

NIH Public Access

Author Manuscript

J Biomed Mater Res A. Author manuscript; available in PMC 2013 October 21.

Published in final edited form as:

J Biomed Mater Res A. 2009 February ; 88(2): 503-519. doi:10.1002/jbm.a.31886.

Adsorbed serum albumin is permissive to macrophage attachment to perfluorocarbon polymer surfaces in culture

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Abstract

Monocyte/macrophage adhesion to biomaterials, correlated with foreign body response, occurs through protein-mediated surface interactions. Albumin-selective perfluorocarbon (FC) biomaterials are generally poorly cell-conducive due to insufficient receptor-mediated surface interactions, but macrophages bind to albumin-coated substrates and also *preferentially* to highly hydrophobic fluorinated surfaces.

Bone marrow macrophages (BMMO) and IC-21, RAW 264.7 and J774A.1 monocyte/macrophage cells were cultured on FC surfaces. Protein deposition onto two distinct FC surfaces from complex and single-component solutions was tracked using fluorescence and time-of-flight secondary ion mass spectrometry (ToF-SIMS) methods. Cell adhesion and growth on protein pre-treated substrates were compared by light microscopy. Flow cytometry and integrin-directed antibody receptor blocking assessed integrins critical for monocyte/macrophage adhesion *in vitro*.

Albumin predominantly adsorbs onto both FC surfaces from 10% serum. In cultures pre-adsorbed with albumin or serum-dilutions, BMMO responded similar to IC-21 at early time points. Compared to Teflon[®] AF, plasma-polymerized FC was less permissive to extended cell proliferation. The ₂ integrins play major roles in macrophage adhesion to FC surfaces: antibody blocking significantly disrupted cell adhesion. Albumin-mediated cell adhesion mechanisms to FC surfaces could not be clarified. Primary BMMO and secondary IC-21 macrophages behave similarly on FC surfaces, regardless of pre-adsorbed protein biasing, with respect to adhesion, cell morphology, motility and proliferation.

Keywords

perfluorocarbon; foreign body reaction; integrin blocking; cell line; serum proteins

INTRODUCTION

Macrophages are key mediators of host inflammatory responses to surgically placed biomedical devices, implicated in an abnormal wound healing response commonly observed around implanted materials known as "foreign body reactions" (FBR).^{1–3} Macrophage cells

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represent a highly adaptable, dynamic cell population with the ability to respond to activating cues of both chemical and physical nature present in their local environment. Foreign body giant cells, the fusion product of monocytes and macrophages and one cellular hallmark of the FBR to implantable materials,⁴ are frequently found at surfaces of biomaterial implants, and material surface chemistries have been shown to influence fusion.⁵ Modulating events that facilitate progression of the FBR at implant sites remains of substantial interest toward improved biomedical device performance and host integration. To date, little *in vivo* control over this complex reaction to implanted materials has been demonstrated.

This study focused on well-known influences on cell adhesion by surface chemistry and the adsorbed protein layer.^{5–7} *In vivo*, biomaterials surfaces are instantly and continuously bombarded with thousands of different host proteins,^{8–12} and surface chemistry is recognized to exert an influence on adsorbed protein composition, exchange dynamics and structural conformations.^{13–15} *In vitro*, surface-denatured proteins lead to increased monocyte adhesion,¹⁶ the first step in a sequence of events that may ultimately result in a FBR *in vivo*. As the predominant plasma protein,^{8,9} albumin's surface interaction has been studied extensively^{17–26} due to albumin's propensity to reduce many biological interactions with surfaces, including reduced thrombogenicity of polymers,¹⁸ decreased bacterial adhesion and device-centered infections,¹⁸ reduced platelet adhesion^{27–29} and limited cell adhesion.³⁰

Albumin has been shown to bind strongly and selectively, to perfluorinated (i.e., plasma deposited and poly(tetrafluoroethylene) or PTFE) surfaces,²⁴ out-competing cell-adhesive extracellular matrix (ECM) proteins like fibronectin in adsorption from binary solutions and serum dilutions, even when fibronectin bulk concentration is biased 10-100 times higher than found physiologically.³¹ Albumin apparently "masks" adsorbed fibronectin, and low rates of endothelial cell adhesion have been correlated to a limiting adsorbed fibronectin surface density as probed by radiolabeling and anti-fibronectin antibodies.³¹ In fact, poor cell adhesion to perfluorinated surfaces has been reported for multiple different cell types. ^{32–36} Thus, increased density of surface-adsorbed albumin on these surfaces correlates to reduced mammalian cell adhesion³⁰ for numerous cell types with important consequences in vivo. Surfaces that "select" albumin from complex biological milieu, such as perfluorocarbons,^{24,31} are predicted to be poorly supportive of cellular adhesion given that most cells are not known to have specific receptors to actively engage albumin in adhesion. Notably, hepatocytes, ³⁷ vascular endothelial cells, ³⁸ and monocytes, macrophage and dendritic cells^{39,40} have been reported to be exceptions. For example, the FcRN receptor, identified as an albumin-binding protein present on monocyte, macrophage and dendritic cells,⁴⁰ binds exclusively at pHs below 6.5 but is not a known cell-adhesive receptor physiologically.³⁹ Hence, in the context of the FBR, highly albumin-adsorbed perfluorinated surfaces might be predicted to elicit reduced inflammatory cell activation if cell-surface adhesion is inhibited. Surprisingly, in vitro cultures of murine primary- and secondaryderived monocyte-macrophage cells have been shown to adhere, grow and proliferate proficiently on hydrophobic (i.e., polystyrene) and perfluorocarbon (e.g., Teflon® AF, plasma-polymerized FC, fluorinated ethylene propylene) substrates.^{41–44}

This study sought to 1) examine and compare aspects of cell adhesion behavior for murine cells of primary- (BMMO) and secondary- (IC-21, J774A.1, RAW 264.7) origins when grown on perfluorinated (i.e., fluorocarbon, FC) surfaces *in vitro*, and 2) to propose a mechanism by which initial monocyte/macrophage cell adhesion events occur on FC surfaces in the context of establishing a FBR to biomaterials. For practical purposes and to allow an appropriate equivalence comparison between primary and secondary derived macrophages, this study focused on responses of BMMO and IC-21 cells exclusively. Of the

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three secondary cell lines initially tested, IC-21 was selected as a best match for comparison to primary BMMO because: 1) IC-21 represents the most physiologically relevant cell line within this group,⁴⁵ and 2) mature IC-21 cell size, morphology, adhesion and spreading behaviors were shown to be very similar to BMMO. To contrast (monocyte-) macrophage cell behavior on FC surfaces, NIH 3T3 fibroblasts, an attachment-dependent cell type but non-adhesive to fluorocarbon surfaces *in vitro*,^{36,44} was used as a control. The ₂ integrin family, vital to development of the inflammatory response *in vivo*⁴⁶ and known to interact with specific ECM proteins as well as denatured proteins at surfaces,¹⁶ was hypothesized to be the primary mediator of any observed initial macrophage adhesion to FC surfaces exposed to biological milieu *in vitro*. Further, as macrophages are known to secrete fibronectin,^{47–50} this was also studied as it may facilitate monocyte and macrophage adhesion to FC surfaces via endogenous ECM production, deposition and cell-based remodeling of the pre-existing adsorbed protein layer.

