

Supporting Information

Rapid, Interface-Driven Domain Orientation in Bottlebrush Diblock Copolymer Films During Thermal Annealing

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Table of Contents

1. Materials and characterization methods	2
2. Thermal Properties of PS-b-PLA Bottlebrush Block Copolymers (DSC)	2
3. Details of Thermal Annealing Process	2
4. Quantification of <i>in situ</i> Reflection Properties via Image Analysis	2
5. Image Processing and Orientation Mapping of Scanning Electron Micrographs	2
6. UV-Vis-Nir Diffuse+Specular Reflection Spectra for All Samples.....	2
7. Volume Fraction Estimation.....	2
8. SEM cross-sectional images of an initial ‘pre-pressed’ BBCP film.....	2
9. Supplemental References.....	2

1. Materials and characterization methods

All reactions were performed in an argon-filled glovebox ($O_2 < 2$ ppm, $H_2O < 0.5$ ppm) at room temperature using oven-dried glassware. THF, toluene, and DCM was dried using a commercial solvent purification system. rac-Lactide {Aldrich}, sec-butyllithium solution {1.3 M in cyclohexane/hexane (92/8), ACROS Organics} and ethylene oxide solution (2.5-3.3 M in THF, Aldrich) was used as received. 1,8-diazabicyclo[5.4.0]undec-7-ene (DBU) {Aldrich} was distilled over CaH_2 and storage under argon at -20 °C. Styrene was pass through neutral alumina plug and stored under argon at -20 °C. $[(H_2IMes)(3-Br-py)_2(Cl)_2Ru=CHPh]$, G3 was synthesized according to literature.¹ exo-5-Norbornene-2-carboxylic acid and exo-5-Norbornene-2-methanol was synthesized according to literature.² NMR characterization of identical materials, but a different synthesis batch can be found in our prior publication.³

Conventional Size Exclusion Chromatography (SEC) was performed using a Tosoh Ecosec HLC-8320GPC at 40 °C with THF (HPLC grade) as the eluent. This SEC is equipped with both a refractive index and UV detector (detector set to 266 nm). The SEC is fitted with a guard column (6.0 mm ID x 4.0 cm x 5 μ m), and two analytical columns (7.8 mm ID x 30 cm x 5 μ m; TSKgel GMH_{HR}-H). The reference flow rate is 0.5 mL min⁻¹ while the analytical column is at 1.0 mL min⁻¹. Polystyrene standards (16 points ranging from 200 Mw to 2.1 million Mw) were used as the general calibration. An additional calibration was created for specifically for linear polylactic acid and only used for linear polylactic acid (12 points ranging from 500 Mw to 10,000 Mw).

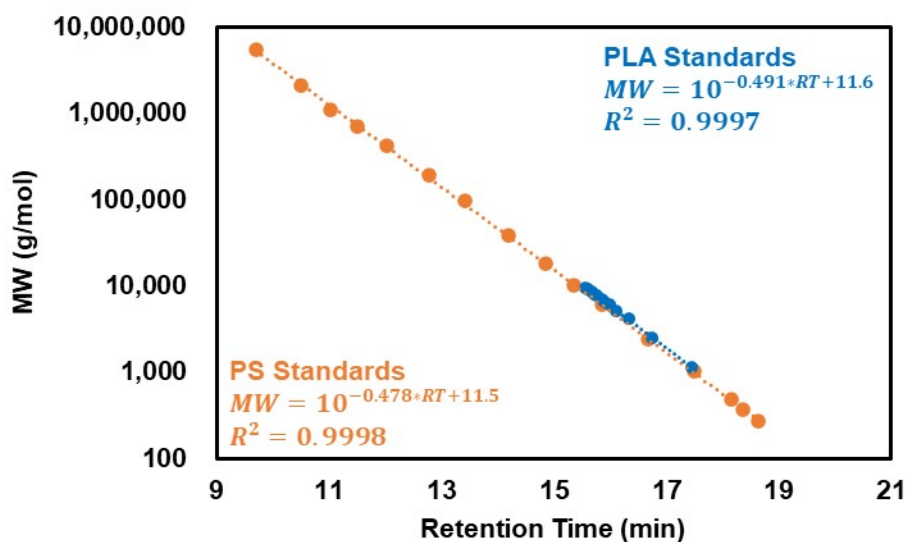


Figure S1. Conventional size exclusion chromatography calibration curves with polystyrene and polylactic acid standards.

Size Exclusion Chromatography (SEC) was performed using a Tosoh Ecosec HLC-8320GPC at 40 °C with THF (HPLC grade) as the eluent. This SEC is equipped with a refractive index, UV detector, and LenS3 Multi-Angle Light Scattering Detector. The SEC is fitted with a guard

column (6.0 mm ID x 4.0 cm x 5 μm), and two analytical columns (7.8 mm ID x 30 cm x 13 μm ; TSKgel Alpha-M). The reference flow rate is 0.3 mL min⁻¹ while the analytical column is at 0.6 mL min⁻¹. The detectors were calibrated with a narrow polystyrene standard ($M_w = 99,000$ Da). Polymer solutions were prepared at a known concentration (ca. 1 mg/mL) and an injection volume of 20 μL was used. dn/dc values for the bottlebrush polymers were obtained for each injection by assuming 100% mass elution of the bottlebrush from the columns (the injection mass was adjusted according to the known wt% from conventional SEC).

Synthesis procedures

The synthesis methodology used in this manuscript to produce the diblock bottlebrush polymer has been reported in several prior publications with extensive characterization and detailed procedures.^{3–5} We provide the key characterization here, and direct readers to prior publications for more details and characterization of the methodology.

Characterization data

Table S1. Characterization data for macromonomers.

Chemistry	$M_{n, \text{GPC}}$ (g/mol) ^a	M_w/M_n^a
PS	4,500	1.03
PLA	4,200	1.06
^a Calculated from conventional GPC calibration respect to PS standards or PLA standards.		

Table S2. Characterization data for PS-PLA diblock bottlebrush (PS: 4.5 kg/mol; PLA: 4.2 kg/mol, 50 wt% PS).

Targeted N_{bb}	M_n^a (kg/mol)	M_w/M_n^a	wt % ^b				Block length PS:PLA	$M_{w, LS}^c$ (kg/mol)
			Diblock BB	Block 1	PLA arm	PS arm		
400	602	1.04	93	6	>1	1	194:229	1800
^a Calculated from conventional GPC calibration respect to PS standards ^b See SI section of reference ⁵ for calculation details. ^c Determined from triple detect GPC								

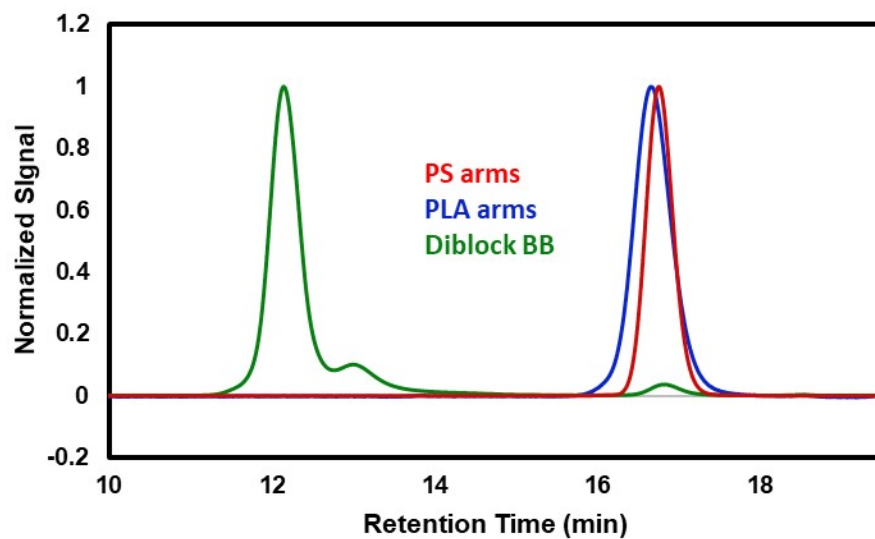


Figure S2. RI-GPC traces for the synthesis of PS-b-PLA diblock bottlebrushes. (The small second hump on the green trace around 13 min is PS-BB cause by catalyst death upon addition of PLA brushes. The small peak at 16.5 min corresponds solely to unfunctionalized PS arms.)

