

Supplementary Information II

Blocky Bromination of Syndiotactic Polystyrene via Post-Polymerization Functionalization in the Heterogeneous Gel State

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```
% This code simulates copolymers with specified degrees of functionalization  
%and random or blocky microstructures and then calculates (1) the sequence  
%length and frequency of consecutive units; (2) the probability that a unit  
%exists in a crystallizable segment; and (3) the prevalence of unique triad  
%sequences in the copolymer chain
```

```
% This code was created using MATLAB® R2017a programming software
```

```
% Authors: Alexandria M. Noble & Kristen F. Noble
```

```
%% Variables
```

```
chain_length = 1442; % number of units in one chain
```

```
amt = 0.55; % fraction of inaccessible styrene units (for a random copolymer  
amt = 0)
```

```
keep_size = 5; % units in the inaccessible block (Note: for the purpose of  
demonstrating how the inaccessible block is created the below code is  
written for a keep_size of 5, however the code can be modified for any  
desired keep_size by following the format of  
Block_InaccessibleFraction.m, shown below. This work used a %keep_size  
of 53.
```

```
r = 1000; % defines the iterative process that generates r number of chains
```

```
ts = 3; % defines that triad sequences will be counted (Note: the code can  
be modified for any desired sequence (e.g., pentads, heptads, etc.) by  
following the format of TriadSequenceCounting.m, shown below.
```

```
l = 26; % units in one crystallizable segment
```

```
percent_functionalization = transpose(0.0:0.02:0.40); % Simulated degrees of  
functionalization (0-40% at intervals of 2%)
```

```
% The copolymer chain will contain 1's to represent styrene (s) units and 0's  
to represent brominated (b) styrene units.
```

```
%% Outputs
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```
keep = []; % stores the fixed styrene indices to create the inaccessible  
fraction
```

```
chain_matrix = []; % stores all chains for one degree of functionalization
```

```
store_avg = []; % stores the average prevalence for each degree of  
functionalization
```

```
store_sd = []; % stores the standard deviation of the prevalences for each  
degree of functionalization
```

```
P = []; % stores the probability that a 1 exists in a segment length of j  
consecutive 1 units
```

```

store_P = []; % stores the probability that a 1 exists in a crystallizable
segment for each chain
Table = []; % stores the average and standard deviation of the probability
that a 1 exists in a crystallizable segment for each degree of
functionalization

for vt = 1:size(percent_functionalization,1) % for each degree of
functionalization
store_P = [];
for uu = 1:r % generates r number of chains and performs the following
on each chain

chain = transpose(ones(1,chain_length)); % creates a chain of 1's of
length chain_length
q = round(amt*floor(chain_length)); % defines the number of
inaccessible 1's by the predefined amount (amt)

fixed_styrene = randsample(chain_length,q); % selects the
inaccessible 1's by random chance

%% run Blocky_InaccessibleFraction.m % code below

%% Blocky_InaccessibleFraction.m
% This code establishes the monomer units in the inaccessible fraction
% Note that for the purpose of demonstrating how the inaccessible block is
% created the below code is written for a keep_size of 5, however the code can
% be modified for any desired keep_size by following the format below. %This
% work used a keep_size of 53.
% This code was created using MATLAB® R2017a programming software

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%% -- Begin Code Blocky_InaccessibleFraction -- %%
keep = []; % stores the fixed styrene indices to create the
inaccessible fraction

for k = 1:size(fixed_styrene,1) % repeat until k is the size of the
fixed styrene matrix (based on the predetermined percent
functionalization and chain length)
while size(keep,1) < q % while the size of the keep matrix is
less than the number of inaccessible 1's (q) established above
if keep_size == 5
if fixed_styrene(k)-1 < 1 % prevents the styrene index
from being less than 1 (if fixed_styrene = 1 then keep 1,2,3)
keep =
[keep;fixed_styrene(k);fixed_styrene(k)+1;fixed_styrene(k)+2];
elseif fixed_styrene(k)-1 < 2
keep = [keep;fixed_styrene(k)-
1;fixed_styrene(k);fixed_styrene(k)+1;fixed_styrene(k)+2];
elseif fixed_styrene(k)+1 > chain_length % prevents the
styrene index from being greater than the chain length (if chain_length
= 1442 and fixed_styrene = 1442 then keep 1440,1441,1442)
keep = [keep;fixed_styrene(k)-2;fixed_styrene(k)-
1;fixed_styrene(k)];
elseif fixed_styrene(k)+2 > chain_length

```

