

1 Segmentation of shot peening patterns using **k**-means 2 clustering

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5 Abstract

6 Shot peen forming is a prominent manufacturing process for shaping aircraft
7 wing skins, which mostly relies on manual intervention, where operators use
8 their expertise to select appropriate peening parameters and apply them in
9 a specific pattern. This paper introduces a crucial advancement towards au-
10 tomating shot peen forming—a segmentation strategy designed to partition
11 peening patterns into uniformly treated zones. Unlike existing optimization
12 methods that necessitate predefined peening treatments, our segmentation
13 strategy automatically identifies optimal treatment parameters for each seg-
14 ment. Our approach comprises a novel clustering algorithm, which divides
15 the pattern into segments, and a noise filtering algorithm that eliminates
16 excessively small segments. The clustering algorithm is a unique adaptation
17 of the k-means method, which considers interconnected centroids due to the
18 coupling of the effects of the top and bottom treatments of the part. The
19 filtering algorithm leverages cellular automata principles. Both algorithms
20 underwent numerical testing using 200 randomly generated test cases. The
21 results indicate that the segmentation strategy consistently maintained forming
22 error within an acceptable range, and remarkably, reduced the forming
23 error in 67 out of 200 cases. This segmentation strategy can seamlessly inte-
24 grate with existing shape optimization tools and a peening treatment library,
25 leading to a fully automated shot peen forming system. The source code for
26 our algorithms is publicly available on GitHub, fostering accessibility and

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²⁷ further research in this domain.

²⁸ *Keywords:* Segmentation, Clustering, Filtering, k-means, Peen forming

1. Introduction

Fabricating aircraft and rockets requires forming large metal plates with high precision. These plates constitute, for example, wing skins, fuselage shells or fuel tank segments. The outlined components often exhibit a nonuniform double curvature, meaning that they are locally curved in two directions and that the local radii of curvature vary over the plate. A cost-effective shaping method that delivers such forms is shot peen forming. This treatment consists in projecting a stream of rigid shot towards the surface of the plate. The shot stream plastically deforms the surface and induces bending of the component (Baughman, 1970). The stream width is smaller than the plate size, and the treatment can be applied to both sides of the plate with a variable intensity. This means that the developed curvatures can be locally controlled by altering the peened segments and the peening parameters. Thus, in the case of wing skins, peen forming is sequentially applied to produce a chord-wise curvature, a span-wise curvature, and a dihedral bend angle. It can also be applied for local stretching and bending of stiffened components (Ramati et al., 1999).

Shot peen forming is still largely performed manually because it relies on the operator’s intuition and expertise in shaping. The disadvantages of the manual process with respect to the automated one include a long and costly trial-and-error design phase, expensive staff training, and health risks for the operators (Frieese, 2006; Miao et al., 2022). However, an essential element of process automation is a numerical simulation tool that can be integrated with a peening robot. Such a tool can enhance repeatability and eliminate the necessity for the trial-and-error search of the optimal pattern, which sometimes takes several months. Several simulation tools were developed in-house by industrial companies, but they are not publicly accessible (Wüstefeld et al., 2006). Moreover, in the public scientific literature, no method exists for selecting shot peen forming treatments based on a given peening pattern. Here, we propose a novel clustering-based method for automatically selecting a peening treatment, which can be combined with the pattern optimization method published earlier (Sushitskii et al., 2022b). No comparison is made with other peening pattern segmentation methods in the literature, because none exists that we are aware of.

Numerical simulation of shot peen forming involves the solution of the *forward* and of the *inverse* problems (Sushitskii et al., 2022b). The forward

65 problem solution means computing the deformation resulting from a pre-
66 defined peening treatment. The inverse problem solution means defining a
67 necessary treatment to achieve a predefined target shape. Hence, it is a nu-
68 merical inverse problem solver which is necessary for industrial companies to
69 accelerate the design phase.

70 The solution to the inverse problem in peen forming is a map of local peening
71 parameters, which we term as *regimes* throughout the paper. These regimes
72 are mapped over the initial geometry of the treated plate, which is called
73 the *peening pattern*. For instance, Figure 1 shows two peening patterns that
74 allow shaping a flat square plate into a (a) target spherical shape: (b) one
75 with continuously varying intensity or an infinity of regimes; (c) and one
76 discrete regimes. The first steps towards numerical resolution of the inverse
77 problem were made by VanLuchene et al. (1995); VanLuchene & Cramer
78 (1996), who simulated the effect of peening using the peening-induced forces.
79 Later, two Kopp & Schulz (2002) and Wang et al. (2008) presented two
80 inverse problem solvers operating directly with the peening parameters . In
81 all these cases, the peening parameters were experimentally related to the
82 peening-induced forces (VanLuchene & Cramer, 1996) or curvatures (Kopp
83 & Schulz, 2002; Wang et al., 2008). The inverse problem can also be solved
84 through formulating the peening pattern in terms of the peening-induced
85 residual stresses (Essa et al., 2015). In turn, the residual stresses can be
86 related to the peening parameters through direct impact simulations (Gariépy
87 et al., 2012; Wang et al., 2022) or through local analytical impact models
88 coupled with global numerical simulations (Xiao et al., 2019).

89 The inverse problem solvers relying on the peening-induced forces (VanLuch-
90 ene et al., 1995; VanLuchene & Cramer, 1996) and stresses (Essa et al., 2015)
91 as loads lose their accuracy in case of large peening-induced rotations. Thus,
92 the large rotations provoke geometrically nonlinear behavior of the plate,
93 which breaks the experimentally developed relations between these loads and
94 the peening parameters. On the other hand, a direct empirical relation be-
95 tween the peening parameters and the induced curvatures (Kopp & Schulz,
96 2002; Wang et al., 2008) does not hold for complex peening patterns, because
97 the local curvatures are influenced by the curvatures in the neighboring zones.