2. Thermal Properties of PS-b-PLA Bottlebrush Block Copolymers (DSC)

As-prepared PS-b-PLA bottlebrush block copolymers exhibited glass transition temperatures (2nd heat) at 95.93 °C (PS block) and 49.15 °C (PLA block) as shown below.

Measurement Parameters.

A 5.5 mg sample was measured in an Aluminum pan (DSC Consumables Tzero) and scanned on a TA instruments Discovery 2500 Differential Scanning Calorimeter (DSC) at the University of Illinois Materials Research Laboratory. A Heat-Cool-Heat cycle was performed by heating from ambient to 150°C, cooling to 25°C, and heating to 150°C. Heating was performed at 10°C/min with 2-minute isotherms at endpoints.

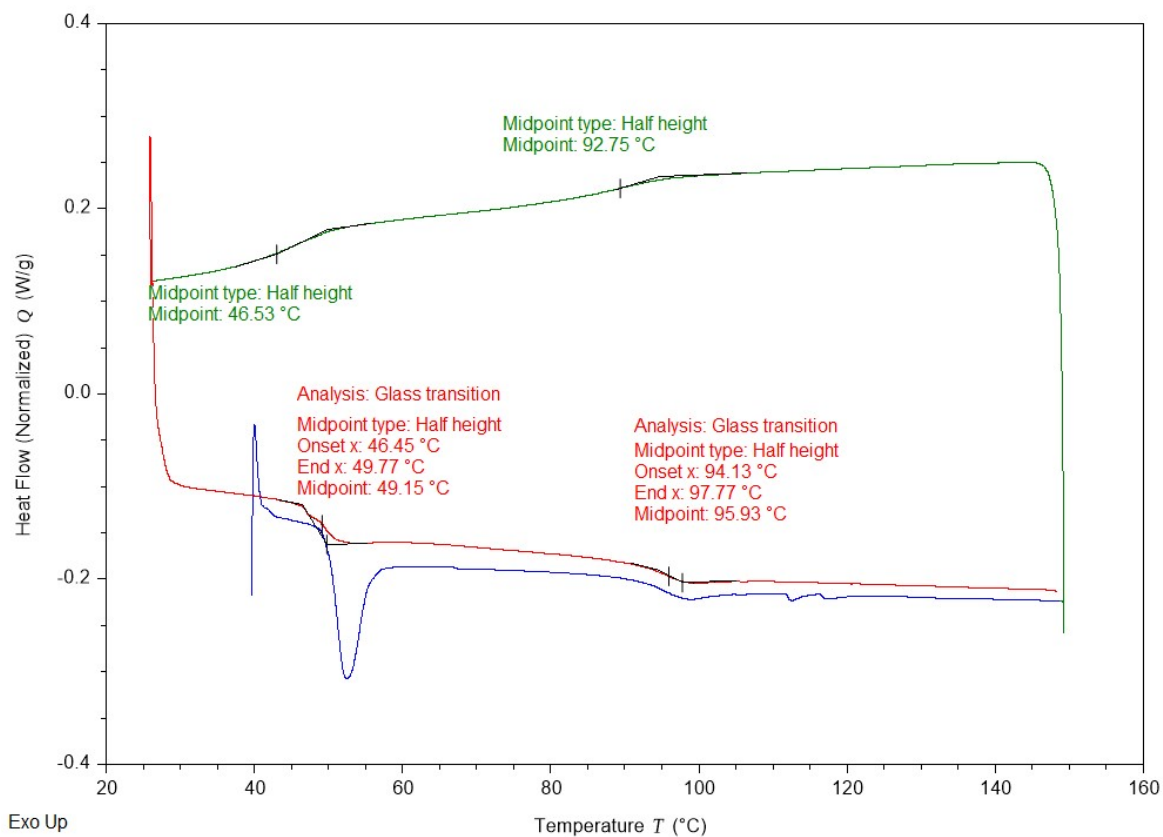


Figure S3. Heat-Cool-Heat data for the synthesized PS-b-PLA Bottlebrush Block copolymer.

3. Details of Thermal Annealing Process

A cartoon of the thermal annealing protocol is provided in **Figure S4** below.

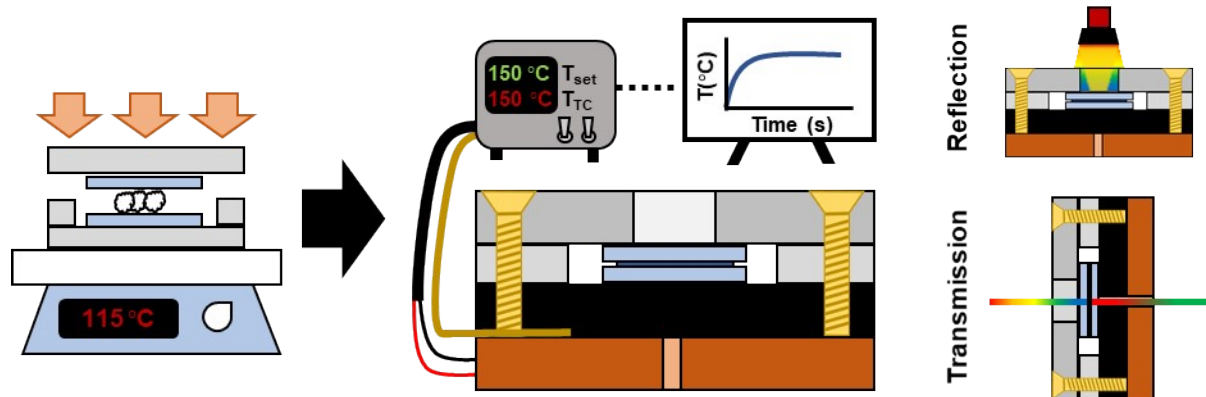


Figure S4. Cartoon of the thermal annealing scheme for two series of films. Films were prepared by first compressing purified BCCP powder between glass slides at 115 °C with film thickness defined by either external (metal) or internal (photoresist) spacers. During annealing, samples were installed in a custom heating/compression cell at room temperature and allowed to heat to 150°C using a closed-loop temperature controller and resistive heater, with a thermocouple embedded below the sample as drawn. In situ monitoring of optical properties was performed in either reflection or transmission mode. Reflection properties were probed using a microscope camera in normal incidence configuration under a ring light. Transmittance was measured by mounting the heating cell within a UV-Vis spectrometer.

The annealing cell was fabricated by the University of Illinois School of Chemical Sciences Machine Shop staff. The cell was designed to accommodate substrates with dimensions of 1" x 1". Prior to assembly, a T-type thermocouple was sealed into a machined groove in the bottom plate with thermal paste and covered with Kapton tape. A photo of the assembled cell is provided in **Figure S5**, with original mechanical drawings following.

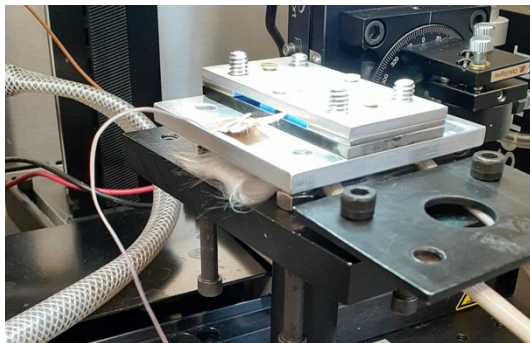


Figure S5. Photograph of the annealing cell in reflection mode.

