

Automatic geomorphological mapping using ground truth data with coverage sampling and random forest algorithms

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21

Abstract

Marine geomorphological maps are useful to understand seafloor structure for 22 example in the context of ecological studies, resources management or con-23 servation planning. Although techniques to build such maps are increasingly 24 sophisticated, manual techniques are still largely used. Automated approaches 25 are needed to get reproducible maps in a reasonable time. This work provides 26 statistical learning approaches based framework to build automatically geomor-27 phological maps. We used bathymetric data to build Digital Bathymetric Model 28 (DBM) and compute terrain attributes characteristic of seafloor geomorphol-29 ogy. Then, we used clustering based algorithms to select automatically ground 30

truth locations from a reference geomorphological map manually made. Finally 31 a supervised classification model random forest based was used to build predic-32 tive models for seafloor geomorphology typologies. Subsequently we studied the 33 effect of DBM resolution, sample size and sampling method of the ground truth 34 locations, in the quality of map production via a series of simulations. Results 35 showed that the proposed framework allowed to build efficiently relevant seafloor 36 geomorphological maps. The best compromise between the sampling effort and 37 the quality of the resulting maps was obtained with 100 m DBM resolution, 200 38 data points sample size and using a complexity-dependent sampling method and 39 led to a map matching at 90% the reference one. 40

41 **Keywords:** geomorphological map, spatial modeling, random forest classification,

42 digital bathymetric model, terrain attributes, lidar data

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43 Declarations

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- ⁴⁵ All accepted principles of ethical and professional conduct have been followed during
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67 1 Introduction

Geomorphological maps are georeferenced delineation of morphological structure and 68 surface composition of a studied land and/or seafloor (Otto and Smith, 2013; Dramis 69 et al, 2011; Pavlopoulos et al, 2009). Marine geomorphological maps are particularly 70 crucial for resource management, conservation efforts, hazard assessment, protected 71 area management, effective marine research campaign planning and various marine-72 related industrial works (Kienholz, 1978; Bishop et al, 2012; Fukunaga et al, 2019; 73 Browne et al. 2010). Such maps provide valuable information on the composition and 74 structure of the seabed which, among other usage, is needed to generate habitat maps 75 through the identification and delineation of different benchic ecosystems like coral 76 reefs, seagrass beds or various deep-sea communities (Pandian et al, 2009; Dramis 77 et al, 2011; Wabnitz et al, 2008). 78

To produce geomorphological maps, different approaches are used (Siart et al, 2009; 79 Hugenholtz et al, 2013). Widely used imagery techniques involve studying the pat-80 terns, textures, shapes, and color variations present in the imagery. This can be done 81 manually by digitizing or tracing the features on the imagery or through automated 82 or semi-automated image segmentation and classification techniques in Geographical 83 Information System (GIS) using available image analysis tools. While imagery can be 84 a valuable tool, it has a limited use for mapping deeply submerged geomorphological 85 features due to water opacity. In such conditions, only acoustic approaches can pro-86 vide usable data which are generally completed by punctual carefully located ground 87 truthing observations using for example scuba-divers or submarine-divers observation, 88 Remote Operated Vehicle (ROV) or Automatic Underwater Vehicle (AUV) picture 89 or videos, drop cameras or seabed sampling (Wynn et al, 2014; Locker et al, 2010). 90 Subsequent treatments, required to generate maps with such data, will be to first pro-91 pose a geomorphologic category for each ground truthing point (ie. Typology), then 92 delineate surfaces of homogeneous typologies. Depending on whether ground truthing 93 points are defined as categories or as quantitative data, interpolation methodologies 94 within the surface to be mapped might take different forms. 95

Many semi-automated or automated approaches have been also proposed in recent 96 decades to achieve objective, automated and repeatable approach to extract mean-97 ingful information using vast quantities of data (Summers et al, 2021). Object-Based 98 Image Analysis (OBIA) which use bathymetric derivatives or a combination of bathy-99 metric derivatives and backscatter to automatically segment the seafloor (Masetti et al, 100 2018; Argyropoulou et al, 2016; Lacharité et al, 2018; Koop et al, 2021; Dekavalla 101 and Argialas, 2017) are widely used. Bathymorphon / geomorphon-based classification 102 (Jasiewicz and Stepinski, 2013; Sowers et al, 2020; Ahn et al, 2023; Novaczek et al, 103 2019) and fuzzy logic scheme (Schmidt and Hewitt, 2004; Lucieer and Lucieer, 2009; 104 Janowski et al, 2021) have been also investigated. The past 10 years, machine learning 105 (Maschmeyer et al, 2019; Misiuk et al, 2021; Janowski et al, 2022; Sklar et al, 2024) 106 and deep learning models (Behrens et al, 2018; Azarafza et al, 2023; Arhant et al, 107 2023) has increasingly been used for geomorphological mapping. Furtherthemore, the 108 performance of these statistical learning methods has also been investigated in com-109 parison with manually ones (Van der Meij et al, 2022; Diesing et al, 2014). In recent 110

¹¹¹ years, although these methods have proven their effectiveness, they are often combined

with underwater imagery which superseded expert manual interpretation and are par-112 ticularly costly for large scale mapping (Van der Meij et al, 2022; Cui et al, 2021; 113 Galvez et al, 2022; Misiuk and Brown, 2023; Breyer et al, 2023). As alternative, this 114 work provides a clustering based algorithm for an optimal ground truth sampling and 115 a learning-based approach for automatic geomorphological mapping. It focuses on the 116 classical situation where experts use morphological map to define surfaces and ground 117 truth measure to define typologies. But to that respect, further considerations need to 118 be addressed: (i) the quality of bathymetric data required, (ii) the typology definition, 119 and (iii) the methodology to use for creating geomorphological maps with such data. 120 (i) Bathymetry is recognized as essential when mapping geomorphology (Wilson et al, 121 2007; Lecours et al, 2016; Fukunaga et al, 2019). It is particularly useful for identify-122 ing and delineating submerged landforms that are not visible in aerial images. Modern 123 bathymetric systems can provide high-resolution data, capturing fine-scale details of 124 the seafloor morphology. Their accuracy is often higher compared to aerial images, as 125 it directly measures the water depth and seafloor elevation. However, the selection of 126 the appropriate technology is crucial to ensure the acquisition of accurate and reliable 127 data suited for an optimal geomorphological map production. More precisely equip-128 ment and material setup will depend on factors such as the scale and coverage needed, 129 water depth, desired resolution, sampling effort and budgetary considerations. Com-130 promise between the sampling effort and the quality of the data have to be made but 131 little tools are available to choose the most optimal setup. Indeed very high-resolution 132 data may be necessary for detailed local studies, while coarser resolution data might 133 suffice for larger-scale regional or global analyses. In the present research, some exam-134 ples to help on such decisions are proposed. 135

(ii) A typology is a description of one geomorphological category of seabed (in our 136 case). However, to make a geomorphological map usable, comparable and understand-137 able by a large international community, categories definition of typologies need to 138 form a consensus. Our study is based on the Millennium Coral Reef Mapping Project 139 (MCRMP) typologies. Initiated by the Institute for Marine Remote Sensing - Univer-140 sity of South Florida (IMaRS/USF) in 2001 and continued since 2003 by reasearchers 141 of the Institut de Recherche pour le Développement (IRD), the MCRMP has proposed 142 a multi-level hierarchical structure (Andréfouët et al, 2004). MCRMP typologies were 143 widely used as they enable several sites around the world to be compared on a the-144 matically rich, homogeneous basis (Andréfouët and Dirberg, 2006). These typologies 145 provide a description of coral reef geomorphology distinguishing reef units such as 146 slopes, flats, passes, terraces, lagoons, channels, etc. 147

(iii) Several approaches can be chosen to generate geomorphological maps from pre-148 viously described data. The most classical approach is to manually draw typologies 149 envelopes using GIS softwares (Minár and Evans, 2008; Otto et al, 2018). However, 150 it is tedious and poorly replicable if temporal changes need to be monitored. Auto-151 mated approaches are of several kind. This work focuses on statistical learning based 152 approaches for their ability to efficiently process and analyze large volumes of spatial 153 data, to learn complex relationships present in the data leading to improved accu-154 racy in produced maps (Stepinski et al, 2007; Siqueira et al, 2022; Van der Meij et al, 155 2022). More precisely four steps are followed: 1) generate a Digital Bathymetric Model 156

(DBM) from bathymetric data, 2) compute terrain attributes from DBM on the entire
surface studied, 3) train random forest based classification algorithm to match terrain
attributes with ground truthing typologies and 4) predict typologies from the entire
surface studied using the classification model generated.

The chosen methodology to generate geomorphological map was tested on a classi-161 cal tropical reef feature: an atoll mapping from South Western Indian Ocean. Our 162 approach is to propose a sensitivity analysis based on a comparison with an existing 163 expert map while degrading DBM definition as proxy of data acquisition setup and 164 varying the number of ground truth points as well as the methodology to select their 165 locations as proxy of field effort. The ultimate goal is to supply tools to plan mapping 166 field campaign using coverage sampling algorithms, codes to semi-automate geomor-167 phological mapping procedure using random forest algorithm and propose metrics to 168 evaluate the quality of the map generated across different resolutions. Recommenda-169 tions to choose the DBM resolution, ground truth size and sites selection are also 170 provided. 171

¹⁷² 2 Materials and methods

173 2.1 Data

For this work, bathymetric data (depth measurements) collected between 2009 and 2010 on Geyser atoll with a surface area of approximately 268 km² are used (Figure 1A). A set of 48.10⁶ data points were collected by LIDAR 1 m resolution calibrated. The depth range captured by this tool often reaches down to -30 m but due to exceptional very good water clarity conditions, bathymetric records on Geyser range between -50 and 4 meters (see their distribution in Figure S1). This data is available on the Hydrography and Oceanography Service of the Navy website ¹.

An expert geomorphological map of the Geyser atoll available here ² is also used. This
 scale-free map was produced by manual contouring, resulting from expert interpreta tion of hyperspectral images. 11 geomorphological typologies have been identified on

 $_{184}$ Geyser (Roos et al, 2017).

