ABSTRACT

This paper is the first in a series in which we perform an extensive comparison of various galaxy-based cluster mass estimation techniques that utilize the positions, velocities and colours of galaxies. Our primary aim is to test the performance of these cluster mass estimation techniques on a diverse set of models that will increase in complexity. We begin by providing participating methods with data from a simple model that delivers idealized clusters, enabling us to quantify the underlying scatter intrinsic to these mass estimation techniques. The mock catalogue is based on a Halo Occupation Distribution (HOD) model that assumes spherical Navarro, Frenk and White (NFW) haloes truncated at $R_{200}$, with no substructure nor colour segregation, and with isotropic, isothermal Maxwellian velocities. We find that, above $10^{14} M_{\odot}$, recovered cluster masses are correlated with the true underlying cluster mass with an intrinsic scatter of typically a factor of 2. Below $10^{14} M_{\odot}$, the scatter rises as the number of member galaxies drops and rapidly approaches an order of magnitude. We find that richness-based methods deliver the lowest scatter, but it is not clear whether such accuracy may simply be the result of using an over-simplistic model to populate the galaxies in their haloes. Even when given the true cluster membership, large scatter is observed for the majority non-richness-based approaches, suggesting that mass reconstruction with a low number of dynamical tracers is inherently problematic.

Key words: methods: numerical – methods: statistical – galaxies: clusters: general – galaxies: haloes – galaxies: kinematics and dynamics – cosmology: observations.

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1 INTRODUCTION

Deducing the masses of the largest gravitationally bound structures in the Universe, galaxy clusters, remains a complex problem that is at the focus of current and future cosmological studies. The characteristics of the galaxy cluster population provide crucial information for studies of large scale-structure (e.g. Bahcall 1988; Einasto et al. 2001; Yang et al. 2005b; Papovich 2008; Willis et al. 2013), constraining cosmological model parameters (see Allen, Evrard & Mantz 2011, for a review) and galaxy evolution studies (e.g. Goto et al. 2003; Postman et al. 2005; Martínez, Coenda & Muriel 2008). Despite the wealth of information clusters can provide, deriving strong constraints from cluster surveys is a non-trivial problem due to the complexity of estimating accurate cluster masses. The use of cluster surveys as a dark energy probe provides greater statistical power than other techniques (Dark Energy Task Force; Albrecht et al. 2006). However, enabling this statistical precision requires significant advances in treating the systematic uncertainties between survey observables and cluster masses. Clusters can be detected across several different wavelength regimes using various techniques. They are identified in optical and infrared light as overdensities in the number counts of galaxies (e.g. Abell 1958; Zwicky, Herzog & Wild 1968), while colour information improves the contrast by selecting the red galaxies that dominate in these systems (e.g. Gladders & Yee 2005; Koester et al. 2007; Szabo et al. 2011; Ascaso, Wittman & Benítez 2012). At X-ray wavelengths, the hot intra-cluster medium produces bright extended sources (e.g. Forman et al. 1972; Böhringer et al. 2000; Rosati, Borgani & Norman 2002; Vikhlinin et al. 2009), while at millimetre wavelengths, inverse Compton scattering of photons from this gas results in characteristic distortions in the cosmic microwave background (e.g. Sunyaev & Zeldovich 1972; Carlstrom, Holder & Reese 2002; Vanderlinde et al. 2010; Planck Collaboration et al. 2013; Hasselfield et al. 2013). Finally, distortions of images of faint background galaxies through weak gravitational lensing offers perhaps the most direct measure of the huge masses of these systems (e.g. Applegate et al. 2012).

Despite these diverse methods of detecting clusters, no cluster observable directly delivers a mass. The cluster mass function is one key method to constrain the dark energy parameter. Ongoing and future dark energy missions plan to consider cluster counts in their analyses. Hence, it is crucial to be able to measure cluster masses as accurately as possible. Follow-up spectroscopy is of great importance to all group/cluster surveys, providing the kinematics of cluster galaxies, which is one of a few mass proxies that is directly related to cluster mass (by providing a direct measure of the dark matter potential well). This series of papers examines various observable–mass relations by testing an extensive range of galaxy-based cluster mass estimation techniques with the aim of calibrating follow-up mass proxies.

Galaxy-based mass estimation techniques commonly follow three general steps: first identify the cluster overdensity, second deduce cluster membership, and, finally, using this membership, estimate a cluster mass. Common optical cluster finding methods include using the Huchra & Geller (1982) Friends-Of-Friends (FOF) group-finding algorithm (e.g. Berlind et al. 2006; Li & Yee 2008; Tempel, Tago & Liivamägi 2012; Jian et al. 2013) and methods based upon Voronoi tessellation (e.g. Marinoni et al. 2002; Lopes et al. 2004; van Breukelen & Clewley 2009; Soares-Santos et al. 2011). Also widely used are red-sequence filtering techniques (e.g. Gladders & Yee 2000; Murphy, Geach & Bower 2012; Rykoff et al. 2013) and methods that rely on the bright central galaxy (BCG) to identify the presence of a cluster (e.g. Yang et al. 2005a; Koester et al. 2007). Cluster catalogues are also constructed using the positions and magnitudes of galaxies to search for over-densities via the matched filter algorithm (e.g. Postman et al. 1996; Kepner et al. 1999; Olsen et al. 1999; Menanteau et al. 2009).

Once the over-densities are identified, many methods select an initial cluster membership using the groups obtained via the FOF algorithm (e.g. Muñoz-Cuartas & Müller 2012; Pearson et al., in preparation; Tempel et al. 2012), whilst others select galaxies within a specified volume in phase space (e.g. von der Linden et al. 2007; Wojtak et al. 2009; Gifford & Miller 2013; Mamon, Biviano & Boué 2013; Sifon et al. 2013; Pearson et al., in preparation) or within a certain region of colour–magnitude space where cluster galaxies are known to reside (e.g. Saro et al. 2013). Once the initial set of member galaxies is chosen, it is common to iteratively refine membership using either the estimated velocity dispersion, radius and colour information, or even a combination of these properties. Deducing which galaxies are members of a cluster is non-trivial, and unfortunately the inclusion of even quite small fractions of interloper galaxies that are not gravitationally bound to the cluster can lead to a strong bias in velocity dispersion-based mass estimates (e.g. Lucey 1983; Borgani et al. 1997; Cen 1997; Biviano et al. 2006; Wojtak et al. 2007). For this reason, methods often employ careful interloper removal techniques, for example, by modelling interloper contamination when performing density fitting, by using the Gapper technique or via iterative clipping as described above. Many methods then follow the classical approach of applying the virial theorem to the projected phase space distribution of member galaxies (e.g. Zwicky 1937; Yahil & Vidal 1977; Evrard et al. 2008), assuming that the system is in equilibrium. Other methods utilize the distribution of galaxies in projected phase space, assuming a Navarro, Frenk and White (NFW) density profile (Navarro, Frenk & White 1996; Navarro, Frenk & White 1997) to obtain an estimate of cluster mass. The number of galaxies associated with a cluster above a given magnitude limit (the richness) is also used as a proxy for mass (e.g. Yee & Ellingson 2003; Andreon & Hurr 2010). In addition, the more recently developed caustic method identifies the projected escape velocity profile of a cluster in radius–velocity phase space, delivering a measure of cluster mass (e.g. Diaferio & Geller 1997; Diaferio 1999; Gifford & Miller 2013).

The aim of this paper is to perform a comprehensive comparison of 23 different methods that employ variations of the techniques described above by deducing both the mass and membership of galaxy clusters from a mock galaxy catalogue. In order to simplify the problem, the clusters are populated with galaxies in a somewhat idealized manner, with cluster locations that are specified a priori; in this way, the basic workings of the various algorithms can be tested under optimal conditions, without the potential for confusion from more complex geometries or misidentified clusters.

