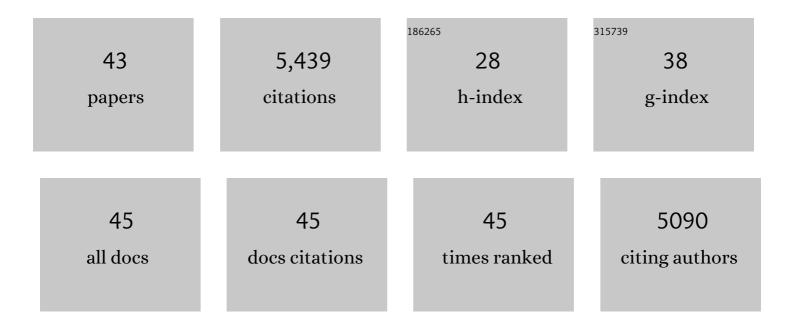
## **Robert Karlsson**

List of Publications by Year in descending order

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| #  | Article   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Kinetic analysis of monoclonal antibody-antigen interactions with a new biosensor based analytical system. Journal of Immunological Methods, 1991, 145, 229-240.  | 1.4 | 1,080     |
| 2  | Experimental design for kinetic analysis of protein-protein interactions with surface plasmon resonance biosensors. Journal of Immunological Methods, 1997, 200, 121-133.   | 1.4 | 524       |
| 3  | Biospecific interaction analysis using surface plasmon resonance detection applied to kinetic, binding site and concentration analysis. Journal of Chromatography A, 1992, 597, 397-410.                          | 3.7 | 431       |
| 4  | SPR for molecular interaction analysis: a review of emerging application areas. Journal of Molecular<br>Recognition, 2004, 17, 151-161.   | 2.1 | 391       |
| 5  | Analyzing a kinetic titration series using affinity biosensors. Analytical Biochemistry, 2006, 349, 136-147.  | 2.4 | 352       |
| 6  | Biosensor Analysis of the Interaction between Immobilized Human Serum Albumin and Drug<br>Compounds for Prediction of Human Serum Albumin Binding Levels. Journal of Medicinal Chemistry,<br>2000, 43, 1986-1992. | 6.4 | 288       |
| 7  | Kinetic and Concentration Analysis Using BIA Technology. Methods, 1994, 6, 99-110.  | 3.8 | 251       |
| 8  | Detection of antigen—antibody interactions by surface plasmon resonance. Application to Epitope<br>Mapping. Journal of Molecular Recognition, 1990, 3, 208-214.   | 2.1 | 218       |
| 9  | Analysis of macromolecular interactions using immobilized ligands. Analytical Biochemistry, 1992, 201, 197-210.   | 2.4 | 195       |
| 10 | Biosensor Analysis of Drug–Target Interactions: Direct and Competitive Binding Assays for<br>Investigation of Interactions between Thrombin and Thrombin Inhibitors. Analytical Biochemistry,<br>2000, 278, 1-13. | 2.4 | 133       |
| 11 | SPR Biosensor Studies of the Direct Interaction between 27 Drugs and a Liposome Surface:Â<br>Correlation with Fraction Absorbed in Humans. Journal of Medicinal Chemistry, 2000, 43, 2083-2086.                   | 6.4 | 133       |
| 12 | Analysis of active antibody concentration. Separation of affinity and concentration parameters.<br>Journal of Immunological Methods, 1993, 166, 75-84.  | 1.4 | 130       |
| 13 | Biomolecular interaction analysis: affinity biosensor technologies for functional analysis of proteins. Current Opinion in Chemical Biology, 1997, 1, 378-383.  | 6.1 | 127       |
| 14 | Screening antibody–antigen interactions in parallel using Biacore A100. Analytical Biochemistry, 2006,<br>353, 181-190.   | 2.4 | 108       |
| 15 | Thermodynamic analysis of protein interactions with biosensor technology. , 1998, 11, 204-210.  |     | 95        |
| 16 | Affinity analysis of non-steady-state data obtained under mass transport limited conditions using<br>BIAcore technology. , 1999, 12, 285-292.   |     | 95        |
| 17 | Kinetic studies of small molecule interactions with protein kinases using biosensor technology.<br>Analytical Biochemistry, 2005, 340, 359-368.   | 2.4 | 94        |
| 18 | Kinetic analysis of the interaction between protein a domain variants and human Fc using plasmon resonance detection. Journal of Molecular Recognition, 1995, 8, 270-278.   | 2.1 | 80        |