¹⁸⁵ 2.2 General methodology

Here an automated scheme using bathymetry and some field typologies verifications 186 also called ground truth to generate reproducible geomorphological maps is used (see 187 Figure S2 in the supplemental materials). The methodology is the following: 1) Using 188 bathymetric data, a Digital Bathymetric Model (DBM) is created at a given resolu-189 tion. 2) From the DBM, terrain attributes corresponding to raster layers which have 190 191 the potential to influence seafloor geomorphology are computed. 3) Using a given coordinate sampling method, a given number of ground truth locations is drawn. 192 By overlaying these locations with the expert map, geomorphological typologies are 193 attributed to each ground truth points. 4) Using the training set of coupled terrain 194 attributes and ground truth typologies, a recursive feature elimination algorithm based 195

 $[\]label{eq:linear} ^{1} \mbox{https://data.shom.fr/donnees} ^{2} \mbox{https://sextant.ifremer.fr/geonetwork/srv/api/records/21232c12-e409-4136-a24a-78c346518cfa}$



Fig. 1 Geyser is an atoll in the Western Indian ocean, specifically located in the northern Mozambique channel between Mayotte and the Gloriosos Islands and covering approximately 268 km². A Bathymetry data (in meters) collected by LIDAR technology and provided by Hydrography and Oceanography Service of the Navy (SHOM). B Geomorphologic structure identified for the Geyser atoll, as a part of the EPICURE project (Roos et al, 2017).

on a random forest classifier is used to select most relevant covariates. The latter are then used to train a random forest based predictive model for geomorphologic typologies. This model is finally used to predict geomorphologic typologies on the whole study site. 5) The quality of the map production was then evaluated using two types of performance criteria: the model performance (Balanced accuracy) and the matching between the generated map and the expert map (Match and Balanced match).

The robustness of our approach is also evaluated using a sentivity analysis through 202 simulation by varying bathymetric data definition, numbers of ground truth points and 203 methodology to select their locations. This involved in examining five DBM resolutions 204 (5 m, 25 m, 50 m, 100 m and 500 m), six sample sizes (50, 100, 200, 500, 700, 1000) 205 and three sampling methods (two spatial coverage sampling methods denoted SCS-206 KMEANS and SCS-CLARA, and a complexity-dependent sampling method denoted 207 CDS). For each of these combinations $(5 \times 6 \times 3)$, 30 replicates of geomorphological 208 maps are generated by replicating the steps 3-5 described above. 209

7

2.2.1 Digital Bathymetric Model creation 210

The raw bathymetric dataset contains approximately 48.10^6 data points giving lat-211 itude, longitude and depth values. To avoid numerical issues due to redundant 212 observations, this dataset was spatially sub-sampled and data points are kept as homo-213 geneous as possible using the *buffer.points* function in the supplemental materials 214 (Roberts, 2015). The sub-sampled data are such that each retained data point is at 215 least at a distance of 5 m from one another. Following this procedure, the new dataset 216 of approximately 8.10^6 data points obtained is used to generate the DBM which con-217 sists in the creation of a square grid of T^2 cells or rasters. The cell size define the DBM 218 resolution. Different resolutions (5 m, 25 m, 50 m, 100 m, 500 m)³ are created using 219 the *dbm* function proposed in the supplemental materials. The bathymetry or depth 220 value of each cell is calculated by taking the average of all the depths inside the same 221 cell. 222

Furthermore, some deep areas are poorly sampled, due to the lack of signal return 223 on the LIDAR sensor (see white zone in the middle of Figure 1A). To get a depth 224 measure for each ground truth location selected using one of the automatic sampling 225 methodology presented further, depth measure on each raster of the study area were 226 required. Thus for each DBM resolution, missing depth of empty cells in this zone 227 are interpolated using an ordinary kriging model via the *ok.dbm* function provided in 228 the supplemental materials. The kriging method is not decribed in details since the 229 problem of missing data is out of the scope of the paper (Cressie, 1988). In contrast, 230 empirical results provided in the supplemental materials show that it performs better 231 than five others spatial interpolation methods for our data Table S1. This imputation 232 method allows to complete bathymetric data on these cells. 233

2.2.2 Terrain attributes calculation 234

It consists in quantifying predictors for seafloor geomorphology. For each DBM reso-235 lution, terrain attributes were calculated using a moving routine from the DBM. More 236 precisely, let us denote the depth $D_{i,j}$ of a given cell with $(i,j) \in [\![1,T]\!] \times [\![1,T]\!]$ and con-237 sider the depth $D_{i+k,j+l}$ of the neighboring cells with $(k,l) \in \mathcal{D}_3 = [-1,1] \times [-1,1]$. 238 These cells are defining the 3×3 window on which the terrain attributes are 239 defined. Note that it is possible to generalize the study to square window with 240 size greater than 3. Selected terrain attributes can be organised into three groups: 241 Slope (Slope) and orientation (Aspect) measures, terrain variability measures such 242 as Roughness (Roughness), Terrain ruggedness index (TRI) and Vector ruggedness 243 measure (VRM) and curvature and relative position measures such as Profile convex-244 ity (profc), Planform convexity (planc) and Bathymetric position index (BPI). 245

Slope has been widely recognized as an important factor for determining benchic habitat and colonization and has been used in many marine studies (Copeland et al, 2013; 247

Fukunaga et al, 2019; Sterne et al, 2020). Like the magnitude of the steepest drop 248

 $^{^{3}}$ The 5 m resolution is the one used by biologists during field verification campaigns. We then looked for a very high resolution at which the geomorphological maps produced were degraded because they were too pixelated. 500 m seemed to be a good choice. Between these two resolutions, we empirically searched for intermediate resolutions enabling us to obtain "significantly" different maps. Hence the 25 m, 50 m and 100 m resolutions were retained

⁸

in depth, Slope (in degrees) is derived from rates of change in x (longitude) and y (latitude) directions and is calculated as follows, for each cell $(i, j) \in [\![1, T]\!] \times [\![1, T]\!]$,

$$Slope_{i,j} = \frac{180}{\pi} \arctan\left(\sqrt{\left(\frac{\partial D_{i,j}}{\partial x}\right)^2 + \left(\frac{\partial D_{i,j}}{\partial y}\right)^2}\right).$$
 (1)

where

$$\begin{cases} \frac{\partial D_{i,j}}{\partial x} = \left[(D_{i+1,j+1} + 2D_{i+1,j} + D_{i+1,j-1}) - (D_{i-1,j+1} + 2D_{i-1,j} + D_{i-1,j-1}) \right] / 8\Delta x \\ \frac{\partial D_{i,j}}{\partial y} = \left[(D_{i+1,j+1} + 2D_{i,j+1} + D_{i-1,j+1}) - (D_{i+1,j-1} + 2D_{i,j-1} + D_{i-1,j-1}) \right] / 8\Delta x \end{cases}$$

where the real number Δ stands for the cell size of the grid.

The orientation measure (Aspect) gives the exposure of a given area to such water waves and is often used in the calculation of others parameters that directly influence habitat (Wilson et al, 2007). Aspect (in degrees) is the compass direction of the steepest drop in depth and is calculated as follows, for each cell $(i, j) \in [\![1, T]\!] \times [\![1, T]\!]$,

$$Aspect_{i,j} = 180 + \frac{180}{\pi} \arctan\left(\frac{\partial D_{i,j}}{\partial x} + \frac{\partial D_{i,j}}{\partial y}\right).$$
 (2)

The roughness measure (*Roughness*) is a critical factor affecting ecological and physical processes on the reef (Leon et al, 2015; Dartnell, 2000). It corresponds to the difference between the maximum and minimum depth values over a 3×3 window and is defined for each cell $(i, j) \in [\![1, T]\!] \times [\![1, T]\!]$ as follows:

$$Roughness_{i,j} = \max_{k,l \in \mathcal{D}_3} (D_{i+k,j+l}) - \min_{k,l \in \mathcal{D}_3} (D_{i+k,j+l}).$$
(3)

The Terrain Ruggedness Index (TRI), is a terrestrial ruggedness measure (Riley et al, 1999) that was adapted to bathymetry data to highlight morphological heterogeneity (Valentine et al, 2004; Rozycka et al, 2017). It is defined as the mean of the absolute differences between the depth value of a cell and the one of its neighboring cells, for each cell $(i, j) \in [\![1, T]\!] \times [\![1, T]\!]$ as follows:

$$TRI_{i,j} = \frac{\sum_{k,l \in \mathcal{D}_3} |D_{i+k,j+l} - D_{i,j}|}{(3^2 - 1)}.$$
(4)

The Vector Ruggedness Measure (VRM) quantifies terrain ruggedness : slope and aspect are decomposed into 3-dimensional vector components using standard vector analysis in a user-specified moving 3×3 window. The vector ruggedness measure is dimensionless because it involves sine and cosine of the slope and aspect measures and its values range from 0 to 1 corresponding to flat regions to rugged ones. Its mathematical definition is omitted to avoid technicalities and details can be found in

 $_{271}$ (Sappington et al, 2007).