The paper is organized as follows. We describe the mock galaxy catalogue in Section 2, and the mass reconstruction methods applied to this catalogue are described in Section 3. In Sections 4 and 5, we present our results on cluster mass and membership comparisons. We end with a discussion of our results and conclusions in Section 6. Throughout the paper we adopt a Lambda cold dark matter ($\Lambda$CDM) cosmology with $\Omega_0 = 0.25$, $\Omega_{\Lambda} = 0.75$, and a Hubble constant of $H_0 = 73$ km s$^{-1}$ Mpc$^{-1}$, although none of the conclusions depends strongly on these parameters.
2 DATA

This paper forms the initial part (Phase I) of a large comparison programme aimed at studying how well halo masses can be recovered using a wide variety of group/cluster mass reconstruction techniques based on galaxy properties. As the first step, we intentionally use a very clean and straightforward set-up: a simple HOD galaxy mock catalogue built upon a nearby-Universe light-cone. Later stages of this project will involve more sophisticated mock galaxy catalogues using both more advanced HOD models (Skibba et al., in preparation) and semi-analytic modelling (Croton et al. 2006). This paper sets out to determine the simplest-case baseline by using a clean, well-defined data set with idealized substructure, sharp boundaries, spherically symmetric haloes and a strong richness correlation. Given initial estimates for the location of the structures, just how bad can it get?

For Phase I, the data set is the mock galaxy catalogue constructed in Muldrew et al. (2012). We briefly describe the catalogue here, and we refer the reader to the above paper and to Skibba & Sheth (2009) for more details. We begin with the Millennium Simulation (Springel et al. 2005), which tracks the evolution of 2160$^3$ dark matter particles of mass $8.6 \times 10^9 h^{-1} M_{\odot}$ from $z = 127$ to $z = 0$ within a comoving box of side length 500$h^{-1}$ Mpc, with a halo mass resolution of $\sim 5 \times 10^9 h^{-1} M_{\odot}$. The simulation adopts a flat ΛCDM cosmology with the following parameters: $\Omega_0 = 0.25$, $\Omega_m = 0.7$, $\sigma_8 = 0.9$, $n = 1$ and $h = 0.73$. Collapsed haloes at $z = 0$ with at least 20 particles are identified with the SUBFIND (Springel et al. 2001) FOF group-finding algorithm, although consistent results are found with other finders (Muldrew, Pearce & Power 2011; Knebe et al. 2011).

The haloes are populated with galaxies whose luminosities and colours follow the halo-model algorithm described in Skibba et al. (2006) and Skibba & Sheth (2009), which is constrained by the luminosity function, colour–magnitude distribution, and luminosity and colour-dependent clustering (Zehavi et al. 2005) as observed in the Sloan Digital Sky Survey (SDSS; York et al. 2000). An important assumption in this HOD model is that all galaxy properties – their numbers, spatial distributions, velocities, luminosities, and colours – are determined by halo mass alone, again rendering the model as simple as possible. We specify a minimum $r$-band luminosity for the galaxies of $M_r = -19 + 5 \log (h)$, to stay well above the resolution limit of the Millennium Simulation.\(^1\)

Haloes are assigned a ‘central’ galaxy which has the same position and velocity as the halo centre (Skibba et al. 2011). ‘Satellite’ galaxies are assumed to be fainter than this object and follow an NFW density profile (Navarro et al. 1997) that obeys the concentration–mass relation from Macciò, Dutton & van den Bosch (2008), with the population extending out to $R_{200}$ (the radius that encloses a density 200 times the critical density of the Universe), assuming isothermal, isotropic, Maxwellian velocity distributions.

In the model of galaxy colours, central and satellite galaxies have different colour–luminosity distributions. The central galaxy is usually the reddest galaxy in a given halo, though satellites are redder than central galaxies at a given luminosity (van den Bosch et al. 2008; Skibba 2009). Satellites are assumed to follow a particular sequence on the colour–magnitude diagram, which approaches the red sequence with increasing luminosity, consistent with what is found in the SDSS group/cluster catalogues (Skibba 2009). Note that alternative approaches to modelling galaxy colours and colour-dependent clustering have recently appeared in the literature (Gerke et al. 2013; Hearin & Watson 2013; Masaki, Lin & Yoshida 2013; Phleps et al. 2013).

We also allow for the expected scatter in the relation between host halo mass and central galaxy luminosity (Zheng, Coil & Zehavi 2007). The number of satellite galaxies in a halo of given mass, $P(N_{\text{sat}}|M)$, is approximated well by a Poisson distribution (Kravtsov et al. 2004), with a mean HOD that increases approximately linearly with mass, $\langle N_{\text{sat}}|M \rangle = [M - M_0]_\text{HOD}/M_\text{HOD}$, hence we adopt this distribution to populate the haloes.\(^2\) The value of $M_\text{HOD}/M_\text{min} \approx 17$ (where $M_\text{min}$ is the mass corresponding to the luminosity threshold: $M_\text{r} < 19 + 5 \log h$), which determines the critical mass above which haloes typically host at least one satellite, is approximately independent of luminosity, and $\alpha \approx 1$ for most of this luminosity range (Zehavi et al. 2011). $M_0$ determines the shape of the satellite HOD at low halo masses and is typically smaller than $M_\text{min}$. The HOD parameters are described in detail in appendix A of Skibba & Sheth (2009).

A galaxy’s velocity is given by the sum of the velocity of its parent halo plus an internal motion contribution within the halo. The internal motions are well approximated by a Maxwellian distribution (admittedly, ΛCDM haloes have more complex velocity distribution functions; see e.g. Sheth & Diaferio 2001 and Bernal et al. 2013), with velocities that are independent Gaussians in each of the three Cartesian coordinates. The dispersion depends on halo mass and radius through the scaling:

\[ \sigma_{200}^2 = G M_{200}/(2 R_{200}). \] (1)

Note that this yields velocity dispersions that are 7 per cent greater than expected for NFW models with realistic anisotropic velocities (Mamon et al. 2013). This overestimate of velocity dispersion and the assumption that it is independent of radius cause a violation of local dynamical equilibrium. In this phase of the project, we also neglect the effects of galaxy velocity bias (Skibba et al. 2011; Munari et al. 2013; Old, Gray & Pearce 2013).

A $90 \times 90'$ light-cone, 500$h^{-1}$ Mpc deep, is constructed by taking a slice through the zero-redshift simulation cone. To deliver a data set with a sufficient number of cluster-sized haloes, the 1000 groups/clusters are selected by taking the 500 most massive, the next 300 richest and finally the groups/clusters with the most luminous brightest cluster galaxy (BCG) are taken to complete the sample. The mass functions of the selected sample and the full light-cone are shown in Fig. 1 via the solid red and black dashed line, respectively. An example of the underlying galaxy distribution inserted by the HOD model is shown in Fig. 2. Black circles indicate the member galaxies for the largest cluster in the sample. By construction, this spatial distribution is smooth and spherical, lacking any imposed substructure, again to keep the test as simple as possible. The red diamonds indicate galaxies in other haloes. As Fig. 2 demonstrates, there are many small haloes that have been populated with HOD galaxies that are not part of our target list but which form the background galaxy distribution for this initial phase of the project, to give the mass measuring algorithms a simple contaminant to

\(^{1}\) The mass resolution of the simulation is sufficient that haloes that host galaxies as faint as $0.1 L_\odot (M_r = 18 + 5 \log (h))$ are typically resolved with more than $\sim 100$ particles (Springel et al. 2005), which corresponds to a stellar mass threshold of $M_\ast \sim 10^{10} h^{-1} M_\odot$.

\(^{2}\) Note that the true relationship between richness and mass is determined by assumptions about the shape of the halo occupation distribution and its mean as a function of mass. The true number distribution at fixed mass is not Poisson, however, because of the group/cluster selection procedure (see Section 2).
reject in their analysis. Once a light-cone is generated, the internal velocity dispersion of this large object will be added to the Hubble recession, stretching it out along the line of sight, generating the usual ‘Finger of God’ effect.

