



# A Continuous Clustering Method for Vector Fields

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## Abstract

A new method for the simplification of flow fields is presented. It is based on continuous clustering. A well-known physical clustering model, the Cahn Hillard model which describes phase separation, is modified to reflect the properties of the data to be visualized. Clusters are defined implicitly as connected components of the positivity set of a density function. An evolution equation for this function is obtained as a suitable gradient flow of an underlying anisotropic energy functional. Here, time serves as the scale parameter. The evolution is characterized by a successive coarsening of patterns — the actual clustering — and meanwhile the underlying simulation data specifies preferable pattern boundaries. Here we discuss the applicability of this new type of approach mainly for flow fields, where the cluster energy penalizes cross streamline boundaries, but the method also carries provisions in other fields as well. The clusters are visualized via iconic representations. A skeletonization algorithm is used to find suitable positions for the icons.

## 1 Introduction

Nowadays, fast computing hardware and efficient numerical algorithms enable highly detailed and large scientific simulations which deliver enormous amounts of data. Various visualization strategies have been proposed to represent such data in an intuitively understandable way.

The larger and more complex the simulation results become, the stronger is the need for a suitable multiscale visualization approach. Simplified representations of the data, useful to see the global pattern, can be further gradually refined for further insight. Moreover, different viewers need different representations. Numerical experts might want to see the raw data in full detail, technological experts might want to see certain features such as vortices, whereas the management might need a simplified presentation.

Clustering, well-known from statistics, is such a multiscale approach. Data are grouped in successively larger sets of strong internal correlation. Many techniques are available for scattered and scalar data, e.g. based on wavelet or Fourier analysis [11, 25]. However, for vector data, only few multiscale visualization methods are available. The most ubiquitous vector field simplification method is still regular subsampling, which is well-known to produce aliases (see e.g. Fig. 1). Turk [23] uses an energy minimizing approach to place equally distributed streamlines at a user prescribed resolution on the screen. Selected streamline drawings are furthermore considered by Jobard et al. [9].

Recently, two approaches for clustering vector data have been proposed. In both approaches a hierarchical clustering tree is produced and the resulting clusters are visualized with arrows. Heckel et al. [8] start from scattered points with vector data. Initially all points are stored in a single cluster, which is recursively split in a top-down manner. At each step, the cluster with the strongest discrepancy between streamlines generated by the original field and its

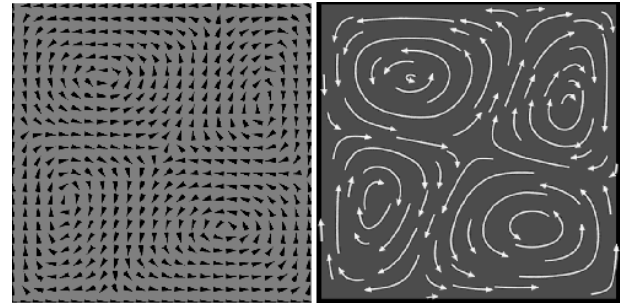


Figure 1: Vector field visualization: hedgehog plot (left), proposed clustering method (right).

approximation by the cluster is bisected with a plane, using principal component analysis. The resulting clusters are guaranteed to be convex, as a result of this bisection approach. However, accurately representing complex fields with convex clusters may require a large cluster count.

Telea and Van Wijk [21] use a bottom-up approach. Initially, each data point is a cluster, next these clusters are merged. In each step the most similar clusters are merged, according to a measure of the difference in position and orientation of the vectors that represent the clusters. The cluster shapes are constrained only indirectly by adapting the weights of the various terms in the error measure. However, this method is sometimes sensitive to the mentioned weight tuning.

Here we propose a continuous clustering method based on diffusion. The main difference to the other approaches is, that no boolean merging or splitting decisions have to be made. Instead a suitable diffusion process continuously enhances strong correlations in the cluster sets. In contrast with Heckel's method, these clusters are not necessarily convex. Hence, curved flow fields can be represented more effectively, as Fig. 1 — obtained with our method — shows. Our approach is motivated by a well-known physical model for phase separation in binary alloys which can be understood as a clustering of material in order to decrease the free energy of the physical system. As major application we consider clustering on flow fields. The method is related to multi-scale image processing methodology which leads to second order parabolic equations, whereas our model here will be a fourth order problem. Perona and Malik [15] have introduced a continuous diffusion model which allows the denoising of images together with edge enhancing. The recovery of lower dimensional structures in images is analyzed by Weickert [24], who introduced an anisotropic nonlinear diffusion method where the diffusion matrix depends on the so called structure tensor of the image. Preußer and Rumpf presented an efficient implementation for large scale image data [17] and used an anisotropic diffusion approach for flow visualization [18].

In detail the aims of our method are

- to extract a collection of nicely shaped subsets of the physical domain, where each of them is being characterized by a

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strong correlation in the underlying physical data and they all together are supposed to cover an approximately fixed fraction of the domain,

- to consider not only one such representation, but a scale of them ranging from fine granularity in the subdivision to very few and coarse cluster sets.

This multiscale should enable the exploration of complicated simulation data and the visual perception of correlations in such data sets at different resolutions. In our model, the clusters will be represented implicitly by a function evolving in time. Concerning the graphical representation we straightforwardly could use a color coded representation of this function on the physical domain as a texture. In the last decade dedicated to flow visualization a variety of such texturing methods has been presented. We mention here the spot noise technique by Van Wijk [5], the line integral convolution method by Cabral and Leedom [2], several improvements and modifications of this method [26, 7, 19], and the already mentioned nonlinear anisotropic diffusion method [18]. Here we use the actual clustering as a precomputing step and pipe the output into an iconic representation approach. Thus, the distinct subsets at any scale have to be represented by suitable graphical icons. This allows a further reduction of graphically represented data, while maintaining and strengthening the informational content.

