Primordial mass segregation in simulations of star formation?

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Accepted 2014 November 11. Received 2014 November 11; in original form 2014 August 4

ABSTRACT
We take the end result of smoothed particle hydrodynamics (SPH) simulations of star formation which include feedback from photoionization and stellar winds and evolve them for a further 10 Myr using N-body simulations. We compare the evolution of each simulation to a control run without feedback, and to a run with photoionization feedback only. In common with previous work, we find that the presence of feedback prevents the runaway growth of massive stars, and the resulting star-forming regions are less dense, and preserve their initial substructure for longer. The addition of stellar winds to the feedback produces only marginal differences compared to the simulations with just photoionization feedback. We search for mass segregation at different stages in the simulations; before feedback is switched on in the SPH runs, at the end of the SPH runs (before N-body integration) and during the N-body evolution. Whether a simulation is primordially mass segregated (i.e. before dynamical evolution) depends extensively on how mass segregation is defined, and different methods for measuring mass segregation give apparently contradictory results. Primordial mass segregation is also less common in the simulations when star formation occurs under the influence of feedback. Further dynamical mass segregation can also take place during the subsequent (gas-free) dynamical evolution. Taken together, our results suggest that extreme caution should be exercised when interpreting the spatial distribution of massive stars relative to low-mass stars in simulations.

Key words: methods: numerical – stars: formation – stars: kinematics and dynamics – open clusters and associations: general.

1 INTRODUCTION
Understanding how and where stars form is one of the central pillars of astrophysics; star-forming regions either form bound clusters (Lada & Lada 2003; Kruijssen 2012, and references therein) or (more usually) disperse into the Galactic disc and they are also the environment in which planetary systems are believed to form (e.g. Haisch, Lada & Lada 2001).

Hydrodynamical simulations (e.g. Bonnell, Clark & Bate 2008; Offner, Hansen & Krumholz 2009; Girichidis et al. 2011), radiation–hydrodynamical simulations (e.g. Peters et al. 2010; Bate 2012; Dale, Ercolano & Bonnell 2012; Hansen et al. 2012; Krumholz, Klein & McKee 2012), and radiation–magnetohydrodynamical simulations (e.g. Myers et al. 2014) of star formation make predictions for the outcome of star formation in dense, or clustered, environments. Ideally, we would like to compare the outcome of simulations of star formation to observations of young star-forming regions to search for similarities in stellar mass functions (Bonnell et al. 1997; Bate 2009; Krumholz et al. 2012), binary and multiplicity properties (Delgado-Donate et al. 2004; Goodwin, Whitworth & Ward-Thompson 2004; Offner et al. 2010; Bate 2012), and in the spatial distributions of stars (Schmeja & Klessen 2006; Girichidis et al. 2012), including mass segregation (Moeckel & Bonnell 2009; Maschberger & Clarke 2011; Kirk, Offner & Redmond 2014; Myers et al. 2014).

One drawback of hydrodynamical simulations is that they do not follow the full dynamical evolution of a star-forming region, either until it forms a cluster, or disperses altogether. This can be remedied slightly by evolving the simulation using a pure N-body code (e.g. Moeckel & Bate 2010; Moeckel et al. 2012; Parker & Dale 2013). Whilst this approach cannot accurately model the gas left over from star formation, recent studies (e.g. Offner et al. 2009; Smith et al. 2011; Kruijssen et al. 2012) suggest that the removal of gas does not strongly affect the subsequent evolution of the cluster, due to high local star formation efficiencies (SFE). For this reason, the classical picture of a cluster becoming unbound following gas removal (Tutukov 1979; Lada, Margulis & Dearborn 1984; Goodwin & Bastian 2006) may not be valid.

In a previous paper (Parker & Dale 2013), we took the outcome of five pairs of smoothed particle hydrodynamics (SPH) simulations and evolved them forward in time using an N-body integrator.

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In each pair, one simulation formed stars under the influence of photoionization feedback, and the other was a control run without feedback. The differences in the SPH calculation were largely limited to differences in the mass functions; the run without feedback formed fewer stars, but they had higher masses and higher stellar densities. The runs with feedback were less dense, and as a result the clusters that formed retained structure for longer, due to their longer relaxation times (Parker & Dale 2013).

In this paper, we take recent simulations by Dale et al. (2014) which include a further source of feedback – namely stellar winds as well as photoionization feedback – and follow their dynamical evolution for a further 10 Myr using $N$-body simulations. As in Parker & Dale (2013), we determine the evolution of their spatial distributions, local stellar density, and fraction of bound stars. We also look for mass segregation, both before and during the subsequent $N$-body evolution. Recently, Kirk et al. (2014) and Myers et al. (2014) have uncovered evidence for primordial mass segregation in their hydrodynamical and radiation–magnetohydrodynamic simulations, respectively – see also Maschberger & Clarke (2011) who find a similar result in an analysis of the Bonnell et al. (2008) simulation of star formation.

Observationally, mass segregation has been found in some young star clusters (e.g. the Orion nebula Cluster – Hillenbrand & Hartmann 1998; Allison et al. 2009a), but not in other regions, including those that contain massive (>8 M$_\odot$) stars (Wright et al. 2014) and those that do not (Kirk & Myers 2011; Parker et al. 2011; Parker, Maschberger & Alves de Oliveira 2012). This begs the question of whether mass segregation is likely to be a primordial outcome of star formation (Bonnell & Davies 1998; Kirk et al. 2014; Myers et al. 2014), whether it is dynamical (Allison et al. 2009b, 2010; Parker et al. 2014), or some combination of the two (Mocek & Bonnell 2009). If none, or very little, mass segregation occurs in simulations of massive star formation, then the most likely scenario is likely to be that it is a predominantly dynamical process.

The paper is organized as follows. In Section 2, we provide a description of the SPH simulations from Dale et al. (2012), Dale, Ercolano & Bonnell (2013), Dale et al. (2014), and the set-up of the subsequent $N$-body simulations. We describe our results in Section 3, we provide a discussion in Section 4, and we conclude in Section 5.