In summary, the model for this simplified initial test generates data where clusters are spherically symmetric, there is no internal substructure, no galaxy velocity bias, no large-scale streaming motions, galaxies follow isotropic orbits and have effectively zero size so there is no blending of objects on the sky.

3 MASS RECONSTRUCTION METHODS

In this section, we present the halo mass reconstruction methods used in this comparison project as listed in Table 1, which also summarizes some basic properties of each method. The following subsections provide brief descriptions of each method, and are headed by an identifying acronym used throughout this paper, as well as giving the names of the developers who participated in this project and the type of method involved. The two main steps performed by each method, the initial galaxy selection and the mass estimation, are separated into broad classes (which are specified in parentheses in the subsection titles). For the procedure of deducing the initial member galaxy sample, methods are categorized as either FOF (star), red sequence (diamond) or phase space (circle) based. The mass estimation procedures are classed as either richness (magenta), phase space (black), radius (blue), abundance matching (green) or velocity dispersion (red) based. Further details can be found in the paper references that are provided in each description, and a more extensive summary of the method characteristics is provided in the Appendix.

3.1 PCN (Pearson & Ponman, phase space, richness)

All PCx methods are based on a cylindrically selected galaxy sample. Starting with the halo positions, galaxies are initially selected from a 5 Mpc radius cylinder about each halo with a depth of ±1000 km s$^{-1}$. The velocity depth is then iterated with a robust 3σ clipping using galaxies within 1 Mpc. To derive masses, the PCN method uses an aperture richness of each cluster as discussed in Pearson et al. (in preparation). Richness is defined as the number of galaxies above a threshold absolute magnitude within 1 Mpc, subtracting an interloper contribution estimated using galaxies in a background annulus of radii 3–5 Mpc. Mass is then estimated using a $M_{500}$–richness relation calibrated on a sample of clusters with SDSS galaxy data and X-ray estimates for $M_{500}$ from Sun et al. (2009) and Sanderson & Ponman (2010). The estimated $M_{500}$ is converted to $M_{200}$ for this project using the mass–concentration relation of Duffy et al. (2008). We estimate statistical errors through bootstrap resampling the observed mass proxy and systematic errors by propagating errors on our calibration relation.

3.2 PFN (Pearson & Ponman, FOF, richness)

All PFx methods are based on an FOF-selected cluster sample. We apply an FOF analysis using the scheme of Eke et al. (2004) which utilizes the positions and velocities of galaxies. The linked clusters are matched to the given cluster centre positions. All linked galaxies are assumed to be cluster members, so we do not include any corrections for interloper contamination. For the PFN method, masses are derived based on the FOF richness of each cluster as discussed in Pearson et al. (in preparation), and calibrated against X-ray masses using the same sample as described for the PCN method, and is converted to $M_{200}$ in the same way. Statistical errors are estimated from Poisson errors propagated through the calibrated mass relation; systematic errors are derived from calibrating uncertainties as for PCN.
ties, relative to the central halo, smaller (in absolute value) than the virial radius, \( R_{200} \), which contains information on the potential, density, and velocity anisotropy profiles of the halo. This is deduced by selecting all galaxies within 1 Mpc and 1333 km s\(^{-1}\) from the halo centre, with a linear relation between \( \log R_{200} \) and \( \log N(1 \text{ Mpc}, 1333 \text{ km s}^{-1}) \) deduced from a robust linear fit to the mock clusters analysed by CLE (see Section 19). The membership is deduced by selecting all galaxies within \( R_{200} \) and with velocities, relative to the central halo, smaller (in absolute value) than \( 2.7 \sigma_{los}(R) \) (computed from an NFW model, as in the CLE method). See Mamon et al. (in preparation).

### 3.3 NUM (Mamon, phase space, richness)

The radius \( R_{200} \) is estimated using the richness measured in a rectangular area of projected phase space within 1 Mpc and 1333 km s\(^{-1}\) from the halo centre, with a linear relation between \( \log R_{200} \) and \( \log N(1 \text{ Mpc}, 1333 \text{ km s}^{-1}) \) deduced from a robust linear fit to the mock clusters analysed by CLE (see Section 19). The membership is deduced by selecting all galaxies within \( R_{200} \) and with velocities, relative to the central halo, smaller (in absolute value) than \( 2.7 \sigma_{los}(R) \) (computed from an NFW model, as in the CLE method). See Mamon et al. (in preparation).

### 3.4 ESC (Gifford & Miller, phase space)

The caustic technique utilizes the radius–velocity phase space information of galaxies in clusters, as well as their dispersion, to estimate the escape velocity profile of the host haloes. The mass profile is inferred by integrating the square of the escape velocity profile multiplied by a parameter \( F_p \) which contains information on the potential, density, and velocity anisotropy profiles of the halo along with fundamental constants. \( F_p \) is treated as approximately constant (see Diaferio 1999 and Serra et al. 2011) with a value of 0.65 as found in Gifford, Miller & Kern (2013). Member galaxies are identified as those within the escape velocity envelope in radius–velocity phase space and within the estimated \( R_{200} \) of the halo. This technique is described in both Gifford & Miller (2013) and Gifford et al. (2013).

### 3.5 MPO (Mamon, phase space)

Starting from the sample of members obtained with the CLN algorithm, the virial radius, \( R_{200} \), total mass scale radius, \( R_p \), red and blue galaxy population scale radii, \( R_{\text{red}} \) and \( R_{\text{blue}} \), and the velocity anisotropies at the virial radius of these red and blue populations are computed using the Bayesian MAMPOSSt method (Mamon et al. 2013). This method jointly fits the positions of the red and blue galaxies in projected phase space. Here, it is assumed that the system is spherically symmetric and that the total mass distribution follows the NFW model, while the red and blue galaxy populations follow NFW models, each with its scale radius. The red and blue populations are assumed to have isotropic orbits at the centre, but increasingly radial or tangential beyond this (with different free outer anisotropies, but a transition scale fixed to be the scale radius of the tracer). The 3D velocities are assumed to be Gaussian at all radii.

### 3.6 MP1 (Mamon, phase space)

MP1 is like MPO, but is colour-blind: a single tracer population is assumed.

### 3.7 RW (Wojtak, phase space)

In this method, the halo mass \( M_{200} \) is derived from the distribution of galaxies in phase space. It is assumed that the galaxies follow a combination of a spherical NFW model (where number follows mass) with a distribution function of energy and angular momentum derived from ACDM haloes (Wojtak et al. 2008), forcing here the inner and outer anisotropies to match those of ACDM haloes, and a constant projected density background term that is kept as a free parameter. See Wojtak et al. (2009) for details. The membership is determined by restricting to galaxies within \( v_{\text{los}} < \sqrt{\Phi(R)/\Phi_1} \), where \( R \) is the projected distance of the galaxy.