MATERIALS AND METHODS

Culture surface preparation for Time-of-Flight Secondary Ion Mass Spectrometry Analysis

Polystyrene (PS) dishes were cut into pieces (~ $1 \text{ cm} \times 1 \text{ cm}$), labeled to indicate "bottom" and affixed to the bottom of PS petri dishes using double-sided tape, or left unfixed to freefloat in protein milieu. All Teflon® AF culture surfaces were prepared by evaporation deposition; fluorocarbon content was subsequently confirmed via x-ray photoelectron spectroscopy (XPS) analysis as described previously.³² Briefly, sufficient Teflon[®] AF (DuPont Fluoroproducts) solution (0.1 wt.% in FC-40 solvent, 3M Corp.) was added to these petri dishes to completely cover all fixed PS pieces. Coated PS surfaces were placed in a vacuum oven overnight at 65°C to remove residual solvent, and plates were misted with 70% ethanol, dried, and treated with biosafety cabinet UV light for 20 minutes immediately before culture, a process shown to sterilize while remaining benign to cell culture surface chemistry.⁴⁴ Teflon[®] AF samples were also reserved as controls. Following sterilization, samples were immersed in either a 3 mg/ml boyine serum albumin (BSA) solution (fraction V, OmniPur®, Sigma, 98% pure by gel electrophoresis) or 10% fetal bovine serum (FBS) solution, both in sterile Dulbecco's phosphate buffered saline with Ca^{2+} and Mg^{2+} (PBS⁺⁺), and incubated at 37°C for 24 hours. Samples were removed, rinsed twice with PBS++, three times with 18 M "Nanopure-grade" ASTM I water and dried under a stream of nitrogen. These samples were subjected to surface analysis as described below.

Preparation of fluoropolymer culture surfaces using plasma deposition

All plasma-prepared fluorocarbon (pp-FC) surfaces employed in these studies were gifts of Dr. E. Fisher and Dr. G. Malkov (Colorado State University, USA), prepared and characterized as previously described.⁴⁴ This method produces a robust (100 nm) thick film of crosslinked perfluorocarbon highly enriched in both $-CF_2$ and $-CF_3$ groups. Briefly, all films were deposited in a home-built inductively coupled radio frequency (RF) (13.56 MHz) plasma reactor.⁵¹ Pulse duty cycle was varied using the internal pulse generator of an RF power supply. Peak applied RF pulse power (*P*) was kept constant at 300W for FC films. For all experiments, C_3F_8 (Air Products, 99%) gas flow was kept constant at 10.0 sccm (standard cubic centimeters per minute), resulting in a reactor pressure of ~200 mTorr.

Time-of-Flight Secondary Ion Mass Spectrometry (ToF-SIMS) analysis

A Physical Electronics 7200 instrument with an 8 keV Cs⁺ ion source, a reflectron time-offlight mass analyzer, chevron-type multi-channel plates, and a time-to-digital converter was used for data acquisition. Data was acquired for the Teflon[®] AF samples after incubation (as described above) with 3 mg/ml BSA or 10% FBS; other protein spectra for comparison were collected previously.^{21,52,53} Positive secondary ions mass spectra were acquired over a mass

range from m/z = 0 to 1000. Negative ion ToF-SIMS spectra were not considered here due to a lack of unique peaks for the different amino acids.⁵⁴ Each spectrum analyzed an area of 100 µm × 100 µm, and the total ion dose used to acquire each spectrum was less than 1 × 10¹² ions/cm². Mass resolution (m/m) of the secondary ion peaks in the positive spectra was typically between 3000 and 5000. The ion beam was moved to a different spot on the sample for each acquisition. Positive spectra were calibrated to the CH₃⁺, C₂H₃⁺, C₃H₅⁺, and C₅H₁₀N⁺ peaks before any further analysis. At least two replicates were prepared for each sample type, with three spectra acquired from each replicate.

Principal Components Analysis (PCA) of ToF-SIMS data

PCA was used to analyze the positive ToF-SIMS data from protein-adsorbed surfaces using scripts written at NESAC/BIO (University of Washington, USA) for MATLAB (The MathWorks, Inc.). All spectral processing (peak selection, normalization, PCA, etc.) was done as described previously.²¹ Briefly, each amino acid was assigned a distinct mass peak, and the abundance of each peak, or amino acid, respectively, was used as input for the dataset. PCA yields an output of both a "scores" and a "loadings" plot.⁵⁵ The scores plot shows the relationship among samples and the loadings plot shows the relationship between the new principal components (PCs) and the original ToF-SIMS peaks. Wagner *et al*²¹ demonstrated that proteins present at a surface can be identified through unique amino acid fragmentation patterns in the ToF-SIMS positive ion spectra. Here, the ToF-SIMS data from BSA and 10% FBS adsorbed onto Teflon[®] AF surfaces was projected into the data set of Wagner *et al*²¹. In addition to PCA, other multivariate analysis methods such as discriminant analysis and neural networks have been used to classify different absorbed protein films.^{53,56}

