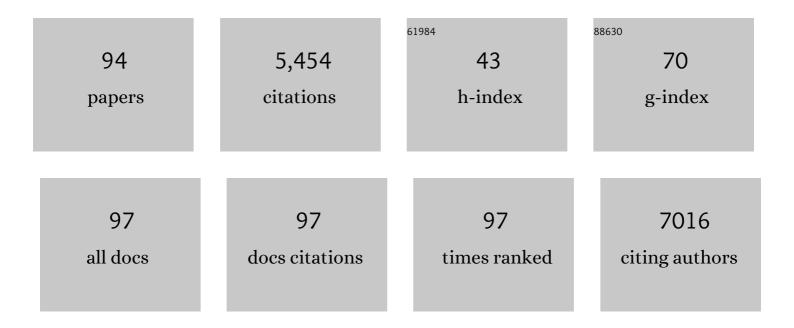
## **Richard A Kammerer**

List of Publications by Year in descending order

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| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | COMP-Ang1: A designed angiopoietin-1 variant with nonleaky angiogenic activity. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 5547-5552.   | 7.1  | 236       |
| 2  | Structural basis of tubulin tyrosination by tubulin tyrosine ligase. Journal of Cell Biology, 2013, 200, 259-270.  | 5.2  | 189       |
| 3  | Stabilization of short collagen-like triple helices by protein engineering. Journal of Molecular<br>Biology, 2001, 308, 1081-1089.   | 4.2  | 177       |
| 4  | Cortexillins, Major Determinants of Cell Shape and Size, Are Actin-Bundling Proteins with a Parallel<br>Coiled-Coil Tail. Cell, 1996, 86, 631-642.   | 28.9 | 172       |
| 5  | An autonomous folding unit mediates the assembly of two-stranded coiled coils. Proceedings of the National Academy of Sciences of the United States of America, 1998, 95, 13419-13424.   | 7.1  | 166       |
| 6  | Exploring amyloid formation by a de novo design. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 4435-4440.  | 7.1  | 166       |
| 7  | Crystal structure of a naturally occurring parallel right-handed coiled coil tetramer. Nature<br>Structural Biology, 2000, 7, 772-776.   | 9.7  | 155       |
| 8  | Microtubule minus-end regulation at spindle poles by an ASPM–katanin complex. Nature Cell Biology,<br>2017, 19, 480-492.   | 10.3 | 147       |
| 9  | Designed angiopoietin-1 variant, COMP-Ang1, protects against radiation-induced endothelial cell<br>apoptosis. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101,<br>5553-5558.        | 7.1  | 134       |
| 10 | Oligomerization and Multimerization Are Critical for Angiopoietin-1 to Bind and Phosphorylate Tie2.<br>Journal of Biological Chemistry, 2005, 280, 20126-20131.  | 3.4  | 134       |
| 11 | Subdomain-Specific Localization of Climp-63 (P63) in the Endoplasmic Reticulum Is Mediated by Its<br>Luminal α-Helical Segment. Journal of Cell Biology, 2001, 153, 1287-1300.   | 5.2  | 127       |
| 12 | Molecular basis of coiled-coil formation. Proceedings of the National Academy of Sciences of the<br>United States of America, 2007, 104, 7062-7067.  | 7.1  | 116       |
| 13 | The coiled-coil trigger site of the rod domain of cortexillin I unveils a distinct network of interhelical and intrahelical salt bridges. Structure, 2000, 8, 223-230.   | 3.3  | 114       |
| 14 | A distinct 14 residue site triggers coiled-coil formation in cortexillin I. EMBO Journal, 1998, 17,<br>1883-1891.  | 7.8  | 113       |
| 15 | Spectraplakins Promote Microtubule-Mediated Axonal Growth by Functioning As Structural<br>Microtubule-Associated Proteins and EB1-Dependent +TIPs (Tip Interacting Proteins). Journal of<br>Neuroscience, 2012, 32, 9143-9158. | 3.6  | 104       |
| 16 | Structural basis for recognition of synaptic vesicle protein 2C by botulinum neurotoxin A. Nature, 2014, 505, 108-111.   | 27.8 | 103       |
| 17 | Tenascin-C Hexabrachion Assembly Is a Sequential Two-step Process Initiated by Coiled-coil α-Helices.<br>Journal of Biological Chemistry, 1998, 273, 10602-10608.  | 3.4  | 99        |
| 18 | Configurational entropy elucidates the role of salt-bridge networks in protein thermostability.<br>Protein Science, 2007, 16, 1349-1359.   | 7.6  | 99        |

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|----|--|------|-----------|