## 2 INITIAL CONDITIONS

In Parker & Dale (2013), we used as our starting conditions the results of SPH simulations of star formation in a parameter space of molecular clouds presented in Dale et al. (2012, 2013). In these simulations, the influence of photoionizing radiation from O-type stars was included according to the algorithm presented in Dale, Ercolano & Clarke (2007) and Dale et al. (2012). Dale et al. (2012, 2013) also present a control run of each simulation in which feedback was switched off. The SPH study has since been extended to include momentum-driven stellar winds as described in Dale & Bonnell (2008), in addition to photoionization. The simulations are terminated after feedback from the O-type stars has been active for 3 Myr, since this is the time when these stars will begin to expire as supernovae.

The principal results of the new SPH study are described in Dale et al. (2014). In general, the additional influence of winds on top of photoionization is modest, except at very early times, when the winds aid the expanding H II regions in clearing dense gas from the deep potential wells in which the O-stars are situated. Nevertheless, there are differences in the numbers and distributions of stars formed in the dual-feedback simulations when compared to the ionization-only runs.

In Table 1, we summarize the results from the SPH studies, including the simulations with stellar winds from the new study by Dale et al. (2014). We list each simulation triplet; the run without feedback first (‘a’), the run with photoionization feedback only (‘b’), and the run with photoionization and stellar winds (‘c’). For each triplet, we list the initial cloud virial ratio, $\alpha_\text{vir}$, cloud mass, $M_\text{cloud}$, the number of stars formed at the end of each simulation, $N_\text{stars}$, the final stellar mass in the simulation, $M_\text{region}$, and the spatial structure.

### Table 1. A summary of the five different triplets of SPH simulations used as the input initial conditions of our $N$-body integrations.

The values in the columns are: the simulation number, the corresponding Run ID from Dale et al. (2012, hereafter D12), Dale et al. (2013, hereafter D13), or Dale et al. (2014, hereafter D14), the type of feedback in the SPH simulation (none, photoionization only, or photoionization and stellar winds), the paper reference, the initial virial ratio of the original clouds $\alpha_\text{init}$ (to distinguish bound from unbound clouds), the initial radius of the cloud in the SPH simulation ($R_\text{cloud}$), the initial mass of the cloud ($M_\text{cloud}$), the number of stars that have formed at the end of the SPH simulation ($N_\text{stars}$), the mass of this star-forming region ($M_\text{region}$), the $Q$-parameter in the SPH simulation at the time feedback is initiated in the feedback runs ($Q_\text{init}$), and the final $Q$-parameter in the SPH simulations ($Q_\text{fin}$).

<table>
<thead>
<tr>
<th>Sim. no.</th>
<th>Run ID</th>
<th>Feedback</th>
<th>Ref.</th>
<th>$\alpha_\text{SPH}^{\text{init}}$</th>
<th>$R_\text{cloud}$</th>
<th>$M_\text{cloud}$</th>
<th>$N_\text{stars}$</th>
<th>$M_\text{region}$</th>
<th>$Q_\text{SPH}^{\text{init}}$</th>
<th>$Q_\text{SPH}^{\text{fin}}$</th>
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<td>578</td>
<td>3207 M$_\odot$</td>
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</tr>
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<td>Photoionization</td>
<td>D12</td>
<td>0.7</td>
<td>5 pc</td>
<td>10 000 M$_\odot$</td>
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<td>2205 M$_\odot$</td>
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<tr>
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<td>Photoionization + wind</td>
<td>D14</td>
<td>0.7</td>
<td>5 pc</td>
<td>10 000 M$_\odot$</td>
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<td>2186 M$_\odot$</td>
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<tr>
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<td>77</td>
<td>594 M$_\odot$</td>
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</tr>
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</table>
as measured by the \( Q \)-parameter (Cartwright & Whitworth 2004) at the point feedback was switched on, and at the end of the SPH simulation.

### 2.1 N-body evolution

We take the final states of five triplets of simulations from Dale et al. (2012), Dale et al. (2013), and Dale et al. (2014) and assume that the combination of the first supernova and stellar winds instantaneously removes any remaining gas both from the feedback and non-feedback simulations, and evolve the resulting gas-free systems with an N-body code.

We evolve the clusters using the fourth-order Hermite-scheme integrator \texttt{kira} within the \textsc{starlab} environment (e.g. Portegies Zwart et al. 1999, 2001). We take the masses, positions, and velocities of the sink particles from the SPH simulations and place these directly into the N-body integrator. In the majority of the SPH runs, the stars are in virial equilibrium, or slightly subvirial, at the end of the simulation (i.e. the initial conditions for the N-body integrator). However, the initial conditions for simulations UF, UP, and UQ were globally unbound, so that one might expect the stars and clusters formed to be in an unbound configuration. In fact, some parts of the globally unbound clouds become bound due to high-velocity gas flows colliding and radiating away kinetic energy, which tends to occur in the dense areas of the clouds where most of the stars form. Therefore, in practice, these unbound clouds form stars that are roughly in virial equilibrium (with virial ratios ranging from 0.4–0.7), apart from Run UF (with feedback), which is highly supervirial, with a virial ratio of 1.9.

The simulation triplets are then evolved for 10 Myr, without a background gas potential. The simulations contain several stars with masses >20 \( \text{M}_\odot \) which are likely to evolve over the 10 Myr duration of the N-body integration. For this reason, we use the \texttt{SeBa} stellar evolution package in the \textsc{starlab} environment (Portegies Zwart & Verbunt 1996, 2012), which provides look-up tables for the evolution of stars according to the time-dependent mass–radius relations in Eggleton, Fitchett & Tout (1989) and Tout et al. (1996). Typically, \texttt{SeBa} updates the evolutionary status of stars on shorter time-scales than the timestep in the \texttt{kira} integrator, although for extremely close systems a lag of up to one timestep can occur.

### 3 RESULTS

In this section, we describe the N-body evolution of spatial structure, local surface density, and the fraction of bound stars, before using three different methods to search for mass segregation. A summary of the evolution of structure, density, and bound fraction, as well as the mass-loss due to stellar evolution, is presented in Table 2.