Fluorescence labeling of proteins for detection of surface adsorption

Fibronectin (from bovine plasma, Sigma) and BSA solutions were fluorescently labeled and subsequently purified per manufacturer's instructions using Alexa Fluor 555[®] and Alexa Fluor 647[®] protein labeling kits (Invitrogen). Protein and dye concentrations were determined by optical density using UV-vis spectroscopy. Subsequently, samples of pp-FC, Teflon[®] AF, negative control CodelinkTM activated hydrogel-coated slides (Amersham Biosciences) or glass coverslips (positive control) were either completely covered with dyelabeled protein solution (depending on the substrate size: small samples were completely immersed, while wax pencil "wells" were drawn on larger slides), or spotted with 30 µl drops (confined within a wax pencil-defined well). Samples were exposed to: 1) single component solutions of fluorescently labeled albumin or fibronectin, or 2) a mixture of both dye-labeled proteins (each with a different fluorescent label) in a range of sample concentrations (albumin, from 1.5 µg/ml to 1.5 mg/ml; fibronectin, from 1.9 µg/ml to 1.9 mg/ml) or 3) a mixture of both proteins with only one dye-labeled protein in the same range of sample concentrations (vida supra). Coverslips were transferred to petri dishes and incubated at 37°C, 98% humidity for 24 hours. No differences in 30 µl spot sizes were discernable after the 24 hour incubation period (i.e., negligible evaporation) when the remaining solution was rinsed from the surface using PBS⁺⁺, further rinsed copiously with PBS⁺⁺, dried under nitrogen and affixed to glass slides using double-sided tape (edges only). Samples were scanned for fluorescence signal using a Perkin Elmer ScanArray Express[™] Microarray Scanner employing appropriate optical filters and wavelengths for each dye employed. Gain and power settings were consistently controlled for sample comparisons. Images were processed for relative fluorescence intensity using ImagePro[™] software, and quantified using Quantity One[®] (Bio-Rad) and MS[™] Excel software.

Cell culture

All cell lines were obtained from the American Type Culture Collection (ATCC, Manassas, VA). Cells were cultured in Dulbecco's modified Eagle's medium (DMEM, Mediatech, for J774A.1 and NIH 3T3) or RPMI 1640 (Mediatech, for IC-21 and RAW 264.7) supplemented with 10% FBS (HyClone, Inc.), and 1% penicillin-streptomycin (Life Technologies). Cultures were maintained in T-175 tissue culture polystyrene (TCPS) flasks (NuncTM) under standard conditions: incubation at 37° C, 98% humidity and 5% CO₂. Cells were dissociated from culture flasks by incubation with Ca²⁺- and Mg²⁺- free cell culture grade HBSS (Life Technologies) (NIH 3T3), or by scraping with a rubber policeman (IC-21). Cell concentration and viability was assessed using standard trypan blue (BioWhittaker) dye exclusion assay and a hemacytometer.⁵⁷ All cell line subcultures were 25 beyond the passage number as received from ATCC.

Primary cell harvest

BMMO were prepared from bone marrow cells harvested from the femurs and tibias of C57BL/6 mice.⁵⁸ To differentiate bone marrow stromal precursors into macrophages, bone marrow-extracted cells were cultured in "complete" bone marrow medium: DMEM supplemented with 10% L929 fibroblast-conditioned medium, 2 mM L-glutamine, 0.01% HEPES, 1% penicillin-streptomycin, and 2 mM non-essential amino acids (Sigma-Aldrich). Cells were grown under standard conditions (*vida supra*) with media changes every two days. This method has been shown to reliably produce differentiated macrophages.⁵⁸

Cell culture on model surfaces

TCPS (Falcon®, Becton Dickinson) and suspension culture PS (Corning Inc.) 15×100 mm dishes and 24-well plates were utilized for both control and experimental conditions. Teflon[®] AF fluoropolymer culture surfaces were prepared by coating PS surfaces as described (*vida supra*). Plates were tested for the presence of contaminating endotoxin using a PyrogeneTM Assay kit (Cambrex), and endotoxin levels were determined to be below the kit detection limit (0.02 EU/ml). Plates were subsequently preconditioned with appropriate medium containing 10% FBS for a minimum of six hours before cell seeding unless otherwise indicated. Cells were seeded at concentrations ranging from 5.0×10^4 to 3×10^6 cells per plate in fresh medium unless otherwise indicated. Initial seeding densities varied slightly for each cell type, due to surface-dependent differences in cell adhesion and growth rates, and to create roughly equivalent cell culture density time endpoints whenever possible.

Protein pre-conditioning and cell culture on select surfaces

FC surfaces were incubated at 37°C for 24 hours with one of the following solutions: PBS⁺⁺, 3 mg/ml BSA in PBS⁺⁺, 0.3 mg/ml fibrinogen (bovine fraction 1, 75% clottable, ICN Biomedical), 10% FBS in PBS⁺⁺, 10% heat inactivated (HI, 56°C, 1 hour) FBS in PBS⁺⁺, 100% FBS or (cell specific) medium containing 10% FBS. At 24 hours, the protein solution or serum was removed by aspiration and cells were immediately seeded in an appropriate cell culture medium containing 10% FBS. Cell culture conditions proceeded as described above.

Phase contrast microscopy

Images of cells on surfaces were obtained with either a Nikon Eclipse TS100 or a Nikon TMS inverted microscope using Nikon objectives. A Kodak DC290 camera was used to capture field images that were subsequently processed using Adobe Photoshop 6.0 (Adobe Systems, Inc.).

Flow cytometry

Cells were pelleted by centrifugation $(300 \times g, 5 \text{ minutes})$ and resuspended in 100 µl of staining solution (1% FBS, 0.01% NaN₃ in PBS⁺⁺) with 1 µg of anti-CD16/32 (clone 93) mAb (eBioscience, F_c receptor block). Cell F_c receptors were blocked for 15 minutes at 4°C. Subsequently, cells were rinsed twice and resuspended in staining solution without antibodies. Cells were transferred to a 96-well plate for subsequent staining with 0.1 µg/ml of one of the following fluorescently conjugated (allophycocyanin, APC or fluorescein isothiocyanate, FITC) mAb per experiment: anti-CD11b (clone M1/70, APC), anti-IgG2a (clone eBR2a, FITC), anti-CD18 (clone M18/2, FITC), all eBioscience). Cells were stained for 30 minutes at 4°C, rinsed twice and resuspended in 500 µl staining solution without antibodies. Cell suspensions were analyzed on a FACSCaliburTM flow cytometer (BD) and data analysis was performed using Dako Cytomation Summit v 4.0 software.