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**Table 1.** Summary of the participating cluster mass estimation methods. Listed is an acronym identifying the method, an indication of the scheme used to undertake member galaxy selection and an indication of the method used to convert this membership list to a mass estimate. Note that acronyms denoted with a star indicate that the method did not use our initial 1000 object target list but rather matched these locations at the end of their analysis. Please see Tables A1 and A2 in the Appendix for more details on each method.

| Method   | Initial galaxy selection       | Mass estimation      | Reference                                      |
|----------|--------------------------------|----------------------|------------------------------------------------|
| PCN      | Phase space                    | Richness             | Pearson et al. (in preparation)                |
| PFN*     | FOF                            | Richness             | Pearson et al. (in preparation)                |
| NUM      | Phase space                    | Richness             | Mamon et al. (in preparation)                  |
| ESC      | Phase space                    | Phase space          | Gifford & Miller (2013)                        |
| MPO      | Phase space                    | Phase space          | Mamon et al. (2013)                            |
| MP1      | Phase space                    | Phase space          | Mamon et al. (2013)                            |
| RW       | Phase space                    | Phase space          | Wojtak et al. (2009)                           |
| TAR*     | FOF                            | Phase space          | Tempel et al. (2014)                           |
| PCO      | Phase space                    | Radius               | Pearson et al. (in preparation)                |
| PFO*     | FOF                            | Radius               | Pearson et al. (in preparation)                |
| PCR      | Phase space                    | Radius               | Pearson et al. (in preparation)                |
| PFR*     | FOF                            | Radius               | Pearson et al. (in preparation)                |
| HBM*     | FOF                            | Abundance matching   | Muñoz-Cuártas & Müller (2012)                 |
| MVM*     | FOF                            | Abundance matching   | Muñoz-Cuártas & Müller (2012)                 |
| AS1      | Red sequence                   | Velocity dispersion  | Saro et al. (2013)                             |
| AS2      | Red sequence                   | Velocity dispersion  | Saro et al. (2013)                             |
| AvL      | Phase space                    | Velocity dispersion  | von der Linden et al. (2007)                   |
| CLE      | Phase space                    | Velocity dispersion  | Mamon et al. (2013)                            |
| CLN      | Phase space                    | Velocity dispersion  | Mamon et al. (2013)                            |
| SG1      | Phase space                    | Velocity dispersion  | Sifon et al. (2013)                            |
| SG2      | Phase space                    | Velocity dispersion  | Sifon et al. (2013)                            |
| PCS      | Phase space                    | Velocity dispersion  | Pearson et al. (in preparation)                |
| PFS*     | FOF                            | Velocity dispersion  | Pearson et al. (in preparation)                |
3.8 TAR (Tempel et al., phase space)

TAR groups/clusters are based upon the conventional FOF group finding algorithm, where the linking-length is calibrated based on the mean distance to nearest galaxy in the plane of the sky. For the current data set $d_1 = 0.44 h^{-1} \text{Mpc}$ and $d_2 = 440 \text{km s}^{-1} = 4.4 h^{-1} \text{Mpc}$ (assuming $d_1/d_2 = 10$). More details of the group finding algorithm are explained in Tago et al. (2008, 2010) and Tempel et al. (2012). The masses of groups/clusters are estimated by applying the virial theorem to the sphere of radius $R_{200}$:

$$M = \frac{3 R_G \sigma_v^2}{G} = 7.0 \times 10^{12} \frac{R_G}{\text{Mpc}} \left(\frac{\sigma_v}{100 \text{km s}^{-1}}\right)^2 M_\odot,$$

(2)

where $\sigma_v$ is the 1D velocity dispersion. The gravitational radius $R_G$ is estimated from the rms projected radius. For that we assume an NFW profile and find the theoretical relationship between these two parameters. Since the concentration parameter of the NFW profile depends on the halo mass (we use the mass–concentration relation from Macciò et al. 2008), we find the final mass iteratively. See Tempel et al. (2014) for more details of the method.

3.9 PCO (Pearson & Ponman, phase space, radius)

Using the galaxy membership of PCN, the galaxy overdensity profiles of clusters are modelled and fitted as described in Pearson et al. (in preparation). A projected NFW profile (Bartelmann 1996) plus a uniform background term to allow for interloper contamination is fitted to all galaxies within 5 Mpc. From the fitted NFW profile a radius $R_{500}$ is found, within which the cumulative number density is 500/ $\Omega_m$ times the mean cosmic number density of galaxies. This number density is estimated from the SDSS luminosity function of Blanton, Hogg & Bahcall (2003) where galaxies are counted above a threshold luminosity of $L_r = -19$. The mass $M_{500}$ within $R_{500}$ is then deduced from $R_{500}$. These overdensity masses have been calibrated against the X-ray masses described under the above a threshold luminosity of $L_r = -19$. The mass $M_{500}$ within $R_{500}$ is then deduced from $R_{500}$. These overdensity masses have been calibrated against the X-ray masses described under the PCN method, and as a result a linear scaling is applied to determine the final $M_{500}$ estimate, which is then extrapolated to $M_{200}$. Error analysis is as for PCN.

3.10 PFO (Pearson & Ponman, FOF, radius)

Using the linked galaxy membership of PFN, the galaxy overdensity profiles of clusters are modelled and fitted as described in Pearson et al. (in preparation). We fit a projected NFW (Bartelmann 1996) profile, assuming that the linked galaxy membership is subject to no interloper contamination. $M_{200}$ is then derived from the fitted profile as for PCO.

3.11 PCR (Pearson & Ponman, phase space, radius)

Using the galaxy membership within 1 Mpc, as derived for PCN, this method is based on the rms radius of each cluster as discussed in Pearson et al (in preparation). Note, however, that since we have no way of knowing which galaxies are interlopers, we are unable to make any statistical allowance for them (in contrast to the PCN method). As for PCN, we apply a relation calibrated on X-ray derived masses to estimate $M_{500}$, which is then extrapolated to $M_{200}$. Error analysis is as for PCN.

3.12 PFR (Pearson & Ponman, FOF, radius)

The method is the same as PCR, except that it is applied to the FOF-selected galaxy membership described for PFN.

3.13 HBM (Muñoz-Cuartas, FOF, abundance matching)

HBM is based upon an ellipsoidal FOF method with linking lengths adapted according to the estimated halo mass. The linking length along the line of sight is controlled by the (theoretical) velocity dispersion of the halo. Cluster masses are determined by abundance matching between the cluster $r$-band luminosity function and the theoretical halo mass function of Warren et al. (2006). The centre of the halo is set to the galaxy with the largest $r$-band luminosity. The method is described in detail in Muñoz-Cuartas & Müller (2012).

3.14 MVM (Müller, FOF, abundance matching)

MVM is the same as HBM with the difference that the centre is assumed to lie at the centre of stellar mass, while the virial theorem is used to compute $M_{200}$. The procedure is described in more detail in Muñoz-Cuartas & Müller (2012).

3.15 AS1 (Saro, red sequence, velocity dispersion)

AS1 was developed to study possible systematics affecting follow-up dynamical mass estimation of high-redshift massive galaxy clusters. By construction, it assumes that the centre of the cluster is known, along with an initial estimate of $R_{200}$ from other observables. It also assumes an intrinsic scatter of $\approx 30$ per cent in mass at fixed velocity dispersion, mainly driven by the triaxial properties of DM haloes. As the simulated clusters in this work are spherical, this is largely overestimated. As the total estimated errors on individual clusters mass could be larger than $\approx 60$ per cent, it does not iterate to solve for $R_{200}$, but is focused more on obtaining an average unbiased mass estimation for an ensemble of clusters. Since, for the purpose of this work, no initial $R_{200}$ was given, it assumes a fiducial value of 1 Mpc for all the mass range. Galaxies must lie within 0.1 mag in colour from a model given by Song et al. (2012), which has proven to be a good fit to the observational data and they must also lie within 4000 km s$^{-1}$ from the cluster centre. A final clipping of $\pm 3 \sigma$ is then performed to remove interlopers and provide a robust estimate of the velocity dispersion. A scaling relation, provided in Saro et al. (2013), is then used to convert the velocity dispersion into $M_{200}$. The model is cosmologically dependent at a background level and assumes a cosmology of $\Omega_M = 0.3$, $\Omega_\Lambda = 0.7$ and $h_0 = 0.7$. More details of this method are described in Saro et al. (2013).

3.16 AS2 (Saro, red sequence, velocity dispersion)

AS2 follows the same procedure as AS1 but the estimated velocity dispersion is corrected by taking into account the number of galaxies as described by equation (6) in Saro et al. (2013). Note: the values of the constants $a$ and $b$ of the relation $\log M_{200} = a + b \log \sigma_v$ employed by the ASx methods can be found in Table 2.