The Curvature position may also be linked to the nature of the seabed. It helps to 272 delimit regions of distinct habitat by identifying boundaries in the character of the 273 terrain ((Wilson et al, 2007)). Bathymetric Position Index (BPI), the marine version 274 of the topographic position index, quantifies where a location on a bathymetric surface 275 is relative to the overall seascape (Mata et al, 2021). It provides an indication of 276 whether any particular pixel forms part of a positive (e.g., crest) or negative (e.g., 277 trough) feature of the surrounding terrain (Lundblad et al, 2006; Wilson et al, 2007). 278 It is calculated using the following formula, for each cell $(i, j) \in [1, T] \times [1, T]$: 279

$$BPI_{i,j} = D_{i,j} - \frac{\sum_{k,l \in \mathcal{D}_3} |D_{i+k,j+l} - D_{i,j}|}{(3^2 - 1)}.$$
(5)

According to (Evans, 1980), Profile convexity (Profc) is the rate of change of Slope 280 and Plan convexity (*Planc*) is the rate of change of Aspect. Negative values in the 281 *Profc* indicate the surface is upwardly convex whereas, positive values indicate that 282 the surface is upwardly concave. Positive values in the *Planc* means the surface is lat-283 erally convex and negative values indicate that the surface is laterally concave. Several 284 methods exist for numerical approximations of these metrics, based on a quadratic 285 form representation f of the surface (Florinsky, 1998; Horn, 1981; Evans, 1980; Zeven-286 bergen and Thorne, 1987). Numerical implementation of (Zevenbergen and Thorne, 287 1987) method's is used in this study; details can be found in (Florinsky, 1998). 288 All the computed terrain attributes, depth and geographic coordinates (longitude, 289

290 latitude) are then stacked to form a multilayer grid of predictors called features or covariates in the sequel Table 1.

Terrain attributes	Reference	Function / R Package					
Slope and Aspect							
Slope	(Horn, 1981)	terrain / raster					
Aspect	(Horn, 1981)	terrain / raster					
Terrain Variability							
Roughness	(Dartnell, 2000)	terrain / raster					
Terrain Ruggedness Index (TRI)	(Wilson et al, 2007)	terrain / raster					
Vector Ruggedness Measure (VRM)	(Ilich et al, 2023)	VRM / MultiscaleDTM					
Curvature and relative position							
Profile Curvature (Profc)	(Zevenbergen and Thorne, 1987)	Curvature / spatialEco					
Planform Curvature (<i>Planc</i>)	(Zevenbergen and Thorne, 1987)	Curvature / spatialEco					
Bathymetric Position Index (BPI)	(Ilich et al, 2023)	BPI / MultiscaleDTM					

²⁹² 2.2.3 Sampling

To build predictive models for geomorphological typologies using a statistical learning approach, a training dataset is required. This one must contain a finite number of ground truth locations and a set of predictive covariates for geomorphological typologies at these locations. In this section, three alternative clustering based sampling methods are proposed to draw automatically these locations inside a given study area. All these methods use cell's centers of a regular grid and choose among the covariates earlier mentioned depending on the choosen algorithm.

Spatial Coverage Sampling using k-means clustering algorithm (SCS-KMEANS)

The basic idea of Spatial Coverage Sampling (SCS) is to draw uniformly sampling loca-302 tions over the study area. It has been shown that SCS on a study area can be achieved 303 by k-means clustering algorithm (Hartigan, 1975). This consists in grouping cell's cen-304 ters of a regular grid on this area using their spatial coordinates as covariates. Note 305 that this regular grid can be the DBM one as long as it does not lead to computional 306 deadlock, otherwise it can be replaced by a raster grid with lower resolution. The final 307 solution of this partition gives the sampling locations and is determined by minimizing 308 a geometric criterion, the mean squared shortest distance between the clusters centroids and the grid cell's centers (Royle and Nychka, 1998; Brus et al, 2006). For the 310 implementation, a k-means algorithm for equal area partitioning is used (Brus, 2019). 311 The scsKM function provided in the supplemental materials can be used to achieve 312 this 313

Spatial Coverage Sampling using CLARA algorithm (SCS-CLARA) 314 K-means clustering approach is time and storage consuming, especially for high DBM 315 resolution. In such case, CLARA algorithm approach could be an alternative. The 316 CLARA algorithm is an extension of the Partitioning Around Medoids (PAM) meth-317 ods (Kaufman and Rousseeuw, 1975) to deal with data containing a large number of 318 objects (more than several thousand observations) in order to reduce computing time 319 and storage problem. Medoids $(M_i)_{i=1}^k$ in a PAM, k being the desired number of clus-320 ters C_i , are cells that minimize their distance to other cell's centers of the cluster. The 321 CLARA algorithm generates $j \in \mathbf{N}^*$ random samples of size n $(n < T^2)$ on individu-322 als, applies a PAM on these samples one after the other, then evaluates the partition 323 quality on each of them by calculating the average global dissimilarity on the complete 324 dataset as follows: 325

$$\sum_{i=1}^{k} \sum_{x_c \in C_i} \frac{d(x_c, M_i)}{T^2}$$
(6)

³²⁶ If this dissimilarity is lower than the previous found one, it considers this solution and ³²⁷ its k medoids as the best current solution. The *scsCLARA* function in the supple-³²⁸ mental materials can be used to choose ground truth locations.

³²⁹ Complexity-Dependent Sampling using CLARA algorithm (CDS)

Terrain morphology can sometime be marked by accidented region where many typologies are observed closed to each other. This is particularly true with coral reef region where for example sand flat, reef slope, reef flat, etc typologies can be close to each

other. In such situations, homogeneous sampling would lead to miss many typolo-333 gies unless a lot of points are requested leading to other problems such as important 334 unbalanced typologies distribution among locations (Brus, 2019). To account for such 335 typologies distribution, ground truth locations are agglomerated around zone of higher 336 ground complexity (i.e. complex terrain). Thus the CDS method is introduced in 337 order to take advantage of these covariates available for each DBM resolution. CDS 338 aims to distribute sampling locations around most heterogeneous or complex areas. 339 It uses DBM's cells as individuals and unlike the SCS which uses the spatial coordi-340 nates, CDS's covariates are chosen among terrain attributes and depth. Depth and 341 Roughness measures are considered in this study. CDS cannot be computed using the 342 k-means clustering algorithm for high DBM resolutions because DBM cells are manda-343 tory to get covariates. This is why the CLARA algorithm is only used. The cdCLARA344 function in the supplemental materials can be used to this end (Figure 2).



Fig. 2 Example of 100 ground truth locations drawn with the different methods: A Spatial Coverage Sampling using K-means clustering algorithm (SCS-KMEANS), B Spatial Coverage Sampling using CLARA algorithm (SCS-CLARA) C and Complexity-dependent sampling using a CLARA algorithm (CDS). Roughness and Depth are used to guide the complexity-dependent sampling.

345

³⁴⁶ 2.2.4 Geomorphology mapping using Random Forest algorithm

To map geomorphological typologies over a whole study area, a supervised classifica-347 tion approach is considered. This work was divided in two steps: a first step to train 348 algorithm and a second step to predict typologies based on trained algorithm. For 349 the training part, each location is generated by one of the previously described sam-350 pling methods and typologies were attributed using the expert map Figure 1B. These 351 located typologies were the target variable, locations coordinates and the correspond-352 ing depth and terrain attributes were used as covariates and both formed the training 353 dataset. In this process, a feature selection scheme random forest based is used to sub-354 set the most relevant covariates of the training set. Then a final model is fitted using 355 the selected covariates. After this first step completed, the second step consisted in 356 using the fitted model to predict the most suitable typology over the whole study area. 357 The generic principle of Random Forest 358

³⁵⁹ The tree based Random Forest (RF) algorithm can be used for a classification task

(Breiman, 2001; Biau and Scornet, 2016). Using the Bootstrap AGGregatING (bagging) principle, RF increases the diversity of the trees by making them grow from different randomly drawn (with replacement) training datasets from the original dataset (Breiman, 1996). At each node of each tree, Rf selects a random subset of features and search for the best split for the node. To classify a new case once the forest is completed, the typology having the most votes over all the trees is retained.

366 Model training

Using a cross-validation scheme with 3 repeats, the training samples are split into 3 367 folds. To train a RF model, three hyperparameters were tuned: the number of trees 368 (Ntrees) of the forest, the number of features used at each node (Mtry) and the min-369 imum number of data points at the terminal node of each tree (nodesize). Indeed, 370 the Ntrees hyperparameter is not tunable in the classical sense but should be set suf-371 ficiently high (Díaz-Uriarte and Alvarez de Andrés, 2006; Oshiro et al, 2012; Probst 372 et al, 2019). The default value of 500 trees is used. The nodesize hyperparameter has 373 been set to 1 for classification task because it generally provides good results (Díaz-374 Uriarte and Alvarez de Andrés, 2006). The Mtry hyperparameter was tuned among 375 fifteen values of hyperparameters chosen automatically by the function tuneLength. 376 A random search optimization strategy which defines a search space as a bounded 377 domain of parameter values and randomly sample points in that domain were used to 378 find the optimal Mtry considering the the resulting Accuracy (Grandini et al, 2020). 379 Models are fitted by repeatedly leaving out one of the folds and performance are deter-380 mined by predicting on the fold left out. The train function of the caret R package 381 were used to this end. 382

383 Terrain attributes selection

By selecting the most relevant terrain attributes as covariates, the risk of over fitting 384 can be reduced and the model's generalization ability improved. Variable or feature 385 importance measures are usually used to rank or select variables. Mean Decrease Impu-386 rity (MDI) and Mean Decrease Accuracy (MDA) are two well-known random forest 387 variable importance measures (Guyon and Elisseeff, 2020; Breiman, 2001; Biau and 388 Scornet, 2016). In this study, the MDA measure also called permutation importance 389 in Breiman's original random forest is chosen since it seems to exhibit less bias than 390 MDI in presence of correlated features (Strobl et al, 2008; Breiman, 2001). Roughly 391 speaking, the MDA measure consists in shuffling values of a given covariate j in the 392 test data or out-of-bag data (that is data excluded from the boostrap sample used to 393 construct the tree) and then computes the difference between the error on the per-394 muted test set and the original test set. More precisely, for each tree t among the ntree 395 trees of the RF, MDA uses the out-of-bag data to compute a prediction error OOB_t . 396 Then, permuting the values of the i^{th} feature in the out-of-bag data, a prediction error 397 OOB_t^j is computed by using the permuted out-of-bag data. The permutation impor-tance of the feature j is thus defined by $MDA_j = \frac{1}{ntree} \sum_{t=1}^{ntree} (OOB_t - OOB_t^j)$. 398 399

⁴⁰⁰ To achieve feature selection via the MDA measure, the backward Recursive Feature ⁴⁰¹ Elimination (RFE) algorithm (Guyon et al, 2002) implemented in the rfe^4 function ⁴⁰² of the *caret* R package is used. This algorithm is based on assessing MDA's impor-

 $_{403}$ tance. MDA is computed by iteratively permuting the values of each input covariate

 $^{{}^{4} \}rm https://topepo.github.io/caret/recursive-feature-elimination.html$

and measuring the resulting drop in prediction accuracy. The feature with the minimum MDA value, representing the least important feature, is systematically removed

from the input set. Subsequently, a new RF model is trained using this reduced set of

⁴⁰⁷ features. This process is continued until the minimal set of input features that yielded

⁴⁰⁸ optimal Accuracy (Grandini et al, 2020) is obtained. To improve the performance of

⁴⁰⁹ feature selection with RFE, a repeated 3-fold cross-validation with 3 repeats is used.

