# S&S<sup>®</sup> Serum Biomarker Chip Displays Specificity & **Reproducibility for 120 Different Human Biomarker Profiles**

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Signal Net Spiked Serum

## Abstract

Abstract We introduce a unipote tool for the determination of relative abundances for human serum biomarkers. Conceptually similar to DNA microarrays, the S&S<sup>®</sup> Serum Biomarker Chip (SRC) is a single capture anthody array that was developed for comparative analysis of serum samples in order to identify differences or similarities in protein expression profiles. The SBC array is manufactured on an introdultose two-pad FAST<sup>®</sup> side in which each pad contains 120 antibodies specific for circulating tumor-related proteins printed in triplicate and arranged as four quadrants per pad. The Universal Linkage System (ULS<sup>®</sup>) protein labeling kit was employed to label serum proteins with either biotion or fluorescein prior to incubation with the SBC array. Data from the array was collected using an Axon scanner to detect fluorescent emitted from Dyomics dves D<sup>®</sup>/<sup>®</sup>47 and DV<sup>®</sup>442 conjusted the Netrentofilm and nut-Fluorescein using an Axon scanner to detect fluorescent emitted from Lyomics diges DPM<sup>46</sup>77 and DPM<sup>46</sup>77 conjugated to Streptavidin and anti-Fluorescent antibody respectively. We show that fluorescently labeled biomarker proteins bind specifically to their capture antibody as determined via competition experiments. The system was tested for reproducibility by running eleven individual experiments and measuring the variation in running eleven individual experiments and measuring the variation in the signal intensity of each target biomarker. The results show that the CV is less than 20% for 105 of the 120 capture biomarker antibiodies. We evaluated array sensitivity by spiking FCS with a cocktail consisting of 26 different biomarker antigens at concentrations of 500, 100, and 25 mg/mL in addition, we were able to observe differences in the labeling patterns when securum samples from individuals with breast, colon, prostate and bladder cancer were compared to age and gender matched controls. The SBC array, when coupled with the ULS protein labeling system, provides the research community with a powerful tool to analyze the relative abundances of 120 individual biomarkers in human serum, requiring significantly less time and money as compared to ELISA techniques.

#### Introduction

Introduction The S&S Serum Biomarker Chip is designed to study relative protein abundance in huma serum samples. This product addresses a growing need for technologies that enable broad molecular profiling of biological samples. The ability to identify multiple serum biomarkers simultaneously has many applications in basic biological research. The use of DNA arrays for profiling mRNA expression in cells has provided valuable information in many biological areas. However, since there is not always a direct correlation between the mRNA level and the expression of the protein, a method that can asay multiple proteins is required for an meningful analysis. Antibody chips provide such a solution and can be used to profile abundance of proteins in samples. The S&S Serum Biomarker associated with human disease states of every major organ. The product enables scientisis to erproduct pattern the relative abund ance of 120 human serum proteins between two samples, such as serum samples from diseased to provide gundanabares Chip is not intended to provide gundanabares. The SFAM Serum Biomarker samples from diseased and healthy individuals. The Serum Biomarker Chip is not intended to provide quantitation such as an ELISA method would provide, but rather is a method to monitor or discover protein abundance changes between biological samples on a broad scale.

abumance changes between hological simples of a broad scale. The S&S Serum Biomarker Chip uses antibody microspots to capture hapten-labeled serum proteins followed by readout using fluorescent reporter molecules. The entire procedure, from sample labeling to slide scanning, lakes one 24 hour day to complete since we recommend an overnight incubation. It is intended to be used with human serum proteins labeled with sink were necessary to control for waraitons in labeling efficiency. A hapten swapping mapping the specific fluerences in ULS<sup>M</sup> labeling efficiencies or differences in antibody-antigen bluding interactions caused either by steric hindrance or solubility issues associated with chemically labeled serum proteins. Here, each pad on the slide is probed with two different serum samples labeled with different haptens, and the second pad is probed with the haptens reversed. The normalized intensity for each element of each pad is calculated as the average of the biotin - and fluorescen-labeled derived calculated as the average of the biotin- and fluorescein-labeled derived intensities from a two-pad experiment. This method is attractive for antibody chips as it takes into account any hapten-specific differences in antigen-antibody interactions.

#### Methods & Results

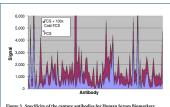
Methods & Kesuits Table 1. The chart below shows all is of the antibody specificity targets present on the Serum Biomarker Chip. Biomarker-specific antibodies were printed on a two-pad S&S FAST's Bide using a Cartesian ProSys 5510 arrayer with SMP3XB pins (TeleChem International, Inc.). Each pad consists of an identical array and each array is comprised of four subarrays. The antibodies were arrayed in triplicate in a proprietary array buffer with approximately Ing of antibody deposited per spot. The antibodies were spotted at a pitch of 0.5mm.

