



Cite this: *Chem. Commun.*, 2016, 52, 3915

Received 21st January 2016,  
Accepted 8th February 2016

DOI: 10.1039/c6cc00611f

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## Role of highly branched, high molecular weight polymer structures in directing uniform polymer particle formation during nanoprecipitation†

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**The new macromolecular architecture, hyperbranched polydendrons, are composed of a broad distribution of molecular weights and architectural variation; however, nanoprecipitation of these materials yields highly uniform, dendron-functional nanoparticles. By isolating different fractions of the diverse samples, the key role of the most highly branched structures in directing nucleation and growth has been explored and determined.**

Polymeric nanostructures have been formed using various techniques including direct synthesis, such as micro/mini-emulsion polymerisation<sup>1,2</sup> and polymerisation induced self-assembly,<sup>3–6</sup> or the post-synthesis self-assembly of polymers of varying architectures, hydrophilicity or hydrophobicity.<sup>7–9</sup> Alternatively, physical fabrication methods such as the “Particle Replication In Non-wetting Templates” (PRINT™) approach<sup>10</sup> may be employed. In many self-assembly routes, a “solvent switch” is used to modify the media environment and drive inter-chain interactions. This can be achieved by changes in pH,<sup>11</sup> addition of salt,<sup>12</sup> removal of a co-solvent or the slow exchange of good solvent with an anti-solvent<sup>13,14</sup> (often water); solution and emulsion approaches have been reported.

A rapid and scale-able solvent switch technique that has gained considerable interest is nanoprecipitation,<sup>15</sup> especially in the formation of degradable nanoparticles for biomedical and nanomedicine applications. In general, polymers are dissolved in a good water-miscible solvent and rapidly added to an aqueous anti-solvent. Descriptions of both spontaneous, energy-barrier-free nucleation (spinodal decomposition)<sup>16</sup> and nucleation-and-growth mechanisms have been reported. In the latter case, providing the polymer is above its thermodynamic solubility limit within the resulting mixed solution, homogeneous nucleation of particles will occur.<sup>16</sup> The particle radii will initially be small,

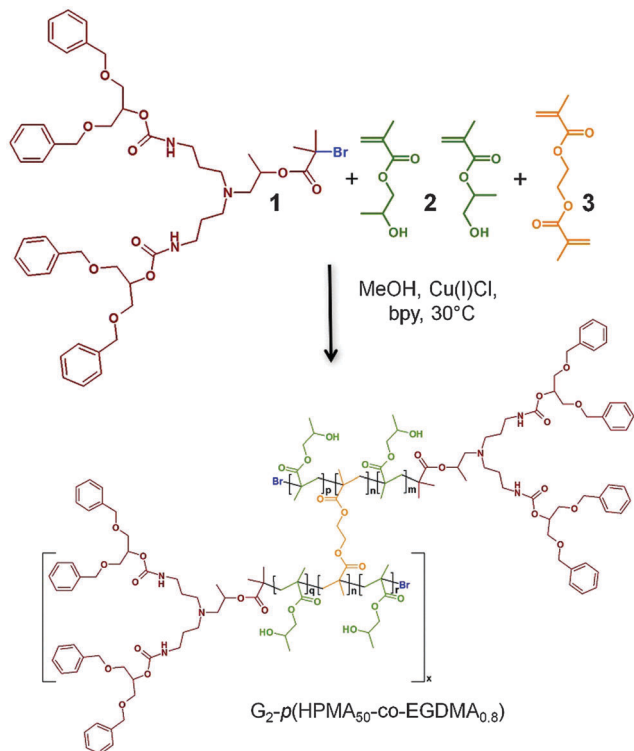
with a free energy of formation that is governed by parameters such as surface tension, particle radius and the free energy difference between media and particle. The free energy will pass through a maximum at a critical nucleus radius leading to rapid growth of larger particles. In either mechanism, growth will continue by depletion of polymer from the surrounding environment or through a process known as diffusion-limited cluster-cluster aggregation, assuming the balance of attractive and repulsive forces will allow interaction and steric/charge stabilisation does not prevent particles from achieving sufficient proximity for effective collision.<sup>15</sup> To ensure rapid polymer nucleation, several groups utilise microfluidic or rapid-mixing techniques that allow distinct separation of the nucleation and growth/aggregation stages.<sup>17,18</sup>