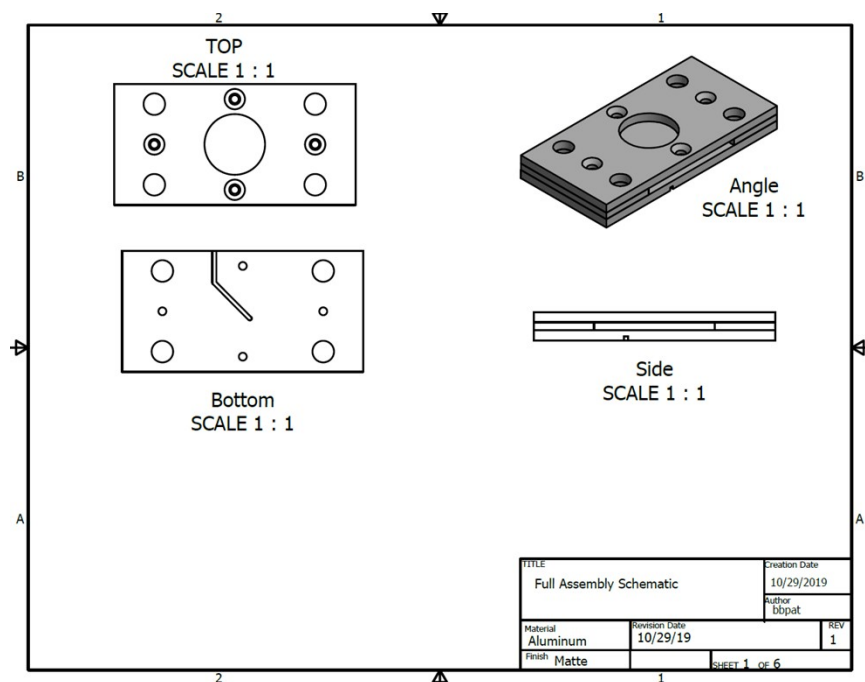


Figure S6. CAD drawing of the cell assembly comprising top and bottom plate and spacers.

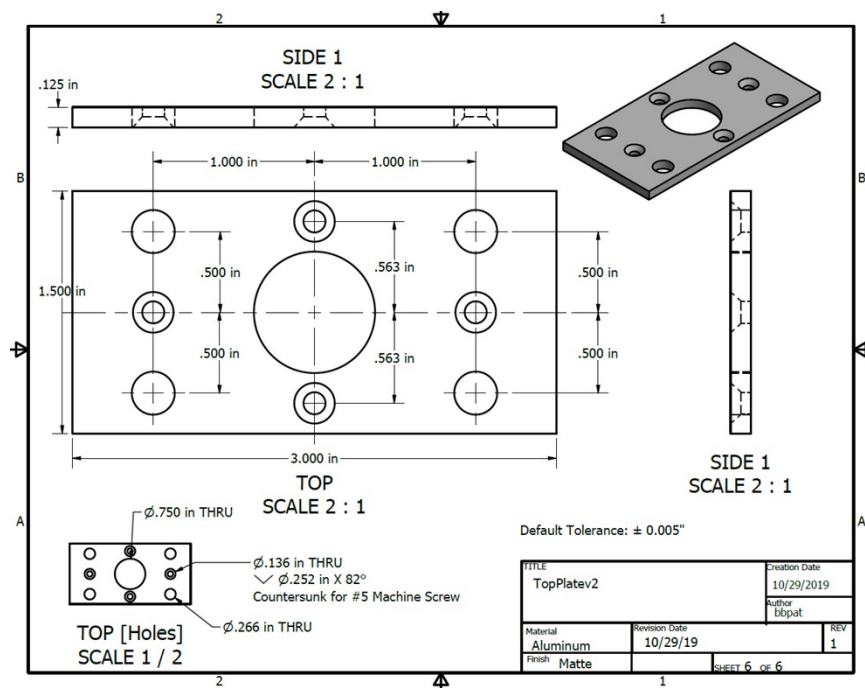


Figure S7. CAD drawing of the top plate of the annealing cell.

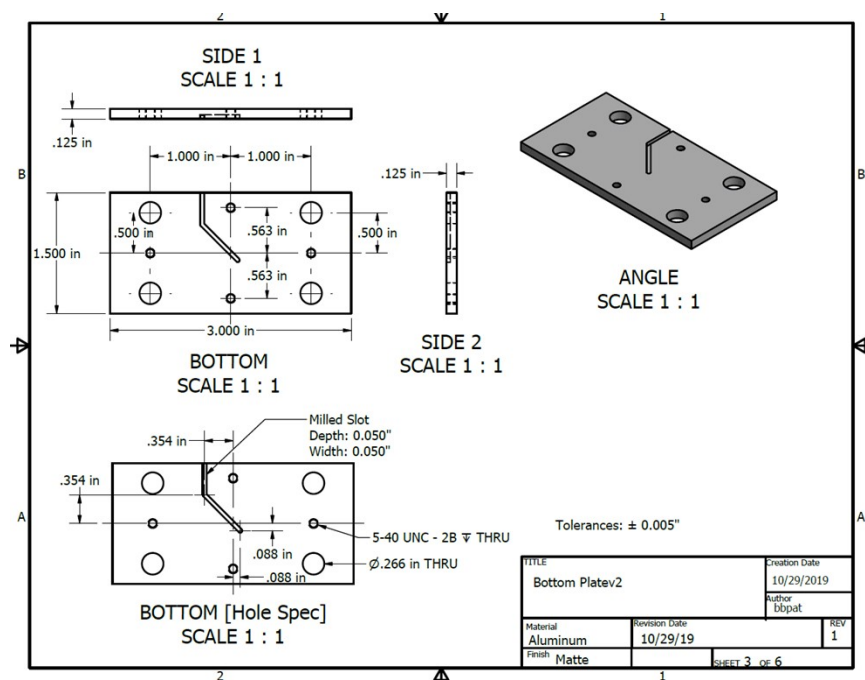


Figure S8. CAD drawing of the bottom plate of the annealing cell (reflection experiments).

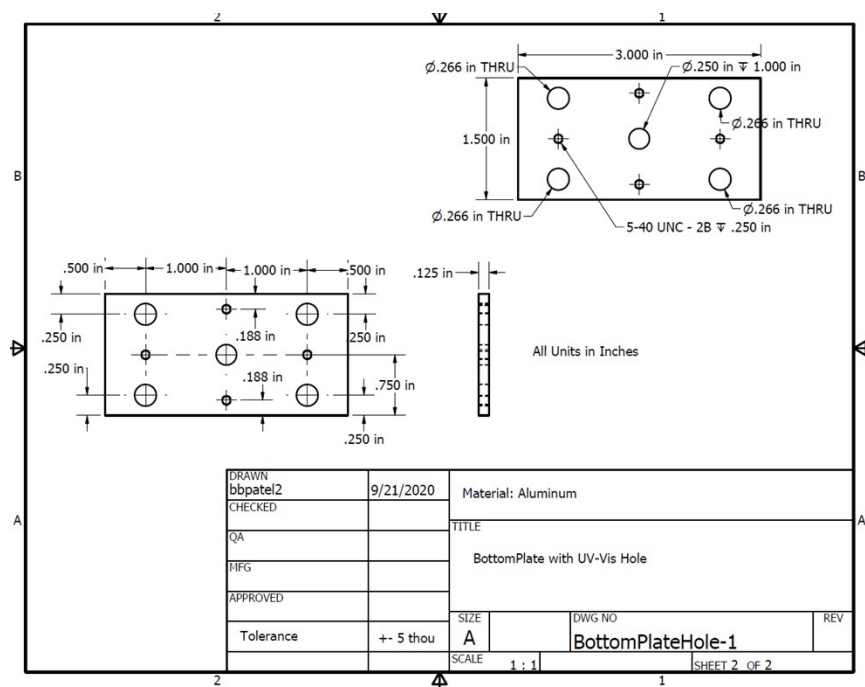


Figure S9. CAD drawing of the bottom plate of the annealing cell (UV-Vis transmission experiments).

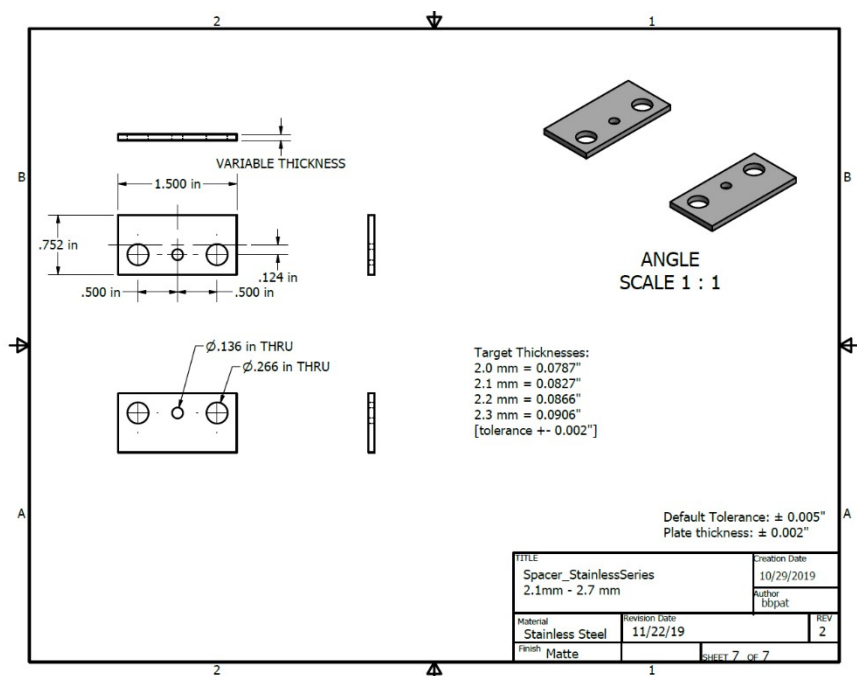


Figure S10. CAD drawing of the series of machined spacers. The 2.0 mm spacer was used for all samples less than $100\ \mu\text{m}$ thick, instead relying on photoresist spacers patterned between the substrates.

