#### INFRAIA H2020



integrated

**N**MR

#### The Call

 Research Infrastructures for translating research on biological structures into innovation in biomedicine. This activity should expand the availability of structural biology services (such as X-ray and neutron scattering, advanced NMR and advanced imaging technologies) to new communities of users, and in particular to scientists with backgrounds other than structural biology, including from SMEs, to benefit translational research in drugs discovery, informed drugs and vaccine design and other fields like biotechnology and biomaterials. Synergies with relevant ESFRI Infrastructures, such as INSTRUCT, EUROBIOIMAGING, EU-OPENSCREEN, and EATRIS, should be duly exploited.

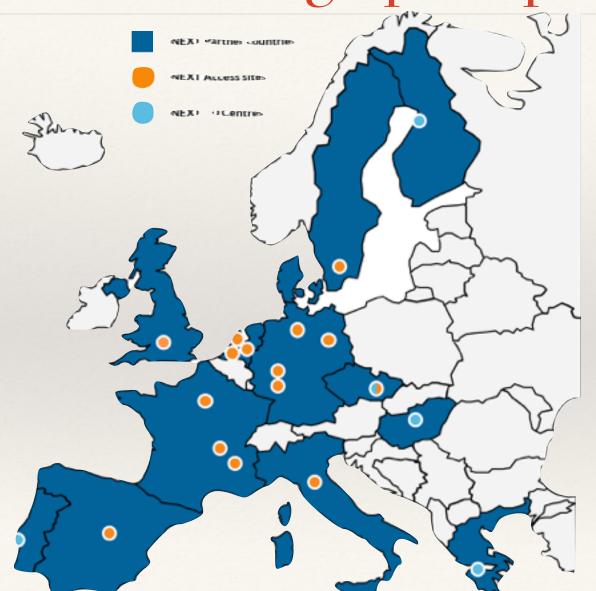


### The Partnership

- \* 23 Partners from 14 Countries
- NMR centres
  - Berlin, Brno, Florence, Frankfurt, Lyon, Utrecht (coordinator)
- Six synchrotron sites
  - Diamond, EMBL-HH, EMBL-GR, ESRF, MAX IV, Soleil
- Five EM facilities
  - \* Brno, Diamond, EMBL-HD, Leiden, Madrid
- Protein interactions in vivo and in vitro
  - Amsterdam (deputy coordinator), EMBL-HD
- Training centres
  - Brno, Budapest, Lisboa, Oulou, Patras, Rehovot
- Research Partners
  - Aarhus, Lund (ESS)



Geographic spread





### Aims

- \* User access to macromolecular X-ray crystallography, solution scattering, NMR, EM and advanced light microscopy.
- Show the possibilities offered by the ESS
- Linking in vitro studies to in-cell interaction studies
- \* Reach out to industry by collaborative research, networking, trainings and showcasing of new opportunities.



### Three Themes of Access

- 1. Structural Audit
- 2. Enhanced Support
- 3. High-End Data Collection



### Structural Audit

- \* A sample of a macromolecule (or complex) is evaluated for
  - \* Purity
  - Stability and aggregation in different buffer formulations
  - Multimeric state by MALLS
  - \* SAXS data collection for low resolution shape determination
  - Crystallisation
  - NMR HSQC (if labeled)
  - \* Utility for EM (if >200kD)
- Results are combined in a Structural Audit Report



# Enhanced Support

- \* Guided access involves training and active participation of guests
- Structure determination by X-ray crystallography, EM, SAXS or NMR
- Ligand and fragment screening
- \* Study of in vitro or in-cell macromolecular interactions



# High-END: "Traditional" Access

- Synchrotron Data Collection
  - Applications for single translation research theme
  - \* One or more group leaders under a common team
  - No Biostruct-type BAGS, but research-theme based
- High-field NMR data collection
  - HSQC characterisation required
- Advanced cryo-EM data collection
  - Only if initial data are available to establish grid quality



# **Enhanced Support Modalities**

- \* X-ray structural analysis (EMBL-GR/ESRF, EMBL-HH)
- NMR Structure analysis (UU, Florence, Frankfurt)
- \* EM sample optimisation (EMBL-HD)
- Ligands and Fragment Screening (EMBL-GR, Diamond)
- Macromolecular Interactions Platform (NKI)
- Advanced Light Imaging (EMBL-HD)



# Training

- \* Integrating X-ray techniques and EM
- Combining X-Rays and Neutrons for Crystallography and Scattering
- Combining solution methods: NMR SAXS Biophysics
- Structural biology for drug development
- The toolkit to study macromolecular interactions
- Integrated methodologies and approaches for Structural Biology
- Training centra in Brno, Budapest, Lisboa, Oulou, Patras, Rehovot



# Joint Research Applications

- 1.Structure-based small molecule inhibitor discovery (ligand screening)
- 2. Integrated study of membrane proteins.
- 3. Study of macromolecules and their complexes in cells.

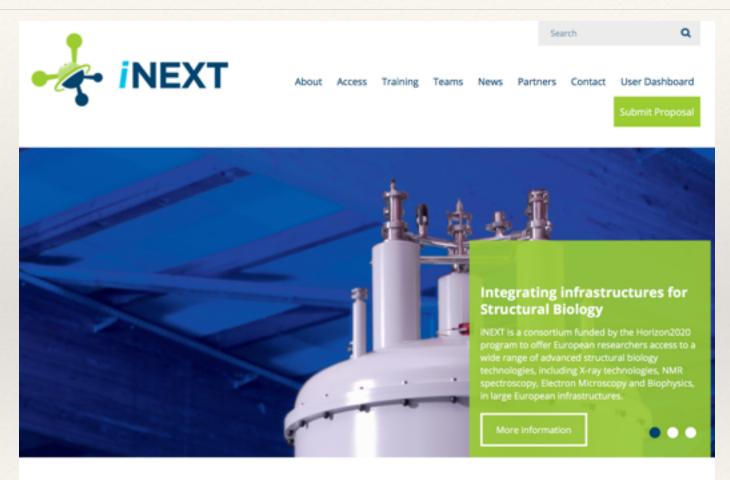


# Networking

- \* Involving and training new scientific communities and the general public.
- Gateways to industry.
- Strengthening the iNEXT community.
- Data Management and links with e-Infrastructures.



# Website: www.inext-eu.org



iNEXT: Infrastructure for NMR, EM and X-rays for Translational Research