Dale et al. (2012, 2013) noted that the absence of feedback in the SPH calculations results in a more top-heavy initial mass function, as the most massive stars do not have their growth regulated by feedback. Photoionization feedback reduces the mass of the star-forming region, and increases the number of stars formed. In general, the addition of stellar winds slightly reduces the number of stars compared to photoionization feedback alone, and also slightly reduces the mass of the region (although there are exceptions, such as Run UF). In all cases, the photoionization+wind models reduce the initial stellar surface density with respect to the runs with photoionization feedback only, and those with no feedback at all.

#### 3.1 Evolution of spatial structure

We examine the evolution of the structure of the simulated regions over the duration of the N-body integration, using the \( Q \)-parameter. The \( Q \)-parameter was pioneered by Cartwright & Whitworth (2004) and Cartwright (2009) and combines the normalized mean edge length of the minimum spanning tree (MST) of all the stars in the region, \( \bar{m} \), with the normalized correlation length between all stars in the region, \( \bar{s} \). The level of substructure is determined by the following equation:

\[
Q = \frac{\bar{m}}{\bar{s}}.
\]
A substructured association or region has $Q < 0.8$, whereas a smooth, centrally concentrated cluster has $Q > 0.8$. The $Q$-parameter has the advantage of being independent of the density of the star-forming region, and purely measures the level of substructure present. The original formulation of the $Q$-parameter assumes the region is spherical, but can be altered to take into account the effects of elongation (Bastian et al. 2009; Cartwright & Whitworth 2009).

In Fig. 1, we compare the evolution of the $Q$-parameter with time in three of the five triplets of simulations (we omit Runs UP and UQ from the plot for clarity). The simulations that formed with feedback are shown by the solid lines; the simulations with ionization feedback only (presented in Parker & Dale 2013) are shown by the thinner lines, and those with ionization feedback and stellar winds are shown by the thicker lines. Simulations with no feedback are shown by the dashed lines. The red lines are Runs J, green lines are Runs I, dark blue lines are Runs UF. Runs with feedback preserve structure for longer as the star-forming region evolves, and two simulations with feedback (I and UF) remain substructured throughout.

The addition of the extra feedback mechanism (stellar winds) does not alter the evolution of the $Q$-parameter significantly with respect to the runs with ionization feedback only, and the results are very similar to those reported in Parker & Dale (2013), namely that star-forming regions which form without feedback lose their structure faster than regions that form with feedback.

### 3.2 Surface densities

Parker & Dale (2013) noted that the regions that form without feedback lose structure faster than their feedback-influenced counterparts due to their higher initial stellar densities. Without the regulating influence of feedback, regions form with slightly higher initial densities and hence shorter local crossing times, which leads to more interactions and the more rapid loss of substructure.

The difference in initial density between the runs with and without feedback is readily apparent when examining the median local surface density. We calculate the local stellar surface density following the prescription of Casertano & Hut (1985), modified to account for the analysis in projection. For an individual star, the local stellar surface density is given by

$$\Sigma = \frac{N - 1}{\pi r_N^2},$$

where $r_N$ is the distance to the $N$th nearest neighbouring star (we adopt $N = 10$ throughout this work; lower $N$ values could bias $\Sigma$ to higher values due to binaries, and higher $N$ values would remove the ‘localness’ from the determination).

In Fig. 2, we show the evolution of the median stellar surface density, $\Sigma_{\text{all}}$, again for the evolution of Runs J, I, and UF. When a region is substructured, it is usually meaningless to define a ‘central’ or ‘core’ density, and Parker & Dale (2013) show that a much better tracer of the true density of a region is the median stellar surface density. The evolution of the runs that form without feedback are shown by the dashed lines, the runs that form with ionization feedback only are shown by the thin solid lines, and the runs that form with ionization feedback and stellar winds (Dale et al. 2014) are shown by the thick lines.

In all sets of simulations, the regions that form without feedback have higher median densities than the simulations that form with feedback. It is these higher densities that facilitate the erasure of...
substructure, as discussed in Parker & Dale (2013) and Parker et al. (2014).

3.3 Bound/unbound stars

We track the number of stars that remain bound as a function of time by calculating the kinetic and potential energies for each star, as detailed in Baumgardt, Hut & Heggie (2002) and Kruijssen et al. (2012). The potential energy of an individual star, $V_i$, is given by

$$V_i = -\sum_{i \neq j} \frac{G m_i m_j}{r_{ij}},$$

where $m_i$ and $m_j$ are the masses of two stars and $r_{ij}$ is the distance between them. The kinetic energy of a star, $T_i$, is given thus:

$$T_i = \frac{1}{2} m_i |\mathbf{v}_i - \mathbf{v}_{ci}|^2,$$

where $\mathbf{v}_i$ and $\mathbf{v}_{ci}$ are the velocity vectors of the star and the centre of mass of the region, respectively. A star is bound if $T_i + V_i < 0$. In Fig. 3, we show the number fraction of stars that remain bound over the N-body integration for three of the five sets of simulations. Again, the simulations that formed with feedback are shown by the solid lines; the simulations that include ionization feedback only are shown by the thinner lines, and the simulations with ionization and stellar winds are shown by the thicker lines. The simulations that formed without feedback are shown by the dashed lines. The colours correspond to the following simulations; red – Run J, green – Run I, dark blue – Run UF.

Simulations that form without feedback have a higher fraction of bound stars at the end of the SPH runs (possibly due to the higher total mass), but the subsequent N-body evolution does not follow a distinct evolutionary path. For example, in Run J without feedback, the final bound fraction of $f_{\text{bound,J}}$ is lower than for the simulations which include feedback. In the other runs, $f_{\text{bound,J}}$ tends to be higher for the simulations without feedback. In the case of Run J, the high initial stellar density (4518 stars pc$^{-2}$) has led to subsequent dynamical interactions unbinding a large fraction of the stars and hence drastically lowered $f_{\text{bound,J}}$ compared to the simulations with feedback.

3.4 Mass segregation

Defining mass segregation has become increasingly difficult due to the many disparate methods which have been promoted in the recent literature. As we will see, different methods which claim to measure mass segregation may actually give very contradictory results.