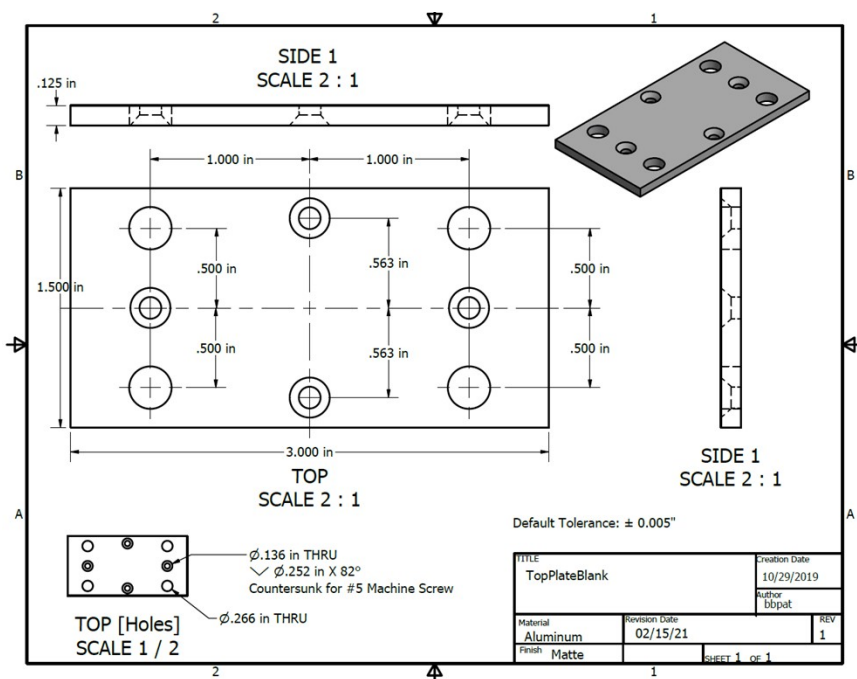


Figure S11. CAD drawing of the 'blank' top plate used for pre-pressing samples before *in situ* annealing.

4. Quantification of *in situ* Reflection Properties via Image Analysis

Matlab code provided below takes as input a folder full of images to be analyzed.

```
%% Software Information
% Name: BCPIImageAnalyzer
% Description: Frame-by-Frame image RGB/HSV Image analysis
% Version: 1.2
% Release Date: 200206
% Version Summary: Added direct calculation of HSV Values from image
% Authors: Bijal Patel, Kush Patel, Justin Kwok

% License Terms

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% All rights reserved.
%
% Developed by: <Diao Lab>
%             <University of Illinois>
%             <http://diao.scs.illinois.edu/Diao\_Lab/Home.html>
%
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% this list of conditions and the following disclaimers.
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% NO EVENT SHALL THE CONTRIBUTORS OR COPYRIGHT HOLDERS BE LIABLE
% FOR ANY
% CLAIM, DAMAGES OR OTHER LIABILITY, WHETHER IN AN ACTION OF
% CONTRACT, TORT
```

% OR OTHERWISE, ARISING FROM, OUT OF OR IN CONNECTION WITH THE
SOFTWARE OR
% THE USE OR OTHER DEALINGS WITH THE SOFTWARE.

%% Step 0 - Initialize

```
disp('Step 0: Initializing...')
clear all
close all
```

%Get current date/time

```
dateTime = datestr(datetime(now,'ConvertFrom','datetime'));
dateTime = strrep(dateTime,":","_");
dateTime = strrep(dateTime,"-","_");
dateTime = strrep(dateTime," ","_");
```

BrightnessThreshold = 1.6;

% Select Folder Path

% ENTER FOLDER PATH WITH ORIGINAL VIDEOS HERE

```
filedir=strcat('[Folder Path Here'],'\');
% or default to userdir
% filedir=strcat(userpath,'\');
```

```
disp('Choose Folder for analysis...')
```

```
[filedir]=uigetdir(strcat(filedir,'*. *'));
```

```
disp('Folder Chosen!')
```

```
filename = input("Enter Sample Name: ','s');
```

```
framerate = str2num(input("Enter Frames per second: ','s'));
```

%Make/choose specific folders for data/frame export

```
expDir = strcat(filedir,'MATLABexport\',filename,"");
```

```
disp('Choose Folder with raw frames...')
```

```
rawFramesDir = strcat(uigetdir(strcat(filedir,'*. *')), "");
```

```
threshFramesDir = strcat(expDir,'ThreshFrames\');
```

```
calcDataDir = strcat(expDir,'calcData\',dateTime,"");
```

```
mkdir(expDir);
```

```
mkdir(threshFramesDir);
```

```
mkdir(calcDataDir);
```

```
logParams = ["1. Export Directory" "2. RawFramesDirectory" "3. ThresholdFramesDirectory"  
"4. CalculatedValuesDirectory"];
```

```
logVals = [expDir rawFramesDir threshFramesDir calcDataDir];
```

```

disp('Step 0: Initializing Complete!')

%% Load Video to RAM
loadToRAM = input("Frame Export Required? (Y/N): ','s');
if (strcmp(loadToRAM,'Y'))
    disp('Importing Video to RAM...')
    obj = VideoReader(strcat(filedir,filenameext));
    vid = read(obj);
    frames = obj.NumberOfFrames;
    framerate = obj.FrameRate;

    logParams = [logParams "Framerate"];
    logVals = [logVals framerate];
    disp('Import Video Complete!')
else
    disp('Skipping Loading Video to RAM')
    frames = size(dir(fullfile(rawFramesDir, '*.jpg')),1);
end

%% Convert Video file to frames and export
if(strcmp(loadToRAM,'Y'))
    disp('Export Raw Frames to File...')
    %Ask user if frames need to be output
    inp = input("    Output Frames? Y/N: ','s');

    %if needs output, do so, else skip
    if (strcmp(inp,'Y'))
        for x = 1:frames
            imwrite(vid(:,:,:,x),strcat(rawFramesDir,filename,'_raw_',num2str(x),'.jpg'));
            fprintf('\tExporting frame: %d out of %d.\n',x,frames);
        end
        disp('Export Raw Frames Complete!')
        beep
    else
        disp('Skipping Frame Export')
    end
else
    disp('Skipping Frame Export')
end

%% Draw mask
disp('Masking')
inp = input("    Draw New Mask? Y/N: ','s');

frames = length(dir(fullfile(rawFramesDir, '*.jpg')));

```

```

if(newAnalysis || strcmp(inp,'Y'))
    disp('Choose Frame to draw mask from.')
    %Choose frame to look at for drawing mask

    maskFrame = chooseFrame(1,frames,rawFramesDir,strcmp(filename),-1);

    %Export Frame
    expTail = strcat(filename,'_maskFrame.jpg');
    expPath = strcat(calcDataDir,expTail);
    saveas(gcf,expPath);

    disp("    Mask Frame chosen! Now Draw Mask");

    %Size of frame image
    imsize=size(imread(strcat(rawFramesDir,filename,'_1','.jpg')));

    %Draw polygon mask
    %mx and my are the vertex positions of your mask
    %left click for points, right click to close loop
    %right click a vertex to create mask
    [mask, mx, my]=roipoly;

    %name image of frame to mask
    im = imread(strcat(rawFramesDir,filename,"_",num2str(maskFrame),'.jpg'));

    %Display mask overlayed onto image
    imshow(imoverlay(mat2gray(im),mask,[0.1 0.8 0.1]));

    %Save masked image
    %Export Frame
    expTail = strcat(filename,'_maskedFrame.jpg');
    expPath = strcat(calcDataDir,expTail);
    saveas(gcf,expPath);

    %Save mask itself
    %Save mask to file

    maskPath = strcat(expDir,'_',dateTime,'_mask.mat');
    save(strcat(calcDataDir,'mask_',dateTime,'.mat'),'mask');

    %uncomment for no mask
    %mask=logical(ones(imsize(1:2)));
else %Load Mask from File
    disp('Choose Mask');