⁴¹⁰ The *createFolds* function of the *caret* package allowed to split data into training and ⁴¹¹ test sets. This function carries a random sampling within geomorphological typologies

in order to balance the classes distributions within the split.

⁴¹³ Once the model is trained, typologies can be predicted across the entire study area. ⁴¹⁴ Typologies predictions are made using the *predict.train* function of the *caret* R

⁴¹⁵ package (Kuhn, 2019).

416 2.2.5 Performance criteria

To assess model's performance, a *Balanced Accuracy* (BA) metric were calculated using the confusion matrix obtained from each model (Grandini et al, 2020). It consists for each typology (class) k, to calculate a Recall score measuring the ability of a model to find all the positive units for this class as follows:

$$Recall_k = \frac{TP_k}{TP_k + FN_k} \tag{7}$$

421 True Positives (TP) are observations predicted to belong to the reference class when

they really do, and False Negatives (FN) are observations predicted to not belong to the reference class when they really do. BA gives an average measure of this concept

⁴²⁴ using the arithmetic mean of Recall across all classes :

$$BA = \frac{\sum_{k=1}^{K} Recall_k}{K} \tag{8}$$

425 where K is the total number of class k.

⁴²⁶ The number of unsampled typologies is used as an indicator of the efficiency of the

sampling method to visit all geomorphologic typologies present in the study area.
To assess the consistency of predictions, the map produced at the different DBM

resolution are compared to Expert map. For this purpose, the expert map is first dicretized into "pixels" size of 5 m resolution (ie. the smallest resolution used in this study). Then the predicted map grid is disaggregated to match the expert map resolution using *disagRast* function in the supplemental materials. To compare two maps, two metrics are calculated using the confusion matrix between the two: *Match* and *Balanced Match*.

 $_{\tt 435}$ $\,$ The Match metric is a simple comparison between the two maps. The Match metric

436 corresponding to the proportion of cells identically labelled is calculated as follows:

$$Match = \frac{N_+}{N_+ + N_-} \tag{9}$$

Where N_+ is the total number of cells where typologies match between the two maps, 437 and N_{-} is the total number of cells that do not match. The *matchRast* function is 438 proposed for the calculation of this metric. However such comparison might hide pre-439 diction problems on some small surface typologies, totally dominated by high surface 440 coverage of some typologies. To take account of such fact, the Balance Match is a 441 sort of ponderated *Match* removing the surface dominant effect that some typologies 442 might have on others. A second metric denoted Balanced Match (BM) inspired by 443 the BA defined in Eq. 8 and using the confusion matrix between a predicted map and 444 the expert map is also calculated. Note that after disaggregating a predicted map, 445 some cells centers around the study site borders may be located outside. These border 446 effects are handled by proportioning BA to the proportion of cells L labelled after the 447 disaggregation of a predicted map. BM is calculated as follows: 448

$$BM = L \cdot \frac{\sum_{k=1}^{K} Recall_k}{K} \tag{10}$$

The balmatchRast function provided in the supplemental materials is used for the BM calculation.

451 **3 Results**

452 3.1 Effect of ground truth sampling methodologies on 453 typologies sampled

Our first methodological assessment was to evaluate the number of missed typologies 454 on the expert map depending on ground truth sampling techniques. Unsurprisingly, 455 the more data points sampled, the greater the chance of sampling all typologies present 456 in the study site, regardless of the sampling method and the resolution of the DBM 457 (Figure 3). Whatever the sampling method and the DBM resolution, small sample 458 size (50 and 100 data points) do not allow to sample all typologies. For these sample 459 sizes, between 1 and 4 typologies are never sampled. It can also be noticed that for 460 500 data points sampled and more, all typologies are sampled no matter the sampling 461 method and the resolution of the DBM. 462

463 3.2 Assessment of model performance based on input data

The Balanced Accuracy criteria helps assessing model performance. The larger the 464 sample size, the more precise these measures (ie. smaller standard errors; Figure 4). 465 For small sample sizes (50 and 100 data points), sampling methods give comparable 466 results or even slightly better results for SCS-KMEANS than others sampling methods. 467 In constrast, from 200 points upwards, complexity-dependent sampling tends to give 468 better results than SCS-CLARA and SCS-KMEANS which give comparable results. 469 This result is confirmed as the sample size increases, with a widening gap among sam-470 pling methodology means and decreasing standard errors (see Figure S3 and Table S3 471 for further details). 472



Fig. 3 Boxplot representing the *Number of missing typologies* metric (y-axis) for different sample size (x-axis), different sampling methods (CDS, SCS-CLARA and SCS-KMEANS) and different bathymetric model resolutions (5 m, 25 m, 50 m, 100 m, 500 m). Each sampling conditions were replicated 30 times and represented as default R boxplot settings

⁴⁷³ 3.3 Assessment of selected terrain attributes

A set of features is selected during the feature selection step of the modeling process
and general outputs are summarized. Only results for 100, 200, 500 data points sample
sizes generated at 100 m DBM resolution using SCS-CLARA method and CDS method
are shown her because they are representative of all results (Figure 5). Geographic
coordinates (*Latitude* and *Longitude*) and *Depth* are almost always selected regardless
the DBM resolution and the sampling method. Terrain variability attributes groups
(*Roughness*, VRM and TRI) that are selected a little more than half of the time.

481 3.4 Evaluation of the quality of produced maps

The *Match* criteria evaluates the consistency of predictions according to the expert map. As previously seen with the *Balanced Accuracy* standard errors values, *Match* errors are non negligible for small sample sizes and decrease when DBM resolutions increase (Figure 6; Figure S3 and Table S4 for further details). In addition, lower *Match* values are recorded at 5 m and 500 m DBM resolutions and higher *Match* values were seen for intermediate DBM resolutions (25 m, 50 m, 100 m) regardless the sample size.

When the sample size and the DBM resolution increase, *Balanced Match* values become more accurate as shown by a decrease of standard errors (Figure 7; Figure S3 and Table S5 for further details). CDS method gives better results than



Fig. 4 Mean (+/- standard error) over 30 draws of the *Balanced Accuracy* metric (y-axis) at different DBM resolutions (x-axis) for different sample size and different sampling methods (CDS, SCS-CLARA and SCS-KMEANS)

others spatial coverage sampling methods for any sample size and any DBM resolution. SCS-CLARA and SCS-KMEANS give comparable results according this criteria.
Comparing results among DBM resolutions, best results are achieved with 50 m and
100 m DBM resolutions.

In Figure 8 two maps generated with data points sampled using CDS methodology are restituted: (A) 50 m DBM resolution using 1000 data points and (B) 100 m DBM resolution using 200 data points. All typologies are sampled and predicted in both cases. Results show that performance criteria measured are better for the (A') case than the (B') case (see (C)). Indeed, some small surface typologies like Drowned subtidal reef flat and inner slope are better predicted and predictions errors on typologies transition areas less important for (A) than for (B).

503 4 Discussion

This methodology enabled us to successfully reconstruct an expert geomorphological 504 map. According to the Number of missing typologies metric, 200 data points are 505 enough to sample almost all the typologies. These locations can be choosen using 506 CDS which performs slightly better than spatial coverage sampling methods (SCS-507 KMEANS and SCS-CLARA). Results between the different DBM resolutions are 508 comparable for small sample sizes considering the *Balanced Accuracy* metric. But, for 509 200 data points and more, 100 m DBM resolution gives clearly better results than oth-510 ers resolutions. The Match metric supports this finding where 50 m and 100 m DBM 511



Fig. 5 Ordered occurence over 30 draws of the selected terrain attributes by the RFE algorithm. Here an example with 100, 200, 500 data points sample sizes (from left to right) generated at 100 m DBM resolution using SCS-CLARA method (top) and CDS method (bottom). The average number of attributes selected is noted above each graph (cf. Terrain attribute section for their definition)

resolutions gives much better results than others DBM resolutions whatever the num-512 ber of data points. Considering the *Balanced Match* criteria which was introduced 513 to contrast the *Match* criteria taking into account typologies surface imbalances, 50 514 m DBM resolution gives a slight better performance than the 100 m DBM resolution, 515 but considering the sampling effort, it would be preferable to use the 100 m DBM. 516 Indeed, while the sampling effort is multiplied by 5 from (B) to (A), the performance 517 recorded is just a little bit better. We have also seen that the precision of the maps 518 produced is sensitive to the resolution of the DBM, the number and locations of the 519 ground truth selection. Thus, reproductible methodologies with associated codes to 520 evaluate their qualities are proposed. In this section, choices on data sampling to gen-521 erate such maps are discussed. Then strong and weak points of the modeling approach 522 and alternative to enhance such work are addressed. 523

⁵²⁴ 4.1 Optimal data acquisition parameters

Creation of a submarine geomorphological map appears, among other consideration, to be constrained between quantity and quality of initial data and cost to acquire and process them. In the present work two kind of data fall in such compromise: the bathymetric data and the ground truth data points.

For the bathymetric data acquisition, high resolution data require advanced technologies, longer survey durations, important storage and sophisticated tools for manipulation which all contribute to higher costs. The results of this study show that a lower DBM resolution do not necessarily lead to the best geomorphological map.