Classically, mass segregation is a signature of the onset of energy equipartition in star clusters, whereby the most massive stars have slower velocities and hence sink to the centre (Spitzer 1969). In this scenario, the cluster is dynamically old, centrally concentrated, and hence has a well-defined radial profile. One can then take different mass bins and compare the density profiles (e.g. Hillenbrand 1997; Pinfield, Jameson & Hodgkin 1998), or look for variations in the slope of the mass function (or luminosity function) with distance from the cluster centre (Carpenter et al. 1997; de Grijs et al. 2002; Gouliermis et al. 2004). A related method is to quantify the variation of the ‘Spitzer radius’ – the rms distance of stars in a cluster around the centre of mass – with luminosity (Gouliermis, de Grijs & Xin 2009).

These methods all require the definition of the cluster centre. This, and the choice of binning can lead to complications and misinterpretation in the data (Ascenso, Alves & Lago 2009). Furthermore, the two main avenues of massive star formation – competitive accretion (Zinnecker 1982; Bonnell, Bate & Zinnecker 1998) and monolithic collapse (McKee & Tan 2003; Krumholz, McKee & Klein 2005) – both predict that the most massive stars should be more centrally located than the lower mass stars, so-called primordial mass segregation. However, as star formation typically occurs in filaments (e.g. Arzoumanian et al. 2011), which usually leads to a hierarchical or substructured spatial distribution of stars, then it becomes almost impossible to define the centre of a star-forming region in order to quantify mass segregation.

In order to avoid the need for defining a centre, several methods have been proposed which compare MSTs of groups of stars (Allison et al. 2009a) or local surface density around all stars (Maschberger & Clarke 2011). Another recently proposed technique is to use an MST to define groups of stars, and then determine whether the most massive stars are closer to the centre of the group than the average stars (Kirk & Myers 2011; Kirk et al. 2014) – the centre is defined as the median position of all stars in the group. We use three of these methods to search for (primordial) mass segregation in the SPH simulations from Dale et al. (2012, 2013, 2014) and in the subsequent N-body evolution (dynamical mass segregation).

3.4.1 Group segregation ratio, $\Omega_{\text{GSR}}$

First, we use the method of Kirk & Myers (2011) and Kirk et al. (2014), in which an MST is constructed for the entire region (see Fig. 4a). The cumulative distribution of all MST branch lengths is then made (Fig. 4b). Two power-law slopes are then fitted to the shortest lengths, and the longest lengths, and the intersection of these slopes defines the boundary of subclustering, $d_{\text{break}}$ (Gutermuth et al. 2009). In Fig. 4(c), the MST of the region is shown, but we have omitted the MST lengths greater than the critical length $d_{\text{break}}$. 

Figure 3. Evolution of the number fraction of bound stars for three of the five triplets of simulations (we omit runs UP and UQ from the plot for clarity, but their behaviour follows the same pattern as runs UF). Simulations with feedback are shown by the solid lines; those with ionization feedback only (presented in Parker & Dale 2013) are shown by the thinner lines, and those with ionization feedback and stellar winds are shown by the thicker lines. Simulations with no feedback are shown by the dashed lines. The red lines are Runs J, green lines are Runs I, and the dark blue lines are Runs UF. There is no correlation between the evolution of the fraction of bound stars and the initial conditions (i.e. feedback versus no feedback).
Mass segregation in star formation?

Figure 4. Mass segregation analysis of stellar groups (as in Kirk & Myers 2011 and Kirk et al. 2014) in the N-body simulation of Run I without feedback at 1 Myr. In panel (a), the MST of the entire star-forming region is shown. In panel (b), the distribution of MST branch lengths is shown, with the power-law fits to the small branches (steep slope) and the long branches (shallow slope) as defined in Gutermuth et al. (2009), Kirk & Myers (2011), and Kirk et al. (2014). The chosen break length \(d_{\text{break}}\) (which defines the groups) is the intersection of these two slopes. In panel (c), the MST lengths exceeding \(d_{\text{break}}\) have been removed, and there are 10 groups containing more than two stars (one is outside the field of view in the plot), and two groups containing more than 10 stars. The 10 most massive stars in the entire simulation are shown by the blue circles, and the most massive star in each of the 10 groups is shown by a red triangle. In panel (d), for each group we show the ratio of the most massive star to the median stellar mass versus the ratio of the distance from the group centre of the most massive star to the group median. The red asterisk symbols are the groups containing 10 or more stars.

The location of the most massive star in each group is shown by the red triangle, and we also show the locations of the 10 most massive stars for the entire region by the large blue circles. We then follow the method described in Kirk & Myers (2011) and Kirk et al. (2014) and look for mass segregation within the groups. If the position of the most massive star in the group \(r_{\text{mm}}\) is closer to the central position than the median value for stars, \(r_{\text{med}}\), the subcluster is said to be mass segregated. This is shown in Fig. 4(d), where we plot the ratio of the highest mass to the median stellar mass in the subcluster against the offset ratio, \(r_{\text{mm}}/r_{\text{med}}\).

In Fig. 4(d), we also distinguish between groups with \(N > 2\) stars, shown by the black crosses, and groups with \(N \geq 10\) shown by the red asterisks. According to the definition in Kirk & Myers (2011) and Kirk et al. (2014), both subclusters with \(N \geq 10\) are mass segregated, because their offset ratios are less than unity. We define a ‘group segregation ratio’, \(\Omega_{\text{GSR}}\), as

\[
\Omega_{\text{GSR}} = \frac{N_{\text{seg}}}{N_{\text{grp}}},
\]

where \(N_{\text{grp}}\) are the number of groups, and \(N_{\text{seg}}\) is the number of these groups that have an offset ratio less than unity. The snapshot shown in Fig. 4 has \(\Omega_{\text{GSR}} = 1\) for groups with \(N \geq 10\), i.e. all of these groups are mass segregated.