Integrin blocking studies

Function-blocking monoclonal antibodies (mAb, all sterile-filtered, azide-free, low endotoxin) directed against the murine integrin 2 (M18/2) chain and a non-specific isotypematched control (Rat IgG2a,) were purchased from eBioscience. BMMO (matured 7 days) and IC-21 cells were pre-incubated for 30 minutes at 4°C with either an integrin-directed mAb or an appropriate isotype control (both 100 µg/ml). Cells were seeded into Teflon[®] AF-coated wells (2 cm²) previously treated for 24 hours with 10% FBS in PBS⁺⁺ at a density of 500 cells/mm² and allowed to adhere for 1 hour. After the 1 hour incubation period, non-adherent cells were removed with two 0.65 ml washes of warm (37°C) PBS⁺⁺. Cells were fixed and stained according to the Wright-Giemsa method⁵⁹ using a commercially available kit (Hema 3[®] Staining System, Fisher). For each sample, adhesion blocking was evaluated by counting the number of cells in three lower-power fields and expressing the result as a percentage of untreated cells (Teflon[®] AF control). For each cell type, 4 independent experiments were performed, and data presented is the average of the 4 trials. Error was reported as standard error of the mean.

Reverse transcriptase-polymerase chain reaction (RT-PCR)

Total RNA from BMMO and IC-21 cells grown on either TCPS or Teflon[®] AF surfaces was extracted and purified at various time points using RNeasyTM kits (Qiagen) per the manufacturer's instructions. First strand cDNA was synthesized from up to 4 µg of total RNA using Superscript IITM reverse transcriptase (Invitrogen) as recommended by the supplier. Both poly dT (Invitrogen) and murine fibronectin-specific primers⁶⁰ (5 - AGCAGTGGGAACGGACCTAC-3, 5 -GTAGGACGTCCCAGCAGC-3, IDT), 100 pmol per reaction, were used to obtain total and fibronectin-specific product cDNA, respectively. Primer-specific PCR amplifications of glyceraldehyde-3-phosphate dehydrogenase (GAPDH) served as "housekeeping" controls for each sample (5 -

AACTTTGGCATTGTGGAAGGGCTC-3, 5 -TGGAAGAGTGGGAGTTGCTGTTGA-3). Primers were either designed using PrimerquestTM software (GAPDH) or as described previously (fibronectin).⁶⁰ All PCR amplifications were performed using an iTaqTM DNA polymerase kit (Bio-Rad) per manufacturer's instructions, on an iCycler thermal cycler (Bio-Rad). Each experiment was performed with 2 separate RNA isolations. PCR products were analyzed on 2% agarose gels, stained with ethidium bromide, and visualized/analyzed using a ChemiDoc XRS system (Bio-Rad) and Quantity One[®] software (Bio-Rad).

RESULTS

ToF-SIMS and PCA indicate albumin is the major protein component adsorbed to Teflon[®] AF substrates from 10% serum or pure albumin solutions

ToF-SIMS and PCA methods previously described have been frequently employed to classify and distinguish surface adsorbed proteins.^{21,52–54} ToF-SIMS data are challenging to interpret due to the highly energetic SIMS ion fragmentation process that produces extensive fragmentation from surface species generating hundreds of peaks in the spectra from a sampling depth of approximately 1-2nm.⁶¹ To facilitate protein identification under these experimental conditions, each amino acid can be assigned one or more characteristic mass peaks; a standard table of mass peaks assigned to specific amino acids has been established.^{21,54}

PCA methods can be used to identify trends in large, complex data sets⁶¹ and have been useful for identifying proteins adsorbed to surfaces,^{21,54} including BSA, fibrinogen, fibronectin, IgG and others (Figure 1A). PCA's mathematical transformations relate a large number of (potentially) correlated variables into a smaller number of uncorrelated variables, i.e. the principal components (PCs). The first PC (x axis) is the PC that accounts for of the largest amount of variability in the data set (for cases reported here, 51%). The second PC (y axis) accounts for the next largest amount of variability in the dataset (here, 17%). In Wagner's model,²¹ as presented here, this allows for qualitative statements to be made about the presence (or absence) of specific protein species on the tested surfaces, the intent of our use of this method here.

The PCA model of Wagner et al²¹ was utilized for analysis of new ToF-SIMS protein data of interest (e.g., albumin) on Teflon[®] AF surfaces pre-treated with either complex (10% FBS) or single component (BSA) protein solutions. Data obtained for BSA were compared to results obtained by Wagner et al^{21} (Figure 1A) for numerous single proteins adsorbed onto a PTFE surface. In this model the ellipses represent the 95% confidence interval for each protein, and most of the proteins (differing in amino acid composition) can be distinguished based on the two principal components shown (Figure 1A). Comparison supports the assertion that BSA present on Teflon[®] AF surfaces exposed to a singlecomponent BSA solution (indicated as $BSA_{Teflon AF}$) is highly similar to BSA results for PTFE and BSA on other surfaces (Figure 1B). In fact, data for all BSA samples shown (0.1 mg/ml BSA solution on mica,²¹ silica (SiO₂),⁵³ and plasma-polymerized poly(*N*-isopropyl acrylamide), (ppNIPAM),⁴⁹ are within the 95% confidence limits of BSA on Teflon[®] AF. In contrast, the 10% FBS (multi-component protein solution) on Teflon[®] AF (FBS_{10%}) ellipse showed a slightly different localization from the BSA samples (Figure 1B). By examining the first principal component (51% of all the variance in the dataset) exclusively, colocalization of the $FBS_{10\%}$ with the $BSA_{Teflon AF}$ and other BSA samples is noted. Examining the second principal component (17% of the overall variance) exclusively, reveals a higher value for the FBS_{10%} compared to the BSA samples. Thus, the analysis requires analysis of the second PC to find differences between the BSA_{Teflon} and FBS_{10%} surfaces.

To evaluate differences between PCA groupings of proteins, a two-dimensional distance between the centers of each protein grouping shown in Figure 1 was calculated. The average normalized distance of BSA_{Teflon AF} to BSA groupings on other substrates was 0.016 ± 0.006 , while the distance between BSA_{Teflon AF} to other proteins was 0.06 ± 0.03 . With the exception of transferrin, all other proteins had distances from BSA_{Teflon AF} that were greater than 0.04 -- four standard deviations higher than the average BSA distance. Thus, the distance between BSA_{Teflon AF} and BSA adsorbed onto other substrates was significantly smaller than that between BSA_{Teflon AF} and all other proteins examined except transferrin.

