SUPPLEMENTARY INFORMATION

Evaluation of Back Scatter Interferometry, a method for detecting protein binding in solution

S. T. Jepsen,^a T. M. Jørgensen,^b W. Zong,^c T. Trydal,^a, S. R. Kristensen^a and H. S. Sørensen^c

^a Dept. of Clinical Biochemistry, Aalborg University Hospital, Hobrovej 18-22, 9000 Aalborg, Denmark.

^b Dept. of Applied Mathematics and Computer Science, Technical University of Denmark, Richard Petersens Plads, Building 324, 2800 Kgs. Lyngby, Denmark

^c Dept. of Photonic Engineering, Technical University of Denmark, Frederiksborgvej 399, 4000 Roskilde, Denmark.

Recorded fringe pattern



Figure 1

Figure 1 shows a typical image of the fringepattern as recorded by the linear CCD. Y-axis is intensity in arbitrary units i.e. high intensity equals the bright spots in the interference pattern.

IgG - Protein A Binding experiments

The IgG Protein A binding experiments was also performed with 2.5 nM Protein A corresponding to that used by; *D. J. Bornhop, J. C. Latham, A. Kussrow, D. A. Markov, R. D. Jones, and H. S. Sørensen, Science, 2007,* **317**, 1732–6 and data from one of several experiments are shown below.

We find no evidence in our data of a measuarable change in refractive index between bound and unbound IgG.



Figure 3

Figure 2 shows end-point data of 2.5 nM Protein A incubated with 17-1150 nM IgG (+) and IgG without addition of Protein A (o). Figure 3 shows the signal difference between data with/without Protein A.