```

        keep = [keep;fixed_styrene(k)-2;fixed_styrene(k)-
1;fixed_styrene(k);fixed_styrene(k)+1];
        else
        keep = [keep;fixed_styrene(k)-2;fixed_styrene(k)-
1;fixed_styrene(k);fixed_styrene(k)+1;fixed_styrene(k)+2];
        end
    end
end
    keep = unique(keep);
end

    %% -- End Code Blocky_InaccessibleFraction -- %%

    q2 = round(percent_functionalization(vt)*floor(chain_length)); %
gives integer value for the percent functionalization (e.g., 40%*length
of chain)
    I2 = transpose(1:1:chain_length); % converts a row of integer values
into a column

    xr = []; % specifies which indices are in the accessible fraction
based on indices in the inaccessible fraction
    for n = 1:size(keep,1)
        xr = [xr;find(I2 == keep(n))];
    end

    I2(xr) = []; % stores the indices not in keep

    rand_bromine = randsample(I2,q2); % selects at random without
replacement indices in the chain

    chain(rand_bromine,1) = 0; % changes specified indices in
rand_bromine to zeros
    chain_matrix(:,uu) = chain;

    x1 = find(chain == 0); % identify bromines
    x2 = find(chain == 1); % identify indices where chain is equal to 1

    chain_f(x1,1)={'b'}; % change all 0's in the chain to 'b' for
bromine
    chain_f(x2,1)={'s'}; % change all 1's in the chain to 's' for
styrene

    %% run TriadSequenceCounting.m %code below

%% TriadSequenceCounting.m
% This code indexes the triad combinations in the chain, calculates the
%frequency and prevalence of the unique triad sequences, and creates a matrix
%of the average and standard deviation of the triad sequence prevalences
%calculated from all generated chains
% Note: this code can be modified for any sequence length (e.g., pentad,
heptad, %etc.)
% This code was created using MATLAB® R2017a programming software

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```

        %% -- Begin Code TriadSequenceCounting -- %%
        tpseq = {}; % stores the triad sequences identified in the loop
below
        for i = 1:chain_length % loop that repeats based on the chain length
            if i <= chain_length-(ts-1) % if the step (i) is less than or
equal to the (chain_length)-ts-1
                indices = i:1:i+(ts-1); % then indices are selected as
shown
                if ts == 3
                    tpseq = [tpseq;
[char(chain_f(indices(1))),char(chain_f(indices(2))),char(chain_f(indic
es(3)))]], {indices}]; % makes triad sequences from the chain
                    end
                end
            end
        end

        if ts == 3 % if calculating the prevalence of triad sequences
            k = 8; % there are 8 possible sequences

            C = {'sss'; 'ssb'; 'sbs'; 'sbb'; 'bss'; 'bsb'; 'bbs'; 'bbb'}; %
creates an array of the 8 possible triad sequences
            C = table(C,C,zeros(size(C,1),1),zeros(size(C,1),1)); % creates
a table to store the triad sequences, their frequency, and their
prevalence
            C.Properties.VariableNames = {'SeqA','SeqB','Freq','Theo_Prev'};

            for i = 1:size(C,1) % for each triad sequence in column SeqA,
identify the unique triad sequences in the chain and count their
frequency (e.g., C.SeqA(i))
                flipped = {fliplr(C.SeqA{i,1})}; % flip C.SeqA in order to
count forward and reverse variants of the sequence

                if strcmp(C.SeqA(i),flipped) % if SeqA equals the flipped
SeqA (e.g., SeqA = 'sss')
                    x = find(strcmp(C.SeqA(i),tpseq(:,1))); % identify the
indices in tpseq where SeqA(i) occurs
                    C.SeqB(i) = C.SeqA(i); % SeqA == SeqB
                    C.Freq(i) = size(x,1); % obtain the size of x, which is
the frequency of each of the symmetric triad sequences without double
counting