| 19 | Op18/stathmin caps a kinked protofilament-like tubulin tetramer. EMBO Journal, 2000, 19, 572-580.  | 7.8  | 92        |
| 20 | Nucleation and propagation of the collagen triple helix in single-chain and trimerized peptides:<br>transition from third to first order kinetics. Journal of Molecular Biology, 2002, 317, 459-470.                                       | 4.2  | 91        |
| 21 | Centriolar CPAP/SAS-4 Imparts Slow Processive Microtubule Growth. Developmental Cell, 2016, 37, 362-376.   | 7.0  | 90        |
| 22 | Heterodimerization of a Functional GABABReceptor Is Mediated by Parallel Coiled-Coil α-Helicesâ€.<br>Biochemistry, 1999, 38, 13263-13269.  | 2.5  | 88        |
| 23 | A conserved trimerization motif controls the topology of short coiled coils. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 13891-13896.  | 7.1  | 88        |
| 24 | Mutations in HPSE2 Cause Urofacial Syndrome. American Journal of Human Genetics, 2010, 86, 963-969.  | 6.2  | 88        |
| 25 | A Novel Receptor-induced Activation Site in the Nipah Virus Attachment Glycoprotein (G) Involved in<br>Triggering the Fusion Glycoprotein (F). Journal of Biological Chemistry, 2009, 284, 1628-1635.                                      | 3.4  | 83        |
| 26 | Collagen Stabilization at Atomic Level. Structure, 2003, 11, 339-346.  | 3.3  | 76        |
| 27 | α-Helical coiled-coil oligomerization domains in extracellular proteins. Matrix Biology, 1997, 15, 555-565.  | 3.6  | 74        |
| 28 | Remorins form a novel family of coiled coil-forming oligomeric and filamentous proteins associated with apical, vascular and embryonic tissues in plants. Plant Molecular Biology, 2004, 55, 579-594.                                      | 3.9  | 74        |
| 29 | SAS-6 engineering reveals interdependence between cartwheel and microtubules in determining centrioleAarchitecture. Nature Cell Biology, 2016, 18, 393-403.  | 10.3 | 73        |
| 30 | All-trans retinol, vitamin D and other hydrophobic compounds bind in the axial pore of the<br>five-stranded coiled-coil domain of cartilage oligomeric matrix protein. EMBO Journal, 1998, 17,<br>5265-5272.                               | 7.8  | 67        |
| 31 | Domain analysis of cortexillin I: actin-bundling, PIP2-binding and the rescue of cytokinesis. EMBO<br>Journal, 1999, 18, 5274-5284.  | 7.8  | 67        |
| 32 | Molecular basis of coiled-coil oligomerization-state specificity. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 19850-19855.   | 7.1  | 66        |
| 33 | Coronin 1 Regulates Cognition and Behavior through Modulation of cAMP/Protein Kinase A Signaling.<br>PLoS Biology, 2014, 12, e1001820.   | 5.6  | 62        |
| 34 | Angiopoietin-1 variant reduces LPS-induced microvascular dysfunction in a murine model of sepsis.<br>Critical Care, 2012, 16, R182.  | 5.8  | 57        |
| 35 | Polymorphism in an Amyloid‣ike Fibrilâ€Forming Model Peptide. Angewandte Chemie - International<br>Edition, 2008, 47, 5842-5845.   | 13.8 | 53        |
| 36 | Evidence That Monoclonal Antibodies Directed against the Integrin Î <sup>2</sup> Subunit<br>Plexin/Semaphorin/Integrin Domain Stimulate Function by Inducing Receptor Extension. Journal of<br>Biological Chemistry, 2005, 280, 4238-4246. | 3.4  | 52        |

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|----|--|-----|-----------|
| 37 | Electrostatic Contributions to the Stability of the GCN4 Leucine Zipper Structure. Journal of<br>Molecular Biology, 2007, 374, 206-219.  | 4.2 | 51        |
