

Francesco Stellato

List of Publications by Year in descending order

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papers

4,294
citations

304368

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264894

42
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docs citations

44
times ranked

4059
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | ARIA – A VUV Beamline for EuPRAXIA@SPARC_LAB. Condensed Matter, 2022, 7, 11. | 0.8 | 5 |
| 2 | Is styrene competitive for dopamine receptor binding?. Biomolecular Concepts, 2022, 13, 200-206. | 1.0 | 0 |
| 3 | Plasma-Generated X-ray Pulses: Betatron Radiation Opportunities at EuPRAXIA@SPARC_LAB. Condensed Matter, 2022, 7, 23. | 0.8 | 5 |
| 4 | Modelling Protein Plasticity: The Example of Frataxin and Its Variants. Molecules, 2022, 27, 1955. | 1.7 | 2 |
| 5 | Cu(II) – Glycerol – Ethylmorpholine Complex Stability Revealed by X-ray Spectroscopy. Journal of Physical Chemistry C, 2021, 125, 1483-1492. | 1.5 | 3 |
| 6 | Zn-Induced Interactions Between SARS-CoV-2 orf7a and BST2/Tetherin. ChemistryOpen, 2021, 10, 1133-1141. | 1.9 | 11 |
| 7 | SARS-CoV-2 Virion Stabilization by Zn Binding. Frontiers in Molecular Biosciences, 2020, 7, 222. | 1.6 | 14 |
| 8 | Dealing with Cu reduction in X-ray absorption spectroscopy experiments. Metallomics, 2019, 11, 1401-1410. | 1.0 | 11 |
| 9 | The Potential of EuPRAXIA@SPARC_LAB for Radiation Based Techniques. Condensed Matter, 2019, 4, 30. | 0.8 | 12 |
| 10 | X-Ray Absorption Spectroscopy Measurements of Cu-ProlAPP Complexes at Physiological Concentrations. Condensed Matter, 2019, 4, 13. | 0.8 | 6 |
| 11 | Design study of a photon beamline for a soft X-ray FEL driven by high gradient acceleration at EuPRAXIA@SPARC_LAB. Nuclear Instruments and Methods in Physics Research, Section A: Accelerators, Spectrometers, Detectors and Associated Equipment, 2018, 909, 294-297. | 0.7 | 3 |
| 12 | Multi-scale theoretical approach to X-ray absorption spectra in disordered systems: an application to the study of Zn(ii) in water. Physical Chemistry Chemical Physics, 2018, 20, 24775-24782. | 1.3 | 10 |
| 13 | Atomic structure of granulin determined from native nanocrystalline granulovirus using an X-ray free-electron laser. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 2247-2252. | 3.3 | 65 |
| 14 | Flow-aligned, single-shot fiber diffraction using a femtosecond X-ray free-electron laser. Cytoskeleton, 2017, 74, 472-481. | 1.0 | 12 |
| 15 | Open data set of live cyanobacterial cells imaged using an X-ray laser. Scientific Data, 2016, 3, 160058. | 2.4 | 7 |
| 16 | A data set from flash X-ray imaging of carboxysomes. Scientific Data, 2016, 3, 160061. | 2.4 | 11 |
| 17 | In cellulo serial crystallography of alcohol oxidase crystals inside yeast cells. IUCr, 2016, 3, 88-95. | 1.0 | 23 |
| 18 | Cu(II) – Zn(II) Cross-Modulation in Amyloid – Beta Peptide Binding: An X-ray Absorption Spectroscopy Study. Journal of Physical Chemistry B, 2015, 119, 15813-15820. | 1.2 | 16 |