98 Another strategy for the peen forming simulation involves formulating the
99 peening pattern in terms of local eigenstrains (Mura, 1987; Korsunsky, 2005).
100 The eigenstrain represents the plastic strain induced by the treatment, which
101 triggers deformation when introduced in the component. The advantage of

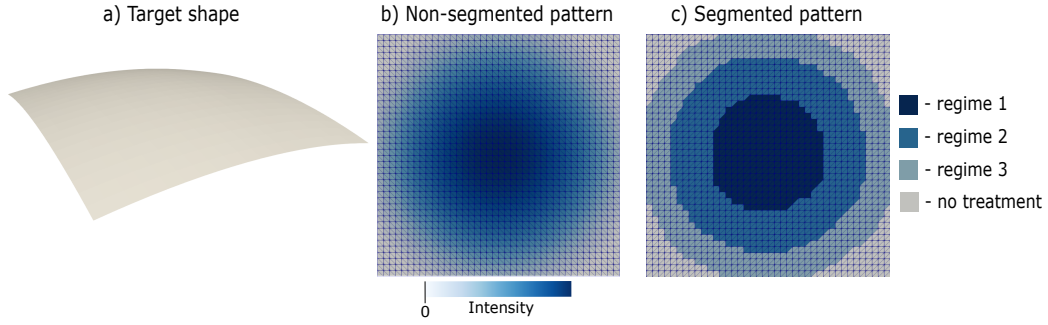


Figure 1. Examples of a non-segmented and of a segmented peening pattern. Both patterns make a flat square plate deform into a spherical patch (a) when applied from both sides of the plate. The non-segmented pattern (b) prescribes a gradual variation of the peening intensity over the plate, which is hardly reproducible with peening equipment. The segmented pattern (c) is reproducible because it consists of uniformly treated segments. Both patterns are discretized with triangular finite elements, and the effect of their application can be simulated using the eigenstrain approach. For this, one must establish in advance a one-to-one correspondence between the peening regimes or the intensities, and the induced eigenstrains.

eigenstrains is that they are independent of the plate geometry, unlike the peening-induced forces, stresses or curvatures (Zhang et al., 2008; Coratella et al., 2015). Moreover, the eigenstrains can be assumed constant during reconfiguration of the component, which was proven by means of numerical simulations and experiments with 76×19 mm shot peened metal coupons (Chen et al., 2014; Chaise et al., 2012). With the eigenstrain approach, the inverse problem can be solved using a numerical optimization algorithm (Miao et al., 2022), an artificial neural network (Siguerdidjane et al., 2020) or the theory of non-Euclidean plates (Sushitskii et al., 2022b). In turn, a one-to-one correspondence between the peening regimes and the induced eigenstrain can be established using experiments (Zhang et al., 2008), small-scale numerical simulations (Wang et al., 2002; Ohta & Sato, 2021), or a combination of both (Korsunsky, 2006). It is, however, imperative to hold the plate flat during peening and to ascertain the absence of stresses and strains in the initial state of the plate, because the established correspondence can be altered otherwise (Miao et al., 2022).

The eigenstrain approach also allows to simulate other forming processes that induce plastic strain in metal components, such as laser peen forming (Luo et al., 2020) or English wheel (Fann, 2022). Moreover, eigenstrains allow to represent a peen formed plate as a laminate subjected to incompatible

expansion or contraction of its layers (Wang et al., 2006), so this approach bridges the gap between peen forming and fabrication of laminated shape-shifting structures. For example, the inverse problem solver presented by Miao et al. (2022) works both for peen forming and for patterned multilayer films (Pajot et al., 2006), and the solver examined by Sushitskii et al. (2022b) uses the results developed in the field of 4D printed elastic sheets (van Rees et al., 2017, 2018).

The numerical inverse problem resolution involves the discretization of the initial geometry with shell finite elements. The computed peening pattern is therefore a discrete map. In the general case, the eigenstrains prescribed to each element vary from one element to another, as it is illustrated in Figure 1 b). On the other hand, the practical conditions of physically applying the peening treatment limit this spatial variation. Thus, the maximal eigenstrain that can be prescribed to a segment is limited by the equipment capacity. In addition, the peening parameters cannot be varied gradually over the plate. Moreover, each modification of peening parameters during treatment slows down the shaping process, which is especially the case for changing the media (Fauchaux, 2019). These reasons suggest that the pattern must consist of segments formed by multiple elements that are prescribed with the same eigenstrain. For a simpler and easier process, the segmented peening pattern must involve the fewest possible peening regimes while preserving the precision of the final shape.

Figure 1 c) presents an example of a segmented pattern. In practice, such segmented patterns are usually applied using masks (Kirk, 1999; Hornauer & Kohler, 1990). If the peening equipment admits only uniform treatment of the component during one cycle, then a separate mask is produced for each peening regime involved.

In particular, the problem of segmentation of the eigenstrain pattern is addressed by Miao et al. (2022); Siguerdidjane et al. (2020). The presented inverse problem solvers operate with one available eigenstrain magnitude, so the peening pattern is a set of treated and untreated segments. Miao et al. (2022) achieved this effect by limiting the overall treated area. This condition forces the optimization algorithm to assign the maximal eigenstrain magnitude to the treated elements and thus penalizes the intermediate eigenstrain values. On the other hand, the neural network presented by Siguerdidjane et al. (2020) naturally operates with one fixed eigenstrain magnitude, which is set during the training phase. Thus, the neural network computes a treat-

159 ment probability for each element, and the elements with high probabilities
160 are then assigned with the fixed eigenstrain magnitude. Although the ex-
161 amined solvers provide ready-to-use patterns, the use of only one eigenstrain
162 magnitude limits the range of achievable target shapes.

163 The inverse problem solver presented by [Sushitskii et al. \(2022b\)](#) does not
164 restrict the variation of the eigenstrain assigned to each element. The division
165 of the pattern into segments is done during post-processing with the help of
166 grouping. This approach allows any number of regimes in the pattern, which
167 means that it is flexible in terms of target shapes. However, the original and
168 the grouped patterns induce different final shapes, so the peening regimes
169 have to be thoroughly adjusted to minimize this difference.

170 The strategies for segmentation of the peening pattern presented by [Sushit-](#)
171 [skii et al. \(2022b\)](#); [Miao et al. \(2022\)](#); [Siguerdidjane et al. \(2020\)](#) share a
172 common drawback: they require having predefined peening regimes. This
173 suggests that if the current peening regimes do not allow to achieve the tar-
174 get shape according to the simulations, then the best suitable regimes must
175 be found by trial-and-error. Thus, if the segmentation is embedded in the
176 inverse problem solver ([Miao et al., 2022](#); [Siguerdidjane et al., 2020](#)), then
177 each new trial means restarting the inverse problem solver. Otherwise, if the
178 segmentation is executed after the inverse problem resolution by means of
179 grouping (see [Sushitskii et al., 2022b](#)), then the grouping algorithm must be
180 relaunched with a new combination of peening regimes, which are defined
181 manually. Hence, the necessity for having pre-defined peening regimes elon-
182 gates the inverse problem solution and makes it less accurate, because the
183 trial-and-error solution is not guaranteed to be the optimal.