The ingredients of our continuous clustering strategy are the following:

- We formulate an *evolution problem* for a function which implicitly describes the set of clusters. The evolution can be interpreted as the gradient flow with respect to an appropriate energy.
- There are *two energy contributions*. The first one leads to the *nucleation of cluster sets* on the physical domain. The second one gives rise for a *successive coarsening of the clusters*.
- Depending on the underlying physical data *surface segments are weighted depending on their location and orientation*. I. e. they are considerably penalized if they are oriented in cross direction to the correlation. Otherwise their energy contribution is kept small.
- On any scale a skeletonization method is used to reduce the informational content of the cluster sets to their essence, which is to be further visualized.
- Finally geometric icons are selected to represent the extracted skeleton information graphically, e. g. arrows in case of vector data.

Let us emphasize that the actual physical data enter the clustering method only via the anisotropic surface energy and the evolved function is solely used to define the cluster sets without any further physical meaning.

As application we mainly consider flow fields, where the concept of correlation along streamlines is near at hand. Nevertheless the methodology is not restricted to flow visualization and thus presented here for more general data.

The organization of the paper is as follows. In Section 2 we outline the physical model of phase separation in binary alloys which motivates this work. Then in the following Section 3 we expand this model and interpret it in terms of a multiscale cluster analysis. A finite element discretization is described in Section 4. In Section 5 the skeletonization approach is outlined and we discuss graphical icons in case of vector data.

Finally in Section 6 we discuss the results and in Section 7 we draw conclusions.

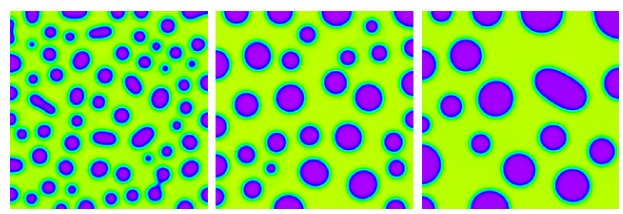


Figure 2: Three timesteps of the original Cahn Hillard phase separation.

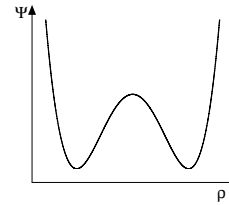


Figure 3: Chemical energy as function of concentration

## 2 Reviewing a physical clustering model

Before we discuss our model of continuous clustering on simulation data we will review in this section a physical model for clustering in metal alloys, which goes back to Cahn and Hilliard [3]. The Cahn–Hilliard model was introduced to describe phase separation and coarsening in binary alloys. Phase separation occurs when a uniform mixture of the alloy is quenched below a certain critical temperature underneath which the uniform mixture becomes unstable. As a result a very fine microstructure of two spatially separated phases with different concentrations develops. In later stages of the evolution on a much slower time scale than the initial phase separation the structures become coarser: either by merging of particles or by growing of bigger particles at the cost of smaller ones. This coarsening can be understood as a clustering, where the system mainly tries to decrease the surface energy of the particles which leads to coarser and coarser structures during the evolution. In the basic Cahn–Hilliard model this surface energy will be isotropic. There are no preferred directions of the interfaces. Hence the particles tend to be ball shaped (cf. Fig. 2). We now shortly outline the basic ideas of the Cahn–Hilliard model. For more details we refer to the review papers by Elliott [6] and Novick–Cohen [14]. The Cahn–Hilliard model is based on a Ginzburg–Landau free energy which is a functional in terms of the concentration difference  $\rho$  of the two material components. The Ginzburg–Landau free energy  $E$  is defined to be

$$E(\rho) := \int_{\Omega} \left\{ \Psi(\rho) + \frac{\gamma}{2} |\nabla \rho|^2 \right\},$$

where  $\Omega$  is a bounded domain. The first term in the free energy,  $\Psi(\rho)$ , is the chemical energy density and typically has a double well form. In this paper we take

$$\Psi(\rho) = \frac{1}{4} (\rho^2 - \beta^2)^2$$

with a constant  $\beta \in (0, 1]$  (cf. Fig. 3). We note that the system is locally in one of the two phases if the value of  $\rho$  is close to one of the two minima  $\pm\beta$  of  $\Psi$ . Now, the diffusion equation for the concentration  $\rho$  is given by

$$\frac{\partial \rho}{\partial t} = \Delta w$$

on  $\mathbb{R}^+ \times \Omega$ . In the equation above we denote by  $w$  the local chemical potential difference which is given as the variational derivative  $\frac{\delta E}{\delta \rho}$  of  $E$  with respect to  $\rho$  (cf. Section 3). Thus, we obtain

$$w = -\gamma \Delta \rho + \Psi'(\rho).$$

The system has to be supplemented with boundary and initial conditions. Here we request  $\frac{\partial w}{\partial \nu} = \frac{\partial}{\partial \nu} \rho = 0$ , where  $\nu$  is the outer normal on  $\partial \Omega$ , and  $\rho(0, \cdot) = \rho_0(\cdot)$  for some initial concentration distribution  $\rho_0$ . We remark that with these boundary conditions mass is conserved and that the Ginzburg–Landau free energy is a Lyapunov functional, i.e. we have

$$\frac{d}{dt} \int_{\Omega} \rho(x, t) dx = 0 \quad \text{and} \quad \frac{d}{dt} E(\rho(t)) \leq 0.$$

Starting with a random perturbation of a constant state  $\bar{u}_0$ , which has a values in the unstable concave part of  $\Psi$ , we observe the following: In the beginning the chemical energy decreases rapidly whereas the gradient energy increases. This is due to the fact that during phase separation  $\rho$  attains values which are at large portions of the domain close to the minima of the chemical energy  $\Psi$ . Since regions of different phase are separated by transition zones with large gradients of  $\rho$ , the gradient energy increases during phase separation. In the second stage of the evolution — the actual clustering — when the structures become coarser, the total amount of transition zones decreases. Correspondingly the amount of gradient energy becomes smaller again.