The FBS_{10%} distance from the BSA samples was 0.05 ± 0.01 , while the average normalized distance of FBS_{10%} to the remaining proteins was 0.08 ± 0.04 . Although this analysis cannot distinguish between proteins *per se* (e.g., transferrin can not be separated from BSA using just the first two PCs), the location of the FBS_{10%} relative to BSA and the other proteins indicates that BSA from the FBS solution is a major component of protein film adsorbed onto the Teflon[®] AF surface from this complex solution.

Fluorescence scanning detection of surface-adsorbed albumin and fibronectin

Results of fluorescence mapping experiments of protein adsorption to various surfaces are shown in Figure 2. Results presented are for equivalent protein concentrations and fluorescence labels by species across all experimental trials (i.e., 4 μ g/ml fibronectin-Alexa Fluor 555[®] and 37 μ g/ml albumin-Alexa Fluor 647[®]). Significantly, this fibronectin concentration is roughly equivalent to that in 10% FBS (i.e., 10% physiological concentration), whereas the albumin concentration is considerably less than encountered in either circumstance (i.e., 1% of that in 10% FBS, 0.1% of physiological concentration) due to fluorescence signal saturation with albumin at physiological concentration and in 1:10 or 1:100 dilutions.

The highly hydrophilic Codelink[™] hydrogel control surfaces showed minimal protein adsorption (background levels) compared to both glass and highly hydrophobic FC surfaces (Figure 2). As a hydrated, uncharged polyacrylamide-based coating, this commercial microarraying surface is designed to repel serum proteins.⁶² Protein adsorption was not significantly different on the Codelink[™] surface based on the protein species (albumin versus fibronectin, Figure 2 insert), very close to background fluorescence (unlabeled protein, Figure 2 insert), reflecting minimal protein adsorption detected on the Codelink[™] surface. Fluorescence intensity assays indicate that albumin inundates FC surfaces, even well below physiological (e.g., at 1/1000th) or standard cell culture condition (1/100th) levels of protein. Fibronectin is barely detectable on FC samples exposed to binary solutions of fibronectin and albumin (data not shown), consistent with previous reports.^{17,19,31,32} Collectively, results of PCA and fluorescence scanning experiments indicate that monocytes/macrophages must *initially* encounter and interact with an albumin-rich FC surface *in vitro* in culture.

Cell adhesion, growth and proliferation on control and FC substrates

Adhesion, growth and proliferation of (monocyte-) macrophage and fibroblast immortalized cell lines was compared with primary-derived murine BMMO cultures on FC (Teflon[®] AF and pp-FC) using light microscopy techniques. Teflon[®] AF has been employed in previous studies employing monocyte/macrophage,^{43,44,63} neuronal⁶⁴ and endothelial⁶⁵ cells due to its solubility and nearly 100% light transmission, allowing facile microscopic evaluation of cell cultures. Based on our preliminary findings related to cell adhesive behavior,^{43,44,63} both FC surface chemistries utilized were predicted to yield similar protein adsorption and cell adhesive patterns, regardless of macrophage cell origin.

Cells were tracked live and after fixation/staining using the Wright-Giemsa technique.⁵⁹ As expected, all cells efficiently colonized TCPS control surfaces (Figure 3A, C, E, G, I). On FC surfaces, all (monocyte-) macrophage cells tested also achieved proficient growth and proliferation, although cell adhesion was less than 100% (Figure 3B, D, F and H, spherical cell morphologies are non-adherent). Adherent cells were not easily removed by rinsing and exhibited motile phenotypes, often displaying lengthy filopodia (up to hundreds of microns from the cell body) which appeared to be "probing" the surface in search of adhesive sites.⁶³ BMMO (Figure 4) and IC-21⁶³ cell populations on FC surfaces showed a mixture of actin-

based cytoskeletal features (e.g., filopodia, membrane ruffling, Figure 4D and F), often with numerous features on a single cell.

A temporal series of phase contrast photomicrographs for BMMO cell adhesion, growth and proliferation on pp-FC surfaces pre-conditioned with 3 mg/ml BSA, 100% FBS or 10% FBS is shown in Figure 4. By 24 hours, some cells had attached and adopted characteristic adherent macrophage morphologies: astral shapes with short filopodia (Figure 4A–C). At Day 8, adherent macrophages had approximately doubled in size (Figure 4D–F, note scale bar difference) and filopodia and membrane ruffles were observed features of the majority of cells, comparable to previous observations on model biomaterial surfaces.^{44,63} Similar surface coverage and growth patterns were observed for each test condition (Figure 4G–I). At Day 12, a substantial portion of each pp-FC surface was covered with cells (Figure 4J, L), although some areas were more sparsely populated (Figure 4K). By Day 19, individual cells adopted macrophage morphologies typically observed on FC surfaces:^{44,63} lengthy filopodia, membrane ruffling, unusual shapes and clustered growth with overlapping regions (Figure 4M–O). Generally, BMMO morphologies on pp-FC surfaces were consistent with those previously observed for the IC-21 macrophage cell line on Teflon[®] AF surfaces.⁶³

For all time points, cell adhesion results for FC surfaces exposed to 3 mg/ml BSA or 10% FBS were comparable, suggestive of albumin biasing of the protein-adsorbed surface, regardless of whether a complex mixture of serum proteins (e.g., FBS) or a single component BSA solution was used to condition the FC surface prior to cell culture. Results shown in Figure 3 (pp-FC) indicate that this surface supported cell growth and proliferation, but when compared to Teflon[®] AF (Figure 5), a significant difference in BMMO response was observed. Although numerous non-adherent cells were observed on each FC surface (i.e., spherical cells with a haloed appearance), Teflon[®] AF was more permissive to BMMO attachment and spreading; this behavior was observed from 8–36 days of culture. On pp-FC, BMMO grew and proliferated until approximately Day 21 of culture, when cells began to detach and die.

In contrast, attachment-dependent NIH 3T3 fibroblast cells (controls) were unable to colonize identically treated pp-FC surfaces (Figure 3J and Figure 6D–I) at any time point. On PS surfaces (Figure 6A–C) cell proliferative kinetics were approximately equivalent for the 3 mg/ml BSA and 10% FBS samples. PS surfaces preconditioned with 100% FBS (Figure 6B) showed substantial improvement in cell proliferation at early time points (through 72 hours). Eventually, the cell populations on all three PS pre-conditioned surfaces reached 100% confluence (data not shown), and large confluent "sheets" of fibroblasts delaminated easily from PS, suggesting that cell-cell contacts were stronger than cell-surface interactions on this surface. On Teflon[®] AF surfaces, BSA and 10% FBS pre-conditioning treatments (Figure 6D and F) also exhibit similar non-adhesive results, with only 100% FBS (Figure 6E) supporting limited NIH 3T3 cell adhesion. The pp-FC surfaces were non-supportive of cell adhesion regardless of pre-conditioning (Figure 6G–I) and this was maintained until NIH 3T3 cultures were terminated (Day 6 post-seeding).