184 Conversely, the problem of segmentation of a peening pattern is similar to the
185 problem of clustering ([Rokach & Maimon, 2005](#); [Xu & Tian, 2015](#)). Cluster-
186 ing algorithms split a set of points into groups based on the point coordinates.
187 In terms of the peening pattern, this means splitting the mesh into segments
188 based on the eigenstrain values assigned to each finite element. Moreover,
189 clustering algorithms are able to find a centroid for each cluster. With regard
190 to the peening pattern, this allows to find the mean eigenstrain in the given
191 segment and to homogenize the eigenstrain in this segment. Each eigenstrain
192 value corresponds to a peening regime, so a clustering algorithm is able to
193 prescribe the best suitable peening regime for each segment. This technique
194 allowed, for example, to compute the pattern shown in Figure 1 (c) starting
195 from the non-segmented pattern presented in Figure 1 (b).

196 An optimal clustering algorithm for the peen forming case must minimize
 197 the difference between the original and the segmented patterns in order to
 198 reduce the difference in the developed shapes. This can be accomplished
 199 using the **k**-means clustering algorithm (Lloyd, 1982), which groups points
 200 in space based on their proximity. This algorithm is also advantageous in
 201 terms of the computation speed and adaptability to sparse point sets (Rokach
 202 & Maimon, 2005; Xu & Tian, 2015).

203 We present in this paper a numerical strategy for segmenting the peening
 204 pattern without predefined peening regimes, which is designed as a post-
 205 processing tool for the inverse problem solver. The segmentation strategy
 206 relies on a novel modified **k**-means clustering algorithm, which is adapted
 207 for the peen forming case but can be applied to solve clustering problems
 208 in other domains. The clustering algorithm is followed by a noise filtering
 209 algorithm that corrects local clustering errors and thus makes the pattern
 210 fully applicable. In turn, the practical applicability of the segmented patterns
 211 is examined by Sushitskii et al. (2022a).

212 2. Theoretical background

213 2.1. The bilayer eigenstrain formulation of the peening pattern

214 The eigenstrain $\boldsymbol{\varepsilon}$ prescribed to each finite element by an eigenstrain-based
 215 inverse problem solver is constant in the in-plane direction inside the ele-
 216 ment, but it varies along the through-thickness coordinate z . We assume
 217 that the treated material is isotropic and not pre-stressed, so that $\varepsilon_{xx}(z) =$
 218 $\varepsilon_{yy}(z) = \gamma(z)$ and $\varepsilon_{zz}(z) = -2\gamma(z)$, where x and y are the in-plane lagrangian
 219 coordinates.

220 A precise measurement of the through-thickness eigenstrain profile $\gamma(z)$ is
 221 challenging due to the stochastic nature of the shot projection (Badreddine
 222 et al., 2014). Thus, the impacts superimpose with each other and are not
 223 necessarily normal to the surface. The shot are not perfectly spherical (Kirk,
 224 1999), and the peening parameters can fluctuate over time due to technical
 225 reasons. Finally, the impacts harden and heat the material (Rouquette et al.,
 226 2009), which alters its response to the subsequent indentations (Benallal
 227 et al., 1989).

228 In order to overcome these complexities in the framework of peen forming
 229 simulation, we idealize the through-thickness eigenstrain profile in the way

230 that it induces the same final shape as the original profile (when assuming
 231 plate theory, see [Sushitskii et al., 2022b](#)). In the general case of treatment
 232 from both sides, the simplest idealized profile involving the fewest variables
 233 is the bilayer profile γ_{bi} . This profile assumes that the plate consists of two
 234 equally thick layers assigned with the eigenstrains ε^t and ε^b as:

$$\gamma_{bi}(z) = \begin{cases} \varepsilon^t & \text{for } 0 < z < \frac{h}{2}, \\ \varepsilon^b & \text{for } -\frac{h}{2} < z < 0, \end{cases} \quad (1)$$

235 where h is the total plate thickness and z is the through-thickness coordinate
 236 having its origin at the mid-surface level. Figure 2 illustrates the idealization
 237 of the through-thickness eigenstrain profile.

238 The bilayer eigenstrain profile is locally described with two variables $(\varepsilon^t, \varepsilon^b)$,
 239 and each of the two peening regimes applied from the top- and bottom-sides
 240 influences both ε^t and ε^b . Thus, the treatment from the top side induces
 241 positive ε^t and negative ε^b , while the treatment from the bottom side induces
 242 negative ε^t and positive ε^b . In case of simultaneous treatments from both
 243 sides, the values of ε^t and ε^b are superpositions of contributions made by the
 244 top and the bottom side peening regimes.

245 Under the assumptions of plate theory, the final shapes induced by the pro-
 246 files $\gamma(z)$ and γ_{bi} coincide if the local forces and moments induced by the two
 247 profiles are equal, meaning that the conditions

$$\int_{-h/2}^{h/2} \gamma(z) dz = \frac{h}{2} (\varepsilon^t + \varepsilon^b), \quad (2)$$

$$\int_{-h/2}^{h/2} \gamma(z) z dz = \frac{h^2}{8} (\varepsilon^t - \varepsilon^b) \quad (3)$$

249 are met. Therefore, Eqns. (2, 3) allow to determine ε^t and ε^b element-
 250 wise and thus to reduce any through-thickness eigenstrain profile $\gamma(z)$ to the
 251 bilayer formulation.