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|----|--|------|-----------|
| 19 | Imaging single cells in a beam of live cyanobacteria with an X-ray laser. <i>Nature Communications</i> , 2015, 6, 5704. | 5.8 | 156 |
| 20 | Expression, purification and crystallization of CTB-MPR, a candidate mucosal vaccine component against HIV-1. <i>IUCr</i> , 2014, 1, 305-317. | 1.0 | 6 |
| 21 | Conformation sequence recovery of a non-periodic object from a diffraction-before-destruction experiment. <i>Optics Express</i> , 2014, 22, 8085. | 1.7 | 11 |
| 22 | Explosion dynamics of sucrose nanospheres monitored by time of flight spectrometry and coherent diffractive imaging at the split-and-delay beam line of the FLASH soft X-ray laser. <i>Optics Express</i> , 2014, 22, 28914. | 1.7 | 13 |
| 23 | Copper-zinc cross-modulation in prion protein binding. <i>European Biophysics Journal</i> , 2014, 43, 631-642. | 1.2 | 15 |
| 24 | High-throughput imaging of heterogeneous cell organelles with an X-ray laser. <i>Nature Photonics</i> , 2014, 8, 943-949. | 15.6 | 156 |
| 25 | Visualizing a protein quake with time-resolved X-ray scattering at a free-electron laser. <i>Nature Methods</i> , 2014, 11, 923-926. | 9.0 | 173 |
| 26 | Serial time-resolved crystallography of photosystem II using a femtosecond X-ray laser. <i>Nature</i> , 2014, 513, 261-265. | 13.7 | 403 |
| 27 | Serial crystallography on <i>in vivo</i> grown microcrystals using synchrotron radiation. <i>IUCr</i> , 2014, 1, 87-94. | 1.0 | 204 |
| 28 | Room-temperature macromolecular serial crystallography using synchrotron radiation. <i>IUCr</i> , 2014, 1, 204-212. | 1.0 | 221 |
| 29 | Natively Inhibited <i>Trypanosoma brucei</i> Cathepsin B Structure Determined by Using an X-ray Laser. <i>Science</i> , 2013, 339, 227-230. | 6.0 | 393 |
| 30 | Crystallographic data processing for free-electron laser sources. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2013, 69, 1231-1240. | 2.5 | 122 |
| 31 | Femtosecond free-electron laser x-ray diffraction data sets for algorithm development. <i>Optics Express</i> , 2012, 20, 4149. | 1.7 | 56 |
| 32 | Time-resolved protein nanocrystallography using an X-ray free-electron laser. <i>Optics Express</i> , 2012, 20, 2706. | 1.7 | 219 |
| 33 | Zn induced structural aggregation patterns of β^2 -amyloid peptides by first-principle simulations and XAS measurements. <i>Metalomics</i> , 2012, 4, 156-165. | 1.0 | 33 |
| 34 | Lipidic phase membrane protein serial femtosecond crystallography. <i>Nature Methods</i> , 2012, 9, 263-265. | 9.0 | 135 |
| 35 | Self-terminating diffraction gates femtosecond X-ray nanocrystallography measurements. <i>Nature Photonics</i> , 2012, 6, 35-40. | 15.6 | 292 |
| 36 | In vivo protein crystallization opens new routes in structural biology. <i>Nature Methods</i> , 2012, 9, 259-262. | 9.0 | 193 |

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|----|---|-----|-----------|
| 37 | High-Resolution Protein Structure Determination by Serial Femtosecond Crystallography. <i>Science</i> , 2012, 337, 362-364. | 6.0 | 758 |
| 38 | Radiation damage in protein serial femtosecond crystallography using an x-ray free-electron laser. <i>Physical Review B</i> , 2011, 84, 214111. | 1.1 | 156 |
| 39 | Zinc modulates copper coordination mode in prion protein octa-repeat subdomains. <i>European Biophysics Journal</i> , 2011, 40, 1259-1270. | 1.2 | 36 |
| 40 | Identifying the structure of the active sites of human recombinant prolidase. <i>European Biophysics Journal</i> , 2010, 39, 935-945. | 1.2 | 30 |
| 41 | An XAS study of the sulfur environment in human neuromelanin and its synthetic analogs. <i>European Biophysics Journal</i> , 2010, 39, 959-970. | 1.2 | 9 |
| 42 | Cu Involvement In Prion Oligopeptide Stability: Experiments And Numerical Simulations. <i>Biophysical Journal</i> , 2009, 96, 590a. | 0.2 | 0 |
| 43 | Identifying the Minimal Copper- and Zinc-binding Site Sequence in Amyloid- β^2 Peptides. <i>Journal of Biological Chemistry</i> , 2008, 283, 10784-10792. | 1.6 | 184 |
| 44 | Metal binding in amyloid β^2 -peptides shows intra- and inter-peptide coordination modes. <i>European Biophysics Journal</i> , 2006, 35, 340-351. | 1.2 | 104 |