3.4.2 Mass segregation ratio, \(\Lambda_{\text{MSR}}\)

We then use the \(\Lambda_{\text{MSR}}\) mass segregation ratio pioneered by Allison et al. (2009a). We find the MST of the \(N_{\text{MST}}\) stars in the chosen...
subset and compare this to the MST of sets of \( N_{\text{MST}} \) random stars in the region. If the length of the MST of the chosen subset is shorter than the average length of the MSTs for the random stars, then the subset has a more concentrated distribution and is said to be mass segregated. Conversely, if the MST length of the chosen subset is longer than the average MST length, then the subset has a less concentrated distribution, and is said to be inversely mass segregated (see e.g. Parker et al. 2011). Alternatively, if the MST length of the chosen subset is equal to the random MST length, we can conclude that no mass segregation is present.

By taking the ratio of the average (mean) random MST length to the subset MST length, a quantitative measure of the degree of mass segregation (normal or inverse) can be obtained. We first determine the subset MST length, \( l_{\text{subset}} \). We then determine the average length of sets of \( N_{\text{MST}} \) random stars each time, \( (l_{\text{average}}) \). There is a dispersion associated with the average length of random MSTs, which is roughly Gaussian and can be quantified as the standard deviation of the lengths \( (l_{\text{average}}) \pm \sigma_{\text{average}} \). However, we conservatively estimate the lower (upper) uncertainty as the MST length which lies 1/6 (5/6) of the way through an ordered list of all the random lengths (corresponding to a 66 per cent deviation from the median value, \( (l_{\text{average}}) \)). This determination prevents a single outlying object from heavily influencing the uncertainty. We can now define the ‘mass segregation ratio’ \( (\Lambda_{\text{MSR}}) \) as the ratio between the average random MST pathlength and that of a chosen subset, or mass range of objects:

\[
\Lambda_{\text{MSR}} = \frac{(l_{\text{average}}) + \sigma_{\text{average}}/\text{subset}}{l_{\text{subset}} - \sigma_{\text{average}}/\text{subset}}.
\]

\( \Lambda_{\text{MSR}} \) of \( \sim 1 \) shows that the stars in the chosen subset are distributed in the same way as all the other stars, whereas \( \Lambda_{\text{MSR}} > 1 \) indicates mass segregation and \( \Lambda_{\text{MSR}} < 1 \) indicates inverse mass segregation, i.e. the chosen subset is more sparsely distributed than the other stars.

In Fig. 5, we show \( \Lambda_{\text{MSR}} \) as a function of the \( N_{\text{MST}} \) stars in the subset for Run I without feedback after 1 Myr of evolution. The four most massive stars are strongly mass segregated, and this also extends to the 10 most massive stars, although the significance is more marginal.

### 3.4.3 Local density ratio, \( \Sigma_{\text{LDR}} \)

The local surface density of massive stars compared to the median surface density of the full region was pioneered by Maschberger & Clarke (2011) as a way of defining mass segregation but minimizing the effects of outliers in the distribution. In this definition, the massive stars have no knowledge of each other, but if their surface density distribution can be shown to be inconsistent with the median surface density distribution of the entire region (by means of a Kolmogorov-Smirnov (KS)test), then the region is said to be mass segregated. In Fig. 6, we show \( \Sigma \) versus \( m \) for every star for Run I without feedback after 1 Myr of \( N \)-body evolution.

Küpper et al. (2011) and Parker et al. (2014) took the ratio of the median surface density of the 10 most massive stars (the red line in Fig. 6) to the region median (the blue dashed line in Fig. 6) to define a ‘local surface density ratio’, \( \Sigma_{\text{LDR}} \):

\[
\Sigma_{\text{LDR}} = \frac{\Sigma_{\text{subset}}}{\Sigma_{\text{all}}}.
\]

The massive stars in this simulation have a higher median surface density than the median value in the region (\( \Sigma_{\text{LDR}} = 3.08 \), and a KS test returns a \( p \)-value of \( 1.35 \times 10^{-2} \) that the two subsets share the same parent distribution).

### 3.4.4 Evolution over 10 Myr

We show the evolution of this simulation (Run I, no feedback) over the full 10 Myr of \( N \)-body evolution in Fig. 7. According to all three measures, this star-forming region has primordial mass segregation. However, the most massive stars evolve and so the subset of the 10 most massive stars in the simulation is not constant. For this reason, the primordial mass segregation disappears according to \( \Lambda_{\text{MSR}} \) and \( \Sigma_{\text{LDR}} \), until dynamical evolution causes a ‘re-segregation’

---

**Figure 5.** Mass segregation as defined by \( \Lambda_{\text{MSR}} \) (Allison et al. 2009a) in the \( N \)-body simulation of Run I without feedback at 1 Myr. \( \Lambda_{\text{MSR}} = 1 \) (i.e. no mass segregation) is indicated by the red horizontal line.

**Figure 6.** The \( \Sigma-m \) plot (Maschberger & Clarke 2011) for the \( N \)-body simulation of Run I without feedback at 1 Myr. The median surface density in the region is 10 stars pc\(^{-2}\) and is shown by the dashed blue line. The median surface density of the 10 most massive stars is 30 stars pc\(^{-2}\), as shown by the solid red line.
3.5 Structure versus mass segregation

For a large set of purely N-body simulations, Parker et al. (2014) showed that plotting spatial structure against mass segregation measurements can distinguish between the initial conditions of star-forming regions. In the following section, we show $Q - \Omega_{GSR}$, $Q - \Lambda_{MSR}$, and then $Q - \Sigma_{LDR}$.

3.5.1 $Q - \Omega_{GSR}$

In Fig. 9, we show the $Q$ parameter plotted against the group mass segregation ratio $\Omega_{GSR}$ for the simulations without feedback (Fig. 9a) and those with ionization and stellar winds (Fig. 9b). We show the data at 0, 2.5, 5, 7.5, and 10 Myr, and we also plot the values from the SPH runs before feedback is switched on, apart from Run UF, which does not contain enough stars at that stage for there to be a group containing 10 or more stars. The colour scheme is as follows; red – Runs J, green – Runs I, dark blue – Runs UF, cyan – Runs UP, and magenta – Runs UQ. (Note that Runs I and UQ both have $Q = 0.42$ and $\Omega_{GSR} = 1$, so only the cyan star symbol is visible in the plot.)