252 2.2. The general approach to **k**-means clustering of the eigenstrain pattern

253 In terms of clustering, the variables $(\varepsilon^t, \varepsilon^b)$ act as coordinates of points on a
 254 plane, where each point corresponds to the peening state of a finite element in
 255 the peening pattern. Typical of a **k**-means method, our clustering algorithm
 256 uses a predefined number of centroids, or in other words, a predefined number

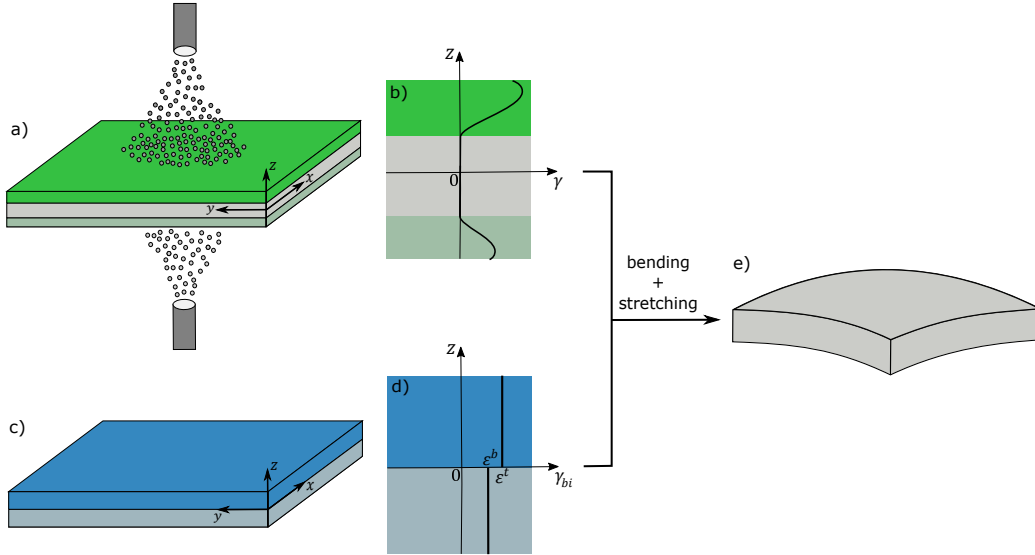


Figure 2. Idealization of the through-thickness eigenstrain profile. In this example, we suppose that the plate is uniformly peened from both sides and that the top side peening regime is more intense. a) The treatment induces eigenstrains in the outer layers of the plate (bright green and pale green), while the internal layer (gray) is not affected. b) The induced eigenstrain prescribes an in-plane expansion to the outer layers, and the expansion magnitude γ is non-uniform along the through-thickness lagrangian coordinate z . c) To idealize the eigenstrain profile, we represent the plate as a bilayer consisting of equally thick layers (bright blue and pale blue). d) We prescribe a uniform eigenstrain to both layers and thus obtain an idealized through-thickness eigenstrain profile γ_{bi} , which is described with two variables: $(\varepsilon^t, \varepsilon^b)$. The values of ε^t and ε^b are computed in the way that γ_{bi} induces the same final shape as $\gamma(z)$ (see Eqns. 2, 3), in the sense of plate theory. e) The final shape induced by γ_{bi} and $\gamma(z)$ is stretched with respect to the initial state and convex towards the side where a more intense treatment was applied.

of peening regimes. When the points are clustered, the algorithm assigns each finite element with the eigenstrains corresponding to the centroid of the cluster containing this element. This strategy allows, in particular, to constrain the maximal eigenstrain in the pattern by imposing limits on the coordinates of the centroids. The cluster centroids are related to the peening regimes, so the clustering algorithm outputs top- and bottom-side peening patterns formulated in terms of peening regimes.

The **k**-means clustering is an iterative process. On each iteration, it attributes the points to the nearest centroids in terms of squared Euclidean distance and then recomputes the centroid positions as the mean points of the corresponding clusters. The iterations stop when all centroids no longer change their positions. The result of clustering is dependent on the initial guess of the centroid positions, which are assigned randomly (Rokach & Maimon, 2005). For this reason, the algorithm must be relaunched several times with different initial centroid positions. Each trial, in turn, comprises several iterations. At the end of each trial, the algorithm evaluates the quality of clustering by computing the sum of squared Euclidean distances between the points and the centroids of clusters to which the points are assigned. The trial providing the lowest sum of distances is chosen as the best clustering. The number of trials is chosen the balance between the computational cost and the desired accuracy. For example, performing 100 trials of the **k**-means algorithm allows to decrease the clustering error by 60% with respect to a single trial on average (Fränti & Sieranoja, 2019). However, the number of all possible partitions of a given set of points is finite, so, if the algorithm is guaranteed to try all partitions in a certain number of trials, then further increasing of this number does not bring any positive effect.

3. Methodology

3.1. The clustering algorithm

We characterize a peening regime with a couple $(\varepsilon^{(1)}, \varepsilon^{(2)})$, where, by convention, $\varepsilon^{(1)}$ is positive and $\varepsilon^{(2)}$ is negative. In other words, we describe the forming effect of a treatment using the bilayer model, where the treated layer expands by $\varepsilon^{(1)}$ and the other layer contracts by $\varepsilon^{(2)}$. Hence, if a plate is treated from the top side with regime i inducing $(\varepsilon_i^{(1)}, \varepsilon_i^{(2)})$ and from the bottom side with regime j inducing $(\varepsilon_j^{(1)}, \varepsilon_j^{(2)})$, then their combination results

Table 1

Formation of the cluster centroids. Each centroid represents a combination of peening regimes applied from the top- and bottom-sides. N available peening regimes form $(N+1)^2$ centroids because the absence of treatment is considered as an additional void regime numbered as 0.

Centroid	Peening regime (top)	Peening regime (bottom)
00	0	0
01	0	1
02	0	2
$\vdots$	$\vdots$	$\vdots$
ij	i	j
$\vdots$	$\vdots$	$\vdots$
NN	N	N

291 in $(\varepsilon_i^{(1)} + \varepsilon_j^{(2)}, \varepsilon_i^{(2)} + \varepsilon_j^{(1)})$. An unclustered pattern consists of such eigen-
 292 strain combinations assigned to each element, but the aim of the clustering
 293 algorithm is to find the best suitable regimes. Consequently, the clustering
 294 algorithm must split the combinations into contributions made by the top
 295 and bottom peening regimes independently.