Fig. 6 Mean (+/- standard error) over 30 draws of the *Match* metric (y-axis) at different DBM resolutions (x-axis) for different sample size and different sampling methods (CDS, SCS-CLARA and SCS-KMEANS)

The 100 m DBM resolution considered in this study appears to strike a good balance 533 between detailed geomorphological maps obtained (Figures 4, 6, 7) with low DBM 534 resolutions (5 and 25 m) and large scale geomorphological maps considering (500 m 535 DBM resolution). Indeed, maps at 5 m, 25 m and 50 m DBM resolutions provide a 536 reasonably detailed view of the seafloor and allow the identification of important fea-537 tures but contain a significant amount of noise or small-scale variability that may not 538 be relevant in constructing a geomorphological map. On the other hand, maps of 500 539 m resolution provide very generalized view of the seafloor representing a significantly 540 coarser scale focusing on major landforms and regional-scale geomorphological pat-541 terns but missing important features required to build such maps (eg. reef patches, 542 pass or canyons ...). The 100 m resolution was also considered suitable in previous 543 studies for various applications, such as regional planning, environmental assessments, 544 and natural hazard evaluations (Curie et al, 2007; Dong et al, 2019). 545

Ground truth data acquisition is often acquired by specialists through scubadiving or 546 snorkling (shallow) or using submarine, ROV or drop camera systems allowing to view 547 and characterise seafloor typologies on specific location. In that respect, the number 548 of location is directly correlated to cost of data gathering and therefore need to be 549 balanced. Furthermore, if the number of data point is directly linked to survey cost, 550 the location of sampling on an heterogeneous seafloor structure could be related to 551 data quality (ie. enhanced typology sampling diversity above oversampling of common 552 typologies). To test such compromises, effects number of data points and methodolo-553 gies to locate them (SCS vs CDS approach) on overall map quality production are 554

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Fig. 7 Mean (+/- standard error) over 30 draws of the *Balanced Match* metric (y-axis) at different DBM resolutions (x-axis) for different sample size and different sampling methods (CDS, SCS-CLARA and SCS-KMEANS).

studied. The choice between homogeneous distribution vs seafloor complexity depen-555 dent approach came from the fact that various typologies might agglomerate around 556 location of complex structures (e.g. barrier reef, patch reef, ...). Figure 2 illustrates this 557 point and show that CDS method amplify sampling in area of important typologies 558 diversity. This explains why Number of missing typologies criteria results obtained 559 with CDS are better than homogeneous sampling methods (Figure 3). However, such 560 approach has the inconvenient that a bathymetric dataset is required prior planning 561 ground truth sampling campaign. Another point of attention is that attributes used 562 as covariates for CDS algorithms should be carefully selected. This study according to 563 the context of used data (coral reef habitats) focuses on metrics promoted in literature 564 (Adev. 1966; Adev and Macintyre, 1973; Battistini, 1975; Minnery et al. 1985) and 565 on empirical results from this work in the supplemental materials with a supervised 566 classification of geomorphologic typologies based on the terrain attributes presented 567 in Table 1. Seafloor depth and roughness were retained as the most relevant. 568

Evaluating effect of bathymetric data points density and number and location of 569 ground truth points was done by comparing produced map to an existing expert 570 manually made map. The *Match* criteria were introduced to see how realistic are pre-571 dictions, regardless the number of sampled typologies. Thus the large geomorphologic 572 units like lagoons (deep, intermediate and shallow), subtidal and deep subtidal reef 573 flats are generally better predicted than the small surface typologies like Pass, Inner 574 slope, Outer slope and Drowned subtidal reef flat typologies. Indeed, such typologies 575 are under-represented in the study area and represent together no more than 10% of 576 the surface. Even evaluating the predictability of such typologies was difficult using 577



Fig. 8 A Predicted map with a 50 m DBM and 1000 sampled locations using CDS methodology. B Predicted map with a 100 m DBM and 200 sampled locations using CDS methodology. Corresponding match / mismatch maps (A' and B') in comparison to the expert one. C Performance criteria measured.

the *Match* metric, hence the use of *Balanced Match* metric pondering typologies by their surface. Using both these metrics, optimal map construction was obtained for 200 ground truth data points obtained using CDS methodology and 100 m DBM resolution (Figure 8).

However, some limitation of the methodology used is to be noted. SCS-CLARA and 582 CDS sampling methods are particularly useful for high dimensional data. The CLARA 583 clustering algorithm can be used on large DBM grid cells data to generate ground truth 584 locations. However, SCS-KMEANS although suitable for spatial coverage sampling, 585 lead to computational deadlock for high DBM resolutions. Furthermore, it can not 586 be used when there is missing data areas in the studied surface. Bathymetric imputa-587 tion of poorly sampled zones was done using ordinary kriging method that performed 588 much better than others spatial interpolation methods (cf. Table S1, an example on 589 the supplemental materials). 590

⁵⁹¹ 4.2 Modeling choices

Traditional approaches to build geomorphologic maps are often time-consuming, laborintensive and has a limited coverage and scale. Remote Sensing techniques although allowing wide coverage and high-resolution, require expertise in image interpretation (Gao, 2009; Gilvear and Bryant, 2016). GIS approaches allow for the integration and analysis of diverse data types support data visualization and complex spatial analysis. But they also require specialized software and expertise (Guzzetti et al, 1999; Napieralski and Li, 2007). In addition, interpretation and analysis heavily rely on data

quality. Besides these techniques, semi-automated and automated approaches through 599 machine learning and deep learning are increasingly used to identify complex pat-600 terns and features for landform classification, feature detection, or segmentation. Deep 601 learning models require large amounts of labeled training data (e.g., images, point 602 clouds) to effectively learn complex patterns and relationships. Training such models 603 involves adjusting millions of parameters through backpropagation and optimization 604 algorithms (e.g., stochastic gradient descent) and their tuning involves finding the right 605 network architecture and hyperparameters. They are often considered black boxes due 606 to their complex architectures, making it challenging to interpret the features that 607 influence predictions (Li et al, 2020). The statistical learning approach used in this 608 study can handle large volumes of data, automate analysis processes and discover com-609 plex patterns and relationships in the data. Interpretability of used models may also 610 be a challenge but less than deep learning ones. In addition, it can work effectively 611 with smaller datasets which was crucial in our objectives. 612

RF model for geomorphologic units clustering is chosen because RF exhibits com-613 plex and non-linear relationships between the features and the dependent variable but 614 also on features among themselves, handling effectively these relationships by using 615 an ensemble of decision trees. It also provide valuable information as feature impor-616 tance measure helping identify the most informative feature which made it preferable 617 to others clustering techniques in many cases. RF was used for supervised classifi-618 cation problems in many recent studies, compared different algorithms has shown 619 effectiveness of RF based algorithms (Zeraatpisheh et al, 2017; Giaccone et al, 2022). 620 It has shown its robustness specifically when data contain uncertainties and handles 621 high-dimensional data efficiently avoiding overfitting and reducing computational com-622 plexity. 623

The variable selection step in the modeling procedure is crucial in the proposed methodology. Terrain attributes calculation is not specifically time-consuming but the relevance of each of them depends on the data and the geomorphological features that are mapped. This study demonstrated that the terrain attributes, although literature based choosen initially, have not all, a great explanatory power on geomorphologic feature. For each generated sample, the number of features selected is also counted and 6 terrain attributes are generally retained.

5 Conclusion

A statistical learning based approach is proposed to automatically map the geomor-632 phology of a study site using bathymetric data and some ground truth data points. 633 On the one hand, tools to help geomorphologists to plan field campaigns in advance 634 through an optimal DBM resolution and an automated sampling methodology to 635 636 achieve field verifications are provided. On the other hand, a flexibility in the proposed methodology allowing the usage of terrain attributes as much as desired since the fea-637 ture selection will help to keep only the most relevant ones is preferred. In addition, 638 statistical and computational tools to compare geomorphological maps produced at 639 different resolutions are provided. The methodology reproducibility is made possible 640 by a set of reusable R scripts. 641

⁶⁴² In the future, an application of the methodology using others data available in oth-⁶⁴³ ers sites is planned. An investigation on others sampling methodology for e.g. which ⁶⁴⁴ would take into account the presence of non sampling sites inside a study area is also ⁶⁴⁵ being considered.

646 6 Supplement information

Additional figures and tables supporting this manuscript can be found after theReference section.

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Fig. S1 Distribution of the raw bathymetric data collected on Geyser.

Table S1 Bathymetry imputation of empty grid cells of a 50 m DBM resolution. The Root Mean Squared Error (RMSE) and its Standard Deviation (SD) were calculated using 5-folds cross-validation scheme for five (05) spatial interpolation models : Inverse Distance Weighted (IDW), Nearest Neighboor (NN), Ordinary Kriging (OK), Universal Kriging (UK), Spatial GAM (GAM)

Method	RMSE	SD (RMSE)
IDW	0.80	0.02
NN	0.82	0.02
OK	0.69	0.01
UK	0.71	0.01
GAM	1.34	0.01

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Fig. S2 Statistical learning approach to build geomorphological maps using bathymetric data and typology field verification samples.



Fig. S3 Boxplot of *Balanced Accuracy* (A), *Match* (B) and *Balanced Match* (C) metrics (y-axis) for different sample size (x-axis), different sampling methods (CDS, SCS-CLARA and SCS-KMEANS) and different bathymetric model resolutions (5 m, 25 m, 50 m, 100 m, 500 m). Each sampling conditions were replicated 30 times and represented as default R boxplot settings.