Studies of the impact of polymer architecture and molecular weight on nanoprecipitation outcomes are very rare.<sup>15</sup> We have reported a new complex polymer architecture – “hyperbranched polydendrons”<sup>19–21</sup> – that overcomes the complexity of dendrimer synthesis by branching between linear–dendritic hybrid polymer chains during propagation (Scheme 1); weight average molecular weights  $>10^6$  g mol<sup>−1</sup>, representing  $>90$  conjoined primary linear–dendritic chains, have been achieved. Hyperbranched polydendrons contain considerable variation in: (1) the number of branched chains (Fig. 1), (2) the position of branched points (Fig. 1) and, (3) molecular weight; a significant concentration of un-branched primary chains (linear–dendritic hybrid polymers) is also present. Despite this diversity, the complex mixtures nanoprecipitate to form stable, highly uniform nanoparticle dispersions. Each particle is surrounded by dendron functionality and outcomes are tunable to allow the multi-valency benefits of dendrimers to be achieved at sizes unobtainable by dendrimer synthesis alone (50–200 nm). Hyperbranched polydendrons mimic our studies of branched polymers without dendron chain-ends,<sup>22,23</sup> forming more uniform nanoprecipitates with increased stability and reduced size compared to particles generated from their chemically-identical linear primary chain building blocks. As such, they are ideal candidates for detailed examination of the role of the very highly branched, very high molecular weight fraction that is present within the molecular weight distribution.

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† Electronic supplementary information (ESI) available: Additional experimental details, NMR spectra, GPC chromatograms, SEM images of nanodispersions, graphs of nanoprecipitate characterisation. See DOI: 10.1039/c6cc00611f



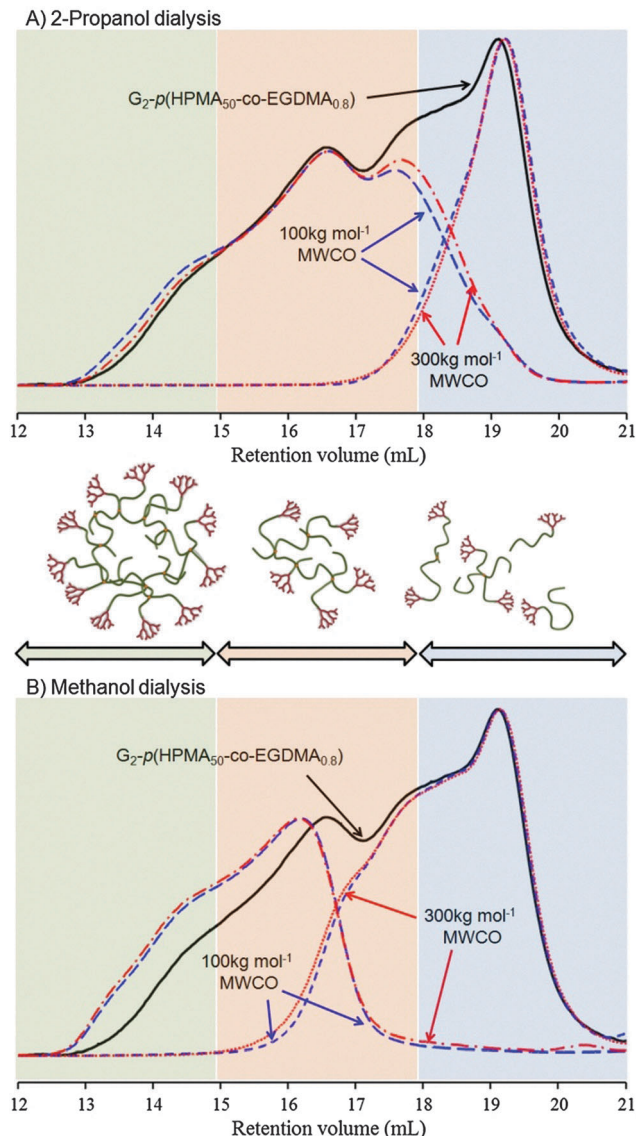


**Scheme 1** Cu-catalysed ATRP synthesis of hyp-polydendrons using HPMA, **2** (both isomers shown), EGDMA, **3**, and a  $G_2$  dendritic initiator, **1**.

