DNA damaging and carcinogenic potential. Although the sensitivity of this battery of genotoxicity assays is high, the specificity is low ("false positive hit"), especially with the

mammalian cell-based assay, compared with *in vivo* data. Understanding the mode of genotoxic action (MoA: aneugen or clastogen) of a genotoxic chemical is another important requirement. The proposed gH2AX/pH3 method permits the measurement of early biomarkers not only able to predict DNA damage, but also differentiate between numerical and structural chromosome damage. This assay is applicable to almost all cell types including cell of human origin. The assay uses innovative techniques with a higher throughput than usual assays and potentially amenable to automation. The method is less prone to irrelevant positive results as compared to some of the current assays because it is able to also discriminate the misleading cytotoxic chemicals.

### **Lab equipment**

Classical cellular biology equipment.

To analyze the two endpoints (gH2Ax and pH3) different methods are suitable (In Cell Western, FACS, HCA...).

### **Method status**

History of use

Internally validated

Published in peer reviewed journal

Currently submitted for further validation by an external party (e.g. OECD, EURL ECVAM,...)

## **PROS, CONS & FUTURE POTENTIAL**

### **Advantages**

- a) The gH2AX/pH3 method has been reported to be more predictive of genotoxicity (for specificity and sensitivity) than the commonly used assays (MNvit and Ames) and could detect efficiently different types of genotoxic compounds at low concentrations.
- b) This assay is faster than MNvit test because it did not require two cell cycles completion.
- c) This assay is the first one to permit to easily discriminate the genotoxic mode of action very powerfully (clastogens, aneugens and misleading cytotoxic chemicals) in any human cell type. The use of cell lines with different metabolism capacities enabled differentiation between directly from bioactivated genotoxins.
- d) The use of multi well plates allows high-throughput format for both cell treatment and parameter measurement.
- e) The biomarkers and the antibodies used for the activity measurement are not protected by any patent or license and the technologies are relatively inexpensive to perform.

### **Challenges**

The gH2AX/pH3 genotoxic method is highly suitable for the detection of genotoxic properties of substances amenable to solution. Chemicals are routinely dissolved in aqueous solvent such as water or organic solvents such as DMSO. For hydrophobic substances like oil-derived chemicals, DMSO extracts can be tested. As with conventional *in vitro* genotoxicity assays, the gH2AX/pH3 biomarkers do not easily allow the evaluation of gasses. It is hoped that new advances in this area will enable such investigations. The method is able to detect genotoxins that require metabolic activation, providing that metabolically competent cells are used or external metabolic activation system is added (e.g., S9 fraction). Since the gH2AX signal is expected to in part result from DNA replication or transcription blocking lesions induced by genotoxins, cells in a proliferative

state may be more appropriate than cells in a confluence state.

## **Modifications**

The use of high content analysis platform permits to detect in parallel different biomarkers. So it is possible to detect other biomarker of interest than only phosphorylated histone H2AX and H3.

## **Future & Other applications**

The gH2AX/pH3 test method is not aimed to replace or delete an existing method. The gH2AX/pH3 method has the unique possibility to provide mechanistic insights into the mechanism of genotoxicity by combining two relevant endpoints in a single assay. gH2AX/pH3 method can be particularly useful in an Adverse Outcome Pathway (AOP) and in the development of an Integrated Approach to Testing and Assessment (IATA). The mechanistic information that is provided by the gH2AX/pH3 method assay can be applied to translate the Molecular Initiating Events (MIE) and cellular responses that are activated upon chemical exposure to carcinogenicity hazards for humans. Discussion will begin this year at OECD for a official test guideline.

## **REFERENCES, ASSOCIATED DOCUMENTS AND OTHER INFORMATION**

### **References**

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## Associated documents

[Kopp\\_ArchTox2019review.pdf](#)

## Other remarks

Since 2008 and the first experiment conducted in our lab, for each cell line, in each experiment, the same benchmark positive control is included. The assay has been done by ten different experimenters with consistently high reproducibility (superior to 95%), independent of the cell line used. The gH2AX/pH3 method was transferred successfully to five academic laboratories (RIKILT, Netherlands; Laboratory of Toxicological Control of Pesticides, Greece; Institut für Toxikologie, Deutschland; IPBS-CNRS UMR5089, France; INSERM U1220-INRA U1416, France). An inter-laboratory trial of the gH2AX/pH3 method was performed with seven different private companies (Litron, Pfizer, Servier, Orion, Sanofi-Aventis, Bayer and Roche Pharma) using 84 chemicals.

The validation has been performed largely according to OECD Guidance document 34 on the validation and international acceptance of new test methods for hazard assessment. An overall concordance between companies of 92% was achieved with a sensitivity of 92% and a specificity of 96%. (see first paper in the References section)

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