DBM: 5 m							
	Sample Size						
Method	50	100	200	500	700	1000	_
SCS-KMEANS SCS-CLARA CDS	$\begin{array}{c} 3.867 \ (0.681) \\ 5.033 \ (0.964) \\ 3.067 \ (0.944) \end{array}$	$\begin{array}{c} 0.967 \ (0.718) \\ 2.133 \ (0.937) \\ 0.967 \ (0.964) \end{array}$	$\begin{array}{c} 0 \ (0) \\ 0.633 \ (0.615) \\ 0.167 \ (0.379) \end{array}$	$\begin{array}{c} 0 & (0) \\ 0 & (0) \\ 0 & (0) \end{array}$	$\begin{array}{c} 0 \ (0) \\ 0 \ (0) \\ 0 \ (0) \end{array}$	$\begin{array}{c} 0 & (0) \\ 0 & (0) \\ 0 & (0) \end{array}$	
DBM: 25 m							
			Sample Size				_
Method	50	100	200	500	700	1000	
SCS-KMEANS SCS-CLARA CDS	$\begin{array}{c} 6 \ (0.643) \\ 5.233 \ (0.898) \\ 3.3 \ (1.179) \end{array}$	$\begin{array}{c} 3.1 (0.885) \\ 2.367 (0.964) \\ 1.467 (0.9) \end{array}$	$\begin{array}{c} 0.533 \ (0.629) \\ 0.567 \ (0.626) \\ 0.3 \ (0.466) \end{array}$	$\begin{array}{c} 0 & (0) \\ 0 & (0) \\ 0 & (0) \end{array}$	$\begin{array}{c} 0 \ (0) \\ 0 \ (0) \\ 0 \ (0) \end{array}$	$\begin{array}{c} 0 \ (0) \\ 0 \ (0) \\ 0 \ (0) \end{array}$	_
DBM: 50 m							_
			Sample Size				_
Method	50	100	200	500	700	1000	_
SCS-KMEANS SCS-CLARA CDS	5.933 (0.74) 5.067 (0.868) 2.833 (1.053)	$\begin{array}{c} 3.167 \; (0.791) \\ 2.1 \; (1.125) \\ 0.933 \; (0.828) \end{array}$	$\begin{array}{c} 0.6 \ (0.724) \\ 0.567 \ (0.679) \\ 0.233 \ (0.43) \end{array}$	$\begin{array}{c} 0 \ (0) \\ 0 \ (0) \\ 0 \ (0) \end{array}$	$\begin{array}{c} 0 \ (0) \\ 0 \ (0) \\ 0 \ (0) \end{array}$	$\begin{array}{c} 0 \ (0) \\ 0 \ (0) \\ 0 \ (0) \end{array}$	_
DBM: 100 m							
			Sample Size				
Method	50	100	200	500		700	1000
SCS-KMEANS SCS-CLARA CDS	5.867 (0.681) 5 (0.947) 2.867 (1.252)	$\begin{array}{c} 3.033 \ (0.85) \\ 2.2 \ (0.805) \\ 0.2 \ (0.407) \end{array}$	$\begin{array}{c} 0.567 \ (0.728) \\ 0.7 \ (0.702) \\ 0.133 \ (0.346) \end{array}$	0.033 (0.	0 (0) .183) 0 (0)	$\begin{array}{c} 0 & (0) \\ 0 & (0) \\ 0 & (0) \end{array}$	$\begin{array}{c} 0 & (0) \\ 0 & (0) \\ 0 & (0) \end{array}$
DBM: 500 m							_
	Sample Size						
Method	50	100	200	500	700	1000	-
SCS-KMEANS SCS-CLARA CDS	5.467 (0.73) 4.8 (1.031) 3.133 (1.042)	$\begin{array}{c} 2.733 \ (0.785) \\ 2.367 \ (1.129) \\ 1.433 \ (0.817) \end{array}$	$egin{array}{c} 0.467 & (0.507) \ 0.4 & (0.563) \ 0.233 & (0.43) \end{array}$	$\begin{array}{c} 0 & (0) \\ 0 & (0) \\ 0 & (0) \end{array}$	$\begin{array}{c} 0 \ (0) \\ 0 \ (0) \\ 0 \ (0) \end{array}$	$\begin{array}{c} 0 \ (0) \\ 0 \ (0) \\ 0 \ (0) \end{array}$	_

Table S2 Mean (standard deviation) over 30 draws of the *Number of missing typologies* metric (y-axis) at different DBM resolutions (x-axis) for different sample size and different sampling methods (CDS, SCS-CLARA and SCS-KMEANS)

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DBM: 5 m						
	Sample Size					
Method	50	100	200	500	700	1000
SCS-KMEANS	0.58(0.147)	0.534(0.132)	$0.541 \ (0.076)$	0.605(0.037)	0.64(0.033)	0.659 (0.025)
SCS-CLARA CDS	$0.482 (0.156) \\ 0.467 (0.11)$	$0.44 (0.105) \\ 0.5 (0.072)$	$0.547 (0.06) \\ 0.559 (0.064)$	$0.582 (0.044) \\ 0.654 (0.047)$	$0.621 (0.027) \\ 0.693 (0.03)$	0.657 (0.035 0.722 (0.023
DBM: 25 m						
			Samp	le Size		
Method	50	100	200	500	700	1000
SCS-KMEANS	0.606(0.133)	0.567(0.111)	$0.578\ (0.081)$	0.666~(0.034)	$0.694\ (0.033)$	0.714 (0.023)
SCS-CLARA	0.58 (0.166)	0.553 (0.118)	0.571 (0.051)	0.653 (0.043)	0.691 (0.034)	0.71 (0.032
CDS	0.467 (0.122)	0.525 (0.095)	0.589 (0.06)	0.684 (0.041)	0.71 (0.032)	0.753 (0.03
DBM: 50 m						
	Sample Size					
Method	50	100	200	500	700	1000
SCS-KMEANS	$0.644 \ (0.154)$	0.57(0.109)	$0.571 \ (0.062)$	$0.657 \ (0.034)$	0.697 (0.029)	0.725(0.02)
SCS-CLARA	0.56 (0.165)	0.513(0.095)	0.583(0.063)	0.659(0.036)	0.695(0.025)	0.724 (0.03
CDS	0.443 (0.123)	0.518 (0.079)	0.595 (0.059)	0.697 (0.033)	0.735 (0.025)	0.775 (0.02)
DBM: 100 m						
			Samp	le Size		
Method	50	100	200	500	700	1000
SCS-KMEANS	0.608(0.167)	0.543(0.097)	0.594(0.081)	0.663(0.034)	0.703(0.027)	0.723(0.02)
SCS-CLARA	0.54(0.11)	0.498(0.099)	0.566(0.084)	0.667(0.038)	0.688(0.027)	0.713 (0.02
CDS	0.511 (0.133)	0.535 (0.076)	0.617 (0.046)	0.718 (0.035)	$0.756\ (0.027)$	0.781 (0.02
DBM: 500 m						
			Samp	le Size		
Method	50	100	200	500	700	1000
SCS-KMEANS	0.576(0.153)	0.535(0.096)	0.555(0.064)	0.66(0.04)	0.705(0.034)	0.784 (0.029
SCS-CLARA	0.549(0.146)	0.505(0.127)	0.547 (0.082)	0.664(0.02)	0.65(0)	0.71 (
CDS	0.505(0.082)	0.542(0.071)	0.604(0.059)	0.659(0.016)	0.69(0)	0.69 (

Table S3 Mean (standard deviation) over 30 draws of the Balanced Accuracy metric (y-axis) at different DBMresolutions (x-axis) for different sample size and different sampling methods (CDS, SCS-CLARA and SCS-KMEANS)

DBM: 5 m						
	Sample Size					
Method	50	100	200	500	700	1000
SCS-KMEANS SCS-CLARA CDS	$\begin{array}{c} 0.74 \ (0.026) \\ 0.734 \ (0.037) \\ 0.766 \ (0.031) \end{array}$	0.793 (0.03) 0.805 (0.019) 0.818 (0.014)	$\begin{array}{c} 0.837 \ (0.009) \\ 0.839 \ (0.008) \\ 0.846 \ (0.01) \end{array}$	0.865 (0.006) 0.865 (0.007) 0.878 (0.008)	$\begin{array}{c} 0.874 \ (0.006) \\ 0.876 \ (0.008) \\ 0.888 \ (0.007) \end{array}$	0.886 (0.006) 0.884 (0.006) 0.9 (0.005)
DBM: 25 m						
			Sampl	e Size		
Method	50	100	200	500	700	1000
SCS-KMEANS SCS-CLARA CDS	$\begin{array}{c} 0.753 \ (0.022) \\ 0.758 \ (0.031) \\ 0.771 \ (0.033) \end{array}$	$\begin{array}{c} 0.812 \ (0.02) \\ 0.815 \ (0.027) \\ 0.832 \ (0.027) \end{array}$	0.857 (0.009) 0.863 (0.008) 0.869 (0.011)	$\begin{array}{c} 0.888 \ (0.007) \\ 0.891 \ (0.006) \\ 0.902 \ (0.01) \end{array}$	$\begin{array}{c} 0.898 \ (0.005) \\ 0.902 \ (0.006) \\ 0.915 \ (0.006) \end{array}$	$\begin{array}{c} 0.909 & (0.006) \\ 0.91 & (0.008) \\ 0.923 & (0.006) \end{array}$
DBM: 50 m						
			Sampl	e Size		
Method	50	100	200	500	700	1000
SCS-KMEANS SCS-CLARA CDS	$\begin{array}{c} 0.755 \ (0.023) \\ 0.76 \ (0.023) \\ 0.794 \ (0.029) \end{array}$	$\begin{array}{c} 0.801 \ (0.025) \\ 0.815 \ (0.03) \\ 0.852 \ (0.015) \end{array}$	$\begin{array}{c} 0.858 \ (0.009) \\ 0.863 \ (0.013) \\ 0.88 \ (0.01) \end{array}$	$\begin{array}{c} 0.891 \ (0.008) \\ 0.896 \ (0.005) \\ 0.91 \ (0.005) \end{array}$	$\begin{array}{c} 0.902 \ (0.005) \\ 0.905 \ (0.006) \\ 0.917 \ (0.004) \end{array}$	0.912 (0.005) 0.912 (0.005) 0.927 (0.005)
DBM: 100 m						
			Sampl	e Size		
Method	50	100	200	500	700	1000
SCS-KMEANS SCS-CLARA CDS	$\begin{array}{c} 0.753 \; (0.024) \\ 0.744 \; (0.031) \\ 0.788 \; (0.028) \end{array}$	$\begin{array}{c} 0.801 \ (0.026) \\ 0.813 \ (0.023) \\ 0.845 \ (0.017) \end{array}$	0.857 (0.008) 0.861 (0.012) 0.875 (0.01)	$\begin{array}{c} 0.89 \ (0.005) \\ 0.893 \ (0.005) \\ 0.91 \ (0.004) \end{array}$	$\begin{array}{c} 0.9 \ (0.003) \\ 0.902 \ (0.007) \\ 0.918 \ (0.004) \end{array}$	$\begin{array}{c} 0.91 \ (0.003 \\ 0.909 \ (0.004 \\ 0.926 \ (0.005 \end{array})$
DBM: 500 m						
	Sample Size					
Method	50	100	200	500	700	1000
SCS-KMEANS SCS-CLARA CDS	$\begin{array}{c} 0.758 \ (0.021) \\ 0.746 \ (0.023) \\ 0.785 \ (0.018) \end{array}$	$\begin{array}{c} 0.793 \ (0.019) \\ 0.802 \ (0.016) \\ 0.817 \ (0.014) \end{array}$	$\begin{array}{c} 0.832 \ (0.008) \\ 0.836 \ (0.01) \\ 0.851 \ (0.007) \end{array}$	$\begin{array}{c} 0.862 \ (0.004) \\ 0.869 \ (0.005) \\ 0.88 \ (0.002) \end{array}$	$0.87 (0.002) \\ 0.87 (0) \\ 0.88 (0)$	0.88(0) 0.88(0) 0.89(0)