296 The clustering algorithm starts by initializing the points that are to be clus-
 297 tered on the coordinate plane $(\varepsilon^t, \varepsilon^b)$. Each finite element e generates one
 298 point with coordinates $(\varepsilon_e^t, \varepsilon_e^b)$, where $(\varepsilon_e^t, \varepsilon_e^b)$ is the eigenstrain prescribed to
 299 this element by the inverse problem solver. Therefore, the elements and the
 300 points are directly associated. Subsequently, the clustering algorithm initial-
 301 izes the *void regime* $(0, 0)$ corresponding to the absence of treatment, and it
 302 also randomly initializes N peening regimes in terms of $(\varepsilon^{(1)}, \varepsilon^{(2)})$ under the
 303 following constraints:

$$\begin{cases} 0 < \varepsilon^{(1)} < \varepsilon_{max}^{(1)}, \\ \varepsilon_{min}^{(2)} < \varepsilon^{(2)} < 0, \end{cases} \quad (4)$$

304 where $\varepsilon_{max}^{(1)}$ and $\varepsilon_{min}^{(2)}$ are, correspondingly, the maximal positive and the
 305 minimal negative eigenstrains achievable with the peening equipment. These
 306 values must be experimentally characterized in advance.

307 Afterwards, the clustering algorithm initializes the cluster centroids, that

are computed as all possible combinations of the peening regimes. Hence, there are $\mathbf{k} = (N + 1)^2$ centroids initialized, and each of them represents a cluster. A centroid formed by regime i applied from the top side and by regime j applied from the bottom side is marked as ij (Table 1). The void regime is numbered as 0, so the centroids marked as $i0$ and $0i$ correspond to treatment with regime i only from the top side and only from the bottom side, respectively. Figure 3 a) shows the points and the centroids initialized on the plane.

The next stage is the deletion of all centroids that lie above the line $\varepsilon^t = \varepsilon^b$. Indeed, the centroids ij and ji are induced by the same peening regimes applied conversely, so they are symmetrical about the line $\varepsilon^t = \varepsilon^b$. However, the algorithm aims at optimizing the peening regimes, and the range of regimes is the same for both sides of the plate. Consequently, the algorithm deletes one centroid from each pair of symmetrical centroids.

The symmetry principle also applies to the points corresponding to the elements. Thus, the eigenstrains $(\varepsilon_e^t, \varepsilon_e^b)$ and $(\varepsilon_e^b, \varepsilon_e^t)$ represent the same treatment combination applied from different sides. However, the points cannot be deleted like the centroids because all points contain information on the given pattern. Accordingly, the clustering algorithm reflects all points that lie above the line $\varepsilon^t = \varepsilon^b$ across this line instead of deletion. With this, the points lying on either side of the line $\varepsilon^t = \varepsilon^b$ are fused and clustered simultaneously. Figure 3 b) traces the reflected points and the preserved centroids.

The iterative process begins next. Similarly as during the standard k-means clustering, each point, i.e., element, is assigned to the cluster represented by the closest centroid, which is shown in Figure 3 c). However, the centroids cannot be relocated freely for every iteration because they are coupled. Indeed, the relocation of centroid ij means altering the eigenstrains induced by both regimes i and j . This, in turn, induces a simultaneous relocation of all centroids formed by regimes i and j in combination with another regime. For this reason, the clustering algorithm does not relocate centroids but adjusts the peening regimes directly. The new centroid positions are then computed as a function of the new peening regimes.

The peening regimes are adjusted using the notion of *contributions*. Let us consider a point $(\varepsilon_e^t, \varepsilon_e^b)$ generated by element e , which is assigned to cluster ij on the current iteration. The centroid of this cluster represents the combination of treatments with regimes i and j . In turn, point e also

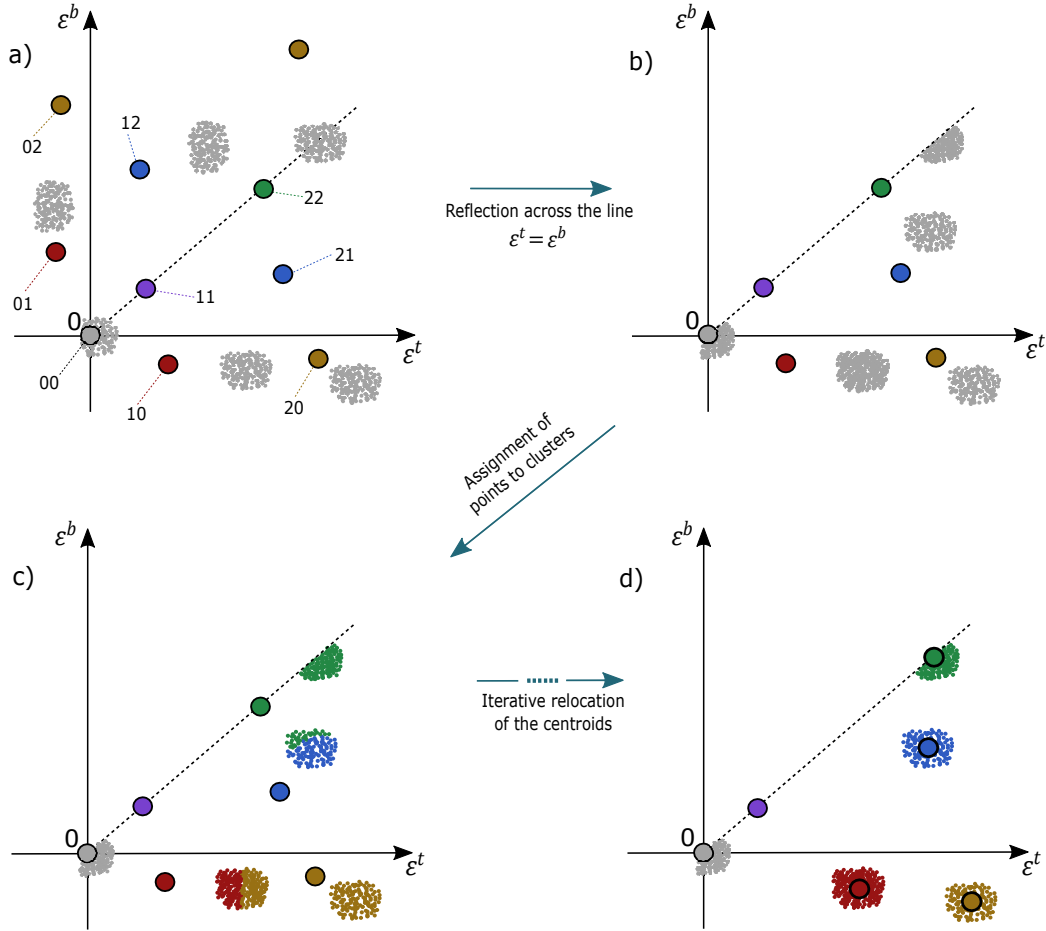


Figure 3. Example of clustering with $N = 2$ available peening regimes.