| 38 | GAS2-like proteins mediate communication between microtubules and actin through interaction with end-binding proteins. Journal of Cell Science, 2014, 127, 2672-82.  | 2.0 | 51        |
| 39 | Characterization of the Matrilin Coiled-coil Domains Reveals Seven Novel Isoforms. Journal of<br>Biological Chemistry, 2002, 277, 19071-19079.   | 3.4 | 50        |
| 40 | Selective Chain Recognition in the C-terminal α-Helical Coiled-coil Region of Laminin. Journal of<br>Molecular Biology, 1995, 250, 64-73.  | 4.2 | 48        |
| 41 | An Intrahelical Salt Bridge within the Trigger Site Stabilizes the GCN4 Leucine Zipper. Journal of<br>Biological Chemistry, 2001, 276, 13685-13688.  | 3.4 | 47        |
| 42 | Interaction of filamin A with the integrin β7cytoplasmic domain: role of alternative splicing and phosphorylation. FEBS Letters, 2004, 569, 185-190.   | 2.8 | 47        |
| 43 | Structural Basis of Formation of the Microtubule Minus-End-Regulating CAMSAP-Katanin Complex.<br>Structure, 2018, 26, 375-382.e4.  | 3.3 | 47        |
| 44 | A Distinct Seven-residue Trigger Sequence Is Indispensable for Proper Coiled-coil Formation of the<br>Human Macrophage Scavenger Receptor Oligomerization Domain. Journal of Biological Chemistry,<br>2000, 275, 11672-11677.      | 3.4 | 46        |
| 45 | Modulation of Agrin Function by Alternative Splicing and Ca2+ Binding. Structure, 2004, 12, 503-515.   | 3.3 | 45        |
| 46 | The Oligomerization Domain of the Asialoglycoprotein Receptor Preferentially Forms 2:2<br>Heterotetramers in Vitro. Journal of Biological Chemistry, 1996, 271, 31996-32001.   | 3.4 | 44        |
| 47 | The laminin-binding domain of agrin is structurally related to N-TIMP-1. Nature Structural Biology, 2001, 8, 705-709.  | 9.7 | 41        |
| 48 | The Angiopoietin-like Factor Cornea-derived Transcript 6 Is a Putative Morphogen for Human Cornea.<br>Journal of Biological Chemistry, 2002, 277, 686-693.   | 3.4 | 41        |
| 49 | De novo design of a two-stranded coiled-coil switch peptide. Journal of Structural Biology, 2006, 155, 146-153.  | 2.8 | 41        |
| 50 | Laminin chain assembly is regulated by specific coiled-coil interactions. Journal of Structural Biology, 2010, 170, 398-405.   | 2.8 | 41        |
| 51 | Heteronuclear NMR assignments and secondary structure of the coiled coil trimerization domain from cartilage matrix protein in oxidized and reduced forms. Protein Science, 1997, 6, 1734-1745.                                    | 7.6 | 40        |
| 52 | Role of dimerization and substrate exclusion in the regulation of bone morphogenetic protein-1 and<br>mammalian tolloid. Proceedings of the National Academy of Sciences of the United States of America,<br>2009, 106, 8561-8566. | 7.1 | 40        |
| 53 | What are oligomerization domains good for?. Matrix Biology, 2000, 19, 283-288.   | 3.6 | 39        |
| 54 | Short Linear Sequence Motif LxxPTPh Targets Diverse Proteins to Growing Microtubule Ends.<br>Structure, 2017, 25, 924-932.e4.  | 3.3 | 37        |

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|----|---|-----|-----------|
| 55 | NMR structure of a parallel homotrimeric coiled coil. Nature Structural and Molecular Biology, 1998, 5, 687-691.  | 8.2 | 36        |
| 56 | Interaction of mammalian end binding proteins with CAP-Gly domains of CLIP-170 and p150glued.<br>Journal of Structural Biology, 2012, 177, 160-167.   | 2.8 | 36        |
| 57 | <sup>15</sup> N backbone dynamics of the Sâ€peptide from ribonuclease A in its free and Sâ€protein bound<br>forms: Toward a siteâ€specific analysis of entropy changes upon folding. Protein Science, 1998, 7,<br>389-402.  | 7.6 | 35        |
