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In-line Fourier-Transform Infrared Spectroscopy as a
Versatile Process Analytical Technology for Preparative Protein Chromatography

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10 Abstract

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Fourier-Transform Infrared Spectroscopy (FTIR) is a well-established spec-11 troscopic method in the analysis of small molecules and protein secondary struc-12 ture. However, FTIR is not commonly applied for in-line monitoring of protein 13 chromatography. Here, the potential of in-line FTIR as a Process Analyti-14 cal Technology (PAT) in downstream processing was investigated in three case 15 studies addressing the limits of currently applied spectroscopic PAT methods. 16 A first case study exploited the secondary structural differences of monoclonal 17 antibodies (mAbs) and lysozyme to selectively quantify the two proteins with 18 Partial Least Squares Regression (PLS) giving Root Mean Square Errors of 19 Cross Validation (RMSECV) of 2.42 g/l and 1.67 g/l, respectively. The corre-20 sponding Q^2 values are 0.92 and, respectively, 0.99, indicating robust models in 21 the calibration range. Second, a process separating lysozyme and PEGylated 22 lysozyme species was monitored giving an estimate of the PEGylation degree of 23 currently eluting species with RMSECV of 2.35 g/l for lysozyme and 1.24 g/l for 24 PEG with Q^2 of 0.96 and 0.94, respectively. Finally, Triton X-100 was added 25 to a feed of lysozyme as a typical process-related impurity. It was shown that 26 the species could be selectively quantified from the FTIR 3D field without PLS 27 calibration. In summary, the proposed PAT tool has the potential to be used 28 as a versatile option for monitoring protein chromatography. It may help to 29 achieve a more complete implementation of the PAT initiative by mitigating 30 limitations of currently used techniques. 31

³² Keywords: chromatography, proteins, process analytical technology (PAT),

³³ fourier-transform infrared spectroscopy (FTIR), downstream processing

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