Comments from the SAXS Community

M.V. Petoukhov

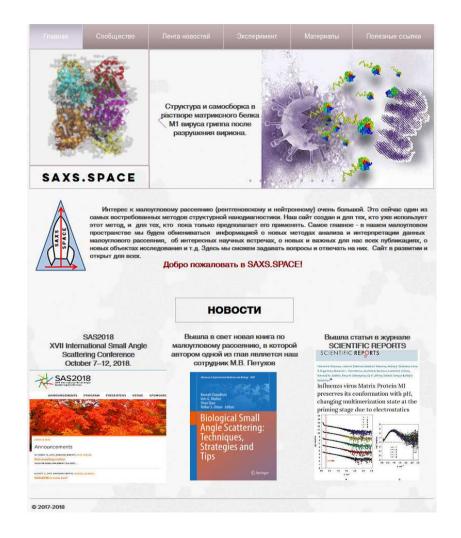
SAXS: Current Objects

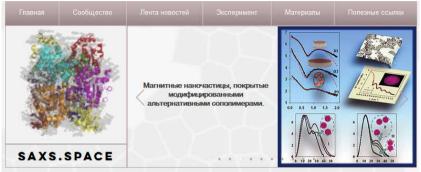
- Coat proteins of plant viruses capsid formation
- Matrix protein M1 interaction with lipid and core
- Study of in cellulo crystallization (in intact living cells)
- Self-assembly processes of biomedically relevant supramers (glycolipids, glycopeptides, oligoglycines)
- Study of transmembrane proteins (APP, ErbB) and their interaction with liposomes
- Structural characterization of dendrimeric macromolecules
- "Conventional" proteins and macromolecular complexes

SAXS: Wishlist for Infrastructure

- Large objects (~10² nm) adequate s-range, small beamstop size
- Study of processes possibility of time-resolved measurements (at least sub-second resolution)
- Phase separation in heterogenious systems inline size-exclusion chromatography setup
- Anomalous SAXS (study of metalloproteins, nanocomposites etc) – tunable wavelength
- Temperature-controlled sample environment
- Robotic sample changer
- State-of-the-art automation, software for data analysis, interpretation and model building

SAXS.SPACE





2017

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