```

```

[maskfilenameext, maskfiledir] = uigetfile(strcat(filedir, '*.*'));
maskPath = strcat(maskfiledir, maskfilenameext);
load(maskPath);
save(strcat(calcDataDir, 'mask_', dateTime, '.mat'), 'mask');
end

logParams = [logParams "MaskFrame"];
logVals = [logVals maskFrame];

disp('Get Mask Complete!')

%% Identify first frame to calculate
disp('Choose first frame to calculate values for (NOT tZERO)')

%Choose frame for t=0
calcStartFrame = chooseFrame(maskFrame, frames, rawFramesDir, strcat(filename), mask);
%Export start Frame
expTail = strcat(filename, '_MaskedCalcStart.jpg');
expPath = strcat(calcDataDir, expTail);
saveas(gcf, expPath);

disp('Frame for calculation start chosen!')

%% Identify frameTzero time correction
disp('Choose frame for time zero...')

%Choose frame for t=0
frameZero = chooseFrame(maskFrame, frames, rawFramesDir, strcat(filename), mask);
%Export Masked Frame
expTail = strcat(filename, '_MaskedTzero.jpg');
expPath = strcat(calcDataDir, expTail);
saveas(gcf, expPath);

disp('tzero Frame Chosen!')

%% Evaluate mean RGB intensity in masked region
fstart=calcStartFrame;%number of starting frame
fspace=1;%number of frames to increment by

numFramestoCalc = frames-fstart; %number of frames to calc
%Preallocate arrays
Int_RGB = zeros(1,numFramestoCalc);
frameAvg_Gint = zeros (1,numFramestoCalc);
frameAvg_Rint = zeros (1,numFramestoCalc);
frameAvg_Bint = zeros (1,numFramestoCalc);

```

```

frameAvg_H = zeros (1,numFramestoCalc);
frameAvg_S = zeros (1,numFramestoCalc);
frameAvg_V = zeros (1,numFramestoCalc);

%Folder for thresholded images

for currentFrame=fstart:fspace:frames
    %Read in frame as RGB
    RGB_frame = imread(strcat(rawFramesDir,filename,'_',num2str(currentFrame),'.jpg'));
    %Convert to HSV
    HSV_frame = rgb2hsv(RGB_frame);

    %extract R, G, and B map values separately
    R=double(RGB_frame(:,:,1));
    G=double(RGB_frame(:,:,2));
    B=double(RGB_frame(:,:,3));

    %extract H, S, and V map values separately
    H=double(HSV_frame(:,:,1))*360;
    S=double(HSV_frame(:,:,2));
    V=double(HSV_frame(:,:,3));

    %NaN values outside of region of interest
    R(mask==0)=nan;
    G(mask==0)=nan;
    B(mask==0)=nan;

    %NaN values outside of region of interest
    H(mask==0)=nan;
    S(mask==0)=nan;
    V(mask==0)=nan;

    %'unwrap' H circle so red values dont appear on both top and bottom
    %of axis
    for hValIndex = 1:size(H)
        if (H(hValIndex)>320)
            H(hValIndex) = H(hValIndex) -360;
        end
    end

    %Get avg RGB values for this frame within mask
    frameAvg_Rint(currentFrame-fstart+1)= nanmean(nanmean(R));
    frameAvg_Gint(currentFrame-fstart+1)= nanmean(nanmean(G));
    frameAvg_Bint(currentFrame-fstart+1)= nanmean(nanmean(B));

    %Mean intensity calculated from R,G,B.

```



```

Int_RGB(currentFrame-fstart+1)=mean([frameAvg_Rint(currentFrame-
fstart+1),frameAvg_Gint(currentFrame-fstart+1),frameAvg_Bint(currentFrame-fstart+1)]);

%NAN overexposed pixels (white) from light source reflection
%White pixels are those above threshold value
[lengthX, lengthY, rgb] = size(RGB_frame);
for column = 1:lengthY
    for row =1:lengthX
        if (sum([R(row,column) G(row,column) B(row,column)]) > BrightnessThreshold*
3*Int_RGB(currentFrame-fstart+1))
            R(row,column)= nan;
            G(row,column)= nan;
            B(row,column)= nan;
            H(row,column)= nan;
            S(row,column)=nan;
            V(row,column)=nan;
        end
    end
end

%Recalculate avg RGB values for this frame within mask
frameAvg_Rint(currentFrame-fstart+1)= nanmean(nanmean(R));
frameAvg_Gint(currentFrame-fstart+1)= nanmean(nanmean(G));
frameAvg_Bint(currentFrame-fstart+1)= nanmean(nanmean(B));

%Recalculate Mean intensity from R,G,B.
Int_RGB(currentFrame-fstart+1)=mean([frameAvg_Rint(currentFrame-
fstart+1),frameAvg_Gint(currentFrame-fstart+1),frameAvg_Bint(currentFrame-fstart+1)]);

%get avg_hsv for this frame within mask
frameAvg_H(currentFrame-fstart+1) = nanmean(nanmean(H));
frameAvg_S(currentFrame-fstart+1) = nanmean(nanmean(S));
frameAvg_V(currentFrame-fstart+1) = nanmean(nanmean(V));

%Export Thresholded Frame
imwrite(uint8(cat(3,R,G,B)),fullfile(threshFramesDir, strcat(num2str(currentFrame),".jpg")));
fprintf('\tOn frame: %d out of %d processed.\n',currentFrame-fstart+1,frames-fstart);
end

logParams = [logParams "Pixel Threshold"];
logVals = [logVals BrightnessThreshold];

disp('Step 6: Calc Mean RGB/HSV Complete!')
%% Step 7 - Plotting Data
disp('Step 7: Plotting Results...')

```

```

%Time management
%frame number array
xFrame=fstart:fspace:frames;

%time array
%calc time with first frame at t = 0s
xTime=(xFrame-frameZero)./framerate;%seconds
xvar=xTime;

%Plotting mean RGB vs time
subplot(2,3,1);
plot(xvar,Int_RGB,'k')
title('Mean RGB Intensity')
xlabel('Time elapsed(s)')
ylabel('Intensity (Arb)')
set(gcf,'color','w');
box on

%Plotting R, G, B, individually and mean vs frame
subplot(2,3,4);
hold on
plot(xFrame,Int_RGB,'k','Linewidth',1.5)
plot(xFrame,frameAvg_Rint,'r','Linewidth',1.5)
plot(xFrame,frameAvg_Gint,'g','Linewidth',1.5)
plot(xFrame,frameAvg_Bint,'b','Linewidth',1.5)

%add thick vertical lines every 10/100/1000 frames
if (frames > 1000)
    for i=0:100:frames
        xline(i,'k');
    end

elseif (frames > 10000)
    for i=0:1000:frames
        xline(i,'k');
    end
else
    for i=0:10:frames
        xline(i,'k');
    end
end

title('RGB Separated Intensity Values')
xlabel('Frame')
ylabel('Intensity (Arb)')
set(gcf,'color','w');