a) At the beginning, the clustering algorithm traces the points (gray dots) that are to be clustered at coordinates (ϵ^t, ϵ^b) . Each point represents a finite element, and its coordinates reflect the eigenstrain assigned to the element by the inverse problem solver. The clustering algorithm randomly initializes the peening regimes and traces the cluster centroids (colored squares), each of which is a combination of two peening regimes for all possible permutations. The coordinates of centroid ij represent the eigenstrain induced by the application of regime i from the top side and of regime j from the bottom side.

b) Next, the algorithm eliminates the symmetry by reflecting the points that lie above the line $\epsilon^t = \epsilon^b$ across this line. The centroids that lie above this line are deleted.

c) Subsequently, each point is assigned to the closest centroid (cluster) in terms of squared Euclidean distance.

d) The algorithm iteratively adjusts the peening regimes, derives new centroid positions and reassigns the points until the stabilization of the centroids. The centroid $(0, 0)$ is fixed at the origin during all iterations. In this example, no points were attributed to cluster 11 (purple), which corresponds to the treatment with regime 1 from both sides.

345 represents a combination of treatments, but the contributions made by the
 346 top and bottom peening are not defined. However, the clustering algorithm
 347 derives these contributions, which have coordinates $(\varepsilon_{e,i}^{(1)}, \varepsilon_{e,i}^{(2)})$ and $(\varepsilon_{e,j}^{(1)}, \varepsilon_{e,j}^{(2)})$,
 348 and uses them to adjust regimes i and j , respectively. The contributions are
 349 derived in the way that they are as close as possible to the eigenstrains
 350 induced by regimes i and j .

351 We formulate the expression for the coordinates of the contributions using the
 352 method of Lagrange multipliers (Hoffmann et al., (2004)). These coordinates
 353 must minimize the sum of squared Euclidean distances f to the regimes i
 354 and j :

$$f = (\varepsilon_{e,i}^{(1)} - \varepsilon_i^{(1)})^2 + (\varepsilon_{e,i}^{(2)} - \varepsilon_i^{(2)})^2 + (\varepsilon_{e,j}^{(1)} - \varepsilon_j^{(1)})^2 + (\varepsilon_{e,j}^{(2)} - \varepsilon_j^{(2)})^2. \quad (5)$$

355 In addition, the combination of contributions $(\varepsilon_{e,i}^{(1)}, \varepsilon_{e,i}^{(2)})$ and $(\varepsilon_{e,j}^{(1)}, \varepsilon_{e,j}^{(2)})$ must
 356 equal $(\varepsilon_e^t, \varepsilon_e^b)$, which leads to:

$$\begin{cases} \varepsilon_{e,i}^{(1)} + \varepsilon_{e,j}^{(2)} = \varepsilon_e^t, \\ \varepsilon_{e,i}^{(2)} + \varepsilon_{e,j}^{(1)} = \varepsilon_e^b. \end{cases} \quad (6)$$

357 With this, the Lagrangian function $\mathcal{L}$ takes the form:

$$\begin{aligned} \mathcal{L} = & (\varepsilon_{e,i}^{(1)} - \varepsilon_i^{(1)})^2 + (\varepsilon_{e,i}^{(2)} - \varepsilon_i^{(2)})^2 + (\varepsilon_{e,j}^{(1)} - \varepsilon_j^{(1)})^2 + (\varepsilon_{e,j}^{(2)} - \varepsilon_j^{(2)})^2 \\ & - \lambda_1 (\varepsilon_{e,i}^{(1)} + \varepsilon_{e,j}^{(2)} - \varepsilon_e^t) - \lambda_2 (\varepsilon_{e,i}^{(2)} + \varepsilon_{e,j}^{(1)} - \varepsilon_e^b). \end{aligned} \quad (7)$$

358 The stationary point of the Lagrangian function is:

$$\begin{cases} \varepsilon_{e,i}^{(1)} = 0.5 (\varepsilon_i^{(1)} - \varepsilon_j^{(2)} + \varepsilon_e^t), \\ \varepsilon_{e,i}^{(2)} = 0.5 (\varepsilon_i^{(2)} - \varepsilon_j^{(1)} + \varepsilon_e^b), \\ \varepsilon_{e,j}^{(1)} = 0.5 (\varepsilon_j^{(1)} - \varepsilon_i^{(2)} + \varepsilon_e^b), \\ \varepsilon_{e,j}^{(2)} = 0.5 (\varepsilon_j^{(2)} - \varepsilon_i^{(1)} + \varepsilon_e^t), \end{cases} \quad (8)$$

359

$$\begin{cases} \lambda_1 = \varepsilon_e^t - \varepsilon_i^{(1)} - \varepsilon_j^{(2)}, \\ \lambda_2 = \varepsilon_e^b - \varepsilon_j^{(1)} - \varepsilon_i^{(2)}. \end{cases} \quad (9)$$

360 The two contributions computed according to Eqns. (8) minimize the sum of
 361 squared Euclidean distances (5) and satisfy conditions (6), simultaneously.
 362 Let $\mathcal{T}_i$ be the set of points assigned to all the centroids formed using regime
 363 i on the current iteration. The algorithm computes the new eigenstrain in-
 364 duced by regime i as the mean of all contributions $(\varepsilon_{e,i}^{(1)}, \varepsilon_{e,i}^{(2)})$ for $e \in \mathcal{T}_i$. The
 365 void regime is, however, fixed at point $(0, 0)$. When all regimes are adjusted,
 366 the algorithm recalculates the centroids and reassigns the points to the clus-
 367 ters. Same as with the standard k -means algorithm, this iterative process is
 368 repeated until stabilization of the peening regimes and, consequently, of the
 369 centroids. The clustering algorithm is summarized as Algorithm 1. Figure 3
 370 d) illustrates the result of clustering.

