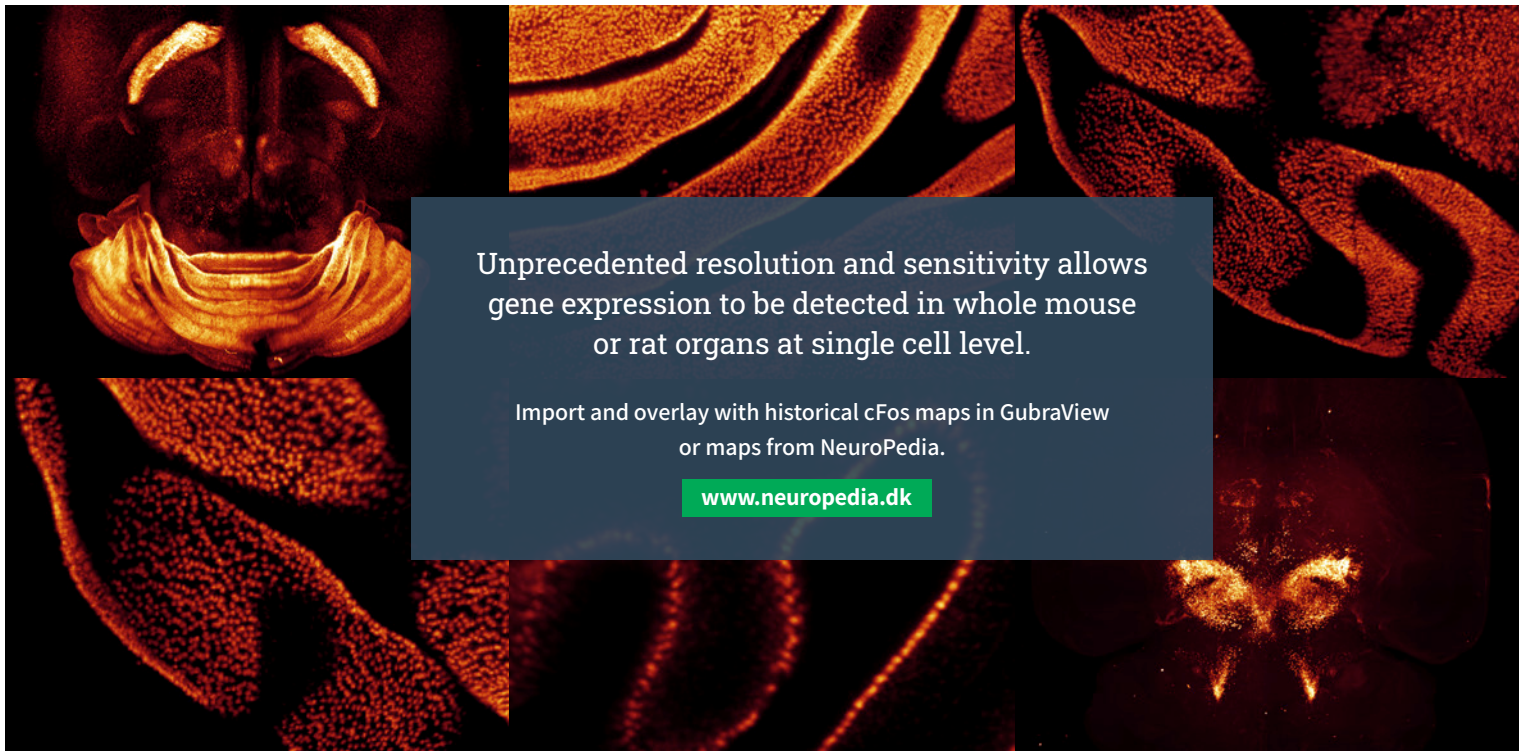# Now it is possible to detect mRNA in intact organs with incredible sensitivity
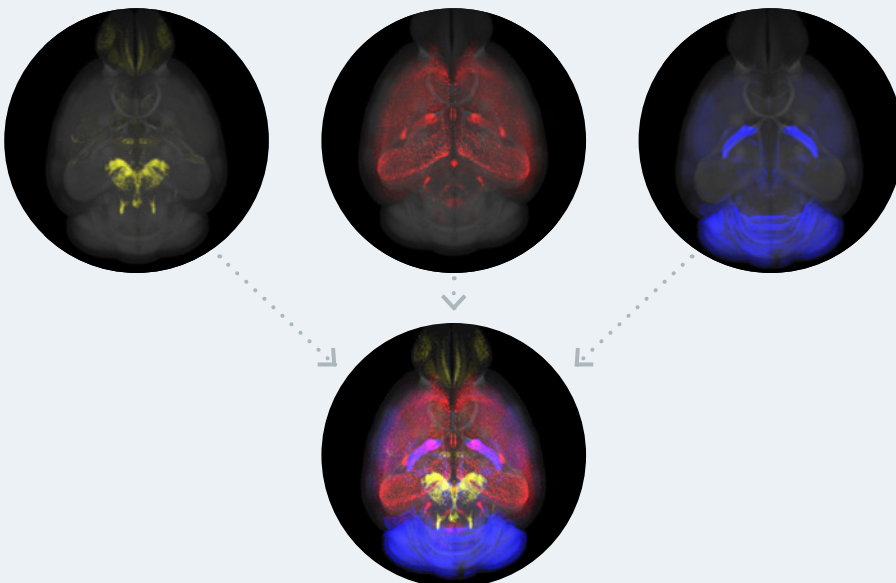

Unprecedented resolution and sensitivity allows gene expression to be detected in whole mouse or rat organs at single cell level.

Import and overlay with historical cFos maps in GubraView or maps from NeuroPedia.

www.neuropedia.dk

Map your ISH data to our reference brain for quantification and anatomical annotation.

Import and overlay with historical cFos maps in GubraView or maps from NeuroPedia.

Preclinical applications:

+ Mapping of receptors and other druggable targets.

+ Detection of viral and non viral payloads in gene therapy.

+ Multiplex up to three genes for quantification of co-expression.

+ Applicable to all species and all genes (if detection limit permits).

+ Immediate early genes for drug responses.

Contact Jacob Hecksher-Sørensen
jhs@gubra.dk