

Electronic Supplementary Material

An antibody-based biomarker discovery method by mass spectrometry sequencing of complementarity determining regions

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Figure S1: Alignment of the variable region of the adalimumab light chain sequence against the IgG germline gene with the highest homology in the IMGT database. The letters in bold below the aligned sequences indicate the mutations in adalimumab from the closest related IgG germline sequence. (<http://imgt.cines.fr/3Dstructure-DB/cgi/DomainGapAlign.cgi>)