Algorithm 1 The clustering algorithm

```
1: for each finite element  $e$  do
2:   Initialize one point with coordinates  $(\varepsilon_e^t, \varepsilon_e^b)$ 
3: end for
4: Initialize the void peening regime  $(0, 0)$ 
5: Randomly initialize  $N$  peening regimes in terms of  $(\varepsilon^{(1)}, \varepsilon^{(2)})$  under constraints (4)
6: Compute the centroids in coordinates  $(\varepsilon^t, \varepsilon^b)$  as all possible combinations of two peening regimes
7: for each centroid that lies above the line  $\varepsilon^t = \varepsilon^b$  do
8:   Delete the centroid
9: end for
10: for each point that lies above the line  $\varepsilon^t = \varepsilon^b$  do
11:   Swap its coordinates  $\varepsilon^t$  and  $\varepsilon^b$ 
12: end for
13: while at least one centroid changes its position do
14:   for each point do
15:     Compute the squared Euclidean distance to each centroid
16:     Assign the point to the cluster formed by the closest centroid
17:     Compute the two contributions using Eqns. (8)
18:   end for
19:   for each regime except for the void regime do
20:     Compute the mean of all contributions belonging to this regime
21:     Assign the mean as the new peening regime
22:   end for
23:   Compute the centroids as the combinations of the adjusted peening regimes
24: end while
25: for each finite element  $e$  do
26:   Assign the element with the eigenstrain corresponding to the centroid of the cluster containing point  $(\varepsilon_e^t, \varepsilon_e^b)$ 
27: end for
```

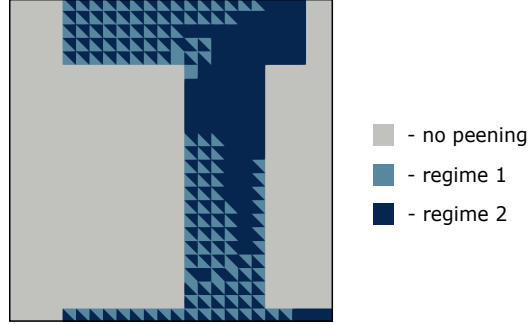


Figure 4. Example of a peening pattern heavily affected by the checkerboard problem after clustering. The pattern is represented on a square plate, which is meshed with triangular finite elements. The elements assigned with different peening regimes of a close intensity are locally mixed, so the proposed pattern is not reproducible in practice.

3.2. Filtering of the peening pattern using cellular automata

The clustering algorithm groups the finite elements based on the eigenstrain prescribed during the inverse problem resolution. After clustering, each finite element in the peening pattern is assigned with a peening regime. The elements assigned with the same regime can be physically situated in different parts of the plate and thus form separate segments. In practice, however, the minimal size of the segments is limited by the shot stream width or the masking resolution. For example, a segment consisting of only one element is not typically reproducible in practice. The presence of such small-scale segments in the clustered pattern is called the *checkerboard* problem, which is illustrated in Figure 4. This problem arises when the eigenstrain prescribed by the inverse problem solver is on the border between two clusters, so the elements assigned with different peening regimes are mixed together after clustering.

We cope with the checkerboard problem with a filtering algorithm, which is executed after clustering for the top and bottom patterns separately. More precisely, we use an image noise filter based on cellular automata (CA) described by [Selvapeter & Hordijk \(2009\)](#). The filter is initially conceived for images consisting of square pixels that are assigned with a gray level, but we reformulate it for the case of triangular meshes where each element is assigned with a peening regime. The filtering algorithm scans the elements and notes the regimes assigned to each element and to its three neighbors, i.e., to its triangular Von Neumann neighborhood ([Zawidzki, 2011](#)). If the algo-

394 rithm detects that the element is mostly surrounded by elements prescribed
 395 with another regime, then it assigns the element with the regime that con-
 396 stitutes the majority. In case when none of the peening regimes constitutes
 397 a majority in the local neighborhood, then the element is assigned with the
 398 regime corresponding to its first neighbor. The same principle applies to
 399 the edge elements that have only two neighbors. The corner elements are
 400 strictly assigned with the regime corresponding to their single neighbor. The
 401 filtering algorithm is iterative and lasts until the stabilization of the assigned
 402 regimes. All triangular elements are scanned and corrected simultaneously
 403 at each iteration. This filter can also be applied after the grouping of the
 404 peening pattern (Sushitskii et al., 2022b), because the grouped patterns can
 405 also face the checkerboard problem.
 406 The filtering algorithm is summarized as Algorithm 2. We formulate the
 407 algorithm in terms of the peening regime ρ_e assigned to element e . The
 408 regimes assigned to the neighboring elements are denoted as ρ_{e1} , ρ_{e2} , ρ_{e3} .

409 4. Results

410 The clustering and the filtering algorithms were implemented in C++ pro-
 411 gramming language with the help of “Eigen” (Guennebaud et al., 2018) li-
 412 brary for data operations and “libigl” (Jacobson et al., 2018) library for tri-
 413 angular mesh handling. The algorithms were validated numerically using 200
 414 random target shapes. For each test case, we firstly solved the inverse prob-
 415 lem using the algorithm presented by Sushitskii et al. (2022b) and obtained
 416 the *free* peening pattern. Secondly, we applied the clustering algorithm (Al-
 417 gorithm 1) and thus obtained the *clustered* peening pattern. Thirdly, we
 418 applied the filtering algorithm (Algorithm 2) and obtained the *filtered clus-*
 419 *tered* peening pattern. Finally, we simulated the application of each of these
 420 three patterns using the method presented by Sushitskii et al. (2022b) to
 421 evaluate discrepancy between this and that. In other words, we solved the
 422 forward problem for each pattern and thus obtained three final shapes. We
 423 computed the dimensionless error Ω in each case as the Hausdorff distance
 424 between the final shape and the target shape divided by the square root of
 425 the total area of the plate.