### 3 A Multiscale clustering approach

The aim of this section is to derive a continuous clustering model mainly on flow data. Motivated by the Cahn Hilliard model for phase separation and particle coarsening (cf. Section 2), we introduce a cluster mapping  $u : \mathbb{R}_0^+ \times \Omega \rightarrow \mathbb{R}$  which will be the solution of an appropriate evolution problem. Thereby, time will serve as the scale parameter leading from fine cluster granularity to successively coarser clusters. For fixed time  $t$  our definition of the set of clusters  $\mathcal{C}(t)$  is founded on the function  $u$  by

$$\mathcal{C}(t) = \{x \mid u(t, x) \geq 0\}.$$

This set splits up into the actual clusters

$$\mathcal{C}(t) = \bigcup_i \mathcal{C}_i(t)$$

where  $\{\mathcal{C}_i(t)\}_i$  are the connected components of  $\mathcal{C}(t)$ .

Now we study the evolution problem which controls the quantity  $u$ . We suppose this evolution to be a suitable clustering model, if for the induced  $\mathcal{C}(t)$

- the number of clusters generically decreases in time,
- the shape of the cluster components strongly corresponds to correlations in the data field,
- the volume fraction covered by  $\mathcal{C}(t)$  is approximately constant in  $t$ , i. e.  $\frac{|\mathcal{C}(t)|}{|\Omega|} \approx \Theta$  for  $\Theta \in (0, 1)$ .

We pick up the physical Cahn-Hilliard model and consider a double well separation potential  $\Psi(u)$  and define a separation energy  $E_s = \int_{\Omega} e_s(u) dx$  with energy density  $e_s(u) = \Psi(u)$ . Under all  $u$  with  $\int_{\Omega} u dx = \bar{u}_0 = \text{const.}$  the energy  $E_s$  is minimal if  $u$  attains only the values  $\pm \beta$ . This leads to a binary decomposition of the domain into two parts, where one part corresponds to  $\{x \mid u(x) = \beta\}$ .

The set  $\{x \mid u(x) = \beta\}$  however can have many connected components and may even be very unstructured. Furthermore there is no mechanism which enforces a successive coarsening and thus a true multiscale of clusters. Therefore, we want to introduce a term penalizing the occurrence of many disconnected cluster components with high interfacial area. Motivated by the Cahn–Hilliard theory of phase transition we choose a gradient energy  $E_{\partial} = \int_{\Omega} e_{\partial} dx$  with local energy density  $e_{\partial}$  that penalizes rapid spatial variations of  $u$ .

In order to have flexibility to choose an anisotropic and inhomogeneous gradient energy, an appropriate definition of an interfacial energy density is given by

$$e_{\partial}(\nabla u) = \frac{\gamma}{2} A \nabla u \cdot \nabla u,$$

where “ $\cdot$ ” denotes the scalar product in  $\mathbb{R}^n$ ,  $\gamma$  is a scaling coefficient and  $A \in \mathbb{R}^{n \times n}$  is some symmetric positive definite matrix that may depend on the space variable and other quantities involved.

In the following we will call the set  $\partial\{x \mid u(x) = 0\}$  the interface. The orientation of the interface can be described by the normal to the interface which, in the case that  $\nabla u \neq 0$ , is given by

$$\nu = \frac{\nabla u}{\|\nabla u\|}.$$

We remark that the interface between the set of positive and negative values of  $u$  is perpendicular to  $\nu$ . For  $A = \text{Id}$  all gradients of  $u$  and hence, all interfaces are penalized equally independent of their orientation. With respect to our clustering intention we consider an anisotropic energy density which strongly depends on the orientation of the local interface and thereby on the direction of  $\nabla u$ .

Let us assume  $v : \Omega \rightarrow \mathbb{R}^n$  to be some vector field on the domain  $\Omega$ . Typically such a field induces a flow on  $\Omega$  with streamlines which are solution of the ordinary differential equation  $\dot{x} = v(x)$ . Now, a natural clustering should emphasize the coherence along the induced streamlines. Thus, cross streamline interfaces have to be penalized significantly by the gradient energy. We choose

$$A := B(v)^T \begin{pmatrix} 1 & 0 \\ 0 & \alpha(\|v\|) \text{Id}_{n-1} \end{pmatrix} B(v)$$

where  $\text{Id}_{n-1}$  is the identity mapping in  $\mathbb{R}^{n-1}$  and for given  $r \in \mathbb{R}^n$  the mapping  $B(r) \in SO(n)$  is a coordinate rotation with  $B(r)r = \|r\|e_1$ . Since interfaces that cross streamlines have to have larger energy we choose a positive  $\alpha$  with  $\alpha \leq 1$ .

Now we define the first variation of the energy

$$w = \frac{\delta E}{\delta u},$$

which is defined on  $\Omega$  by

$$\int_{\Omega} \frac{\delta E}{\delta u} \zeta dx := \frac{d}{d\epsilon} E(u + \epsilon \zeta) \Big|_{\epsilon=0}.$$

We obtain  $w = \Psi'(u) - \gamma \text{div}(A \nabla u)$ .