```

```

hold off
box on

%Plotting R, G, B, individually and mean vs time
subplot(2,3,[2 3]);
hold on
plot(xvar,Int_RGB,'k','Linewidth',1.5)
plot(xvar,frameAvg_Rint,'r','Linewidth',1.5)
plot(xvar,frameAvg_Gint,'g','Linewidth',1.5)
plot(xvar,frameAvg_Bint,'b','Linewidth',1.5)
title('RGB Separated Intensity Values')
xlabel('Time elapsed (s)')
ylabel('Intensity (Arb)')
set(gcf,'color','w');
box on
hold off
set(gcf, 'Position', [100, 100, 1000, 500])

%Plotting H and V vs time
subplot(2,3,[5 6]);
hold on
yyaxis left
ylabel('Hue')
plot(xvar,frameAvg_H,'k','Linewidth',1.5)
yyaxis right
plot(xvar,frameAvg_S,'b','Linewidth',1.5)
plot(xvar,frameAvg_V,'r','Linewidth',1.5)
title('Hue and Value over time')
xlabel('Time elapsed (s)')
ylabel('Value')
set(gcf,'color','w');
box on
hold off
set(gcf, 'Position', [100, 100, 1000, 500])
legend("Hue","Saturation","Value");

%Save figure
saveas(gcf,strcat(calcDataDir,'Plot.png'));

%Concatenate results into matrix [frame xtime(s) Mean1(Arb) Mean2(Arb) Rint(Arb)
Gint(Arb) Bint(Arb)]
m = [transpose(xFrame) transpose(xTime) transpose(Int_RGB) ...
      transpose(frameAvg_Rint) transpose(frameAvg_Gint) ...
      transpose(frameAvg_Bint) transpose(frameAvg_H) transpose(frameAvg_S) ...
      transpose(frameAvg_V)];

```

```

%Export Text
expTail = strcat(filename, '_Data.txt');
expPath = strcat(calcDataDir, expTail);

%Write data file headers
fid = fopen(expPath, 'w');
fprintf(fid, "Frame, TimeElapsed(s), Mean Intensity, Red Intensity, Green Intensity, Blue
Intensity, Hue (degrees), Saturation (0-1), Value (0-1)\n");
fclose(fid);

%Write data
dlmwrite(expPath, m, '-append', 'delimiter', ',');

disp('Step 7 Complete!')

%% Step 8: Write Log File
disp('Step 8 Writing Log File...!')

%Write Log File

if(writeLog(expDir, logParams, logVals))
    disp('Log File Written');
else
    disp('Log Failed');
end

%% END

disp('Sequence Complete!')

%% Local functions
%Writes parameters to log file
function logged = writeLog(expPath, params, vals)
    %Get current date/time
    dateTime = datestr(datetime(now, 'ConvertFrom', 'datetime'));
    dateTime = strrep(dateTime, ":", "_");
    dateTime = strrep(dateTime, "-", "_");
    dateTime = strrep(dateTime, " ", "_");

    logName = strcat(expPath, '\AnalysisLog_', dateTime, '_txt');
    logFileID = fopen(logName, 'w');
    fprintf(logFileID, 'Log Entry - Generated \n%s\n', dateTime);
    outputs = [params; vals];

```

```

    fprintf(logFileID, '%s\n',outputs);
    fclose(logFileID);
    logged = 1;
end

%Loads parameters from log file
function data = readLog(logFilePath)
    logFid = fopen(logFilePath);
    txt = textscan(logFid,'%s','delimiter','\n');
    fclose(logFid);
    txt = txt{1}; %Pull first element of cell array
    txt = string(txt); %convert to string array
    %Convert to n x 2 array of param-val pairs
    data = [txt(1:2:end) txt(2:2:end)];
end

%Allows user to choose a frame
function pickedFrame = chooseFrame(frame, maxFrames, framesDir, filename,mask)
    currentFrame = frame;
    pickedFrame = 0;
    done = 0;

    while (done ~= 1)
        %display masked, if mask given

        im = imread(strcat(framesDir,filename,'_',num2str(currentFrame),'.jpg'));

        if (mask~= -1)
            imshow(imoverlay(mat2gray(im),mask,[0.0 0.5 0.0]));
        else
            imshow(im);
        end

        cmd = input('    q for done, asd+fg for -(10/5/1)+(1/5/10), frames: ','s');
        switch cmd
            case 'q'
                done = 1;
                pickedFrame = currentFrame;
            case 'd'
                if (currentFrame > 1)
                    currentFrame = currentFrame-1;
                end
            case 's'
                if (currentFrame > 11)
                    currentFrame = currentFrame-10;
                end
            default
                %do nothing
        end
    end
end

```

```

    end
case 'a'
    if (currentFrame > 101)
        currentFrame = currentFrame-100;
    end
case 'f'
    if (currentFrame < maxFrames)
        currentFrame = currentFrame+1;
    end
case 'g'
    if (currentFrame < maxFrames-10)
        currentFrame = currentFrame+10;
    end
case 'h'
    if (currentFrame < maxFrames-100)
        currentFrame = currentFrame+100;
    end
otherwise
    disp('Unrecognized input\n');
end

end
end

```

5. Image Processing and Orientation Mapping of Scanning Electron Micrographs

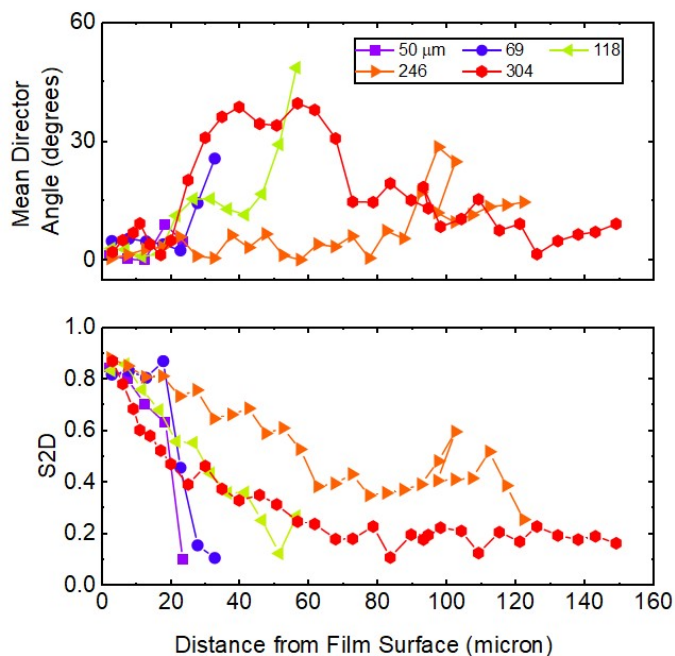


Figure S12. Results of orientation analysis for films characterized by cross-sectional scanning electron microscopy.

Image analysis for cross-sectional scanning electron micrographs was performed in two steps. First, images were pre-processed in FIJI(ImageJ2)⁶ to obtain binarized black and white domains. Then, images were processed using GTFiber2.0 by Persson et al.⁷ to obtain orientation statistics and colorized fiber maps.

Step 1: Image Preprocessing

In brief, image pre-processing consisted of 4 steps:

1. **Enhancing local contrast** to eliminate edge brightening effects from topographical variation under the electron beam.
2. **Applying a band pass filter** to enhance features with periodicity on the order of the domain size
3. **Binarization to black and white**
4. **Removal of off-film regions** for the top and bottom image of the series (if applicable)

For one series of particularly high-resolution images, it was found that small surface features (likely from Au/Pd deposition) were frequently causing misidentification of fibers. For this set, an optional gaussian blur step was added to the macro. **Figure S12** depicts the intermediate steps of this process.

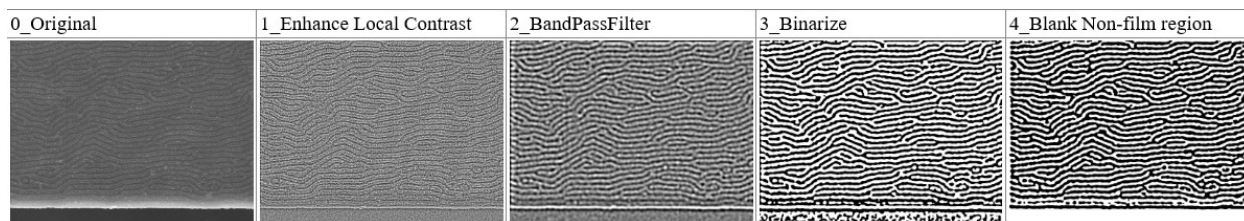


Figure S13. Snapshots of SEM pre-processing steps.