Surface pre-conditioning treatments also included PBS⁺⁺, 0.3 mg/ml fibrinogen (control), 10% HI FBS in PBS⁺⁺ and cell-specific medium containing 10% FBS. Regardless of treatment, BMMO (data not shown) and IC-21 cells both adhered rapidly and spread on FC surfaces, exhibiting great variance in cell size (Supplemental Figure 1). Results for BMMO and IC-21 cell response to the most frequently employed pre-conditioning treatments are shown in Figure 7. When compared to TCPS positive controls (Figure 7, A, C) IC-21 cells on Teflon[®] AF surfaces (Figure 7E, G, I) exhibited more motile phenotypic features. In contrast, BMMO cells were larger with more motile phenotypes on TCPS surfaces at this time point (Figure 7, B, D versus F, H, J). In general, less BMMO adhesion was observed on

Teflon[®] AF surfaces than TCPS, whereas IC-21 adhesion was comparable on both surfaces regardless of protein pre-conditioning (Supplemental Figure 2).

A majority of IC-21 and BMMO cells present Mac 1 integrin subunits

Flow cytometry experiments were performed to determine levels of expression of $_2$ and $_M$ subunits comprising the Mac-1 (CD11b/CD18) integrin receptor in BMMO and IC-21 populations. Results (Figure 8) indicate that 76 ± 13% of the IC-21 and 80 ± 5% of the BMMO cells express $_2$ (CD18), and 90 ± 1% of the IC-21 and 87 ± 3% of the BMMO populations express $_M$ (CD11b). Integrin $_2$ is often implicated in macrophage adhesion⁶⁶ and extravasation.⁶⁷ Mac-1 ($_M/_2$) is known to interact with numerous ligands, including complement protein C3, fibrinogen, ICAM-1, -2 and -3, VCAM-1 and factor X.⁶⁸ Further, Mac 1 has been implicated in mediating leukocyte adhesion to denatured proteins, including albumin⁶⁹ and denatured albumin, ¹⁶ facilitating interactions of cells with solid substrates.¹⁶ Hence, its presence in the majority of these cells implicates a possible mechanism for albumin-based surface adhesion.

Functional blocking of integrin β_2 reduces cell adhesion on Teflon® AF substrates and leads to altered adherent macrophage morphology

Antibody function-blocking experiments were used to determine the role of $_1$ and $_2$ integrin subunits on macrophage adhesion to FC surfaces. Controls for 1) the surface, Teflon[®] AF (Figure 9A, D) and 2) non-specific binding (isotype matched control, Rat IgG2a, ; Figure 9B, E) were included. mAb-blocking against integrin $_2$ produced significantly reduced cell adhesion for both the IC-21 (reduced 88 ± 6%) and BMMO (reduced 80 ± 9%) cells on Teflon[®] AF surfaces (Figure 9C, F) compared to untreated cells seeded under identical conditions. Further, at one hour post-seeding, $_2$ -blocked seeded cells (Figure 9C, F) exhibit altered morphologies when compared to both those unblocked on Teflon[®] AF (Figure 9A, D) and isotype controls (Figure 9B, E). Post-blocking adherent cell morphology was more spherical with fewer obvious adhesion sites, fewer filopodia and less extensive membrane ruffling, reflecting phenotypic change.

Incomplete blocking of cell adhesion observed in this experiment suggests (an) additional mechanism(s) for cell attachment, given that flow cytometry shows 20–24% of the cell populations lack a ₂ receptor, and blocking efficiency directed against ₂ is less than 100%. Other integrins may contribute to monocyte/macrophage adhesion. Analogous assay of ₁ integrin-mediated adhesion were inconclusive due to high levels of non-specific binding for isotype controls (data not shown). Non-integrin mediated cell attachment mechanisms may include non-specific binding, hydrophobic interactions and interactions (primarily electrostatic in nature) between heparin- (or heparin-like) binding domain and proteoglycans or glycoproteins found on cells/surfaces.⁶⁸ However, that these are specific to macrophage adhesion on FC, and not other cell types shown refractory to FC binding in serum, seems implausible.

Fibronectin transcript expression in cells adherent on control and Teflon[®] AF surfaces

Fibronectin mRNA production in both BMMO and IC-21 (Supplemental Figure 3) cells grown on TCPS and Teflon[®] AF surfaces was confirmed at various time points (up to 24-hours post-seeding). Agarose gel electrophoresis results for fibronectin cDNA amplification (Supplemental Figure 3A, lanes 2–5; B lanes 2–4) yielded a 400 bp amplicon band. Each sample was also positive for GAPDH (data not shown). BMMO cells were positive for fibronectin transcript production at 30 and 60 minutes on both surfaces (Supplemental Figure 3A, lanes 2–5); whereas IC-21 cells expressed fibronectin transcript at 30 minutes and two hours post-seeding on TCPS (Supplemental Figure 3B, lanes 2–3) but not Teflon[®] AF (data not shown). IC-21 cells were positive for fibronectin production on Teflon[®] AF,

but only at 24 hours post-seeding (Supplemental Figure 3B, lane 4). Endogenous macrophage fibronectin production demonstrated at early culture time points (30 minutes in some cases, Supplemental Figure 3) may increase adsorbed matrix protein density sufficiently to increase cell adhesive site density, facilitating cell-surface interactions, producing cell-enhanced remodeling of the pre-existing surface adsorbed protein layer and additional monocyte/macrophage adhesion as cultures progress temporally.