Here we isolate the very high molecular weight fraction of hydrophobic hyperbranched polydendrons and study its role in directing successful nanoprecipitation to allow a greater mechanistic understanding and offer more general opportunities for enhanced nanoprecipitation processes.

The polymers were synthesised using previously reported approaches,<sup>19–21</sup> namely copper-catalysed methanolic atom transfer radical polymerisation at 30 °C, initiated using a new generation 2 ( $G_2$ ) dendron initiator, **1**, and a monomer mixture of 2-hydroxypropyl methacrylate, **2** (HPMA), and ethylene glycol dimethacrylate, **3** (EGDMA) in a 1 : 2 : 3 molar ratio of 1 : 50 : 0.8 (target number average degree of polymerisation for each primary chain of 50 monomer units). This ratio also ensures <1 brancher per primary chain and avoids gelation. The number average molecular weight ( $M_n$ ) and weight average molecular weight ( $M_w$ ) of the resulting branched polymer ( $G_2$ - $p$ (HPMA<sub>50</sub>-co-EGDMA<sub>0.8</sub>)) were determined by triple detection size exclusion chromatography (DMF/0.01 M LiBr eluent; SEC) to be 115 700 g mol<sup>-1</sup> and 1 538 000 g mol<sup>-1</sup> respectively (dispersity ( $D$ ) = 13.3; Fig. 1). The equivalent linear polymerisation in the absence of **3** yielded a polymer ( $G_2$ - $p$ (HPMA<sub>50</sub>)) of  $M_n$  = 12 300 g mol<sup>-1</sup> and  $M_w$  = 15 500 g mol<sup>-1</sup> ( $D$  = 1.26). Branching with **3** does not impact the formation of primary chains;<sup>24</sup> therefore, it is correct to consider the weight average structures within the  $G_2$ - $p$ (HPMA<sub>50</sub>-co-EGDMA<sub>0.8</sub>) as comprising approximately 100 conjoined  $G_2$ - $p$ (HPMA<sub>50</sub>) chains.

This is highly important as  $M_w$  represents the molecular weight of macromolecules at the mean of the weight fraction and, therefore, approximately half of the physical mass of the sample comprises branched structures with a minimum



**Fig. 1** SEC (DMF/0.01 M LiBr eluent) refractive index chromatograms of hyperbranched polydendron samples before (solid black line) and after (red and blue dashed lines) dialysis using either (A) 2-propanol or (B) methanol as the dialysis solvent. Tubing with two different molecular weight cutoffs (MWCO) were utilized in each solvent condition.

of 100 conjoined chains; the  $M_n$  value indicates a large number of linear and lesser branched structures, as expected.

$G_2$ - $p$ (HPMA<sub>50</sub>-co-EGDMA<sub>0.8</sub>) was fractionated using alcoholic dialysis (3 days) with either methanol or 2-propanol, and tubing with two nominal molecular weight cut-off values (MWCO = 100 kg mol<sup>-1</sup> and 300 kg mol<sup>-1</sup>). Two good solvents for  $p$ (HPMA) were studied due to the potential for variable wetting and pore swelling within the tubing; outer solutions were collected and fully replenished each day. Solutions within the dialysis tubing, and the combined solvent fractions collected from the reservoir, were dried and the recovered polymer studied by triple detection SEC (Fig. 1 and Table 1).