The following imagej2 macro code was written to streamline batch processing of images:

```
// Get Directory of files
// Batch Processing part modified from https://imagej.nih.gov/ij/macros/BatchProcessFolders.txt
sourceDir = getDirectory("Choose a Directory ");
setBatchMode(true);
count = 0;
countFiles(sourceDir);
n = 0;
processFiles(sourceDir);
print(count+" files processed");

function countFiles(sourcedir)
{
    list = getFileList(sourceDir);
    for (i=0; i<list.length; i++)
    {
        if (endsWith(list[i], "/"))
            print("subdirectory skipped");
            //countFiles(""+sourceDir+list[i]);
        else
            count++;
    }
}
```



```
}
```

```
function processFiles(sourceDir)
```

```
{
```

```
    list = getFileList(sourceDir);
```

```
    for (i=0; i<list.length; i++)
```

```
    {
```

```
        if (endsWith(list[i], "/"))
```

```
            print("subdirectory skipped");
```

```
            //countFiles(""+sourceDir+list[i]);
```

```
        else
```

```
        {
```

```
            showProgress(n++, count);
```

```
            path = sourceDir+list[i];
```

```
            processFile(path);
```

```
        }
```

```
    }
```

```
}
```

```
function processFile(path)
```

```
{
```

```
    if (endsWith(path, ".tif"))
```

```
    {
```

```
        open(path);
```

```
        //Get root of filename
```

```
        sourceName = File.nameWithoutExtension;
```

```
        print("Processing file:" + sourceName + "in folder:" + sourceDir);
```

```

print("Creating Output Folders");
//Create output folders

ClaheDir = sourceDir+"1_ContrastEnhanced"+File.separator;
File.makeDirectory(ClaheDir);
print(ClaheDir);

FFTBandPassDir = sourceDir+"2_BandPassFiltered"+File.separator;
File.makeDirectory(FFTBandPassDir);

BinarizedDir = sourceDir+"3_Binarized"+File.separator;
File.makeDirectory(BinarizedDir);

InvertedDir = sourceDir+"4_Inverted"+File.separator;
File.makeDirectory(InvertedDir);


print("Begin full processing macro");
print("Step 1: Enhancing Local Contrast");

//if data is too high resolution, smaller blocksize accentuates noise: go between 20-50
//run("Enhance Local Contrast (CLAHE)", "blocksize=50 histogram=256 maximum=1000
mask=*None*");

    run("Enhance Local Contrast (CLAHE)", "blocksize=20 histogram=256 maximum=1000
mask=*None*");


//Optional Blur for very noisy data
//run("Gaussian Blur...", "sigma=8");


saveAs("jpeg", ClaheDir+sourceName+"_CLAHE");
print("Saved jpeg to " + ClaheDir+sourceName+"_CLAHE");


print("Step 2: Running BandPass Filter");

run("Bandpass Filter...", "filter_large=30 filter_small=15 suppress=Vertical tolerance=5
autoscale saturate");

```

```

saveAs("jpeg", FFTBandPassDir+sourceName+"_FFTBandPass");
print("\tSaved jpeg to " + FFTBandPassDir+sourceName+"_FFTBandPass");

print("Step 3: Running Binarization");
setOption("BlackBackground", false);

//if default threshold is 'good enough'
run("Convert to Mask");

//if manual threshold needed, use following line
//setThreshold(0, 112);
//run("Convert to Mask");

//remove speckles if needed
//run("Remove Outliers...", "radius=10 threshold=50 which=Dark");
//run("Remove Outliers...", "radius=10 threshold=50 which=Bright");

saveAs("jpeg", BinarizedDir+sourceName+"_FFTBandPass");
print("\tSaved jpeg to " + BinarizedDir+sourceName+"_FFTBandPass");

print("Step 4: Inverting B/W");
setOption("BlackBackground", false);
run("Invert");
saveAs("tif", InvertedDir+sourceName+"_FIJIProcessed");
print("\tSaved tif to " + InvertedDir+sourceName+"_FIJIProcessed");
print("End full processing macro");
close();
}}

```

6. UV-Vis-Nir Diffuse+Specular Reflection Spectra for All Samples

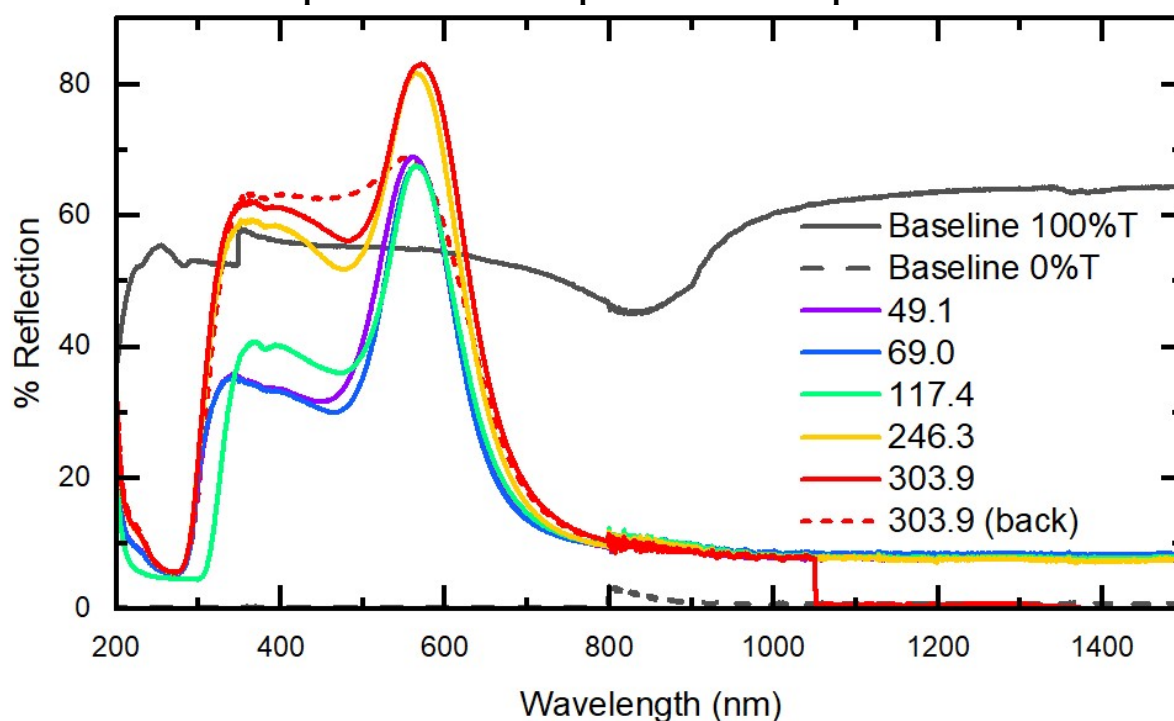


Figure S14. Raw Specular+Diffuse reflection spectra take after samples were annealed for 10 minutes under the microscope camera. Legend contains film thickness in microns.

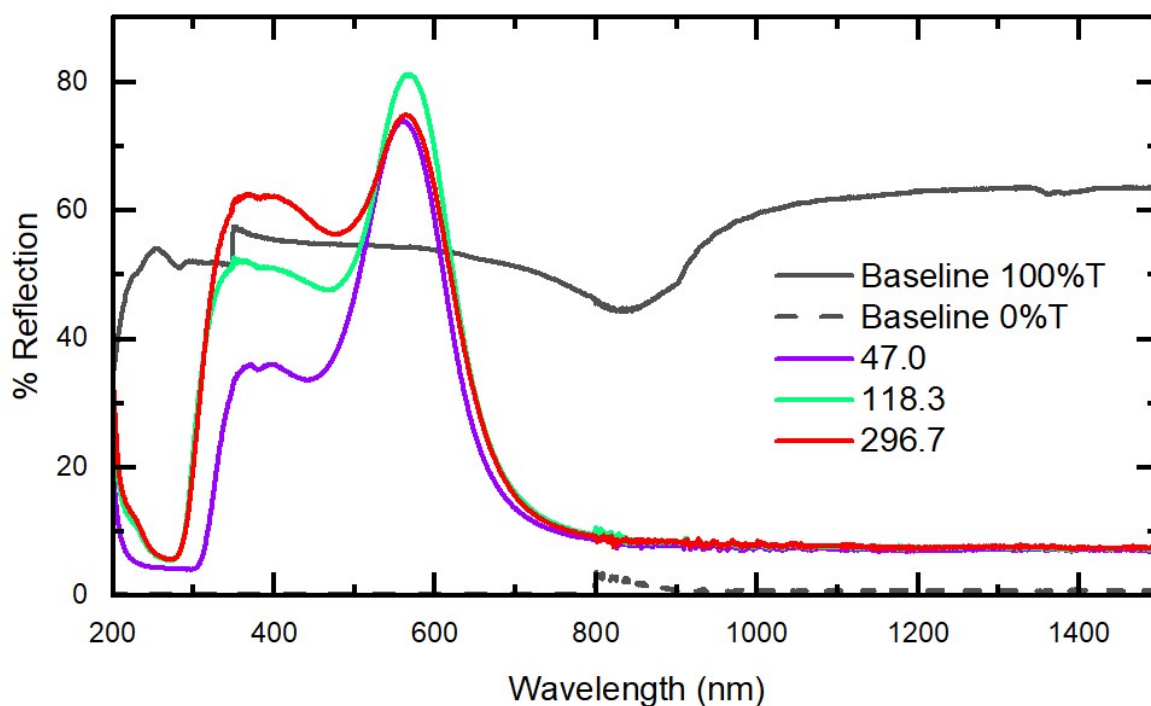


Figure S15. Raw Specular+Diffuse reflection spectra take after samples were annealed for 10 minutes within the UV-Vis transmission instrument. Legend contains film thickness in microns.