Alpha1 antichymotrypsin	Alpha fetoprotein	Angiogenin	Alpha2 macroglobulin
Apolipopeotein	Angiostatin	Bone sialoprotein	Angiopoietin-2
Beta2 microglobulin	CA125	Cathepsin B	Apolipoprotein J
CA15-3	CA50	Ceruloplasmin	Chorionic gonadotropin-beta
CA19-9	Carcinoembryoinc antigen (gp 2)	C-reactive protein	Epidermal growth factor receptor
Carcinoembryonic antigen (gp 4)	Chorionic gonadotropin- alpha	Cyclin-dependent kinase inhibitor 2A	Fibroblast Growth factor- basic
Chendroitin Sulftate	Cytokeratin Fragment 21-1	E-Selectin	Hepatocyte Growth Factor
Chromogranin	Fas ligand	Estrogen Receptor	IgG
Collagen Type I	Haptoglobulin	FAS	IL-13
Complement C4	lgA	Ferritin	IL-17
Eotanin	lgG	Fibroblast Growth factor-7	IL-2 receptor-alpha
Epidermal growth factor	lgM	GM-CSF	IL-8
ErbB2	IL-12p40	løG	Insulin growth factor binding protein 3
G-CSF	IL-1-alpha	IL-10	Interferon-gamma
Hemoglobin	IL1-beta	IL-2	Kallikrein-9
ICAM-1	IL-3	Insulin-like growth factor 1	Myeloperonidase
1gG	IL-4	IP-10	Neuron Specific Enolase
IL12-p70	IL-6	Laninin	PDGF (all isoforms)
IL-2 receptor-beta	Insulin	MCP-1	PDGF (BB isoform only)
IL-5	Kallikrein-14	MCP-2	Placental Alkaline Phosphatase
IL-7	Kallikrein-5	M-CSF	plasminogen activator inhibitor
Kallikrein-12	Lose-density lipoprotein	MIP-1-alpha	Prostatic acid phosphatase
MCP-3	MCP-4	Myoglobin	\$100
MMP-3	MMP-2	PSA (Free)	Serum Albumin
Plasminogen	MMP-9	PSA-Alpha-chemotrypsin complex	Tetranectin
RANTES	Osteopontin	Serum Albumin	Thrombopoietin
Serum Alburnin	PSA (Total)	TGF-alpha	Thrombospondin-1
TAG-72	Serum Albumin	Tyrosinase	TNF-alpha
TGF-beta	Sialyl Lewis X	Urokinase Plasminogen Activator	Tumor-Associated Trypsin Inhibitor
Thyroglobulin	TIMPI	VCAM-1	VEGF-D
TIMP2	TNF-beta	VE-Cadherin	Von Willebrand Factor
VEGF	Transferrin		

Figure 1. Direct labeling of human serum proteins using the Universal Labeling System -detection of Biomarkers spaked into Human Serum. The graph demonstrates that the USP protein Licking negative accessible biodic sequences proteins dynamic strategies and methanism. Two proteins were used for this spaking experiment, epidermal growth factor (EG) and protein sequelic angle of SNA. Differentiation and proteins and the distinct form a horizon sequelic angle of SNA. The Constantian sing methanism. USD strategies and the strategies are also been apprecised by the strategies and and and concentration = 200 angleti). The reaction was processed to remove une reacted ULS should and subsequently included and an SAST dide apprecision the appropriate cop-ture antibuler. The taidet was stated and incubated with respersion copy biologic.

Figure 2. SVPRO' Roby staining of the S&S Serum Biomarker Chip. Serum Biomarker Chips were treated with the protein stain SYPRO' Roby according to the manufacturers instructions. The average signal intensity for each antibody triplicate on both the top and bottom pad were polent and the regression line drawn (Ferson). The side was examed with the Axon 4200A scamer using the blue laser at A&Bs and a FMT setting o '00 mol laser rower of 95%. Data analysis was completed by using Array Vision FSLT.

Reproducibility of the array compares both pads of the slide: Tot y SYPRO\* Ruby demonstrates that nearly identical quantities of an a both the top and bottom pad of each two pad S&S Serum Bioman



alf Serum: Four microarce-biotin. These labeled proteins or r Chip either in the presence or CS. The results clearly show th d FCS proteins suggesting that r who woult of non-specific with ULS Biomarke "cold") F he labele at cair serum (FCS) was tabeted to 7 / n e incubated for 16 hours on the Serum sence of a 100-fold excess of unlabeled the cold FCS failed to inhibit the bindin ch of the binding observed under these reactions of bovine proteins with the

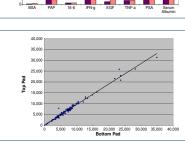
Figure 4. Average signal intensity and %CV of protein expression ratios from multipl Stram Biomarker Chipe. Elsevin individual labeling reactions were performed on the strand strand strand strand strand strand strand strand strand good mundal ogication. These labeling motions were used to prove the SoS Stramo Biomarker Chip, and the relative protein expression ratio for each of the 120 specificiti was determined. The binding raction was detected using both threshold strand Stramer, Spot finding background corrections, and two-channel data sampling was eccompliable with Array Visuar YAST software. The SDC analysis weakbook is a Microsoft. 2006 flate coveres thareaccut data into humer of solves that represent the Microsoft. 2006 flate coveres thareaccut data into microsoft solves that represent Microson Exzer hie that converts morescent data into humercal values that represent the abundance of antigen in Sample A relative to antigen in Sample B. Average signal intensity was determined by taking the average of the sum of the Dy547 signal and the Dy647 signal for each triplicate on the array.