Let us assume that the evolution of the cluster mapping  $u$  is governed by diffusion where the corresponding flux linearly depends on the negative gradient of the first variation of energy. As the simplest model we choose  $\frac{\partial}{\partial t} u - \Delta w = 0$  and end up with the following fourth order parabolic problem:

Find a continuous cluster mapping  $u : \mathbb{R}_0^+ \times \Omega \rightarrow \mathbb{R}$ , such that

$$\frac{\partial}{\partial t} u - \Delta w = 0 \tag{1}$$

$$w = \Psi'(u) - \gamma \text{div}(A \nabla u) \tag{2}$$

with boundary conditions  $\frac{\partial}{\partial \nu} u = \frac{\partial}{\partial \nu} w = 0$  and prescribed initial data  $u(0, \cdot) = u_0(\cdot)$ .

This modified Cahn–Hilliard equation can be interpreted as the  $H^{-1}$  gradient flow for the energy  $E$  (see [14] for a discussion of this fact in the case of the standard Cahn–Hilliard equation). In particular we immediately obtain the Lyapunov property

$$\frac{\partial}{\partial t} E(u) \leq 0. \quad (3)$$

This energy decay is in fundamental accordance to the desired successive pattern coarsening in the evolution. After an initial short period of phase separation it is mainly the interfacial energy contribution which is successively reduced. Furthermore, as in the case of the standard Cahn–Hilliard equation, we obtain that  $\int_{\Omega} u(x, t) dx$  is constant in time, which corresponds to the approximate volume conservation of the generated scale of cluster sets.

In general it does not make sense to consider certain initial data, if no a priori information on the clustering is known. As initial data  $u_0$  we thus choose a constant value  $\bar{u}_0$  plus some small random noise. The constant  $\bar{u}_0$  depends on the volume fraction  $\Theta$  of the domain later on to be covered by the clusters, i.e. by the sets  $\{x \mid u(t, x) \geq 0\}$ . Therefore, we choose

$$\bar{u}_0 = \Theta\beta - (1 - \Theta)\beta.$$

Starting with a random perturbation of this constant first very rapidly cluster patterns will grow without any prescribed location and orientation. This is in order to decrease  $E_s = \int_{\Omega} \Psi(u) dx$  which forces the solution to obtain values close to  $\pm\beta$  in most of the domain  $\Omega$ . After this process the clusters orientate themselves in an anisotropic way to decrease the amount of the anisotropic gradient energy  $E_{\beta}$ . In addition, the cluster becomes coarser and coarser due to the fact that smaller particles shrink and larger ones grow. We remark that in particular one observes that a large particle that is surrounded by smaller ones grow to the expense of the smaller ones. This implies that as time evolves locally only the main features of the clusters will be kept.

Altogether, we obtain a scale  $u(t, \cdot)$  of cluster mappings and induced cluster sets  $\mathcal{C}(t)$ . They represent a successively coarser representation of simulation data and continuously enhances coherences in the underlying simulation data set, where the cluster set  $\mathcal{C}(t)$  will cover a volume of approximate size  $\Theta|\Omega|$ . As already mentioned, the multiscale property comes along with the decay of the energy  $E(u)$  (see (3)).

To summarize, the vector field that is to be represented defines the anisotropy of the energy and therefore governs the diffusion process of  $u$ . Roughly speaking, the vector field determines in which direction an interface between phases is relatively "cheap" (from an energy point of view). As the energy is minimized during the evolution the interfaces will move in such a way that there are mostly "cheap" interfaces (i.e. interfaces where  $\nabla u$  is roughly perpendicular to  $v$ , which means that the interface is roughly parallel to  $v$ ). So for any particle most of its boundary will be aligned with the vector field  $v$ . If not to large the particles themselves will be aligned to the vector field.

## 4 Discretization of the Diffusion Problem

In what follows we briefly discuss the discretization and implementation of the evolution problem for the cluster mapping  $u$  and the set of clusters  $\mathcal{C}(t)$ . For this purpose a finite element discretization in space and some discrete scheme in time are considered. Here, uppercase letters denote discrete quantities which correspond to continuous quantities in lowercase letters. Hence, we consider an ap-

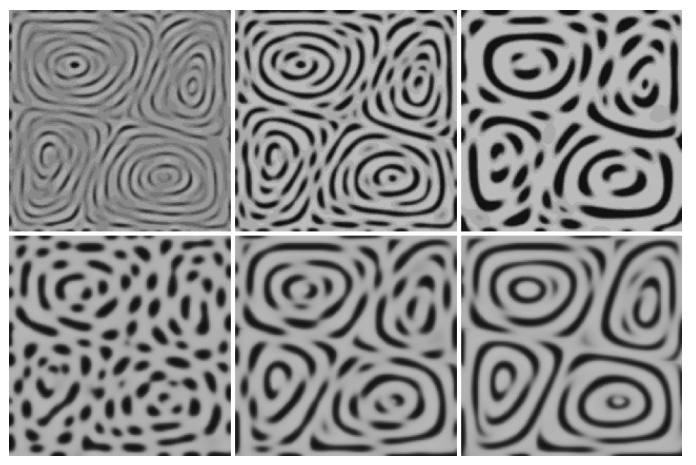


Figure 4: Continuous clustering of a vector field: time evolution (upper row), effect of increasing anisotropy (lower row). The computation is based on a grid of resolution  $257^2$ .

propriate continuous variational formulation for (1), (2), given by

$$\begin{aligned} \left( \frac{\partial}{\partial t} u, \theta \right) + (\nabla w, \nabla \theta) &= 0, \\ (w, \chi) &= (\Psi'(u), \xi) + (A \nabla u, \nabla \xi), \end{aligned}$$

which shall hold for all  $\theta, \xi \in C^\infty(\bar{\Omega})$ , where  $(\cdot, \cdot)$  denotes the  $L^2$  product on the domain  $\Omega$ . For a finite element implementation we now replace the continuous solution and test functions in this formulation by discrete approximations in some finite element space. Here we have restricted ourselves to finite elements on regular adaptive grids  $\mathcal{M}_h$  in 2D and 3D generated by recursive subdivision of elements  $E$ . On these grids we consider the bilinear, respectively trilinear finite element spaces  $V^h$  for the approximation of  $u$  and  $w$  on  $\Omega$ . Numerical integration of the  $L^2$  products is based on the lumped masses product  $(\cdot, \cdot)_h$  [22]. Furthermore we consider a center of mass quadrature rule for the bilinear forms  $(\nabla \cdot, \nabla \cdot)$  and  $(A \nabla \cdot, \nabla \cdot)$ . Especially, we replace  $A$  by the piecewise constant diffusion tensor  $A^h$ , with  $A^h|_E = A(c_E)$ , where  $c_E$  is the element's center of mass.