3.17 AvL (von der Linden, phase space, velocity dispersion)

This method is a relatively simple velocity dispersion estimator, as used for SDSS clusters in von der Linden et al. (2007) and for EDisCS clusters in Milvang-Jensen et al. (2008). Galaxies are iteratively selected to lie within $2.5 \sigma_v$ and $0.8 R_{200}$ – the latter is
estimated from $\sigma_v$ by assuming the virial theorem. These cuts are chosen to make the method relatively insensitive to contamination from nearby structures; $R_{200}$ and the final $\sigma_v$ are corrected for the expected bias from sigma-clipping. Note: the values of the constants $a$ and $b$ of the relation $\log M_{200} = a + b \log \sigma_v$ employed by AvL can be found in Table 2.

3.18 CLE (Mamon, phase space, velocity dispersion)

The initial membership is limited to $R < 3$ Mpc and $|v| < 4000$ km s$^{-1}$. A relative velocity gap technique (Girardi et al. 1993), with gapper coefficient $C = 4$, is initially applied to remove obvious interlopers (keeping the largest subsample). The radius $R_{200}$ is first estimated from the aperture velocity dispersion, where the measured value (using the robust Median Absolute Deviation; see Beers, Flynn & Gebhardt 1990) is matched to the aperture velocity dispersion at $R_{200}$. This is predicted for a spherical single-component NFW model with concentration of $c = 4$, with the Mamon & Łokas (2005) velocity anisotropy profile (with anisotropy radius equal to the scale radius of the NFW model, as found for $\Lambda$CDM haloes by Mamon, Biviano & Murante 2010). The membership is recovered by selecting all galaxies within $R_{200}$ with velocities relative to the central one smaller (in absolute value) than $2.7\sigma_{200}(R)$ (computed from an NFW model with the same velocity anisotropy model, but now assuming a concentration obtained from the $\Lambda$CDM concentration–mass relation of Macciò et al. 2008). The virial radius and membership are iterated, now measuring the aperture velocity dispersion using the unbiased standard deviation. The method is described in the appendix of Mamon et al. (2013). Note: the values of the constants $a$ and $b$ of the relation $\log M_{200} = a + b \log \sigma_v$ employed by CLE can be found in Table 2.

3.19 CLN (Mamon, phase space, velocity dispersion)

CLN is similar to CLE, but now using the output of NUM as input. Note: the values of the constants $a$ and $b$ of the relation $\log M_{200} = a + b \log \sigma_v$ employed by CLN can be found in Table 2.

3.20 SG1 (Sifón, phase space, velocity dispersion)

Both SG1 and SG2 implement the shifting gapper of Fadda et al. (1996) and the velocity dispersion-mass relation of Evrard et al. (2008). All galaxies within 4000 km s$^{-1}$ (rest-frame) of the cluster redshift are binned in projected radial annuli, each of which has at least 15 galaxies and a minimum width of 250 kpc. Galaxies within each bin are ordered by the modulus of the velocity and a main body is defined by finding a gap between two successive velocities of 500 km s$^{-1}$ or more. All galaxies within 1000 km s$^{-1}$ of this boundary are considered halo members. The velocity dispersion is the bi-weight estimate of scale (Beers et al. 1990) of all members. From this velocity dispersion, a mass, $M_{200}$, is estimated from the velocity dispersion–mass relation of Evrard et al. (2008), and the radius, $R_{200}$, is estimated from this mass. A new velocity dispersion is computed using only members within $R_{200}$, and this process is repeated until convergence (usually $\sim 3$ iterations). The full description of the implementation is in Sifón et al. (2013).

3.21 SG2 (Sifón, phase space, velocity dispersion)

SG2 is the same algorithm as SG1 but with different parameters for the shifting gapper method: radial bins have a minimum width of 150 kpc and 10 galaxies; the main body boundary is 300 km s$^{-1}$ and all galaxies within 500 km s$^{-1}$ of this boundary are considered members. Consequently, SG1 and SG2 only differ in the membership selection. Note: the values of the constants $a$ and $b$ of the relation $\log M_{200} = a + b \log \sigma_v$ employed by the SGx methods can be found in Table 2.

3.22 PCS (Pearson & Ponman, phase space, velocity dispersion)

Using the galaxy membership within 1 Mpc, as for PCN, this method is based on the velocity dispersion of each cluster as discussed in Pearson et al. (in preparation). The velocity dispersion is determined using the Gapper estimator (Beers et al. 1990). From the virial theorem, we expect $M \propto \sigma^3$. In practice, both the normalization and power-law index of the relation between mass and velocity dispersion has been calibrated to the X-ray derived masses, extrapolated to $M_{200}$, and errors estimated, as described for PCN. Note: the values of the constants $a$ and $b$ of the relation $\log M_{200} = a + b \log \sigma_v$ employed by PCS can be found in Table 2.

3.23 PFS (Pearson & Ponman, FOF, velocity dispersion)

This algorithm is identical to PCS, except that it uses the FOF-linked galaxy membership of PFN. Note: the values of the constants $a$ and $b$ of the relation $\log M_{200} = a + b \log \sigma_v$ employed by PFS can be found in Table 2.

4 RESULTS: CLUSTER MASS COMPARISON

In this section, we present results comparing the recovered group/cluster masses of the different reconstruction methods to the ‘true’ masses in the catalogues. We first make comparisons when the group members are selected by the algorithms, before examining the simpler case where actual galaxy membership is specified (i.e. the case in which membership is known a priori). The differences between these results will allow us to distinguish between uncertainties due to the mass estimates and uncertainties due to the identification of group and cluster members. We note that supplying the ‘true’ galaxy membership does not necessarily guarantee an improvement, nor should it be expected: the methods have at most one or two free parameters, and are not generally tuned to each specific

| Method | $a$ | $b$ |
|--------|-----|-----|
| AS1    | 1.080 | 2.910 |
| AS2    | 1.080 | 2.910 |
| AvL    | 1.220 | 3.000 |
| CLE    | 1.064 | 3.000 |
| CLN    | 1.064 | 3.000 |
| SG1    | 1.034 | 2.975 |
| SG2    | 1.034 | 2.975 |
| PCS    | 0.668 | 2.280 |
| PFS    | 0.797 | 2.750 |

Table 2. Values of the constants $a$ and $b$ of the relation $\log M_{200}/10^{14} M_\odot = a + b \log (\sigma_v/1000$ km s$^{-1}$) employed by methods that utilize the group/cluster velocity dispersion. Please see Tables A1 and A2 in the Appendix for more details on each method.
data set but are rather run using their default settings. As such, if the default setting assumes a more restrictive or extensive galaxy membership, the calculation of mass from galaxy members selected in a different way is not guaranteed to be reliable. However, the level of scatter will indicate whether the calculation of mass from galaxy members selected in a different way is a serious issue.

Fig. 3 shows the recovered mass versus input mass when the group/cluster membership is not given in advance. The colour of scatter will indicate whether the calculation of mass from galaxy members selected in a different way is a serious issue.
Figure 4. Left hand side: rms error on log mass versus mean recovered mass (in dex) when the true galaxy membership is not known. The dashed black line identifies where the mean of the true mass distribution lies. Right hand side: rms error on log mass versus mean recovered mass (in dex) for three mass bins when the true galaxy membership is not known. Mass groups 1, 2 and 3 represent clusters with ‘true’ $M_{200}$ within the ranges $\log(M_{200}) \leq 14.25$, $14.25 < \log(M_{200}) \leq 14.45$ and $14.45 < \log(M_{200})$, respectively.

The scheme separates the methods into the broad classifications introduced in the last section and are classed as either richness (magenta), phase space (black), radius (blue), abundance matching (green) or velocity dispersion (red) based. Methods where membership is based on an FOF linking method have star-shaped markers, red sequence-based methods have diamond markers and phase space-based methods have circle markers. If returned, the vertical bars at the left hand side of each panel indicate the statistical (red) and systematic (blue) errors estimated internally by the mass reconstruction methods themselves without reference to this plot.