DISCUSSION

Significant to the FBR observed in vivo surrounding implanted fluorinated polymers.^{3,70} in vitro macrophage cultures were found to efficiently adhere to and colonize FC surfaces of two different chemical compositions despite an abundant adsorbed albumin overlayer. This distinct FC adhesive behavior characterized cells of both primary- (BMMO) and secondary-(IC-21) origin. At early time points, BMMO colonization was more successful on standard tissue culture substrates designed to promote cell adhesion, growth and proliferation compared to FC. However, BMMO cells were able to achieve nearly 100% confluent surface coverage on FC substrates as cultures progressed. Both FC substrate cultures were allowed to progress to extended (21-36 day) time points not typically assayed *in vitro*. Some previous reports assay "non-adhesive" albuminated substrates such as FC for very short (~24 hours) time periods,⁷¹ perhaps missing a more long-term efficient cell adhesion to FC surfaces described here. While short-term interactions might be interesting for *in vitro* drug screening assays using macrophages (i.e., for inflammatory mediators), longer-term adhesion in serum is relevant to the FBR scenario. Findings confirm numerous previous reports of "anomalous" monocyte/macrophage growth^{41,42} and motility on hydrophobic surfaces^{16,33,69} that contrast the better-known poor adhesion characteristics typical of numerous cultured attachment-dependent mammalian cell types on FC surfaces.^{15,36,72,73}

Several protein pre-conditioning treatments produced similar effects on observed macrophage adhesion to and colonization of FC substrates, although slightly improved cell adhesion was noted for FC surfaces pre-biased with the known macrophage-adhesive protein^{1,74} fibrinogen. Previous studies show unique behavior for proteins in general, and albumin specifically, on FC surfaces.³⁰⁻³² As supported by ToF-SIMS data, albumin, the most abundant serum protein (55% of total serum mass),⁹ binds to FC surfaces with high affinity and retention, resisting surfactant elution.^{23,24,31} This albumin-binding characteristic has been correlated with improved surface resistance to cell adhesion, related to the ability of albumin to effectively block recognition of adhesive protein motifs present on these surfaces via integrin receptor-mediated interactions.^{24,31,36,75,76} This is significant given that ToF-SIMS and fluorescence staining show that albumin is the primary adsorbed protein on culture surfaces assessed here and correlated with macrophage adhesion. Other potential adsorbed protein mediators of (monocyte-) macrophage adhesion to FC surfaces in vitro include complement protein C3,⁶⁶ fibrinogen,^{77,78} and trace matrix proteins (e.g., fibronectin and vitronectin),^{79,80} each known to interact with cells via specific integrin receptor interactions. However, previous serum-based adsorption studies indicate that negligible serum-adsorbed fibronectin is detectable on PTFE²⁰ or Teflon[®] AF.⁸¹ Fibrinogen should be effectively removed during standard commercial serum preparation.⁸²

Interestingly, despite presumed albumin dominance of FC surfaces in culture media and similarity in general FC surface composition, $pp-C_3F_8$ surfaces were less permissive than Teflon[®] AF surfaces to BMMO cultures, suggesting that variations in FC surface chemistry or architecture translate to altered protein conformations that affect these naïve cells in culture. The two FC surfaces likely exhibit different surface and bulk FC chain arrangements: Teflon[®] AF has amorphous $-CF_2$ - backbone segments dispersed with 2,2-bis (trifluoromethyl)-4,5-difluoro-1,3-dioxide segments that present $-CF_3$ groups dispersed in

the surface zone (see XPS spectra in Ref. 32). pp-C₃F₈ FC is an amorphous, crosslinked thin film,^{83,84} with different plasma preparation conditions producing differing amounts of -CF₂units oriented both in- and out of- plane at the surface, and varying amounts of co-existing – CF₃ terminal groups.⁸³ The reported static aqueous contact angle for pp-FC (112°)⁸¹ suggests a terminating surface layer largely of –CF₃ groups. Hence, these pp-FC surfaces have different mixtures of -CF₂- and –CF₃ chemistry. These FC surface structural differences can translate to non-equivalent adsorbed protein densities²⁴ and/or conformations.^{85,86} Since lower surface energy correlates to higher albumin retention,²⁴ and FC surfaces are known to both sequester albumin and exhibit poor cell adhesion, different adhesive behaviors observed for BMMO cells (the most naïve and perhaps the most biologically-relevant/stimulus-sensitive cell type explored here) on pp-FC versus Teflon[®] AF surfaces may be attributed to changing albumin adsorbed states on surfaces over time in culture.

Previous findings indicate that monocytes and activated macrophage cell lines adhere proficiently to denatured proteins, including albumin,^{16,69} with direct relevance to the scenario described here. Given that: 1) highly hydrophobic surfaces (e.g., FC) have been shown to be albumin selective, 2) albumin is apparently the major protein adsorbed to both FC surfaces, and 3) the mechanism of adhesion to FC surfaces was consistent with findings of Davis,¹⁶ i.e., functional blocking of the Mac-1 2 integrin significantly impaired adhesion of BMMO and IC-21 cells to FC substrates, this suggests that these macrophage cells utilize integrin-mediated binding mechanisms primarily to adhere to denatured proteins present on FC surfaces. Importantly, this interaction may proceed via integrin interactions with protein sites exposed only after albumin surface denaturation or altered conformational state(s) are achieved. Nonetheless, presence of surface-adsorbed fibronectin or other ECM protein density sufficient to accommodate integrin attachment cannot be completely ruled out (except in 100% BSA pre-adsorbed cultures), despite the lack of evidence from fluorescence studies and the strong supporting evidence for predominantly albumin-covered surfaces. While mAb blocking experiments directed toward the 2 integrin Mac-1 receptor subunit reduced cell adhesion substantially for both primary- and secondary-derived cells on FC substrates, some limited adhesion was still observed, indicating that other interactions mediate macrophage adhesion to FC surfaces. Collectively, these studies demonstrate that primary BMMO and secondary-derived IC-21 cells behave very similarly in their adhesive capacity and mechanism, and cell morphology, motility and proliferative behavior on FC surfaces regardless of protein surface biasing. Differences observed between IC-21 and BMMO cells grown to extended culture times (36 days) on pp-FC and Teflon® AF surfaces may reflect important phenotypic differences enhanced by extending culture times. Since many immortalized secondary cells lines are selected for their ability to readily grow in adhesion-dependent cultures, phenotypic differences between primary (i.e., surface naïve) macrophages and those of secondary origin might be attributed to this selection bias.

Lastly, *in vitro* "biocompatibility" testing and cell-based high-throughput drug screening assays (i.e., for inflammatory tests) often employ immortalized, secondary-derived (monocyte-) macrophage cell lines due to their inflammatory phenotype, ready availability, ease of culture, and cost-effectiveness, despite the fact that these cells exhibit varied adhesion patterns, culture requirements, longevity and unknown "equivalence" to primary inflammatory cell phenotypes. Masking agents such as albumin and assay substrate materials including FC are used to prevent undesired cell adhesion to certain assay components in these cultures. This study indicates that this strategy might in fact facilitate unwanted cell adhesion and perhaps undesired activation of these cells, producing assay artifacts not representative of *in vivo* conditions.