Table S4 Mean (standard deviation) over 30 draws of the *Match* metric (y-axis) at different DBM resolutions (x-axis) for different sample size and different sampling methods (CDS, SCS-CLARA and SCS-KMEANS)

$\begin{tabular}{ c c c c c c } \hline Sample Size \\ \hline \begin{tabular}{ c c c c c c c c c c c c c c c c c c c$	DBM: 5 m						
$\begin{tabular}{ c c c c c c c c c c c c c c c c c c c$		Sample Size					
$ \frac{\text{SCS-KMEANS}}{\text{SCS-CLARA}} \underbrace{\begin{array}{c} 0.292 \ (0.035) \\ 0.306 \ (0.051) \\ 0.412 \ (0.071) \\ 0.482 \ (0.048) \\ 0.521 \ (0.056) \\ 0.521 \ (0.056) \\ 0.607 \ (0.029) \\ 0.686 \ (0.026) \\ 0.674 \ (0.024) \\ 0.723 \ (0.023) \\ 0.773 \ (0.023) \\ 0.778 \ (0.025) \\ 0.778 \ (0.024) \\ 0.778 \ (0.024) \\ 0.778 \ (0.024) \\ 0.778 \ (0.024) \\ 0.778 \ (0.025) \\ 0.778 \ (0.025) \\ 0.778 \ (0.025) \\ 0.778 \ (0.025) \\ 0.778 \ (0.025) \\ 0.778 \ (0.025) \\ 0.778 \ (0.021) \\ 0.778 \ (0.021) \\ 0.778 \ (0.021) \\ 0.778 \ (0.022) \\ 0.778 \ (0.021) \\ 0.778 \ (0.015) \\ 0.787 \ (0.011) \\ 0.787 \ (0.011) \\ 0.787 \ (0.011) \\ 0.787 \ (0.011) \\ 0.787 \ (0.011) \\ 0.778 \ (0.021) \\ 0.776 \ (0.021) \\ 0.777 \ (0.011) \\ 0.777 \ (0.011) \\ 0.777 \ (0.011) \\ 0.777 \ (0.011) \\ 0.777 \ (0.011) \\ 0.777 \ (0.011) \\ 0.777 \ (0.011) \\ 0.777 \ (0.012) \\ 0.777 \ (0.016) \\ 0.777 \$	Method	50	100	200	500	700	1000
DBM: 25 m Sample Size Method 50 100 200 500 700 1000 SCS-KMEANS SCS-CLARA 0.306 (0.036) 0.413 (0.057) 0.47 (0.055) 0.479 (0.08) 0.611 (0.027) 0.623 (0.032) 0.699 (0.024) 0.701 (0.025) 0.732 (0.027) 0.73 (0.021) 0.758 (0.028) 0.759 (0.03) DBM: 50 m Sample Size Method 50 100 200 500 700 1000 SCS-KMEANS SCS-CLARA 0.316 (0.037) 0.335 (0.053) 0.444 (0.055) 0.484 (0.074) 0.603 (0.031) 0.669 (0.031) 0.705 (0.022) 0.736 (0.022) 0.767 (0.031) SCS-CLARA 0.336 (0.037) 0.444 (0.053) 0.669 (0.031) 0.710 (0.019) 0.736 (0.024) 0.761 (0.031) DBM: 100 m Sample Size Method 50 100 200 500 700 1000 SCS-KMEANS SCS-CLARA 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) Method 50 100 200 500 700 1000 SC	SCS-KMEANS SCS-CLARA CDS	$\begin{array}{c} 0.292 \ (0.035) \\ 0.306 \ (0.051) \\ 0.412 \ (0.071) \end{array}$	$\begin{array}{c} 0.445 \ (0.062) \\ 0.482 \ (0.048) \\ 0.521 \ (0.056) \end{array}$	$\begin{array}{c} 0.576 \ (0.027) \\ 0.581 \ (0.027) \\ 0.607 \ (0.029) \end{array}$	$\begin{array}{c} 0.662 \ (0.019) \\ 0.649 \ (0.026) \\ 0.686 \ (0.024) \end{array}$	$\begin{array}{c} 0.686 \ (0.026) \\ 0.674 \ (0.024) \\ 0.723 \ (0.023) \end{array}$	$\begin{array}{c} 0.723 \ (0.021) \\ 0.708 \ (0.025) \\ 0.755 \ (0.02) \end{array}$
Method 50 100 200 500 700 1000 SCS-KMEANS CDS 0.306 (0.036) 0.438 (0.066) 0.477 (0.055) 0.479 (0.08) 0.623 (0.032) 0.623 (0.032) 0.701 (0.025) 0.701 (0.025) 0.732 (0.027) 0.73 (0.021) 0.758 (0.038) 0.758 (0.030) CDS 0.413 (0.057) 0.553 (0.054) 0.623 (0.032) 0.701 (0.025) 0.76 (0.016) 0.757 (0.015) DBM: 50 m	DBM: 25 m						
Method501002005007001000SCS-KMEANS CDS0.306 (0.036) 0.438 (0.066) 0.419 (0.087)0.417 (0.055) 0.479 (0.08) 0.553 (0.054)0.611 (0.027) 0.623 (0.032) 0.623 (0.032) 0.726 (0.02)0.732 (0.027) 0.732 (0.021) 0.736 (0.021) 0.758 (0.038) 0.787 (0.015)DBM: 50 mSample SizeMethod501002005007001000SCS-KMEANS DCS-KMEANS 0.316 (0.037)0.444 (0.055) 0.443 (0.046)0.603 (0.031) 0.448 (0.055)0.705 (0.023) 0.615 (0.048)0.736 (0.02) 0.705 (0.025)0.766 (0.031) 0.736 (0.024)Method501002005007001000SCS-KMEANS DCS-CLARA0.316 (0.037) 0.335 (0.053)0.444 (0.055) 0.485 (0.074)0.603 (0.031) 0.615 (0.048)0.711 (0.019) 0.711 (0.019)0.766 (0.021) 0.736 (0.024)DBM: 100 mSample SizeMethod501002005007001000SCS-KMEANS DSCS-CLARA0.309 (0.032) 0.325 (0.049)0.446 (0.063) 0.474 (0.064)0.611 (0.026) 0.607 (0.04)0.731 (0.022) 0.731 (0.025)0.756 (0.021) 0.756 (0.021)DBM: 500 m0.426 (0.071) 0.426 (0.071)0.581 (0.041) 0.581 (0.041)0.666 (0.036) 0.666 (0.036)0.756 (0.015) 0.756 (0.015)0.731 (0.025) 0.731 (0.025)0.756 (0.021) 0.586 (0.021)DBM: 500 m </td <td></td> <td></td> <td></td> <td>Samp</td> <td>le Size</td> <td></td> <td></td>				Samp	le Size		
SCS-KMEANS SCS-CLARA 0.306 (0.036) 0.338 (0.06) 0.47 (0.055) 0.479 (0.08) 0.611 (0.027) 0.623 (0.032) 0.699 (0.024) 0.701 (0.025) 0.732 (0.027) 0.73 (0.021) 0.758 (0.028) 0.759 (0.03) DBM: 50 m Sample Size Method 50 100 200 500 700 1000 SCS-KMEANS CDS 0.316 (0.037) 0.444 (0.055) 0.603 (0.031) 0.705 (0.025) 0.736 (0.02) 0.767 (0.031) SCS-KMEANS CDS 0.316 (0.037) 0.444 (0.055) 0.603 (0.031) 0.711 (0.019) 0.736 (0.024) 0.761 (0.024) SCS-KMEANS CDS 0.335 (0.053) 0.445 (0.074) 0.615 (0.048) 0.711 (0.019) 0.736 (0.024) 0.761 (0.024) DBM: 100 m Sample Size Method 50 100 200 500 700 1000 SCS-KMEANS CDS 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) SCS-KMEANS CDS 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) SCS-	Method	50	100	200	500	700	1000
DBM: 50 m Sample Size Method 50 100 200 500 700 1000 SCS-KMEANS SCS-CLARA CDS 0.316 (0.037) 0.335 (0.053) 0.444 (0.055) 0.448 (0.046) 0.603 (0.031) 0.485 (0.074) 0.705 (0.025) 0.615 (0.048) 0.736 (0.024) 0.711 (0.019) 0.736 (0.024) 0.736 (0.024) 0.767 (0.031) 0.761 (0.024) DBM: 100 m Sample Size Method 50 100 200 500 700 1000 SCS-KMEANS CDS 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) Method 50 100 200 500 700 1000 SCS-KMEANS CDS 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) SCS-CLARA 0.3324 (0.049) 0.474 (0.064) 0.607 (0.04) 0.709 (0.025) 0.731 (0.025) 0.756 (0.021) CDS 0.426 (0.071) 0.581 (0.041) 0.66 (0.036) 0.756 (0.015) 0.779 (0.016) 0.805 (0.022)	SCS-KMEANS SCS-CLARA CDS	$\begin{array}{c} 0.306 \ (0.036) \\ 0.338 \ (0.06) \\ 0.413 \ (0.057) \end{array}$	$\begin{array}{c} 0.47 \; (0.055) \\ 0.479 \; (0.08) \\ 0.553 \; (0.054) \end{array}$	$\begin{array}{c} 0.611 \ (0.027) \\ 0.623 \ (0.032) \\ 0.64 \ (0.032) \end{array}$	$\begin{array}{c} 0.699 \ (0.024) \\ 0.701 \ (0.025) \\ 0.726 \ (0.02) \end{array}$	$\begin{array}{c} 0.732 \ (0.027) \\ 0.73 \ (0.021) \\ 0.76 \ (0.016) \end{array}$	$0.758 (0.028) \\ 0.759 (0.03) \\ 0.787 (0.015)$