Encouragingly, we see a correlation across the input mass range $13 < \log(M_{200}/M_{\odot}) < 15$. There is generally good agreement, at least for the inferred masses of massive galaxy clusters. Nonetheless, one can see substantial scatter, especially at masses of $\log(M_{200}/M_{\odot}) < 14$, typically associated with groups. Although for some methods or mass regimes the masses may be slightly overestimated, the masses of groups and poor clusters appear to be more often underestimated, except for the methods based upon richness (PCN, PFN and NUM), as well as HBM. These biases are also apparent in Fig. B2, which shows the residual recovered mass in dex via $\log(M_{200,\text{Rec}}/M_{200,\text{True}})$. In some cases, these masses may be underestimated by more than an order of magnitude. Phase space methods MPO, MP1, TAR and velocity dispersion methods SGx fail to recover masses for some groups/clusters, but the number of such cases amounts to a very low fraction of the sample. The PCR method fails to recover reliable masses. This method uses the rms radius of the galaxy distribution extracted within a 1 Mpc aperture (and velocity range). However, this parameter is inflated by the presence of interlopers (which can removed statistically when calculating richness, for example) and is reduced by the imposed 1 Mpc aperture. It is noticeable that the PFR method, which is also based on rms radius, but is less affected by

Figure 5. The CDF of recovered halo mass delivered by the 23 methods when the group/cluster membership is not known. The true $M_{200}$ CDF is shown via the thick black dashed line. It is clear that all methods are recovering lower halo masses than that of the ‘true’ mass. A two-sample Kolmogorov–Smirnov (KS) test also demonstrates that the recovered mass distributions are not statistically similar to the true mass distribution (delivering $p$-values for all methods $p < 0.01$).
interlopers and has no restrictive aperture imposed, performs significantly better.

In the left-hand side of Fig. 4, we quantify the error in the estimated masses by calculating the rms of the difference between the recovered mass and the input mass (in dex) and we display this versus the mean of the recovered mass distribution. The black dashed vertical line identifies the mean of the true mass distribution. It is clear that the majority of methods, with the exception of AvL and MVM, are systematically biased to lower halo masses. For the majority of the methods, the rms error on $M_{200}$ is of the order of 0.3 dex (i.e. a factor of 2). Richness-based methods NUM, PFN and PCN produce the lowest rms values indicating lower scatter. Both PCR and HBM are outliers, delivering substantially higher rms values for all methods across all masses. As seen earlier, we find that the majority of methods have a higher mean deviation for groups/clusters in the lowest mass bin, this is highlighted in the right hand side of Fig. 4 where the rms of the difference between the recovered and input masses (in dex) is shown for the three mass bins: log($M_{200}$/M$_\odot$) $\leq$ 14.25, 14.25 $< \log(M_{200}) \leq$ 14.45 and 14.45 $< \log(M_{200})$. The bins are chosen so that there are roughly equal numbers of clusters in each mass bin. Here ‘1’ represents the method with the lowest rms and ‘23’ represents the method with the highest rms. Groups/clusters that are not recovered by the methods are excluded in this calculation. The overall rms ranking calculated for groups/clusters of all masses where the average bias of a given method has been subtracted, Rank$_{\sigma}$, is given in the second to last column. The overall rms ranking calculated without bias subtraction is given in the final column.

| Method | $\log(M_{200}) \leq 14.25$ | $14.25 < \log(M_{200}) \leq 14.45$ | $\log(M_{200}) > 14.45$ | All masses |
|--------|-----------------|-----------------|-----------------|-----------|
| Mean   | σ               | rms             | Rank            | Mean      | σ               | rms             | Rank            | Mean      | σ               | rms             | Rank            | Rank$_{\sigma}$ | Rank |
| PCN    | 0.14            | 0.12            | 0.18            | 2         | 0.10            | 0.07            | 0.12            | 1         | 0.08            | 0.07            | 0.11            | 1         | 2         |
| PFN    | 0.14            | 0.13            | 0.19            | 3         | 0.15            | 0.11            | 0.18            | 3         | 0.17            | 0.08            | 0.19            | 3         | 3         |
| NUM    | 0.14            | 0.11            | 0.17            | 1         | 0.15            | 0.09            | 0.17            | 2         | 0.18            | 0.09            | 0.20            | 4         | 1         |
| ESC    | 0.36            | 0.30            | 0.46            | 17        | 0.28            | 0.18            | 0.33            | 16        | 0.27            | 0.15            | 0.31            | 15        | 16        |
| MPO    | 0.28            | 0.26            | 0.38            | 10        | 0.20            | 0.17            | 0.26            | 5         | 0.17            | 0.15            | 0.23            | 6         | 12        |
| MP1    | 0.28            | 0.27            | 0.39            | 11        | 0.20            | 0.17            | 0.27            | 8         | 0.17            | 0.16            | 0.23            | 7         | 13        |
| RW     | 0.38            | 0.26            | 0.47            | 18        | 0.34            | 0.18            | 0.38            | 20        | 0.33            | 0.15            | 0.36            | 19        | 9         |
| TAR    | 0.23            | 0.26            | 0.35            | 7         | 0.19            | 0.18            | 0.26            | 6         | 0.21            | 0.28            | 0.35            | 18        | 10        |
| PCO    | 0.28            | 0.24            | 0.37            | 8         | 0.23            | 0.18            | 0.29            | 9         | 0.15            | 0.15            | 0.21            | 5         | 6         |
| PFO    | 0.26            | 0.31            | 0.41            | 14        | 0.21            | 0.21            | 0.29            | 11        | 0.13            | 0.13            | 0.19            | 2         | 11        |
| PCR    | 0.68            | 0.57            | 0.89            | 23        | 0.53            | 0.40            | 0.66            | 23        | 0.55            | 0.34            | 0.64            | 22        | 22        |
| PFR    | 0.32            | 0.20            | 0.38            | 9         | 0.33            | 0.16            | 0.37            | 18        | 0.37            | 0.17            | 0.41            | 20        | 14        |
| HBM    | 0.19            | 0.35            | 0.40            | 13        | 0.32            | 0.58            | 0.66            | 22        | 0.31            | 0.56            | 0.73            | 23        | 23        |
| MVM    | 0.29            | 0.27            | 0.40            | 12        | 0.24            | 0.21            | 0.32            | 14        | 0.21            | 0.16            | 0.26            | 11        | 19        |
| AS1    | 0.37            | 0.35            | 0.51            | 21        | 0.27            | 0.23            | 0.35            | 17        | 0.25            | 0.21            | 0.33            | 17        | 21        |
| AS2    | 0.32            | 0.33            | 0.46            | 16        | 0.23            | 0.22            | 0.32            | 15        | 0.22            | 0.20            | 0.30            | 14        | 20        |
| AvL    | 0.25            | 0.22            | 0.34            | 6         | 0.19            | 0.14            | 0.23            | 4         | 0.17            | 0.16            | 0.23            | 8         | 7         |
| CLE    | 0.34            | 0.32            | 0.47            | 19        | 0.23            | 0.18            | 0.29            | 10        | 0.20            | 0.14            | 0.24            | 9         | 17        |
| CLN    | 0.34            | 0.34            | 0.48            | 20        | 0.25            | 0.18            | 0.31            | 13        | 0.23            | 0.18            | 0.29            | 13        | 8         |
| SG1    | 0.35            | 0.29            | 0.46            | 15        | 0.29            | 0.24            | 0.37            | 19        | 0.24            | 0.16            | 0.29            | 12        | 18        |
| SG2    | 0.40            | 0.33            | 0.52            | 22        | 0.39            | 0.22            | 0.45            | 21        | 0.37            | 0.17            | 0.41            | 21        | 15        |
| PCS    | 0.23            | 0.16            | 0.28            | 4         | 0.25            | 0.15            | 0.29            | 12        | 0.28            | 0.13            | 0.31            | 16        | 4         |
| PFS    | 0.23            | 0.19            | 0.30            | 5         | 0.21            | 0.16            | 0.26            | 7         | 0.22            | 0.12            | 0.25            | 10        | 5         |