CONCLUSIONS

The major conclusions of this study are that: 1) macrophages readily adhere to and populate albuminated FC surfaces that are non-permissive to other cell types *in vitro*, 2) cells of both primary and secondary (immortalized) origin respond to FC surfaces very similarly over short (<1 week) culture periods, 3) the Mac-1 $_2$ integrin receptor subunit plays a major role in establishing initial cell adhesive contacts to FC surfaces in the presence of both albumin and serum media, and 4) albumin's use as masking agent in many cell-based assays, drug screening, and as a passivating agent implanted biomaterials might not be advised where macrophage adhesion is undesirable.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

This work was funded by NIH grant EB 000894. ToF-SIMS and PCA experiments were performed at the National ESCA and Surface Analysis Center for Biomedical Problems, funded by NIH grant EB-002027. The authors acknowledge A. García (Georgia Institute of Technology), G. Hagen, P. Gong, P. Wu, M. Gonzalez-Juarerro (Colorado State University), M.S. Wagner (Proctor and Gamble) and D.J. Graham (Assemblon, Inc.) for helpful technical guidance. Y–F. "Jerry" Liu is appreciated for technical assistance with RT-PCR experiments.

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Figure 1.

ToF-SIMS principal component analysis (PCA) performed on BSA- and 10% FBS-treated Teflon® AF surfaces, projected into the model of Wagner *et al.*²¹ A) Wagner's PCA-based model identifying and grouping various proteins adsorbed to PTFE. Abbreviations: collagen (Col), cytochrome C (Cyc), fibrinogen (Fgn), fibronectin (Fn), (gamma)-globulin (Glb), hemoglobin (HB), immunoglobin G (IgG), lysozyme (Lsz), lactoferrin (Ltf), myoglobin (Myg), transferrin (Trf). B) Samples of 3 mg/ml BSA and 10% FBS adsorbed on Teflon® AF substrates were analyzed by ToF-SIMS and projected into Wagner's BSA model for other substrates.



Figure 2.

Fluorescence scanning results for Alexa Fluor®-labeled (as described in Material and Methods) fibronectin and BSA protein solutions (37 µg/ml BSA-Alexa Fluor 647®, and 4 µg/ml fibronectin-Alexa Fluor 555®) exposed to surfaces with a wide range of wettabilities for 24 hours: Codelink[™] and glass controls (very hydrophilic), FC (very hydrophobic). Unlabeled proteins were incubated with FC surfaces as a negative control. Inset: Magnified results for unlabeled protein samples and labeled proteins exposed to hydrophilic Codelink[™] substrates.



Figure 3.

Phase contrast photomicrographs of live cells on control (TCPS) and pp-FC surfaces for primary-derived cells (BMMO) and secondary-derived immortalized cell lines of (monocyte-) macrophage (RAW 264.7, J774A.1, IC-21) and fibroblast (NIH 3T3) origin. Cell seeding density varied, as described in Materials and Methods. Cultures depicted represent 2–4 days post-seeding on each surface and represent the typical sub-confluent growth pattern observed for each cell type (except J). Images are representative of multiple fields (3) and multiple replicates (2) for each test condition.



Figure 4.

Phase contrast photomicrographs of live BMMO grown on uniform pp-FC surfacespreconditioned with a single component BSA (3 mg/ml) solution, 100% FBS or 10% FBS. Cells were seeded at densities of 1500 cells/mm² Times indicated and scale bars are relevant for each row of images. Circles indicate "astral" morphology (A, E), arrows indicate filopodia (D, F, M, O) and bold arrows indicate membrane ruffling (D, F). Multiple cytoskeletal features (i.e., filopodia, membrane ruffling) are common features on many cells (e.g., D and F). Images are representative of multiple fields (3) and multiple replicates (2) for each test condition.



Figure 5.

Phase contrast photomicrographs of live BMMO (Day 8 in culture) on FC surfaces exposed to preconditioning treatments as indicated. Cells were seeded at densities of 600 cells/mm² on Teflon® AF and 1500 cells/mm² on pp-FC. Circles indicate astral morphology (C–E), arrows indicate filopodia (A–D, F) and bold arrows indicate membrane ruffling (D). Scale bar shown in F is relevant for all images. Images are representative of multiple fields (3) and multiple replicates (2) for each test condition.



Figure 6.

Phase contrast photomicrographs of live NIH 3T3 cells 48 hours post-seeding in 10% serum containing media on uniform pp-FC surfaces pre-treated with (3 mg/ml) BSA (A, D and G), 100% (B, E and H) or 10% serum (C, F and I). NIH 3T3 fibroblasts fail to effectively colonize FC surfaces under any of the conditions tested (D–I). Scale bars are relevant for each row. Images are representative of multiple fields (> 5) and multiple plates (2) for each test condition.



Figure 7.

Phase contrast photomicrographs of IC-21 and BMMO cells on control (TCPS) and Teflon® AF surfaces with preconditioning treatments (Teflon® AF only) as indicated. Cells were seeded at a density of 500 cells/mm², fixed and stained after one hour. Scale bars in D and J are equivalent, and relative for all panels except A and B. Images are representative of multiple fields (3) and multiple replicates (2) for each test condition.



Figure 8.

Flow cytometry analysis of integrin $_2$ (CD18) and $_M$ (CD11b) expression on populations of BMMO and IC-21 cells grown on TCPS under standard conditions. A) Histograms showing specific integrin labeling. Black line represents isotype control and dotted line represents fluorescence intensity expressed by binding of isotype control or integrin of interest (CD18, CD 11b) to BMMO or IC-21 cells, respectively. B) Bar graphs representation of data shown in A, indicating the percent of total sampled cell population positive for the integrin indicated. n=3 for each experiment, error bars represent standard error of the mean.



Figure 9.

Phase contrast photomicrographs of results for control and $_2$ blocks performed on IC-21 cells grown on Teflon® AF surfaces preconditioned with 10% FBS. Cells were incubated without antibodies or with either isotype control or blocking antibodies at a concentration of 100 µg/ml for 30 minutes prior to seeding at a density of 500 cells/mm². Cells were fixed and stained after one hour. Scale bars are relevant for each row. Images are representative of multiple fields (3) and multiple replicates (2) for each test condition.