DBM: 50 m Sample Size Method 50 100 200 500 700 1000 SCS-KMEANS SCS-CLARA 0.316 (0.037) 0.444 (0.055) 0.603 (0.031) 0.705 (0.025) 0.736 (0.02) 0.767 (0.031) CDS 0.443 (0.046) 0.577 (0.053) 0.669 (0.031) 0.745 (0.019) 0.736 (0.024) 0.761 (0.024) DBM: 100 m Sample Size Method 50 100 200 500 700 1000 SCS-KMEANS CDS 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) Method 50 100 200 500 700 1000 SCS-KMEANS SCS-CLARA 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) SCS-CLARA 0.324 (0.049) 0.474 (0.064) 0.607 (0.04) 0.709 (0.025) 0.731 (0.025) 0.756 (0.021) DBM: 500 m DBM: 500 m Contract of the state of the							
Method 50 100 200 500 700 1000 SCS-KMEANS SCS-CLARA 0.316 (0.037) 0.335 (0.053) 0.444 (0.055) 0.485 (0.074) 0.603 (0.031) 0.615 (0.048) 0.705 (0.025) 0.711 (0.019) 0.736 (0.024) 0.736 (0.024) 0.761 (0.024) 0.761 (0.024) DBM: 100 m	DBM: 50 m						
Method 50 100 200 500 700 1000 SCS-KMEANS SCS-CLARA CDS 0.316 (0.037) 0.443 (0.046) 0.444 (0.055) 0.485 (0.074) 0.603 (0.031) 0.615 (0.048) 0.705 (0.025) 0.711 (0.019) 0.736 (0.024) 0.736 (0.024) 0.767 (0.031) 0.761 (0.024) DBM: 100 m				Samp	le Size		
SCS-KMEANS SCS-CLARA 0.316 (0.037) 0.335 (0.053) 0.444 (0.055) 0.485 (0.074) 0.603 (0.031) 0.615 (0.048) 0.705 (0.025) 0.711 (0.019) 0.736 (0.02) 0.736 (0.024) 0.767 (0.031) 0.761 (0.024) CDS 0.443 (0.046) 0.577 (0.053) 0.669 (0.031) 0.745 (0.019) 0.736 (0.024) 0.761 (0.024) DBM: 100 m Sample Size Method 50 100 200 500 700 1000 SCS-CLARA 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) SCS-SCLARA 0.309 (0.032) 0.446 (0.064) 0.607 (0.04) 0.709 (0.025) 0.731 (0.022) 0.767 (0.019) SCS-CLARA 0.324 (0.049) 0.474 (0.064) 0.607 (0.04) 0.709 (0.025) 0.731 (0.025) 0.756 (0.021) DBM: 500 m DBM: 500 m	Method	50	100	200	500	700	1000
DBM: 100 m Sample Size Method 50 100 200 500 700 1000 SCS-KMEANS 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.017) SCS-KMEANS 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) SCS-CLARA 0.324 (0.049) 0.474 (0.064) 0.607 (0.04) 0.709 (0.025) 0.731 (0.022) 0.766 (0.021) CDS 0.426 (0.071) 0.581 (0.041) 0.666 (0.036) 0.756 (0.015) 0.779 (0.016) 0.805 (0.022)	SCS-KMEANS	0.316 (0.037) 0.335 (0.053)	0.444 (0.055) 0.485 (0.074)	0.603 (0.031) 0.615 (0.048)	0.705 (0.025) 0.711 (0.019)	0.736 (0.02)	0.767 (0.031) 0.761 (0.024)
DBM: 100 m Sample Size Method 50 100 200 500 700 1000 SCS-KMEANS SCS-CLARA 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) CDS 0.324 (0.049) 0.474 (0.064) 0.607 (0.04) 0.709 (0.025) 0.731 (0.025) 0.756 (0.021) CDS 0.426 (0.071) 0.581 (0.041) 0.666 (0.036) 0.756 (0.015) 0.779 (0.016) 0.805 (0.022) DBM: 500 m E E E E E E	CDS	0.443 (0.046)	0.400(0.014) 0.577(0.053)	0.669 (0.031)	0.745 (0.019)	0.772 (0.017)	0.797 (0.017)
DBM: 100 m Sample Size Method 50 100 200 500 700 1000 SCS-KMEANS 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) SCS-CLARA 0.324 (0.049) 0.474 (0.064) 0.607 (0.04) 0.709 (0.025) 0.731 (0.025) 0.756 (0.021) CDS 0.426 (0.071) 0.581 (0.041) 0.666 (0.036) 0.756 (0.015) 0.779 (0.016) 0.805 (0.022) DBM: 500 m							
Method 50 100 200 500 700 1000 SCS-KMEANS 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) SCS-CLARA 0.324 (0.049) 0.474 (0.064) 0.607 (0.04) 0.709 (0.025) 0.731 (0.025) 0.756 (0.021) CDS 0.426 (0.071) 0.581 (0.041) 0.666 (0.036) 0.756 (0.015) 0.779 (0.016) 0.805 (0.022) DBM: 500 m L L L L L L	DBM: 100 m						
Method 50 100 200 500 700 1000 SCS-KMEANS 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) SCS-CLARA 0.324 (0.049) 0.474 (0.064) 0.607 (0.04) 0.709 (0.025) 0.731 (0.022) 0.756 (0.021) CDS 0.426 (0.071) 0.581 (0.041) 0.66 (0.036) 0.756 (0.015) 0.779 (0.016) 0.805 (0.022) DBM: 500 m				Samp	le Size		
SCS-KMEANS 0.309 (0.032) 0.446 (0.063) 0.611 (0.026) 0.707 (0.018) 0.731 (0.022) 0.767 (0.019) SCS-CLARA 0.324 (0.049) 0.474 (0.064) 0.607 (0.04) 0.709 (0.025) 0.731 (0.025) 0.756 (0.021) CDS 0.426 (0.071) 0.581 (0.041) 0.666 (0.036) 0.756 (0.015) 0.779 (0.016) 0.805 (0.022) DBM: 500 m	Method	50	100	200	500	700	1000
SCS-CLARA 0.324 (0.049) 0.474 (0.064) 0.607 (0.04) 0.709 (0.025) 0.731 (0.025) 0.756 (0.021) CDS 0.426 (0.071) 0.581 (0.041) 0.66 (0.036) 0.756 (0.015) 0.779 (0.016) 0.805 (0.022) DBM: 500 m	SCS-KMEANS	0.309(0.032)	$0.446\ (0.063)$	$0.611 \ (0.026)$	$0.707 \ (0.018)$	$0.731 \ (0.022)$	$0.767 \ (0.019)$
DBM: 500 m	SCS-CLARA	$0.324 (0.049) \\ 0.426 (0.071)$	$0.474 (0.064) \\ 0.581 (0.041)$	0.607 (0.04) 0.66 (0.036)	0.709 (0.025) 0.756 (0.015)	$0.731 \ (0.025) \\ 0.779 \ (0.016)$	$0.756 (0.021) \\ 0.805 (0.022)$
DBM: 500 m							
	DBM: 500 m						
Sample Size		Sample Size					
Method 50 100 200 500 700 1000	Method	50	100	200	500	700	1000
SCS-KMEANS 0.335 (0.041) 0.432 (0.046) 0.566 (0.031) 0.679 (0.018) 0.707 (0.016) 0.737 (0.012)	SCS-KMEANS	0.335(0.041)	0.432(0.046)	0.566 (0.031)	0.679(0.018)	0.707 (0.016)	0.737(0.012)
SCS-CLARA 0.32 (0.043) 0.458 (0.057) 0.569 (0.034) 0.701 (0.015) 0.72 (0) 0.75 (0) CDS 0.428 (0.044) 0.521 (0.046) 0.615 (0.029) 0.726 (0.009) 0.75 (0) 0.76 (0)	SCS-CLARA CDS	$\begin{array}{c} 0.32 \ (0.043) \\ 0.428 \ (0.044) \end{array}$	$0.458 (0.057) \\ 0.521 (0.046)$	0.569 (0.034) 0.615 (0.029)	$\begin{array}{c} 0.701 \ (0.015) \\ 0.726 \ (0.009) \end{array}$	$\begin{array}{c} 0.72 \ (0) \\ 0.75 \ (0) \end{array}$	$\begin{array}{c} 0.75 \ (0) \\ 0.76 \ (0) \end{array}$

Table S5 Mean (standard deviation) over 30 draws of the *Balanced Match* metric (y-axis) at different DBM resolutions (x-axis) for different sample size and different sampling methods (CDS, SCS-CLARA and SCS-KMEANS)