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|----|---|-----|-----------|
| 19 | Rational Design of Thermodynamic and Kinetic Binding Profiles by Optimizing Surface Water Networks<br>Coating Protein-Bound Ligands. Journal of Medicinal Chemistry, 2016, 59, 10530-10548.   | 6.4 | 64        |
| 20 | Identification of Fc Gamma Receptor Glycoforms That Produce Differential Binding Kinetics for Rituximab. Molecular and Cellular Proteomics, 2017, 16, 1770-1788.  | 3.8 | 64        |
| 21 | Fc Gamma Receptor Glycosylation Modulates the Binding of IgG Glycoforms: A Requirement for Stable Antibody Interactions. Journal of Proteome Research, 2014, 13, 5471-5485.   | 3.7 | 61        |
| 22 | Determination of Interaction Kinetic Constants for HIV-1 Protease Inhibitors Using Optical Biosensor<br>Technology. Analytical Biochemistry, 2001, 291, 207-218.  | 2.4 | 58        |
| 23 | Label-Free Primary Screening and Affinity Ranking of Fragment Libraries Using Parallel Analysis of<br>Protein Panels. Journal of Biomolecular Screening, 2008, 13, 202-209.   | 2.6 | 50        |
| 24 | Binding of autoreactive mouse anti-type II collagen antibodies derived from the primary and the<br>secondary immune response investigated with the biosensor technique. Journal of Immunological<br>Methods, 1995, 188, 63-71.                | 1.4 | 49        |
| 25 | A Thermodynamic Characterization of the Binding of Thrombin Inhibitors to Human Thrombin,<br>Combining Biosensor Technology, Stopped-Flow Spectrophotometry, and Microcalorimetry.<br>Analytical Biochemistry, 2002, 300, 152-162.            | 2.4 | 40        |
| 26 | Evaluation of calibration-free concentration analysis provided by Biacoreâ"¢ systems. Analytical<br>Biochemistry, 2016, 510, 88-97.   | 2.4 | 35        |
| 27 | Biosensor Analysis of the Interaction between Drug Compounds and Liposomes of Different<br>Properties; a Two-Dimensional Characterization Tool for Estimation of Membrane Absorption. Journal<br>of Pharmaceutical Sciences, 2005, 94, 25-37. | 3.3 | 33        |
| 28 | Direct and competitive kinetic analysis of the interaction between human IgG1 and a one domain analogue of protein A. Journal of Immunological Methods, 1995, 183, 43-49.   | 1.4 | 32        |
| 29 | Comparison of surface plasmon resonance binding curves for characterization of protein interactions and analysis of screening data. Analytical Biochemistry, 2016, 502, 53-63.  | 2.4 | 31        |
| 30 | Biosensor-based characterization of serum antibodies during development of an anti-IgE<br>immunotherapeutic against allergy and asthma. Journal of Molecular Recognition, 2007, 20, 22-31.  | 2.1 | 30        |
| 31 | Label-free kinetic binding data as a decisive element in drug discovery. Expert Opinion on Drug<br>Discovery, 2006, 1, 439-446.   | 5.0 | 26        |
| 32 | Affinity Measurement Using Surface Plasmon Resonance. , 2004, 248, 389-416.   |     | 25        |
| 33 | Calibration-free concentration analysis of protein biomarkers in human serum using surface plasmon<br>resonance. Talanta, 2015, 144, 801-808.   | 5.5 | 22        |
| 34 | Statistical aspects of van't Hoff analysis: a simulation study. Journal of Molecular Recognition, 2007, 20, 379-385.  | 2.1 | 21        |
| 35 | Biosensor binding data and its applicability to the determination of active concentration. Biophysical<br>Reviews, 2016, 8, 347-358.  | 3.2 | 21        |
| 36 | Elucidating the Origin of Long Residence Time Binding for Inhibitors of the Metalloprotease<br>Thermolysin. ACS Chemical Biology, 2017, 12, 225-233.  | 3.4 | 14        |

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|----|--|-----|-----------|
| 37 | New insights into DNA-binding behavior of Wilms Tumor Protein (WT1) — A dual study. Biophysical<br>Chemistry, 2009, 145, 116-125.  | 2.8 | 11        |
| 38 | Surrogate potency assays: Comparison of binding profiles complements dose response curves for unambiguous assessment of relative potencies. Journal of Pharmaceutical Analysis, 2018, 8, 138-146.  | 5.3 | 10        |
| 39 | Use of Isotachophoresis for Identification of Pollen Allergens. International Archives of Allergy and<br>Immunology, 1982, 68, 222-225.  | 2.1 | 6         |
| 40 | Pre-labeling of diverse protein samples with a fixed amount of Cy5 for sodium dodecyl<br>sulfate–polyacrylamide gel electrophoresis analysis. Analytical Biochemistry, 2015, 484, 51-57.   | 2.4 | 6         |
| 41 | Surface Plasmon Resonance in Binding Site, Kinetic, and Concentration Analyses. , 2013, , 209-221.   |     | 4         |
| 42 | 047 Biospecific interaction analysis using surface plasmon resonance detection. Mapping of binding<br>sites on IGF-II and kinetic analysis of the interaction of IGF-II with IGFBP-1 and the IGF-II receptor.<br>Fresenius' Journal of Analytical Chemistry, 1992, 343, 100-101. | 1.5 | 0         |
| 43 | BlAcore Technology — A Marriage Partner to Mass Spectrometry ?. , 1998, , 295-303.   |     | 0         |