| 58 | Structural Analysis of the Sixth Immunoglobulinâ€Like Domain of Mouse Neural Cell Adhesion<br>Molecule L1 and Its Interactions with αvl²3, αIIbl²3, and α5l²1 Integrins. Journal of Neurochemistry, 1998, 71,<br>2615-2625. | 3.9 | 35        |
| 59 | Structure of the Extracellular Domain of Tie Receptor Tyrosine Kinases and Localization of the Angiopoietin-binding Epitope. Journal of Biological Chemistry, 2006, 281, 28408-28414.                                       | 3.4 | 35        |
| 60 | Role of the nucleotidyl cyclase helical domain in catalytically active dimer formation. Proceedings of the United States of America, 2017, 114, E9821-E9828.  | 7.1 | 35        |
| 61 | Atomic Models of De Novo Designed ccβ-Met Amyloid-Like Fibrils. Journal of Molecular Biology, 2008,<br>376, 898-912.  | 4.2 | 34        |
| 62 | A 35-kDa Protein Is the Basic Unit of the Core from the 2 × 104-kDa Aggregation Factor Responsible for<br>Species-specific Cell Adhesion in the Marine Sponge. Journal of Biological Chemistry, 1996, 271,<br>23558-23565.  | 3.4 | 33        |
| 63 | Characterization of G2L3 (GAS2-like 3), a New Microtubule- and Actin-binding Protein Related to Spectraplakins. Journal of Biological Chemistry, 2011, 286, 24987-24995.  | 3.4 | 31        |
| 64 | Angiopoietin-1 regulates microvascular reactivity and protects the microcirculation during acute endothelial dysfunction: Role of eNOS and VE-cadherin. Pharmacological Research, 2014, 80, 43-51.                          | 7.1 | 31        |
| 65 | Biophysical and Structural Characterization of the Centriolar Protein Cep104 Interaction Network.<br>Journal of Biological Chemistry, 2016, 291, 18496-18504.   | 3.4 | 31        |
| 66 | Structure and disorder in the ribonuclease S-peptide probed by NMR residual dipolar couplings.<br>Protein Science, 2009, 12, 2132-2140.   | 7.6 | 27        |
| 67 | Stabilization of the α-Helical Coiled-coil Domain in Laminin by C-terminal Disulfide Bonds. Journal of<br>Molecular Biology, 1995, 250, 74-79.  | 4.2 | 26        |
| 68 | Structural basis for misregulation of kinesin KIF21A autoinhibition by CFEOM1 disease mutations.<br>Scientific Reports, 2016, 6, 30668.   | 3.3 | 26        |
| 69 | Toward a High-Resolution Structure of Phospholamban:Â Design of Soluble Transmembrane Domain<br>Mutantsâ€. Biochemistry, 2000, 39, 6825-6831.   | 2.5 | 25        |
| 70 | Botulinum neurotoxins: new questions arising from structural biology. Trends in Biochemical Sciences, 2014, 39, 517-526.  | 7.5 | 25        |
| 71 | The unusually stable coiled-coil domain of COMP exhibits cold and heat denaturation in 4–6 M<br>guanidinium chloride. Biophysical Chemistry, 2000, 85, 179-186.   | 2.8 | 24        |
| 72 | A Type IV Translocated Legionella Cysteine Phytase Counteracts Intracellular Growth Restriction by<br>Phytate. Journal of Biological Chemistry, 2014, 289, 34175-34188.   | 3.4 | 24        |

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|----|--|-----|-----------|
| 73 | Structural basis of katanin p60:p80 complex formation. Scientific Reports, 2017, 7, 14893.   | 3.3 | 24        |
| 74 | Crystal structure of the BoNT/A2 receptor-binding domain in complex with the luminal domain of its neuronal receptor SV2C. Scientific Reports, 2017, 7, 43588.   | 3.3 | 23        |
| 75 | Rat GTP cyclohydrolase I is a homodecameric protein complex containing high-affinity calcium-binding sites 1 1Edited by W. Baumeister. Journal of Molecular Biology, 1998, 279, 189-199.   | 4.2 | 21        |