In this plot, $\Omega_{GSR} = 0.5$ (half of the groups are mass segregated) is shown by the vertical dashed line, and the boundary between a substructured and centrally concentrated spatial distribution ($Q = 0.8$) is shown by the horizontal dashed line. Before the feedback mechanisms are switched on, all the groups have their most...
massive member more centrally concentrated than the average star. When no feedback is switched on (Fig. 9a), this behaviour carries forward to the end of the SPH simulation \(t = 0\) Myr in the N-body integration. As the regions (and the massive stars within them) evolve, the mass segregation is gradually lost – so much so that after 10 Myr only one simulation (UQ – the magenta triangle) has \(\Omega_{\text{GSR}} > 0.5\).

Conversely, in the simulations with photoionization and stellar winds, feedback has wiped out the early primordial mass segregation in groups in three out of five simulations [the red (Run J), cyan (Run UP) and magenta (Run UQ) plus signs]. However, the subsequent evolution of these regions leads to most simulations having \(\Omega_{\text{GSR}} > 0.5\) – i.e. most of the groups are segregated in the sense that their most massive member is closer to the group centre than the average star.

### 3.5.2 \(Q - \Lambda_{\text{MSR}}\)

In Fig. 10, we show the \(Q\) parameter plotted against the mass segregation ratio \(\Lambda_{\text{MSR}}\) for the simulations without feedback (Fig. 10a) and those with ionization and stellar winds (Fig. 10b).

As in Fig. 9, we show the data at 0, 2.5, 5, 7.5, and 10 Myr, and we also plot the values from the SPH runs before feedback is switched on. At this point in the SPH run, the number of stars that have already formed can be rather low (<50) and we only plot the SPH measurement for Runs J and UP, which already have enough stars to make the determination of \(\Lambda_{\text{MSR}}\) meaningful. The colour scheme is as follows: red – Runs J, green – Runs I, dark blue – Runs UF, cyan – Runs UP, and magenta – Runs UQ. The large symbols indicate when \(\Lambda_{\text{MSR}}\) deviates significantly from unity (i.e. mass segregation or inverse mass segregation is present).

The example shown in Fig. 7(a) – Run I without feedback – is shown by the green symbols in Fig. 10(a), and the corresponding run with feedback (Fig. 8a) is shown by the green symbols in Fig. 10(b).

Primordial mass segregation as measured by \(\Lambda_{\text{MSR}}\) is found in one simulation (Run I) out of five for the regions without feedback, and in one of five simulations (Run J) with feedback (though not the same simulation). Run J was not mass segregated before feedback was switched on, with \(\Lambda_{\text{MSR}} = 1.2\) – the red star in Fig. 10(a).

Considering the runs that formed without feedback, in addition to the behaviour of Run I which was shown in Fig. 7(a), over 10 Myr of subsequent dynamical evolution Run J (the red symbols) fluctuates between being significantly mass segregated (at 5 and 7.5 Myr), and significantly inversely mass segregated (at 2.5 and 10 Myr). Run UF (the dark blue symbols) dynamically mass segregates and remains so over the full 10 Myr. Run UP (the cyan symbols) dynamically inversely mass segregates after 2.5 Myr and Run UQ (the magenta points) does not significantly mass segregate or inverse mass segregate aside from a brief snapshot at 5 Myr.

The runs that form with feedback in general display no primordial mass segregation. The one simulation that does show primordial mass segregation (Run J) unsegregates, due to stellar evolution – the 10 most massive stars at \(t = 0\) Myr are not the same 10 most massive stars even after only 2.5 Myr. Run UF (the dark blue symbols) dynamically mass segregates until after 7.5 Myr, when dynamical interactions eject two massive stars and the cluster reverts to being unsegregated after 10 Myr. Run UP (the cyan symbols) first becomes inversely mass segregated, before dynamical interactions lead to normal mass segregation.

### 3.5.3 \(Q - \Sigma_{\text{LDR}}\)

In Fig. 11, we show the \(Q\) parameter plotted against the local surface density ratio \(\Sigma_{\text{LDR}}\). Notably, in the simulations without feedback (Fig. 11a), nearly all simulations show primordial mass segregation at \(t = 0\) Myr, whereas none are mass segregated before feedback is switched on in the SPH runs (the star symbols clustered around \(\Sigma_{\text{LDR}} = 1\) in both panels). The subsequent combination of dynamical and stellar evolution generally erases this signature.
Figure 10. Spatial structure $\mathcal{Q}$ versus $\Lambda_{\text{MSR}}$. In panel (a), we show the evolution of all five star-forming regions which form without feedback and in panel (b) we show the evolution of all five regions that form with ionization and stellar wind feedback. The red symbols are Runs J, green symbols are Runs I, dark blue symbols are Runs UF, cyan symbols are Runs UP, and the magenta symbols are Runs UQ. In addition, we show the measurements from the SPH simulations before feedback is switched on (only Runs J and UP have enough stars at that stage to calculate $\Lambda_{\text{MSR}}$). The larger symbols indicate where the deviation of $\Lambda_{\text{MSR}}$ from unity is significant. $\Lambda_{\text{MSR}} = 1$ (no mass segregation) is shown by the vertical dashed line, and the boundary between a substructured and centrally concentrated spatial distribution ($\mathcal{Q} = 0.8$) is shown by the horizontal dashed line.

Figure 11. Spatial structure $\mathcal{Q}$ versus $\Sigma_{\text{LDR}}$. In panel (a), we show the evolution of all five star-forming regions which form without feedback and in panel (b) we show the evolution of all five regions that form with ionization and stellar wind feedback. The red symbols are Runs J, green symbols are Runs I, dark blue symbols are Runs UF, cyan symbols are Runs UP, and the magenta symbols are Runs UQ. In addition, we show the measurements from the SPH simulations before feedback is switched on. The larger symbols indicate where the deviation of $\Sigma_{\text{LDR}}$ from unity is significant. $\Sigma_{\text{LDR}} = 1$ (no mass segregation) is shown by the vertical dashed line, and the boundary between a substructured and centrally concentrated spatial distribution ($\mathcal{Q} = 0.8$) is shown by the horizontal dashed line.

apart from several snapshots in Run J (the red symbols) and Run I (the green symbols).

