

Thiemo C Huuk

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/784997/publications.pdf>

Version: 2024-02-01

11
papers

270
citations

933447

10
h-index

1281871

11
g-index

12
all docs

12
docs citations

12
times ranked

138
citing authors

#	ARTICLE	IF	CITATIONS
1	Model-based integrated optimization and evaluation of a multi-step ion exchange chromatography. Separation and Purification Technology, 2014, 136, 207-222.	7.9	56
2	Simulating and Optimizing Preparative Protein Chromatography with ChromX. Journal of Chemical Education, 2015, 92, 1497-1502.	2.3	44
3	UV absorption-based inverse modeling of protein chromatography. Engineering in Life Sciences, 2016, 16, 99-106.	3.6	33
4	Modeling of complex antibody elution behavior under high protein load densities in ion exchange chromatography using an asymmetric activity coefficient. Biotechnology Journal, 2017, 12, 1600336.	3.5	24
5	A versatile noninvasive method for adsorber quantification in batch and column chromatography based on the ionic capacity. Biotechnology Progress, 2016, 32, 666-677.	2.6	22
6	Calibration-free inverse modeling of ion-exchange chromatography in industrial antibody purification. Engineering in Life Sciences, 2016, 16, 107-113.	3.6	21
7	Protein adsorption on ion exchange adsorbers: A comparison of a stoichiometric and non-stoichiometric modeling approach. Journal of Chromatography A, 2021, 1653, 462397.	3.7	21
8	Analysis of complex protein elution behavior in preparative ion exchange processes using a colloidal particle adsorption model. Journal of Chromatography A, 2021, 1654, 462439.	3.7	16
9	Modeling the Gibbs-Donnan effect during ultrafiltration and diafiltration processes using the Poisson-Boltzmann theory in combination with a basic Stern model. Journal of Membrane Science, 2022, 648, 120333.	8.2	15
10	Adsorption of colloidal proteins in ion-exchange chromatography under consideration of charge regulation. Journal of Chromatography A, 2020, 1611, 460608.	3.7	11
11	Deconvolution of high-throughput multicomponent isotherms using multivariate data analysis of protein spectra. Engineering in Life Sciences, 2016, 16, 194-201.	3.6	7