| 76 | Collagen VI, Conformation of A-domain Arrays and Microfibril Architecture. Journal of Biological Chemistry, 2011, 286, 40266-40275.  | 3.4 | 21        |
| 77 | Contributions of the ionization states of acidic residues to the stability of the coiled coil domain of matrilin-1. FEBS Letters, 1999, 446, 75-80.  | 2.8 | 18        |
| 78 | Angiopoietin-1 enhances neutrophil chemotaxis in vitro and migration in vivo through interaction with CD18 and release of CCL4. Scientific Reports, 2017, 7, 2332.   | 3.3 | 13        |
| 79 | Highâ€Level Production of Phenylacetaldehyde using Fusionâ€Tagged Styrene Oxide Isomerase. Advanced<br>Synthesis and Catalysis, 2021, 363, 1714-1721.  | 4.3 | 12        |
| 80 | Crystal Structure of a Heterotetrameric Katanin p60:p80 Complex. Structure, 2019, 27, 1375-1383.e3.  | 3.3 | 11        |
| 81 | The nuclear protein Waharan is required for endosomal-lysosomal trafficking in <i>Drosophila</i> .<br>Journal of Cell Science, 2010, 123, 2369-2374.   | 2.0 | 10        |
| 82 | The role of the N-terminal amphipathic helix in bacterial YidC: Insights from functional studies, the<br>crystal structure and molecular dynamics simulations. Biochimica Et Biophysica Acta - Biomembranes,<br>2022, 1864, 183825.            | 2.6 | 10        |
| 83 | Thermodynamic and Structural Studies of Carbohydrate Binding by the Agrin-G3 Domain. Biochemistry, 2007, 46, 9541-9550.  | 2.5 | 9         |
| 84 | Crystallization and Preliminary X-Ray Diffraction Analysis of the 190-ÃLong Coiled-Coil Dimerization<br>Domain of the Actin-Bundling Protein Cortexillin I fromDictyostelium discoideum. Journal of<br>Structural Biology, 1998, 122, 293-296. | 2.8 | 7         |
| 85 | Structure of the BoNT/A1 – receptor complex. Toxicon, 2015, 107, 25-31.  | 1.6 | 6         |
| 86 | Nuclear Magnetic Resonance Structures of GCN4p Are Largely Conserved When Ion Pairs Are<br>Disrupted at Acidic pH but Show a Relaxation of the Coiled Coil Superhelix. Biochemistry, 2017, 56,<br>1604-1619.                                   | 2.5 | 6         |
| 87 | Design of a Coiled-Coil-based Model Peptide System to Explore the Fundamentals of Amyloid Fibril<br>Formation. International Journal of Peptide Research and Therapeutics, 2005, 11, 43-52.  | 1.9 | 5         |
| 88 | Synthesis and Evaluation of Biphenyl Compounds as Kinesin Spindle Protein Inhibitors. Chemistry and<br>Biodiversity, 2013, 10, 538-555.  | 2.1 | 5         |
| 89 | 1H, 13C and 15N backbone assignments for the C-terminal globular domain of agrin. Journal of<br>Biomolecular NMR, 2001, 20, 295-296.   | 2.8 | 4         |
| 90 | Crystal structure of the catalytic domain of botulinum neurotoxin subtype A3. Journal of Biological<br>Chemistry, 2021, 296, 100684.   | 3.4 | 4         |

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|----|--|-----|-----------|
| 91 | Structural insights into the interaction of botulinum neurotoxin a with its neuronal receptor SV2C.<br>Toxicon, 2020, 175, 36-43.  | 1.6 | 3         |
| 92 | Structural Basis for the Oligomerization-State Switch from a Dimer to a Trimer of an Engineered Cortexillin-1 Coiled-Coil Variant. PLoS ONE, 2013, 8, e63370.                                      | 2.5 | 2         |
| 93 | Mutations in HPSE2 Cause Urofacial Syndrome. American Journal of Human Genetics, 2010, 87, 309.  | 6.2 | 1         |
| 94 | Homodimerization of coronin A through the Câ€ŧerminal coiled oil domain is essential for<br>multicellular differentiation of <i>DictyosteliumÂdiscoideum</i> . FEBS Letters, 2020, 594, 2116-2127. | 2.8 | 1         |