The different MWCO dialysis tubing had a minimal impact on the separation, presumably due to non-aqueous solvent use;



**Table 1** SEC analysis of  $G_2$ - $p$ (HPMA<sub>50</sub>-EGDMA<sub>0.8</sub>) hyperbranched poly-dendrons fractions after alcoholic dialysis under varying conditions

| Dialysis solvent (tubing MWCO) | Recovery site         | $G_2$ - $p$ (HPMA <sub>50</sub> - <i>co</i> -EGDMA <sub>0.8</sub> ) |                                     |      |
|--------------------------------|-----------------------|---|-------------------------------------|------|
|                                |                       | $M_n^a$ (SEC, g mol <sup>-1</sup> )                                 | $M_w^a$ (SEC, g mol <sup>-1</sup> ) | $D$  |
| —                              | —                     | 115 700   | 1 538 000                           | 13.3 |
| 2-Propanol                     | —                     | —   | —                                   | —    |
| 100 kg mol <sup>-1</sup>       | External <sup>b</sup> | 13 300  | 23 400                              | 1.75 |
|                                | Internal <sup>c</sup> | 178 500   | 1 148 000                           | 6.44 |
| 300 kg mol <sup>-1</sup>       | External <sup>b</sup> | 14 000  | 21 700                              | 1.55 |
|                                | Internal <sup>c</sup> | 136 400   | 955 800                             | 7.01 |
| Methanol                       | —                     | —   | —                                   | —    |
| 100 kg mol <sup>-1</sup>       | External <sup>b</sup> | 28 300  | 77 500                              | 2.74 |
|                                | Internal <sup>c</sup> | 761 200   | 2 326 000                           | 3.06 |
| 300 kg mol <sup>-1</sup>       | External <sup>b</sup> | 25 400  | 99 500                              | 3.92 |
|                                | Internal <sup>c</sup> | 615 800   | 1 990 000                           | 3.23 |

<sup>a</sup> Triple detection SEC using DMF/0.01 M LiBr eluent. <sup>b</sup> Polymer recovered from solution surrounding dialysis tubing. <sup>c</sup> Polymer recovered from inside dialysis tubing. See ESI for dn/dc values.

SEC chromatograms overlaid almost exactly (Fig. 1) and calculated  $M_n$  and  $M_w$  values were very similar (Table 1). Methanolic dialysis allowed enhanced isolation of the high molecular weight fraction with the lowest dispersity values.

Nanoprecipitation of the eight fractions (Table 1) and the linear-dendritic hybrid  $G_2$ - $p$ (HPMA<sub>50</sub>) was conducted from THF solutions (5 mg mL<sup>-1</sup>) using deionised water as the anti-solvent and targeting final nanoparticle concentrations of 1 mg mL<sup>-1</sup> after THF removal. Dynamic light scattering analysis (DLS; Table 2) of the unfiltered samples yielded  $z$ -average diameter ( $D_z$ ) values of 620–770 nm for nanoprecipitates of  $G_2$ - $p$ (HPMA<sub>50</sub>) and the least branched materials fractionated using 2-propanol; similar methanol dialysis fractions generated significant macro-phase separated material. When the high molecular weight fractions isolated using 2-propanol or methanol were nanoprecipitated under identical conditions, considerably more uniform size distributions were observed with  $D_z$  values ranging from 135–205 nm and polydispersity (PDI) values ranging from 0.068–0.095 (Table 2).

To investigate the role of the very high molecular weight species in directing the nanoprecipitation process of highly disperse samples, the high molecular weight fraction of  $G_2$ - $p$ (HPMA<sub>50</sub>-*co*-EGDMA<sub>0.8</sub>) collected using the 100 kg mol<sup>-1</sup> MWCO tubing and methanolic dialysis ( $G_2$ - $p$ (HPMA<sub>50</sub>-*co*-EGDMA<sub>0.8</sub>)<sub>MeOH</sub>) was systematically added at varying weight ratios to the  $G_2$ - $p$ (HPMA<sub>50</sub>) sample prior to nanoprecipitation. This fraction was selected as it possessed the highest  $M_n$  and  $M_w$  values and the lowest  $D$  value of the high molecular weight fractions, thereby allowing the most extreme case of molecular weight and architectural diversity after mixing with  $G_2$ - $p$ (HPMA<sub>50</sub>). Nanoprecipitation of a  $G_2$ - $p$ (HPMA<sub>50</sub>)/ $G_2$ - $p$ (HPMA<sub>50</sub>-*co*-EGDMA<sub>0.8</sub>)<sub>MeOH</sub> mixture containing just 1 wt% branched material generated significantly lower  $D_z$  and PDI values than those seen for  $G_2$ - $p$ (HPMA<sub>50</sub>) alone; 198 nm (PDI = 0.034) versus 670 nm (PDI = 0.215) respectively (Fig. 2A).