Note that though this group/cluster catalogue is derived from the same light-cone, the sample is not exactly the same as those in the majority of the remaining methods are very similar, with typical mass estimation errors of a factor of 2 to 3.
Figure 6. The true cluster mass versus recovered mass when the group/cluster membership is known. The black dotted line represents the 1:1 relation. ‘NR’ in the legend represents groups/clusters that are not recovered because they are found to have very low ($<10^{10} M_\odot$) or zero mass. The black ticks that lie across the 1:1 relation represent the minimum and maximum input group/cluster $M_{200}$. The vertical red bar represents the mean statistical error delivered by methods and the vertical blue bar represents the mean systematic error delivered by methods.

previous figures. In order to maintain a blind set-up for comparison, new clusters from the light-cone were added into the sample. As a result, this sample has, on average, poorer groups/clusters. One can see qualitatively similar correlations as in the previous results, but detailed comparisons indicate interesting differences between them. We immediately see that when the methods are not allowed to restrict the galaxy membership according to their usual scheme, for the majority of methods, the recovered masses are poorer (AvL, SG1,
5 RESULTS: CLUSTER MEMBERSHIP

We now examine the galaxy cluster membership delivered by the various methods and compare the richnesses of the recovered systems. Fig. 8 presents the richness of the recovered groups and clusters compared to the number of member galaxies in the source catalogue.

In general, the recovery of galaxy membership is very good and we find tighter relations with somewhat lower levels of scatter in comparison to the mass recovery results (also highlighted in Fig. C2 in the Appendix). Certain methods tend to miss members of massive clusters, such as both the AS and PC approaches. This deficit is intrinsic to these methods, in that they are deliberately conservative in their membership selection, focusing on the very central regions of each object; the bias that this selection introduces in the recovered mass is then calibrated out of the estimator. Other methods, such as CLE, ESC, PFx, RW, and SGx, are more contaminated by interlopers and consequently have richness estimates that are biased high. As mentioned earlier, this plot also illustrates that the high recovered mass up-scattered clusters seen in the ASx, ESC and SGx methods in Fig. 3 do not seem to be due to line-of-sight contamination by higher mass objects as very few objects have spuriously high numbers of recovered galaxies for these methods.

Most methods have distributions similar in shape to the true sample (as shown in Fig. C1 in the Appendix) although some distributions are, as noted above, offset either to high values due to interloper inclusion or low values due to conservative membership criteria. Finally, Fig. 9 compares the recovered richnesses as a function of recovered mass for the different methods. The three richness-based methods, PFN, PCN, and NUM, have, as expected, very tight relations. For both the input catalogue and these methods, there is a close relationship between richness and cluster mass which may not hold in the real universe. This strong correlation is a consequence of the simple model that we have adopted for this initial part of the comparison project. HBM, a velocity-dispersion-based method, also has a particularly narrow distribution combined with some catastrophic failures due to the mis-matching of a small number of groups/clusters. In contrast, many other methods have more scatter in both recovered number and recovered mass. Moreover, the slope of the \( N(M) \) relation is not well recovered for some methods (e.g. ASx, PCx) because of the variation in completeness as a function of mass seen in Fig. 8 for these methods: although they recover the halo masses to a similar level of accuracy as the other methods, they should not be used for reliable member galaxy determination.

6 DISCUSSION

The initial set-up used for this project was kept deliberately simple. We began with a simulated dark matter halo catalogue, and a model that inserts galaxies via smooth, spherically symmetric NFW distributions centred at the centre of the dark matter potential well and scaled by the mass of the halo. Within the \( z = 0 \) snapshot, haloes of mass above \( 10^{11.5} \, \text{M}_\odot \) (Fig. 1) are populated and a light-cone is then drawn through this distribution to create the ‘observations’ used for this test. Once this baseline study has quantified and minimized the uncertainties intrinsic in mass estimation, we will move on to a more sophisticated cluster model to identify the additional levels of uncertainty that such complexity introduces. Due to the simplicity of the model used for this initial phase and the use of a single cosmology, we cannot comment on the absolute calibration of each model, other than noting that the values have been calibrated to at least approximate reality. The main focus of this paper is to quantify the underlying scatter inherent in cluster mass estimation techniques that use the positions, velocities, and colours of galaxies.
Figure 8. Recovered number of galaxies associated with each group/cluster versus the true number of galaxies when the group/cluster membership is not known. The black dotted line represents the 1:1 relation and the black ticks represent the true minimum and maximum number of galaxies associated with the input group/clusters. ‘NR’ in the legend represents groups/clusters that are not recovered because they are found to have very low ($<10^{10} \, M_\odot$) or zero mass.

There are three general stages involved in galaxy-based cluster mass estimation. The first stage is the identification of a group/cluster overdensity, the second is the selection of galaxies deemed to be group/cluster members, and the third is the estimation of cluster properties based on this membership. These steps are not, in practice, independent from each other. For instance, a cluster mass estimation method based on dynamical properties might be very sensitive to contamination by unrelated field galaxies. As such,
it is perhaps better in such a method to be very conservative with the membership selection at the expense of completeness and then recalibrate the mass estimate based on this incomplete galaxy sample. Conversely, a method based on the volume covered might not be sensitive to interlopers but highly reliant on obtaining a nearly complete galaxy sample.

Following the philosophy of this study of making things as simple as possible, and to aid the inter-comparison of the results of different
methods, we supplied the participants with a list of initial centres (i.e. the first stage of this process) about which to look for structures. We further note that not all methods taking part in this study include this step. The centres of the group/cluster sample correspond with the location of the BCG in all cases and are the ‘true’ location of the halo centre in the DM simulation (the HOD model used places the brightest galaxy at the location of the most bound DM particle in the halo). Some methods (indicated by an asterisk in Table 1) chose not to use this information, and instead used the full galaxy catalogue detecting initial centres themselves. After calculating the properties of the identified groups/clusters, these methods then matched to our supplied coordinates. This is admirable and a more stringent test of these methods. We aim to investigate the issue of initial search location further in subsequent work.

We conclude from this study that, for clusters with masses above $10^{14} \, M_{\odot}$, the uncertainty in the methods seems to be around a factor of 2. Richness-based methods have the smallest uncertainties, but this reliability may be due to the underlying simplicity of the HOD model, which includes no asphericity, dynamical substructure or large-scale velocity distortions. However, we note that low scatter in the richness–mass relation has been observed for photometric samples (e.g. Rozo et al. 2014). Below $10^{14} \, M_{\odot}$, the scatter rises as the number of member galaxies drops, and the uncertainty rapidly approaches an order of magnitude. This level of error has severe implications for studies of cosmology based on cluster masses. Observational challenges such as spectroscopic tests of spherical systems with known locations and no imprecision in the cluster mass estimate. We stress that this experimental design did not improve mass estimates; for the majority of methods, we supplied the participants with a list of initial centres themselves. After calculating the properties of the identified groups/clusters, these methods then matched to our supplied coordinates. This is admirable and a more stringent test of these methods. We aim to investigate the issue of initial search location further in subsequent work.

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The authors contributed in the following ways to this paper: LO, RAS, FRP, & DC designed and organized this project. LO performed the analysis presented and wrote the majority of the paper. LO, ET, SIM, MEG, RP, TP & FRP organized the workshop that initiated this project. MRM & YW contributed to the analysis. The other authors (as listed in Section 3) provided results and descriptions of their respective algorithms. All authors helped proof-read the paper.