When feedback is included (Fig. 11b), no simulations have primordial mass segregation according to $\Sigma_{\text{LDR}}$. Furthermore, following dynamical evolution, only one simulation displays a $\Sigma_{\text{LDR}}$ ratio that is significantly higher than unity – Run UP at 7.5 Myr.

The large fraction of simulations without feedback that display a high $\Sigma_{\text{LDR}}$ at the end of the SPH calculation compared to the simulations with feedback suggests that the absence of feedback leads to higher stellar densities and hence enables the massive stars to acquire a retinue of lower mass stars (as seen in pure $N$-body simulations; Parker et al. 2014). However, once stellar evolution is included, the massive stars lose so much mass (Parker & Dale 2013) that the determination of $\Sigma_{\text{LDR}}$ begins to include stars with local densities comparable to the median value in the region.
4 DISCUSSION

We have expanded upon the work in Parker & Dale (2013) and followed the dynamical evolution of five SPH simulations of star formation which also include stellar winds, as well as photoionization feedback, and compare them to control run simulations which have feedback switched off. The inclusion of stellar winds does not appreciably affect the formation of a region any more so than in the simulations which only include photoionization as a feedback source. The results are similar to those in Parker & Dale (2013); regions which form under the influence of feedback have lower stellar densities than those that do not. The systematically lower densities raise the local crossing time in the regions, and this leads to the regions retaining substructure far longer than the regions which are not influenced by feedback.

We also look for mass segregation in both the non-feedback, and the feedback-influenced star-forming regions. Recent analyses of hydrodynamical simulations of star formation by Kirk et al. (2014) and Myers et al. (2014) have claimed to find primordial mass segregation; the massive stars segregate early on in the calculation and remain segregated throughout.

As discussed in Parker et al. (2014) and Parker & Goodwin (in preparation), different methods to search for mass segregation routinely produce apparently contradictory results, especially for complex spatial distributions. For example, the $\lambda_{\text{MSR}}$ ratio (Allison et al. 2009a) measures the spatial distribution of a subset of massive stars compared to random subsets, whereas the $\Sigma_{\text{LDR}}$ ratio (Küpper et al. 2011; Maschberger & Clarke 2011; Parker et al. 2014) measures the surface density around individual massive stars, and compares this to the surface density around average stars. Finally, the method from Kirk & Myers (2011) and Kirk et al. (2014), which we use to define a ‘group segregation ratio’ – $\Omega_{\text{GSR}}$, measures the distance of the most massive star from the centre of a subcluster to the distance of the median mass star from the centre.

$\lambda_{\text{MSR}}$ provides information on the global spatial distribution of massive stars, whereas $\Sigma_{\text{LDR}}$ provides information on the local density around those stars. As shown in Parker & Goodwin (in preparation), the $\Omega_{\text{GSR}}$ method sometimes reflects the local density of massive stars, but is reliant on dividing up a star-forming region into individual groups. It is not clear whether this assumption is valid; for example, in these simulations there is only one star formation ‘event’ and the massive stars – and associated low-mass stars – are not independent of each other.

This is apparent in Figs 7 and 8, where $\lambda_{\text{MSR}}$ and $\Sigma_{\text{LDR}}$ display similar behaviour, whereas $\Omega_{\text{GSR}}$ does not. More often than not, $\Omega_{\text{GSR}}$ shows a region to be mass segregated on local scales when the global distribution is consistent with a random distribution. We therefore suggest that the claim of primordial mass segregation in clusters by Kirk et al. (2014) has been influenced by the method used to define mass segregation and does not reflect the true spatial distribution from the outcome of star formation.

Furthermore, when we examine the results of several SPH simulations, we see primordial mass segregation in some, but not all, simulations (either four in five, or two in five using $\Omega_{\text{GSR}}$, one in five using $\lambda_{\text{MSR}}$, and either none, or four in five using $\Sigma_{\text{LDR}}$ depending on whether feedback was switched on or not). Interestingly, Myers et al. (2014) find mass segregation according to $\Sigma_{\text{LDR}}$ in their magnetohydrodynamical simulations of star formation which include feedback, whereas we only see high $\Sigma_{\text{LDR}}$ ratios in the simulations without feedback.

In order to compare our results to those of Myers et al., we estimate the $\Sigma_{\text{LDR}}$ ratio from their fig. 17 using their median cluster values (the coloured lines in their fig. 17) and the median of the 10 most massive stars as shown in their figure (using all stars with masses $>$1 M$_\odot$ gives almost identical results). In their control (hydro only) run $\Sigma_{\text{LDR}} = 1.4$, in their run with a ‘weak’ magnetic field ($\Sigma_{\text{LDR}} = 4.4$) and in their run with a ‘strong’ magnetic field $\Sigma_{\text{LDR}} = 4.4$. The presence of a magnetic field may have caused mass segregation in this particular simulation, but the level of mass segregation according to $\Sigma_{\text{LDR}}$ does not increase with increasing magnetic field. As we have seen with the simulations from Dale et al. (2012, 2013, 2014), mass segregation can be random, and more than one simulation is required to ascertain whether the addition of extra physics is the root cause of a different spatial distribution for the most massive stars.

Furthermore, the simulations of Myers et al. (2014) form somewhat smaller systems (‘clumps’) than the full cluster simulations of Dale et al. (2012, 2013, 2014), and their feedback mechanisms also differ (for example, they include protostellar outflows but no photoionization feedback or stellar winds). It is possible that this feedback is weaker in terms of regulating accretion flows, and hence the calculations of Myers et al. (2014) may behave more like the control runs from Dale et al. (2012, 2013, 2014), unless magnetic fields are dominant in governing the spatial distribution of stars. In either case, more simulations that include magnetic fields are required to address this question.

This suggests that either the advent of mass segregation in simulations of star formation is a random event, or that the prescriptions of feedback for the initial conditions differ enough to produce significantly different spatial distributions of stars.

