Supporting information

Additional Supporting Information may be found in the online version of this article at the publishers web-site:

Figure S1. Characterization of CHO cell vesicles. Representative results for DLS (**A**) and Bradford Assay (**B**) of vesicle solutions made from CHO cells.

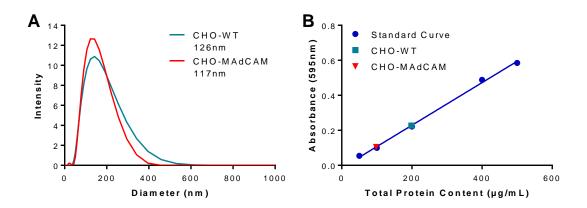


Figure S2. Characterization of human colon tissue VRH. Representative results for DLS (**A**) and Bradford Assay (**B**) of VRH made from colon tissue. Polydispersity index was measured and if greater than 0.28 by DLS, the VRH sample was probe sonicated again on ice for 90sec in a pulsed manner (5sec on, 1sec off), followed by centrifugation and characterization of the sample by DLS and Bradford as described.

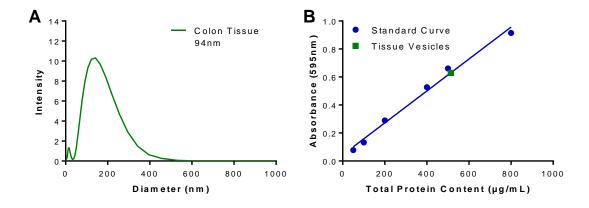


Table S1. Components of the reference and test samples. A pictorial representation of the reference and test samples used in the work-flow.

| Matrix Tastad | Binding Samples | | Control Samples | | |
|--|-----------------|-------------|---------------------|--|------|
| Matrix Tested | Reference | Test | Reference | | Test |
| Soluble F _C Recombinant MAdCAM | ~ | > | | | 100 |
| Recombinant Membrane MAdCAM (CHO cells) | | | | | |
| Native Soluble MAdCAM in Serum | 46 | > | Legend | | |
| | | | | Dilutions done with PBS Dilutions done with serum Dilutions done with homogenate | |
| | | | | | |
| Native Membrane MAdCAM in Tissue | | | | | |
| | | | * | PF-00547659 | |
| Soluble and Membrane MAdCAM in Tissue Soluble MAdCAM in Tissue | 3 | | Isotype Control mAb | | |
| | | | ** | CHO-WT Cell Vesicles | |
| | | | | CHO-MAdCAM Cell Vesicles | |
| | | | | Tissue Vesicles | |
| | | | 6 | MAdCAM Protein | |

Figure S3. Cell based assay protocol diagram. Illustration of the experimental procedure to measure the binding affinity in CHO cell vesicles

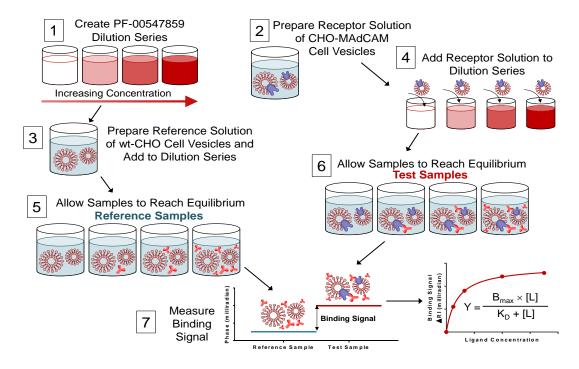
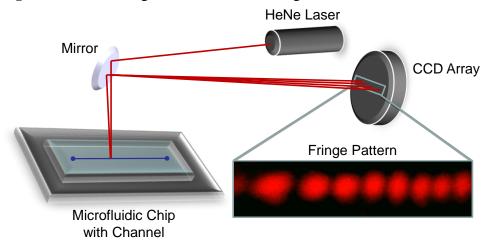


Figure S4. Block diagram of a Backscattering Interferometer.



Measurement of hMAdCAM Concentration in Biological Samples by LC-MS/MS

Briefly, for MAdCAM concentrations the LC system was configured with a WPS-3000 autoinjector and an FLM-3300 flow manager consisting of a temperature controlled column compartment set at 15°C and using 3 valves and 3 pumps: a Loading Pump connected to the antipeptide antibody column, an NCS module supplying the analytical column with nanoflow rates, and Micro Pump washing and equilibrating the reversed-phase trap cartridge. There were three chromatographic separation stages employed: an anti-peptide antibody column, a reverse-phase trap cartridge and a reverse phase column. The loading and elution flow rates on the anti-peptide antibody column typically ranged between 0.3 and 0.6 mL/min, while the flow rate of the analytical column was 600 nL/min. The detection of MAdCAM-derived peptide was performed on an API5500Q mass spectrometer in multiple reaction monitoring (MRM) using a Nanospray III interface (AB Sciex, Toronto, Canada). AB Sciex Analyst software (Version 1.5.2) was used to acquire data and determine peak areas. Peak area ratios (PAR) of the MAdCAM to the internal standard peptide peak area were calculated. Calibration curves were obtained by plotting PAR of the calibration samples versus sample concentration. The curve was fitted with a nonlinear regression applying a 1/concentration weighted model using Labstats (Excel add-in). All sample concentrations were then calculated from their PARs against the calibration curve; a representative calibration curve is shown in Figure S5. The quality of the curve fit was evaluated by back-calculating the MAdCAM concentration of the calibrants. Each assay run was calibrated using an eight or ten point (tissue or serum assay, respectively) standard curve analyzed in duplicates across a MAdCAM quantification range from 2.29 to 5000 pM (tissue assay) and from 2.62 to 10000 pM (serum assay). Additionally, for quality control purposes, human serum and colon tissue homogenate, including their dilutions and spikes with recombinant human MAdCAM.Fc (R & D Systems, MN) were analyzed on multiple testing occasions in replicates (n=3 serum; n=2 tissue). A summary of all the values calculated are shown in Table S2. Coefficients of variation and relative error of the standards must be equal to or less than 20% for a run to be considered acceptable.

Table S2. MAdCAM sample concentrations. Tabulation of sample concentrations measured by LC-MS/MS and used in BSI

| Sample | Concentration by LC-MS/MS (pM) | Dilution to BSI | Concentration in BSI (pM) |
|---------------------|--------------------------------|-----------------|---------------------------|
| Human serum 100% | 999 ± 32 | various | Various |
| CHO-rhMAdCAM VRH | 1719.5 ^a | 50 | 34 |
| Colon tissue VRH | 15 ^a | 349 | 0.043 |
| Colon tissue ILH | 44 ^a | 4 | 11 |

^a no SD was determined due to the rare sample type and availability

Figure S4. Quantification of MAdCAM using LC/MS. Representative results for LC/MS quantification of MAdCAM protein in vesicle solutions made from colon tissue.