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APPENDIX A: PROPERTIES OF THE MASS RECONSTRUCTION METHODS

Table A1. Table illustrating the member galaxy selection process for all methods. The second column details how each method selects an initial member galaxy sample, while the third column outlines the member galaxy sample refining process. Finally, the fourth column describes how methods treat interloping galaxies that are not associated with the clusters.

| Methods | Initial galaxy selection | Member galaxy selection methodology | Treatment of interlopers |
|---------|--------------------------|-------------------------------------|--------------------------|
| PCN     | Within 5 Mpc, 1000 km s\(^{-1}\) | Clipping of ±3 \(\sigma\), using galaxies within 1 Mpc | Use galaxies at 3–5 Mpc to find interloper population to remove |
| PFN     | FOF                      | No                                  | No                       |
| NUM     | Within 1 Mpc, 1333 km s\(^{-1}\) | 1) Estimate \(R_{200}\) by a relationship between \(R_{200}\) and richness deduced from CLE; 2) Galaxies within \(R_{200}\) and with velocities less than 2.7 \(\sigma_{\text{los}}(R)\) are selected | No                       |
| ESC     | Within preliminary \(R_{200}\) estimate and ±3500 km s\(^{-1}\) | Gapper technique                  | Removed in refining by Gapper technique |
| MPO     | Input from CLN            | 1) Calculate \(R_{200}, R_p, R_{\text{red}}, R_{\text{blue}}\) by MAMPOST method; 2) Select members within radius according to colour | No                       |
| MP1     | Input from CLN            | Same as MPO except colour blind    | No                       |
| RW      | Within 3 Mpc, 4000 km s\(^{-1}\) | Within \(R_{200}\), \(|\Phi^{(R)}|^{1/2}\), where \(R_{200}\) obtained iteratively | No                       |
| TAR     | FOF                      | No                                  | No                       |
| PCO     | Input from PCN            | Input from PCN                      | Include interloper contamination in density fitting |
| PFO     | Input from PFN            | Input from PFN                      | No                       |
| PCR     | Input from PCN            | Input from PCN                      | Same as PCN |
| FFR     | Input from PFN            | Input from PFN                      | No                       |
| HBM     | FOF (ellipsoidal search range, centre of most luminous galaxy) | Increasing mass limits, then FOF, loops until closure condition | No                       |
| MVM     | Same as HBM               | Same as HBM                         | No                       |
| AS1     | Within 1 Mpc, 4000 km s\(^{-1}\), constrained by colour–magnitude relation | Clipping of ±3 \(\sigma\) | Removed by clipping of ±3 \(\sigma\) |
| AS2     | Within 1 Mpc, 4000 km s\(^{-1}\), constrained by colour–magnitude relation | Clipping of ±3 \(\sigma\) | Removed by clipping of ±3 \(\sigma\) |
| AvL     | Within 2.5 \(\sigma_{\text{v}}\) and 0.8 \(R_{200}\) | Obtain \(R_{200}\) and \(\sigma_{\text{v}}\) by \(\sigma\)-clipping | No                       |
| CLE     | Within 3 Mpc, 4000 km s\(^{-1}\) | 1) Estimate \(R_{200}\) by aperture velocity dispersion; 2) galaxies within \(R_{200}\) and with velocities less than 2.7 \(\sigma_{\text{los}}(R)\) are selected; 3) Iterate steps 1 and 2 until convergence | Obvious interlopers are removed by velocity gap technique, then further treated in iteration by \(\sigma\) clipping |
| CLN     | Input from NUM            | Same as CLE                         | Same as CLE |
| SG1     | Within 4000 km s\(^{-1}\) | 1) Measure \(\sigma_{\text{gal}}\), estimate \(M_{200}\) and \(R_{200}\); 2) Select galaxies within \(R_{200}\); 3) Iterate steps 1 and 2 until convergence | Shifting gapper with minimum bin size of 250 kpc and 15 galaxies; velocity limit 1000 km s\(^{-1}\) from main body |
| SG2     | Within 4000 km s\(^{-1}\) | 1) Measure \(\sigma_{\text{gal}}\), estimate \(M_{200}\) and \(R_{200}\); 2) Select galaxies within \(R_{200}\); 3) Iterate steps 1 and 2 until convergence | Shifting gapper with minimum bin size of 150 kpc and 10 galaxies; velocity limit 500 km s\(^{-1}\) from main body |
| PCS     | Input from PCN            | Input from PCN                      | Same as PCN |
| PFS     | Input from PFN            | Input from PFN                      | No                       |
Table A2. Table showing the characteristics of the mass reconstruction process of methods used in this comparison. The second, third, fourth and fifth columns illustrate whether a method calculates/utilizes the velocities, velocity dispersion, radial distance of galaxies from cluster centre, the richness and the projected phase space information of galaxies, respectively. If a method assumed a mass or number density profile it is indicated in columns six and seven.

| Methods | Velocities | Velocity dispersion | Radial distance | Richness | Projected phase space | Mass density profile | Number density profile |
|---------|------------|---------------------|----------------|----------|-----------------------|---------------------|------------------------|
| PCN     | Yes        | No                  | No             | Yes      | No                    | No                  | No                     |
| PFN     | Yes        | No                  | No             | Yes      | No                    | No                  | No                     |
| NUM     | Yes        | No                  | No             | Yes      | No                    | No                  | No                     |
| ESC     | Yes        | Yes                 | Yes            | No       | No                    | Caustics            | No                     |
| MPO     | Yes        | No                  | Yes            | No       | Yes                   | NFW                 | Yes                    |
| MP1     | Yes        | No                  | Yes            | No       | Yes                   | NFW                 | Yes                    |
| RW      | Yes        | No                  | Yes            | No       | No                    | NFW                 | No                     |
| TAR     | Yes        | Yes                 | Yes            | No       | No                    | NFW                 | No                     |
| PCO     | Yes        | No                  | No             | No       | No                    | NFW                 | Yes                    |
| PFO     | Yes        | No                  | No             | No       | No                    | NFW                 | Yes                    |
| PCR     | Yes        | No                  | Yes            | No       | No                    | No                  | No                     |
| PFR     | Yes        | No                  | Yes            | No       | No                    | No                  | No                     |
| HBM     | Yes        | Yes                 | Yes            | No       | No                    | NFW                 | No                     |
| MVM     | Yes        | Yes                 | Yes            | No       | No                    | NFW                 | No                     |
| AS1     | Yes        | Yes                 | No             | No       | No                    | No                  | No                     |
| AS2     | Yes        | No                  | Yes            | No       | Yes                   | No                  | No                     |
| AsL     | Yes        | Yes                 | Yes            | No       | No                    | No                  | No                     |
| CLE     | Yes        | Yes                 | No             | No       | No                    | NFW                 | NFW                    |
| CLN     | Yes        | Yes                 | No             | No       | No                    | NFW                 | NFW                    |
| SG1     | Yes        | Yes                 | Yes            | No       | No                    | No                  | No                     |
| SG2     | Yes        | Yes                 | Yes            | No       | No                    | No                  | No                     |
| PCS     | Yes        | Yes                 | No             | No       | No                    | No                  | No                     |
| PFS     | Yes        | Yes                 | No             | No       | No                    | No                  | No                     |
Figure B1. Recovered mass distributions when the group/cluster membership is not known. The black dotted line represents the mean of the true mass distribution and the grey distributions on each subplot represent the true mass distributions.
Figure B2. Mass bias versus true mass when the group/cluster membership is not known. The black dotted line represents a residual of zero. ‘NR’ in the legend represents groups/clusters that are not recovered because they are found to have very low (<10^6 M_☉) or zero mass.
Figure B3. Recovered mass distributions when the group/cluster membership is known. The black dotted line represents the mean of the true mass distribution and the grey distributions on each subplot represent the true mass distributions.
Figure B4. Mass bias versus true mass when the group/cluster membership is known. The black dotted line represents a residual of zero. ‘NR’ in the legend represents groups/clusters that are not recovered because they are found to have very low ($<10^{10} \text{M}_\odot$) or zero mass.
Figure C1. Distributions of the recovered number of galaxies associated with each group/cluster when the membership is not known. The grey distribution on each subplot represents the true richness distribution of the groups/clusters. The black dotted line presents the mean of this input richness distribution.
Figure C2. Richness bias versus true richness when the group/cluster membership is not known. The black dotted line represents a residual of zero. ‘NR’ in the legend represents groups/clusters that are not recovered because they are found to have very low ($<10^{10} M_\odot$) or zero mass.

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