

# DryLab® Publications

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- 230. Computer-Assisted Method Development for Small and Large Molecules**  
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- 229. Computer-assisted UHPLC–MS method development and optimization for the determination of 24 antineoplastic drugs used in hospital pharmacy**, Nicolas Guichard, Szabolcs Fekete, Davy Guillarme, Pascal Bonnabry, Sandrine Fleury-Souverain, *J Pharm Biomed Anal.*, **164**, 5 February, 395–401 (2019)
- 228. Is hydrophobic interaction chromatography the most suitable technique to characterize site-specific antibody-drug conjugates?** Valentina D'Atri, Reinhard Pell, Adrian Clarke, Davy Guillarme, Szabolcs Fekete  
*J Chromatogr. A*, **1586**, 8 February, 149–153 (2019)
- 227. Tuning selectivity in cation-exchange chromatography applied for monoclonal antibody separations, part 1: Alternative mobile phases and fine tuning of the separation**, Evelin Farsang et. Al,  
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- 226. Development of a design of experiments optimized method for quantification of polysorbate 80 based on oleic acid using UHPLC-MS**, Julia Punchman, Dirk-H. Evers, Christel C. Müller-Goymann, Michael E. Herbig,  
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- 225. Structure assisted impurity profiling for rapid method development in liquid chromatography**  
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- 224. Simultaneous optimization of mobile phase composition and pH using retention modeling and experimental design**, Norbert Rácz, Imre Molnár, Arnold Zöldhegyi, Hans-Jürgen Rieger, Róbert Kormány, *J Pharm Biomed Anal.*, **160**, 336–343 (2018)
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- 221. Analysis of recombinant monoclonal antibodies in hydrophilic interaction chromatography: A generic method development approach** B. Bobály, V. D'Atri, A. Beck, D. Guillarme, Sz. Fekete, *J. Pharm. Biomed. Anal.* **145** (2017) 25 October, 24–32
- 220. Computer-Assisted Method Development for Small and Large Molecules**, Sz. Fekete, R. Kormány, and D. Guillarme, *LCGC Special* **30**(2017) Issue 6, 14–21
- 219. Separation of antibody drug conjugate species by RPLC: A generic method development approach** S. Fekete, I. Molnár, D. Guillarme, *J. Pharm. Biomed. Anal.* **137**(2017) 15 April, 60–69
- 218. Separation of Atropisomers by Chiral Liquid Chromatography and Thermodynamic Analysis of Separation Mechanism** L. Zhang, Y. Hu, E. Galella, F. P. Tomasella, W. P. Fish, *J. Pharm. Anal.* **7**(2017) 156–162
- 217. Renewal of an old European Pharmacopoeia method for Terazosin using modeling with mass spectrometric peak tracking** R. Kormány, I. Molnár, J. Fekete, *J. Pharm. Biomed. Anal.* **135**(2017) 20 February, 8–15
- 216. Optimization of non-linear gradient in hydrophobic interaction chromatography for the analytical characterization of antibody-drug conjugates** B. Bobály, G. M. Randazzo, S. Rudaz, D. Guillarme, S. Fekete,  
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