7. Volume Fraction Estimation

The volume fraction of PS was estimated by substituting the following data:

Table S3. Synthetic and chemical properties of polymacromonomers

<i>Species</i>	$M_n(\text{g/mol})$	DP	$\rho\left(\frac{\text{g}}{\text{cm}^3}\right)$	Density Citation
PS-NB	4500	200	1.04	8
PLA-NB	4200	200	1.25	9

into the following equation, as per Dalsin et al¹⁰

$$\Phi_{PS} = \frac{(M_{n,PS-NB} * DP_{PS-NB} / \rho_{PS-NB})}{(M_{n,PS-NB} * DP_{PS-NB} / \rho_{PS-NB}) + (M_{n,PLA-NB} * DP_{PLA-NB} / \rho_{PLA-NB})} = 0.56$$

Here, we approximate the density of the polymacromonomer to be the same as reported value for the bulk amorphous arm species.

8. SEM cross-sectional images of an initial ‘pre-pressed’ BBCP film

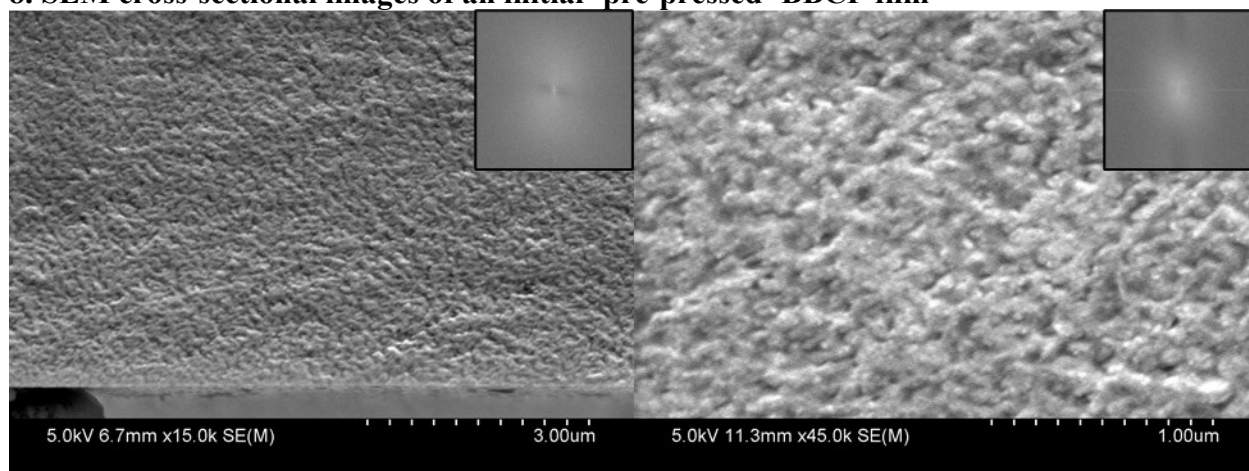


Figure S16. Cross-sectional SEM images near the edge of the pre-pressed BBCP film. Insets contain power spectrum images calculated using ImageJ’s FFT plugin.

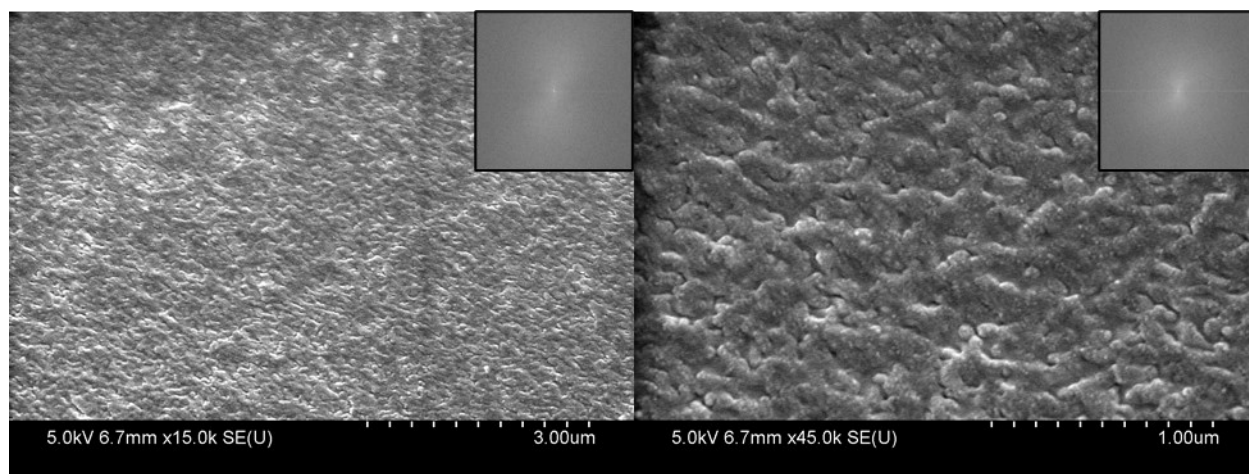


Figure S17. Cross-sectional SEM images near the center of the pre-pressed BBCP film. Insets contain power spectrum images calculated using ImageJ’s FFT plugin.

For comparison, **Figure S18** contains cross-sectional SEM images taken from the edge and center of a well-ordered film after annealing, along with their power spectrum images.

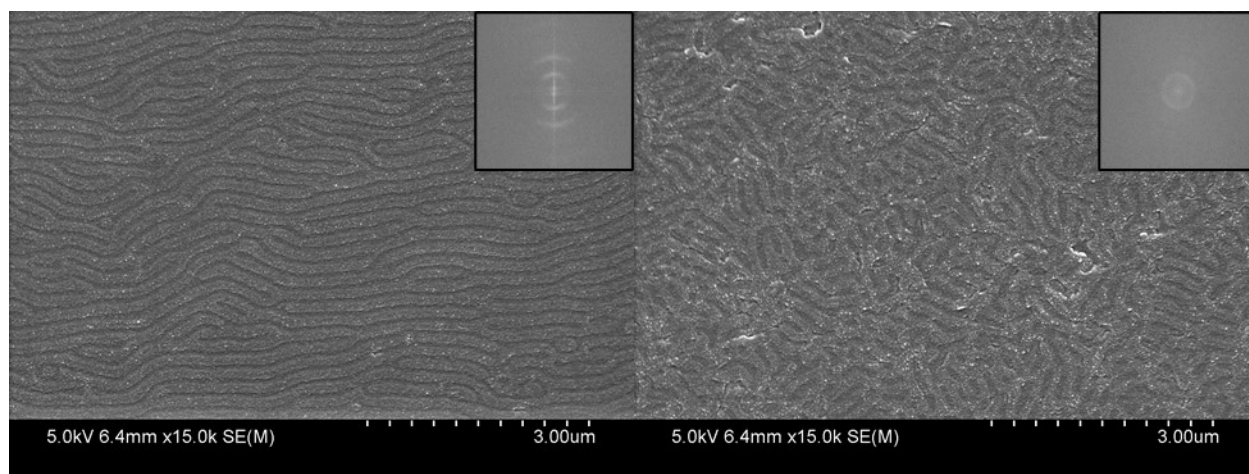


Figure S18. Cross-sectional SEM images taken at the center and edge of an annealed film (69.0 μm thick film). Insets contain power spectrum images calculated using ImageJ's FFT plugin.

9. Supplemental References

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