For the discretization in time we have taken into account two possibilities: a first order implicit Euler scheme and a second order  $\theta$ -splitting scheme (see Bristeau et al. [1] and Müller Urbaniak [13]). Both are known to be strongly A-stable. While we can prove the energy decay property (3) for the implicit Euler scheme, we use the  $\theta$ -splitting for practical computations as it allows larger time steps.

In the case of the implicit Euler scheme the time derivative is discretized by  $\frac{\partial}{\partial t} u((n+1)\tau) \approx \frac{U^{n+1} - U^n}{\tau}$  where  $\tau$  is the selected time step and  $U^h$  an approximation of  $u(n\tau)$ . A brief introduction to the more complicated  $\theta$ -splitting can be found in the appendix.

Finally, we can derive a fully discrete scheme. For the "hat shaped" multilinear basis functions  $\Phi_i$  and the discrete piecewise constant anisotropic diffusion matrix  $A^h$  we define by

$$\begin{aligned} M_h &:= ((\Phi_i, \Phi_j)_h)_{ij}, \\ L_h(A) &:= \left( (A^h \nabla \Phi_i, \nabla \Phi_j) \right)_{ij} \end{aligned}$$

the diagonal lumped mass and the anisotropic stiffness matrix respectively and by  $L_h := L_h(\text{Id})$  the standard stiffness matrix. These global matrices  $M_h$ ,  $L_h$ , and  $L_h(A)$  are assembled in a grid traversal collecting the contributions on all local grid elements as it is standard in finite element programming [4].

If we indicate by a bar coefficient vectors corresponding to finite element functions in the basis  $\{\Phi_i\}_i$ , we obtain the backward Euler discretization

$$M_h \bar{U}^{n+1} + \tau L_h(\bar{\Psi}'(\bar{U}^{n+1})) + \gamma M_h^{-1} L_h(A) \bar{U}^{n+1} = M_h \bar{U}^n$$

with  $U^0 = \mathcal{I}_h u_0$ , where  $\mathcal{I}_h$  is the interpolation on grid  $\mathcal{M}_h$ . By obvious notation  $\bar{\Psi}'(\cdot) := (\Psi'(\cdot))_i$  is the vector of nodal wise derivatives of  $\Psi$ .

In each step of the discrete evolution we have to solve this system of nonlinear equations. In order to do this we apply some Newton scheme which typically converges in a few steps if we consider moderate time steps and pick up the old solution at the old time step as the initial guess for the newton iteration.

The efficiency of our approach is further increased by an adaptive grid refinement and coarsening strategy. Here, we used a heuristic strategy which refines in interfacial regions and coarsens in the pure phases.

In the case of the implicit Euler scheme, it is possible to prove discrete counterparts of the mass conservation and energy decay properties. I. e.

$$\int_{\Omega} U^n dx = \int_{\Omega} \mathcal{I}_h u_0 dx$$

and

$$E_h(U^n) := \int_{\Omega} \left\{ \mathcal{I}_h \Psi(U^n) + A^h \nabla U^n \cdot \nabla U^n \right\} dx$$

is non increasing (discrete Lyapunov property) and thus gives reason for the discrete multiscale property of our method

$$E_h(U^{n+1}) \leq E_h(U^n) \leq \dots \leq E_h(U^1) \leq E_h(\mathcal{I}_h u_0).$$

## 5 Iconic Representation of the Clusters

The clustering method described in Section 3 produces clusters which emphasize the spatial coherence in the data. In what follows concerning the iconic representation we focus to the case of flow data. Nevertheless this exposition might inspire the reader to think of different applications along the same guidelines. For flow data, cluster interfaces tend to be tangent to the streamlines of the underlying vector field, so the clusters' shapes convey local insight in the vector field direction. On the other hand, the physical phase separation model presented in section 2 produces clusters which tend to be evenly distributed over the domain of interest  $\Omega$ .

Consequently, such clusters could be used as a starting point for producing a simplified visualization of the structure of the underlying vector field. For this, we propose to reduce each cluster to one curved arrow icon. For every cluster, the size and spatial position of the icon should reflect the size of the cluster, whereas the curvature and arrow direction should be related to the vector field inside the respective cluster. We have chosen to use the curved arrow icons as they convey several information levels in a simple, easy to understand manner [21], as compared to other, more abstract icons.

The iconic visualization pipeline based on the multiscale clustering proceeds as follows (see also Figs. 5 and 6). First, the clusters  $\mathcal{C}$  are extracted from the Cahn-Hilliard equation solution  $u$ . Next, the skeletons of the clusters are computed as sets of discrete points, as shown further in section 5.2. Next, the center points of the skeletons are detected and used to construct the curved arrows by streamline tracing, as discussed in section 5.4. The reason why we use such an apparently complicated method is, that typically the clusters are large thin often curved structures and streamline tracing is rather sensitive with respect to the choice of the starting point. The rest of this section explains the several steps in detail.