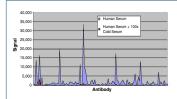
Two **Color Reproducibility**: The WCV of the protein expression ratios for 105 of the 120 specificities on the S88 Serum Biomarker chip is less than 20%. For those antibo-where the %CV is higher than 20% the total signal intensity from both Dy547 and Dy647 is often very low suggesting that these proteins are not present in detectable quantities in the serum sample sterded.

igner 5. Limits of detection and Sensitivity of the Capture Antibodies on the erram Biomarker Chip. A cocktail of 26 different purchised human antigens were pilot into tical cal calcum (FK) at three different concentrations 50,000 and 25 grafin final concentration. Four microliters of this spiked ICS was compared to a nyclate ICS using the two-color system of artipexiding IVS and anti-fluoresci-metry for a borner spiked arcs multiple were labeled with historin-ULS and homescien-ULS for hours in pald A, while the dye way combination was included and homescine-ULS hours in pald A, while the dye way combination was included and historin-ULS and un-spiked arcmit habeled in the hours in part A, while the dye way combination was included in the hourin (LS). A while the dye way combination was included which hourin-ULS) and un-spiked arcmit habeled which hourin-ULS. The interpret of the tice of the neeroge spikel animetary of the graft Chi Cas and the Cas Arcerage signal intensity of the centrel in the dye way combined and the spikel dye the spikel of the spikel Cas and the spikel Cas. Arcerage signal intensity of the centrel in the dye way combined as a spikel arcmit habeled at the spikel Cas and the spikel Cas. Arcerage signal intensity of the centrel in the dye and the spikel arcmit habeled at the spikel Cas and the spikel arcmit habeled at the spikel arcmit habeled at the spikel arcmit and the spikel arcmit habeled at the spikel arcmit and the spikel arcmit and the spikel arcmit and the spikel arcmit arcmit are arcmit at the spikel arcmit and the spikel arcmit and

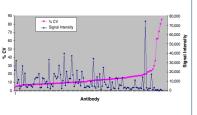
The S&S Serum Biomarker Chip detects purified human antigens spiked into a cor plex protein mixture (FCS). The greater the amount of protein spiked into the FCS the higher the observed ratio of that protein relative to the un-spiked FCS. As the concentration of spiked protein was decreased, the observed ratios decreased as we

Figure 5. Li Serum Bion spiked into f ng/ml final c





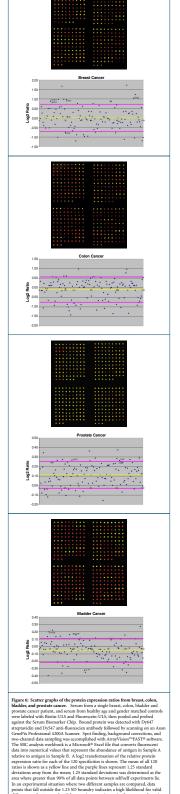
3b. Human Ser -so. somma serum: Four mecotiters of human sera was labeled with ULS-biotin for hours. These labeled proteins were inclusabled for 16 hours on the Serum Biomarker Chip either in the presence or absence of a 100-664 eccess of unlabeled "Codd" hume serum. Unlabeled human serum inhibited nearly all the signal at all the spots on the array. This result demonstrates that the binding of human process to the Serum Nomarke chip is principally the result of specific binding of the antigens with ther array. The serue of the se



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### Summary

- · The S&S Serum Biomarker Chip contains 120 antibodies to human serum biomarkers reproducibly printed in triplicate.
- Using the ULS-labeling system the %CV from multiple experiments was less the 20% for the majority of the elements on the S&S Serum Biomarker Chip
- The S&S Serum Biomarker Chip shows that human antigens bind specifically to the capture antibodies immobilized on the chip's surface.
- Purified human proteins spiked into non-human serum (FCS) were captured and detected by the Serum Biomarker Chip in a dose-dependent manner.
- The S&S Serum Biomarker Chip demonstrated differences in the relative protein abundances between serum samples from normal and cancer patients.



fall outside the 1.25 SD boundry indicates a high likelihood 1 scatter graphs of the protein expression ratios from serum neer patients clearly show that the S&S Serum Biomarker (Ch ences in relative protein abundance between serum samples, cates that equal quantities of a particular protein are present use greater than a perior indicate that the protein is more abund in serum while a value of loss than zero indicates that the second second second second second second second at serum while a value of loss than zero indicates that the second s

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