Further addition of  $G_2$ - $p$ (HPMA<sub>50</sub>-*co*-EGDMA<sub>0.8</sub>)<sub>MeOH</sub> led to a range of nanoparticles with steadily decreasing  $D_z$  values (PDI = 0.034–0.110), with the lowest value (145 nm) achieved from nanoprecipitation in the absence of linear  $G_2$ - $p$ (HPMA<sub>50</sub>). The impact of highly branched, high molecular weight material

**Table 2** Dynamic light scattering

| Polymer sample (dialysis solvent; tubing MWCO) | Recovery site           | $D_z^a$ (nm) | $D_n^a$ (nm) | PDI   |
|--|-------------------------|--------------|--------------|-------|
| $G_2$ - $p$ (HPMA <sub>50</sub> )              | —                       | 670          | 555          | 0.215 |
| Dialysis fractions                             | —                       | —            | —            | —     |
| 2-Propanol                                     | —                       | —            | —            | —     |
| 100 kg mol <sup>-1</sup>                       | External <sup>b</sup>   | 620          | 525          | 0.214 |
|  | Internal <sup>c</sup>   | 190          | 165          | 0.070 |
| 300 kg mol <sup>-1</sup>                       | External <sup>b</sup>   | 770          | 390          | 0.424 |
|  | Internal <sup>c</sup>   | 205          | 175          | 0.095 |
| Methanol                                       | —                       | —            | —            | —     |
| 100 kg mol <sup>-1</sup>                       | External <sup>b,d</sup> | —            | —            | —     |
|  | Internal <sup>c</sup>   | 145          | 115          | 0.075 |
| 300 kg mol <sup>-1</sup>                       | External <sup>b,d</sup> | —            | —            | —     |
|  | Internal <sup>c</sup>   | 135          | 105          | 0.068 |

<sup>a</sup> DLS measured as unfiltered 1 mg mL<sup>-1</sup> aqueous dispersions. <sup>b</sup> Polymer recovered from solution surrounding dialysis tubing. <sup>c</sup> Polymer recovered from inside dialysis tubing. <sup>d</sup> Significant macro-phase separation.

within the nanoprecipitation of linear  $G_2$ - $p$ (HPMA<sub>50</sub>) can be rationalised by consideration of the nucleation process in the mixed solvent/anti-solvent environment. In the absence of branched polymer, the diffusion of good solvent will lead to the media becoming increasingly less able to solvate the linear-dendritic hybrid polymer chains; these will collapse to form small particles comprising single or small numbers of collapsed macromolecules. It is unlikely that these small nuclei will exceed the critical radius ( $r^*$ ), and slow assembly of chains will lead to poorly defined nucleation and growth stages (Fig. 2B); clearly distinct nucleation and growth are key to the formation of narrow size distributions.<sup>15</sup>

Conversely, the population of unimolecular structures comprising > 100 conjoined chains will lead to the rapid formation of considerably larger nuclei, many of which would be expected to exceed  $r^*$  and allow a distinct nucleation stage; the linear-dendritic polymer would require a considerable number of single chains to coalesce in a rapid and concerted multi-body collision to achieve the same outcome. We recently reported a lack of discernible light scattering immediately after addition of linear polymer solutions to anti-solvents, in contrast to the observed strong scattering from branched polymers under identical conditions; this would support the absence of rapid nucleation in lower molecular weight linear materials.<sup>23</sup> Fractionation to generate  $G_2$ - $p$ (HPMA<sub>50</sub>-*co*-EGDMA<sub>0.8</sub>)<sub>MeOH</sub> also increased the number of conjoined chains within the weight average branched structure to > 150 (number average structures contain > 60 chains) (Table 1).