## 5.1 Cluster Extraction

First we extract the clusters  $\mathcal{C}$  from the scalar field  $u$ . For this, we classify all the cells of the discretization of the field  $u$  as cluster outside, border, or inside cells, based on the sign of  $u(x)$ . Moreover, all border and inside cells belonging to a given cluster are labelled by the cluster's ID as presented by e.g. Walsum et al. in [16].

## 5.2 Skeletonization

In the second step, clusters are reduced to their *skeletons*. By skeleton, we understand here a set of points which, if connected, produce a 'spine' which conveys the shape information of the original cluster in a compact manner.

There are numerous skeletonization algorithms [10, 12]. However, many such algorithms produce skeletons with complex, tree-like topologies. As we intend here to use the skeletons only to produce the arrow icons, we prefer simple, polyline-like topologies.

To produce such skeletons, we use a discrete method based on the eikonal equation [20]. Given a boundary curve  $\Gamma$  in two dimensions (or a boundary surface, in 3D) and a function  $T$ , such that  $T = 0$  on  $\Gamma$ , the eikonal equation is  $|\nabla T| = 1$ . If we regard  $\Gamma$  as being the level set (e.g. isoline or isosurface) of the function  $T$ , the above equation describes the evolution in time of  $\Gamma$  in normal direction to  $\Gamma$ . In our case,  $\Gamma$  coincides with the previously detected cluster boundaries.

As presented by Sethian [20], we discretize the above equation on the same grid used to solve the Cahn-Hilliard equation, as follows (for the 2D case):

$$\begin{aligned} \max(D_{ij}^{-x} T, 0)^2 + \min(D_{ij}^{+x} T, 0)^2 + \\ \max(D_{ij}^{-y} T, 0)^2 + \min(D_{ij}^{+y} T, 0)^2 = F_{ij}^{-2}, \end{aligned} \quad (4)$$

where the  $ij$  denotes the current grid point and the operators  $D^+$  and  $D^-$  denote the forward and backward differences at that grid point. On a 2D regular grid of cell size  $h$  we have  $D_{ij}^{+x} T = \frac{T_{i+1,j} - T_{i,j}}{h}$  and  $D_{ij}^{-x} T = \frac{T_{i,j} - T_{i-1,j}}{h}$ , and similarly for the  $y$  axis.

Equation (4) is solved by using the fast marching method, as described in [20]. The fast marching method starts constructing the solution  $T$  from the initially known points with  $T = 0$  on the boundary  $\Gamma$ . At each step, the solution  $T$  is constructed from the point with the smallest computed  $T$  value, by stepping away from the boundary in a downwind direction. The boundary is marched inwards until it collapses into a single line, namely the skeleton points of the cluster  $\Gamma$ .

## 5.3 Reconnection and Center Detection

The skeletonization produces a set of usually disjoint skeleton points. Next, these points are reconnected (based on a closest point strategy) to form a polyline. For every cluster, we then compute the center of its polyline and use it in the next step of the pipeline.

## 5.4 Icon Construction

From the skeleton centers detected in the previous step, streamlines are traced in the vector field until they reach the borders of the clusters within which they evolve. Next, curved arrow geometries are constructed around the extracted streamlines. Finally we discuss the application of the continuous clustering method and the associated curved arrow visualization to various vector fields.

The leftmost image in Fig. 6 shows a solution  $u(t, \cdot)$  of the Cahn-Hilliard process driven by a 3-vortex vector field on a  $64 \times 64$  grid.

The thresholding of the continuous signal  $u$  into clusters is shown in the second image of Fig. 6. The clusters overlaid with the



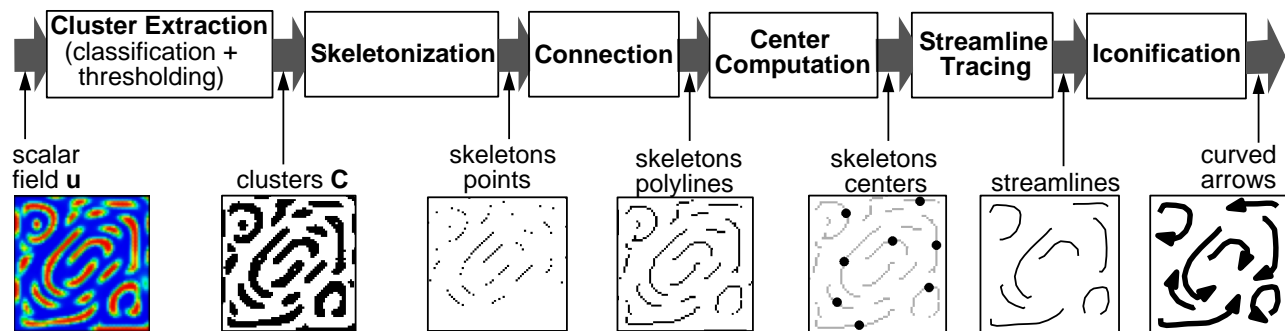


Figure 5: Pipeline for iconic cluster visualization

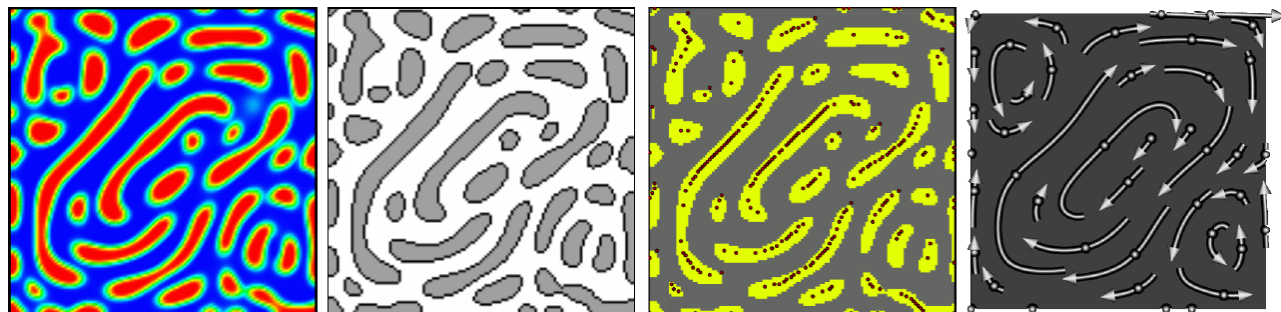


Figure 6: Clustering pipeline, from left to right: diffusion solution, clusters, skeleton points, curved arrow visualization. The underlying grid is of size  $257^2$ .

