

Masaki Okumura

List of Publications by Year in descending order

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Version: 2024-02-01

47
papers

1,221
citations

394390

19
h-index

377849

34
g-index

52
all docs

52
docs citations

52
times ranked

1604
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Synergistic cooperation of PDI family members in peroxiredoxin 4-driven oxidative protein folding. <i>Scientific Reports</i> , 2013, 3, 2456. | 3.3 | 118 |
| 2 | Structures and functions of protein disulfide isomerase family members involved in proteostasis in the endoplasmic reticulum. <i>Free Radical Biology and Medicine</i> , 2015, 83, 314-322. | 2.9 | 94 |
| 3 | Impact of membrane curvature on amyloid aggregation. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2018, 1860, 1741-1764. | 2.6 | 88 |
| 4 | Preparation of Selenoinsulin as a Long-Lasting Insulin Analogue. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 5522-5526. | 13.8 | 80 |
| 5 | High-resolution X-ray analysis reveals binding of arginine to aromatic residues of lysozyme surface: implication of suppression of protein aggregation by arginine. <i>Protein Engineering, Design and Selection</i> , 2011, 24, 269-274. | 2.1 | 75 |
| 6 | Redox-assisted regulation of Ca ²⁺ homeostasis in the endoplasmic reticulum by disulfide reductase ERdj5. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E6055-E6063. | 7.1 | 74 |
| 7 | The Protein Disulfide Isomerase Family: from proteostasis to pathogenesis. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2020, 1864, 129338. | 2.4 | 66 |
| 8 | A PDI-catalyzed thiol-disulfide switch regulates the production of hydrogen peroxide by human Ero1. <i>Free Radical Biology and Medicine</i> , 2015, 83, 361-372. | 2.9 | 59 |
| 9 | Dynamic assembly of protein disulfide isomerase in catalysis of oxidative folding. <i>Nature Chemical Biology</i> , 2019, 15, 499-509. | 8.0 | 58 |
| 10 | Radically Different Thioredoxin Domain Arrangement of ERp46, an Efficient Disulfide Bond Introducer of the Mammalian PDI Family. <i>Structure</i> , 2014, 22, 431-443. | 3.3 | 49 |
| 11 | Acceleration of disulfide-coupled protein folding using glutathione derivatives. <i>FEBS Journal</i> , 2011, 278, 1137-1144. | 4.7 | 44 |
| 12 | Inhibition of the Functional Interplay between Endoplasmic Reticulum (ER) Oxidoreductin-1 \pm (Ero1 \pm) and Protein-disulfide Isomerase (PDI) by the Endocrine Disruptor Bisphenol A. <i>Journal of Biological Chemistry</i> , 2014, 289, 27004-27018. | 3.4 | 38 |
| 13 | One-Dimensional Sliding of p53 Along DNA Is Accelerated in the Presence of Ca ²⁺ or Mg ²⁺ at Millimolar Concentrations. <i>Journal of Molecular Biology</i> , 2015, 427, 2663-2678. | 4.2 | 37 |
| 14 | A chemical method for investigating disulfide-coupled peptide and protein folding. <i>FEBS Journal</i> , 2012, 279, 2283-2295. | 4.7 | 34 |
| 15 | Diverse Structural Conversion and Aggregation Pathways of Alzheimer's Amyloid- β (1-40). <i>ACS Nano</i> , 2019, 13, 8766-8783. | 14.6 | 33 |
| 16 | The Highly Dynamic Nature of ERdj5 Is Key to Efficient Elimination of Aberrant Protein Oligomers through ER-Associated Degradation. <i>Structure</i> , 2017, 25, 846-857.e4. | 3.3 | 25 |
| 17 | Characterization and optimization of two-chain folding pathways of insulin via native chain assembly. <i>Communications Chemistry</i> , 2018, 1, . | 4.5 | 24 |
| 18 | Characterization of the endoplasmic reticulum-resident peroxidases GPx7 and GPx8 shows the higher oxidative activity of GPx7 and its linkage to oxidative protein folding. <i>Journal of Biological Chemistry</i> , 2020, 295, 12772-12785. | 3.4 | 23 |