Decreasing  $D_z$  values with increasing content of  $G_2$ - $p$ (HPMA<sub>50</sub>-*co*-EGDMA<sub>0.8</sub>)<sub>MeOH</sub> (Fig. 2A) supports a nucleation-driven mechanism as increased nucleation density from additional highly branched polymer will lead to a larger number of smaller nanoparticles at constant polymer mass.<sup>25</sup>

The number of nanoparticles formed can be approximated by assuming the polymer density and utilising the number average diameter ( $D_n$ ) values (Table S1, ESI<sup>†</sup>), and it is important to note that the number of chains and chain ends does not vary throughout this study. Previous studies have shown that increasing polymer molecular weight alone will lead to little difference in  $D_z$  values within nanoprecipitations of linear polymer systems, despite considerable variation in solution viscosity,



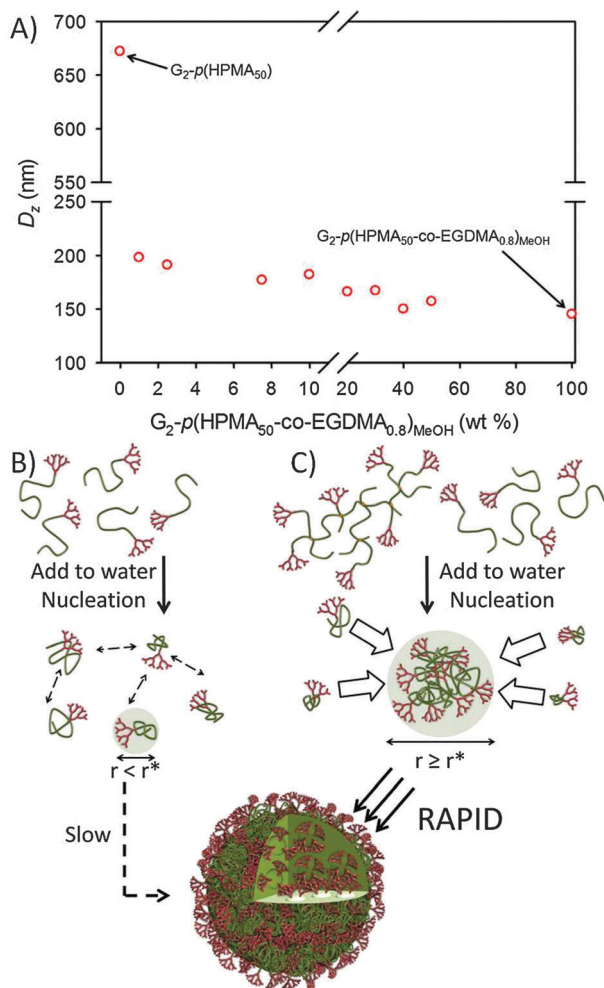


Fig. 2 Effect of addition of highly branched, high molecular weight  $G_2-p(\text{HPMA}_{50}\text{-co-EGDMA}_{0.8})_{\text{MeOH}}$  to  $G_2-p(\text{HPMA}_{50})$ . (A) Observed modification of nanoprecipitate  $D_z$  (DLS) with increasing branched polymer; schematic representation of nucleation from (B) linear-dendritic hybrid polymers with slow growth stage and (C) hyperbranched polydendron generating large nucleus and rapid growth.

with increased particle sizes resulting for some polymers.<sup>26–28</sup> The effects seen here are, therefore, clearly driven by the number of conjoined chains and the architecture of the high molecular weight fraction.

The ability to direct nanoprecipitation outcomes by introducing small fractions of highly branched polymers may have considerable value in ensuring consistency in medical products, decreasing complexity of mixing processes and in controlling nanoparticle sizes and polydispersities. To achieve such results with as little as 1 wt% of branched polymer opens significant opportunities,

offers simplification within large scale processes and provides mechanistic insight.

The authors gratefully acknowledge financial support from the University of Liverpool and the Engineering and Physical Sciences Research Council for a PhD studentship (FL) and grant funding (EP/I038721/1) that underpinned the research.

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