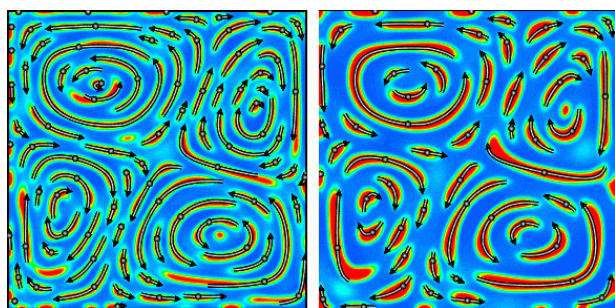


Figure 7: From left to right the results of two successive time steps of the clustering evolution with corresponding icons are shown.

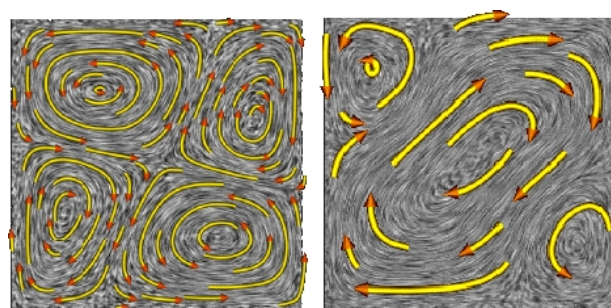


Figure 8: Two different convective vector fields are depicted by the clustering method. On the left a fine representation is shown, whereas on the right a resulting coarse representation is depicted.

extracted skeleton points are shown in the third image of Fig. 6. The rightmost image in Fig. 6 visualizes the vector field with streamline-based icons. The curved arrows, initiated from the skeleton centers depicted as small balls, are clipped by the borders of the clusters into which they evolve.

A similar visualization is shown, for two different clustering time instants, in Fig. 7. The multiscale feature of the clustering is visible in the reduction of the arrow count. An enhancement of the proposed curved arrow visualization is shown in Fig. 8 by the addition of a spot noise textured background. Finally, Fig. 9 shows the proposed method applied on a circular 2D vortex.

## 6 Discussion

In this section, we compare the presented continuous clustering method with the discrete clustering method presented in [21]. Similarly to the method presented here, discrete clustering builds a vec-

tor field multiscale representation by merging neighbouring cells with similar vector values. The time parameter of the Cahn-Hilliard equation is equivalent with the iteration count in the bottom-up discrete cluster merging. The continuous clustering method delivers a continuous scale of successively coarser cluster sets. In contrast, discrete clustering proceeds in distinct steps, where two clusters are merged at each step.

Figure 11 shows the discrete clustering of the two vector fields discussed in the previous section. Regarding the cluster shapes, the continuous clustering explicitly constrains the shape via the minimization of the interfacial energy, in order to obtain vector-aligned, smooth-shaped clusters. In contrast, the discrete clustering does not constrain the cluster shapes in any manner, assuming that their growth to a 'natural' partition of the vector field can be governed only by the inter-cluster similarity function. This can lead however to 'badly' shaped (e.g thin and long) clusters, which are hard to represent by curved arrow icons. In this respect, we see the con-

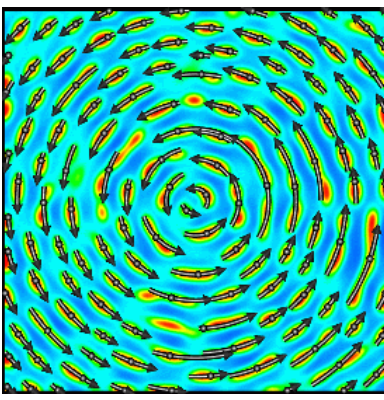


Figure 9: Visualization of a circular flow field by the clustering method.

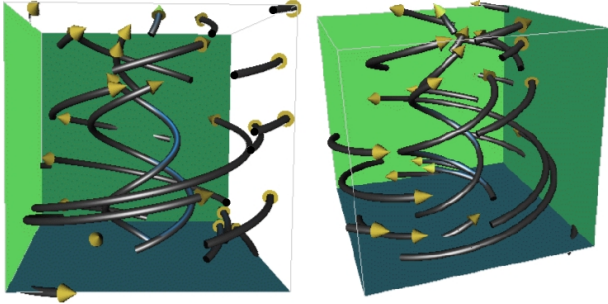


Figure 10: A 3D vectorfield is visualized by the clustering method

trolling of the cluster shape in the continuous clustering method as an advantage. However, discrete clustering always merges the two most resembling clusters, so the intrinsic symmetry of the underlying vector field remains visible in the clustering (see [21] for details). This may be seen as an advantage of the discrete clustering method, see Fig. 11. Finally, the shapes produced by the continuous clustering are not constrained to simple convex ones, as in the method presented by Heckel et al [8]. We have applied the continuous clustering method also to the visualization of 3D fields. Fig. 10 shows the visualization of a 3D circular vortex field from two different viewpoints. The produced arrow icons illustrate the clustering of the data in the center of the domain, where the flow is dominated by a vertical swirling motion, and along the domain's boundary, where the flow mainly rotates in horizontal planes.