In the SPH simulations of Dale et al. (2012, 2013, 2014), the runs without feedback lead to the most massive stars attaining much higher local densities than those with feedback. The reason for this is probably that feedback from the O-stars essentially shuts down accretion on to the main subclusters so they stop growing, both in the sense of the stars they already have not acquiring more mass and in the sense of them having nothing to make new stars with. Therefore, the local SFE and thus local stellar density do not increase as much in the feedback runs as they do in the control runs.

The initial conditions of the SPH runs (in terms of initial cloud mass, density, and virial ratio) do not appear to influence whether mass segregation occurs, and certainly not as much as the presence (or absence) of feedback. If we consider the $\Sigma_{\text{LDR}}$ plot (Fig. 10), in the case without feedback (panel a) Run I is mass segregated, but the more dense version of this simulation (Run J) is not. Conversely, in the presence of feedback (panel b) Run I is not mass segregated, whereas Run J is.

Whilst resolution tests were conducted on some of the SPH simulations presented in Dale et al. (2012), they were mainly directed at showing that the convergence of global properties such as the evolution of the SFE and global ionization fraction was adequate, and did not include any of the simulations discussed here. The principal effect of increasing the resolution in an SPH simulation is to permit the formation of lower mass objects which otherwise cannot be modelled. We did not find that the shape of the mass function (except at the lowest mass end), the total stellar mass, or the...
spatial distribution of stellar mass were substantially affected by the simulation resolution.

We also found no evidence that feedback alters the stellar mass function, so that increasing the simulation resolution would be likely to affect the stellar content of the control and feedback hydrodynamic simulations in very similar ways, by effectively extending their mass functions to lower masses. In reality, some of this additional fragmentation would be prevented by physics which is not currently included in the SPH simulations, such as accretion feedback.

The effect of the presence of an additional population of low-mass stars on the outcome of the N-body simulations is not trivial to quantify. It is very unlikely to qualitatively alter our results (one of which is that it is not always possible to get a consensus from different algorithms on whether mass segregation is present in a single data set), although it may alter the time-scales on which the modelled stellar systems dynamically mass segregate or unsegregate. The time-scale for mass segregation is related to the local crossing time (i.e. local density) and the presence of more low-mass stars would make the region more dense, reduce the crossing time and hence make the region more likely to dynamically mass segregate on faster time-scales. However, very dense clusters can eject massive stars (sometimes dissolving the cluster entirely; Allison & Goodwin 2011; Parker et al. 2014), so it is not clear whether the ‘missing’ low-mass stars in the SPH runs would affect the amount of mass segregation measured.

During the subsequent dynamical evolution via N-body simulations, mass segregation can be erased through dynamical interactions or (more usually) mass-loss from stellar evolution, which has the effect of changing the 10 most massive stars in the bin used to define $\Sigma_{\text{LDR}}$ or $\Lambda_{\text{MSR}}$. Together, the above results suggest that mass segregation need not be primordial, and conversely, that a high-mass cluster which does not display mass segregation after a certain time may have been mass segregated in the past, especially if it contains highly evolved stars. The only limits we can place on mass segregation is that if it is not observed in a low-mass region without evolved stars (Parker et al. 2011, 2012), then it is unlikely that this region was ever mass segregated. Models that explain mass segregation of more massive clusters (such as the Orion nebula Cluster) via the cool collapse of a star-forming region (Allison et al. 2010; Allison & Goodwin 2011; Parker et al. 2014) are still as likely an explanation for the observed mass segregation in these clusters, as opposed to primordial mass segregation.

5 CONCLUSIONS

We have evolved five triplets of hydrodynamical simulations of star formation from Dale et al. (2012, 2013, 2014) further in time using the N-body method. The hydrodynamical simulations follow the formation of a region with (a) no feedback, (b) photoionization feedback only, and (c) photoionization feedback and stellar winds.

We have looked for differences in the spatial distribution of stars, local density, and the fraction of bound stars. We have also used three different methods used to quantify mass segregation – $\Lambda_{\text{MSR}}$ (Allison et al. 2009a), $\Sigma_{\text{LDR}}$ (Maschberger & Clarke 2011; Parker et al. 2014), and the group segregation ratio, $\Omega_{\text{GSR}}$ (Kirk & Myers 2011; Kirk et al. 2014).

In terms of the global evolution of the regions, our conclusions are similar to those in Parker & Dale (2013), where we looked for differences in simulations with no feedback, and simulations with photoionization feedback only. The addition of stellar winds feedback has little effect above the photoionization feedback; the star-forming regions subject to feedback form with lower densities and hence remain substructured for longer than the simulation without feedback. Generally, the simulations that form with feedback contain fewer bound stars at the end of the N-body integration, possibly due to the lower mass of the star-forming regions.

Using three different techniques to search for mass segregation presents a rather confusing picture. When feedback is not switched on, the regions are usually primordially mass segregated according to $\Omega_{\text{GSR}}$ and $\Sigma_{\text{LDR}}$, whereas the $\Lambda_{\text{MSR}}$ measure does not usually measure primordial mass segregation in the simulations. When feedback is switched on, all three measures show no preferential spatial distribution for the most massive stars, suggesting that the inclusion of more realistic physics in the hydrodynamical simulations suppresses primordial mass segregation, although more simulations that include magnetic fields would be highly desirable, and could potentially alter these conclusions.

When evolved for 10 Myr, some simulations dynamically mass segregate. However, a combination of further dynamical evolution, and stellar evolution of the most massive stars, can also cause some regions to unsegregate, and sometimes re-segregate due to different subsets of massive stars being included in the determination.

We conclude that extreme caution should be exercised when interpreting mass segregation in the outcome of hydrodynamical simulations of star formation. Different methods define mass segregation in different ways, and dynamical and stellar evolution can also affect searches for mass segregation. We note that several observed low-mass regions do not display mass segregation, and based on the results presented here, we suggest that mass segregation is not always primordial.

ACKNOWLEDGEMENTS

RJP acknowledges support from the Royal Astronomical Society in the form of a research fellowship. This research was supported by the DFG cluster of excellence ‘Origin and Structure of the Universe’ (JED, BE). We thank Tobias Achenbrenner for the preliminary study in his Bachelorarbeit at USM, which was the genesis of this paper.

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