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|----|--|-----|-----------|
| 19 | Coupling effects of thiol and urea-type groups for promotion of oxidative protein folding. <i>Chemical Communications</i> , 2019, 55, 759-762. | 4.1 | 21 |
| 20 | PDI Family Members as Guides for Client Folding and Assembly. <i>International Journal of Molecular Sciences</i> , 2020, 21, 9351. | 4.1 | 20 |
| 21 | Preparation of Selenoinsulin as a Long-Lasting Insulin Analogue. <i>Angewandte Chemie</i> , 2017, 129, 5614-5618. | 2.0 | 18 |
| 22 | Visualization of structural dynamics of protein disulfide isomerase enzymes in catalysis of oxidative folding and reductive unfolding. <i>Current Opinion in Structural Biology</i> , 2021, 66, 49-57. | 5.7 | 16 |
| 23 | Human ER Oxidoreductin-1 \pm (Ero1 \pm) Undergoes Dual Regulation through Complementary Redox Interactions with Protein-Disulfide Isomerase. <i>Journal of Biological Chemistry</i> , 2016, 291, 23952-23964. | 3.4 | 15 |
| 24 | Energy landscape of polymorphic amyloid generation of β 2-microglobulin revealed by calorimetry. <i>Chemical Communications</i> , 2018, 54, 7995-7998. | 4.1 | 14 |
| 25 | Antipsychotic olanzapine-induced misfolding of proinsulin in the endoplasmic reticulum accounts for atypical development of diabetes. <i>ELife</i> , 2020, 9, . | 6.0 | 14 |
| 26 | Cysteines 208 and 241 in Ero1 \pm are required for maximal catalytic turnover. <i>Redox Biology</i> , 2016, 7, 14-20. | 9.0 | 13 |
| 27 | Effects of positively charged redox molecules on disulfide-coupled protein folding. <i>FEBS Letters</i> , 2012, 586, 3926-3930. | 2.8 | 11 |
| 28 | Chemical Methods for Producing Disulfide Bonds in Peptides and Proteins to Study Folding Regulation. <i>Current Protocols in Protein Science</i> , 2014, 76, 28.7.1-28.7.13. | 2.8 | 9 |
| 29 | Structural stability of amyloid fibrils depends on the existence of the peripheral sequence near the core cross β region. <i>FEBS Letters</i> , 2015, 589, 3541-3547. | 2.8 | 9 |
| 30 | A unique leucine-valine adhesive motif supports structure and function of protein disulfide isomerase P5 via dimerization. <i>Structure</i> , 2021, 29, 1357-1370.e6. | 3.3 | 8 |
| 31 | Ero1-Mediated Reoxidation of Protein Disulfide Isomerase Accelerates the Folding of Cone Snail Toxins. <i>International Journal of Molecular Sciences</i> , 2018, 19, 3418. | 4.1 | 6 |
| 32 | Ca ²⁺ Regulates ERp57-Calnexin Complex Formation. <i>Molecules</i> , 2021, 26, 2853. | 3.8 | 6 |
| 33 | Distinct roles and actions of protein disulfide isomerase family enzymes in catalysis of nascent-chain disulfide bond formation. <i>IScience</i> , 2021, 24, 102296. | 4.1 | 5 |
| 34 | Glutathione Ethylester, a Novel Protein Refolding Reagent, Enhances both the Efficiency of Refolding and Correct Disulfide Formation. <i>Protein Journal</i> , 2012, 31, 499-503. | 1.6 | 4 |
| 35 | Crystallization and preliminary X-ray structural studies of human prouroguanylin. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2008, 64, 531-532. | 0.7 | 3 |
| 36 | Chemical Methods and Approaches to the Regioselective Formation of Multiple Disulfide Bonds. <i>Current Protocols in Protein Science</i> , 2014, 76, 28.8.1-28.8.28. | 2.8 | 3 |

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|----|---|-----|-----------|
| 37 | Crystallization and preliminary crystallographic analysis of the complex between triiodothyronine and the C ₂ fragment of rat protein disulfide isomerase. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012, 68, 476-478. | 0.7 | 2 |
| 38 | Conjugate of Thiol and Guanidyl Units with Oligoethylene Glycol Linkage for Manipulation of Oxidative Protein Folding. <i>Molecules</i> , 2021, 26, 879. | 3.8 | 2 |
| 39 | Functional Interplay between P5 and PDI/ERp72 to Drive Protein Folding. <i>Biology</i> , 2021, 10, 1112. | 2.8 | 2 |
| 40 | Folding Mechanism of a Precursor Protein of a Peptide Hormone Mediated by an Intra-Molecular Chaperone. <i>Biophysical Journal</i> , 2012, 102, 56a. | 0.5 | 0 |
| 41 | Role of Leu66 in the Folding of Uroguanylin Assisted by Intra-Molecular Chaperone. <i>Biophysical Journal</i> , 2012, 102, 56a. | 0.5 | 0 |
| 42 | Acceleration of Disulfide-Coupled Protein Folding by Positively Charged Glutathione Derivatives. <i>Biophysical Journal</i> , 2012, 102, 57a. | 0.5 | 0 |
| 43 | Regulation of Disulfide Coupled Folding of De Novo Designed Precursor Protein. <i>Biophysical Journal</i> , 2014, 106, 472a. | 0.5 | 0 |
| 44 | Positively Charged Redox Agents Accelerate Disulfide Coupled Protein Folding. <i>Biophysical Journal</i> , 2014, 106, 472a. | 0.5 | 0 |
| 45 | Folding Analyses of the Major Folding Intermediate of Prouroguanylin using Deletion Mutants. <i>Biophysical Journal</i> , 2016, 110, 389a. | 0.5 | 0 |
| 46 | Disulfide Selectivity under the Control of Secondary Structure in Protein Folding. <i>Biophysical Journal</i> , 2016, 110, 210a. | 0.5 | 0 |
| 47 | Structural Analyses of a Linker Region of the Amyloid Precursor Protein. <i>Biophysical Journal</i> , 2018, 114, 78a. | 0.5 | 0 |