Finally we present an application of our approach for 2D image processing, where we generate a scale of brush stroke type representations of a greyscale image. Thus, let us consider the intensity of an image as a scalar function  $s : \Omega \rightarrow \mathbb{R}$ . If we intend to release brush stroke along regions of homogeneous values of the scalar quantity  $s$ , we need to energetically favour interfaces which have a tangent space locally perpendicular to  $\nabla s$ . Hence, we choose a corresponding quadratic form with

$$A := B(\nabla s)^T \begin{pmatrix} \alpha & 0 \\ 0 & \text{Id}_{n-1} \end{pmatrix} B(\nabla s),$$

where for given  $r \in \mathbb{R}^n$  the mapping  $B(r) \in SO(n)$  is again a coordinate rotation with  $B(r)r = \|r\|e_1$ , and  $1 \gg \alpha > 0$  (cf. Fig. 12).

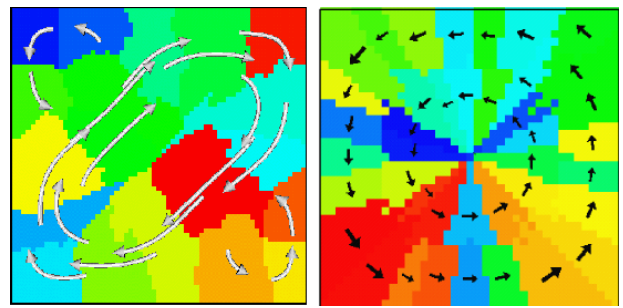


Figure 11: Discrete clustering of a 3-vortex field (left) and a circular vortex field (right).

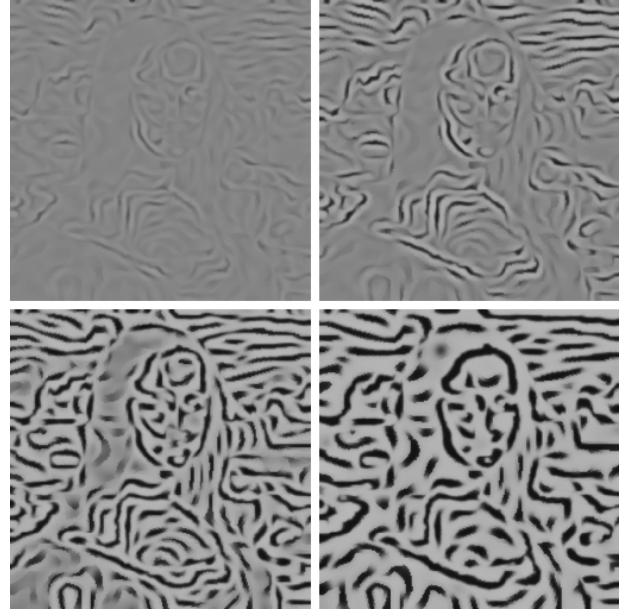


Figure 12: Multiscale brush-stroke representation of a greyscale Mona Lisa image.

## 7 Conclusions

We have presented a new multiscale clustering approach which is based on a continuous model for clustering on scientific data. The approach is motivated by well-known physical clustering models describing the phase separation and coarsening process in metal alloys. As a case study we focused on the clustering of flow fields. Future research could be directed to

- the improvement of the performance of the method with respect to computing time, where parallization or implementation of the underlying diffusion in texture hardware may help to overcome the computational bottleneck,
- the construction of further appropriate interfacial energies for different applications,
- a detailed classification of the skeleton shapes and the selection of appropriate icons, e. g. for saddle points or vortices in flow fields.

## Appendix

Here we briefly outline the implementation of the  $\theta$ -splitting scheme. Due to its strong stability properties it allows much larger timesteps scheme and we have used it in the current implementation of our clustering model. The scheme divides any time step in three substeps (see Figure 13). In each substep the linear operator is split up into two parts with coefficients  $\alpha$  and  $1 - \alpha$  respectively, one of which is taken implicitly the other explicitly. The nonlinear term is taken implicitly in the middle substep only.

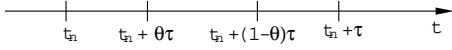


Figure 13:  $\theta$ -splitting, subdivision of time steps

For the parameter  $\alpha \in (0.5; 1]$ ,  $\beta = 1 - \alpha$  and  $\theta \in (0; 0.5)$  the scheme reads as follows:

$$\begin{aligned}
 & \left( M_h + \alpha\theta\tau L_h M_h^{-1} L_h(A) \right) \bar{U}^{n+\theta} \\
 &= \left( M_h - \beta\theta\tau L_h M_h^{-1} L_h(A) \right) \bar{U}^n - \theta\tau L_h \psi'(\bar{U}^n), \\
 & \left( M_h + \beta(1-2\theta)\tau L_h M_h^{-1} L_h(A) \right) \bar{U}^{n+1-\theta} \\
 & \quad + (1-2\theta)\tau L_h \psi'(\bar{U}^{n+1-\theta}) \\
 &= \left( M_h - \alpha(1-2\theta)\tau L_h M_h^{-1} L_h(A) \right) \bar{U}^{n+\theta}, \\
 & \left( M_h + \alpha\theta\tau L_h M_h^{-1} L_h(A) \right) \bar{U}^{n+1} \\
 &= \left( M_h - \beta\theta\tau L_h M_h^{-1} L_h(A) \right) \bar{U}^{n+1-\theta} \\
 & \quad - \theta\tau L_h \psi'(\bar{U}^{n+1-\theta})
 \end{aligned}$$

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