                else % if SeqA does not equal the flipped SeqA (SeqA(i) is
asymmetric)
                    C.SeqB(i) = flipped; % populate column SeqB with the
appropriate variant from SeqA (e.g., 'ssb' = 'bss')
                    x = size(find(strcmp(C.SeqA(i),tpseq(:,1))),1)+...
size(find(strcmp(flipped,tpseq(:,1))),1); % obtain
the size of tpseq for the locations of C.SeqA(i) and the variant in
C.SeqB(i) and add the two size functions together to determine the
overall frequency
                    C.Freq(i) = x ; % records the overall frequency of
asymmetric SeqA(i) and its variant
                end
            end
        end
    end
end

```

```

C.Theo_Prev = C.Freq/size(tpseq,1); % calculates the prevalence of
each triad sequence as the ratio of the frequency to the total number
of triads in the chain (size of tpseq)

```

```

prev(:,uu) = C(:,4); % stores the prevalences for each triad
sequence from all generated chains
avg_prev = mean(prev,2); % calculates the average of the prevalences
sd_prev = std(prev,[],2); % calculates the standard deviation of the
prevalence

```

```

%% -- End Code TriadSequenceCounting -- %%

```

```

% Length and frequency of consecutive units

```

```

string = sprintf('%d',chain); % Converts the chain matrix into a
string with no spaces

```

```

t1=textscan(string,'%s','delimiter','0','multipleDelimsAsOne',1); %
reads the consecutive 1's from the string using 0's as delimiters (to
calculate the segment length of consecutive bromine, the delimiter
should be changed from 0 to 1)
s = t1{:}; % computes the length of each consecutive segment in the
string
data = cellfun('length',s); % assigns the length of each segment
(e.g., 111 = 3, 111111 = 6)

```

```

[number_times segment_length] = hist(data, 1:(chain_length)); %
makes matrices of the segment_lengths and their frequencies in the
string

```

```

Table1 = transpose([segment_length;number_times]); % converts the
above arrays into one matrix of segment length and frequency

```

```

% Calculate the probability that a 1 is from a segment of at least the
crystallizable segment

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```

j = Table1(:,1); % segment lengths of consecutive styrene units
P = []; % stores the probability that a 1 exists in a segment length
of j consecutive 1 units

```

```

for t = 1:size(j,1) % for each segment length
S = Table1(:,1).*Table1(:,2); % calculates the number of 1's in
each segment length (segment length*frequency)
wj = (1-percent_functionalization(vt))*(S/(chain_length*(1-
percent_functionalization(vt)))); % probability that a unit chosen at
random is a 1 and is a member of a sequence of j consecutive units

```

```

if j(t) < 1 % if the segment length is less than that of the
defined crystallizable segment (1) then P = 0
P = [P; 0]; % Probability based on Flory, P. J., Theory of
Crystallization in Copolymers. T. Faraday Soc. 1955, 51 (0), 848-857.
elseif j(t) >= 1

```

```
        P = [P; ((j(t)-1+1)/j(t))*wj(t)]; % if the segment length is
at least that of the defined crystallizable segment (l) then calculates
probability based on Flory, P. J., Theory of Crystallization in
Copolymers. T. Faraday Soc. 1955, 51 (0), 848-857.
```

```
    end
end
```

```
    sumP = sum(P); % sums all of the P's to determine the probability
that a unit chosen at random is a 1 from a crystallizable segment
    store_P = [store_P;sumP]; % stores the probability that a 1 exists
in a crystallizable segment for each chain
```

```
end
```

```
store_avg = [store_avg, avg_prev]; % stores the average of the
prevalences (rows) for each degree of functionalization (columns)
store_sd = [store_sd, sd_prev]; % stores the standard deviation of the
prevalences (rows) for each degree of functionalization (columns)
p1 = 100.*(mean(store_P)); % calculates the average of the store_P's for
each degree of functionalization
p2 = 100.*(std(store_P)); % calculates the standard deviation of the
store_P's for each degree of functionalization
Table = [Table; percent_functionalization(vt).*100,p1,p2]; % stores the
degree of functionalization and the average and standard deviation of
the probability that a 1 exists in a crystallizable segment
```

```
end
```