426 The target shapes were generated as a result of the application of random
 427 peening patterns on a 1×1 m square plate. The plate thickness and the
 428 Poisson’s ratio were randomly assigned in each test case. The plates were

Algorithm 2 The filtering algorithm

```
1: while the eigenstrain is reassigned at least for one element do
2:   for each finite element  $e$  do
3:     if the element has three neighbors  $e_1, e_2, e_3$ , then
4:       if  $\rho_e \neq \rho_{e1}$  and  $(\rho_{e1} = \rho_{e2}$  or  $\rho_{e1} = \rho_{e3})$  then
5:         Assign the value of  $\rho_{e1}$  to  $\rho_e$ 
6:       else if  $\rho_e \neq \rho_{e2}$  and  $\rho_{e2} = \rho_{e3}$  then
7:         Assign the value of  $\rho_{e2}$  to  $\rho_e$ 
8:       else if
9:          $(\rho_e \neq \rho_{e1}$  and  $\rho_e \neq \rho_{e2}$  and  $\rho_e \neq \rho_{e3})$  and
10:         $(\rho_{e1} \neq \rho_{e2}$  and  $\rho_{e1} \neq \rho_{e3}$  and  $\rho_{e2} \neq \rho_{e3})$  then
11:          Assign the value of  $\rho_{e1}$  to  $\rho_e$ 
12:        end if
13:      end if
14:      if the element has two neighbors  $e_1, e_2$ , then
15:        if  $\rho_e \neq \rho_{e1}$  and  $\rho_e \neq \rho_{e2}$  then
16:          Assign the value of  $\rho_{e1}$  to  $\rho_e$ 
17:        end if
18:      end if
19:      if the element has one neighbor  $e_1$ , then
20:        if  $\rho_e \neq \rho_{e1}$  then
21:          Assign the value of  $\rho_{e1}$  to  $\rho_e$ 
22:        end if
23:      end if
24:    end for
25: end while
```

429 meshed with 1152 triangular elements, and the random peening patterns
 430 were generated on this mesh using the algorithm presented by [Sushitskii](#)
 431 [et al. \(2022b\)](#). The first 100 test cases (series 1) involved only one available
 432 peening regime, which was applied in randomly situated rectangular segments
 433 of the plate. The second 100 cases (series 2) involved four available regimes
 434 applied in random segments.

435 Figure [B.8](#) illustrates the numerical validation workflow using one test case
 436 from series 2. The presented clustered pattern (Figure [B.8 d](#)) leads to the
 437 highest Ω in series 2 because the pattern is significantly affected by the
 438 checkerboard problem. The filtered clustered pattern (Figure [B.8 e](#)) contains
 439 irregularly shaped peening segments, but it is not subjected to the checker-
 440 board problem, so it is applicable in practice.

441 Figure [6](#) summarizes the validation results in terms of Ω . The histograms
 442 demonstrate that the dimensionless error Ω did not exceed 0.4% for the final
 443 shapes provided by the free patterns. Both clustering and filtering algorithms
 444 made a slightly negative impact in terms of Ω , but Ω stayed in the same range
 445 of 0.4% for all the test cases. Moreover, Ω remained inferior to 0.1% for more
 446 than 80% of cases in series 1 and for more than 70% of cases in series 2.

447 During the clustering phase, we fixed the number of available regimes as one
 448 for series 1 and as four for series 2. No other information on the peening
 449 regimes was provided to the clustering algorithm, and the centroid positions
 450 were not constrained. The number of trials for the clustering algorithm was
 451 fixed as 100. The influence of clustering in terms of Ω is plotted in Figure [7](#)
 452 a). Thus, the clustered pattern decreased Ω with respect to the free pattern
 453 for 78 cases in series 1 and for 32 cases in series 2. The decrease in Ω in these
 454 cases is explained by the segmented structure of the random patterns that
 455 were used to generate the target shapes. Hence, the clustering removed the
 456 gradual eigenstrain variation from the free patterns and made the clustered
 457 patterns more similar to the original ones.

458 On the other hand, the difference in the influence of clustering on the two
 459 series is explained by the fact the clustered patterns in series 1 were less
 460 subjected to the checkerboard problem. Indeed, the only available peening
 461 regime was situated far from the origin in terms of $(\varepsilon^t, \varepsilon^b)$, so the majority
 462 of elements was assigned to the correct clusters. The main source of error in
 463 this case was the peening regime itself, which was slightly different from the
 464 peening regime applied in the original random pattern. On the other hand,
 465 the regimes in series 2 were situated close to each other, so the quantity of

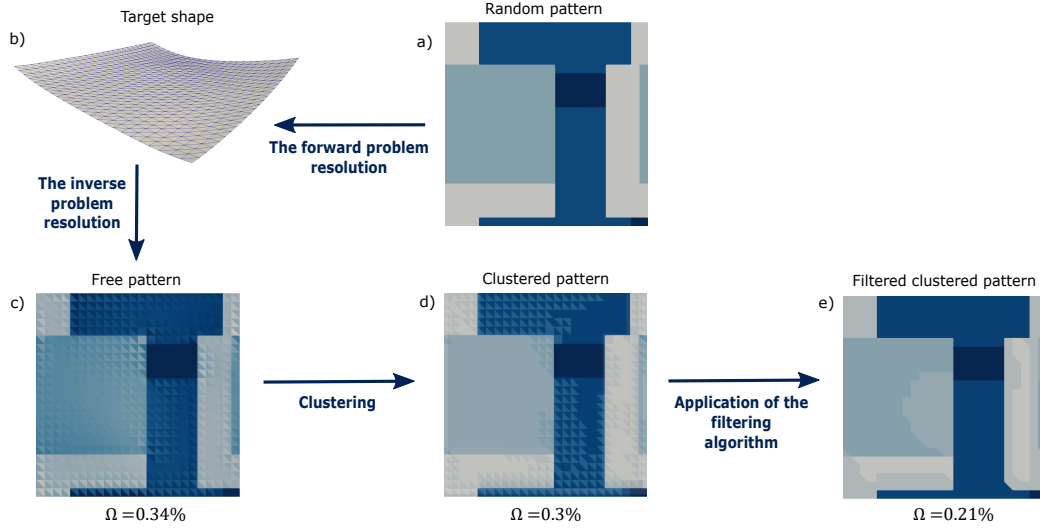


Figure 5. One of the test cases from series 2 that demonstrated a marked checkerboard problem. a) The random peening pattern generated on a 1×1 m sized plate using the algorithm presented by [Sushitskii et al. \(2022b\)](#). Different colors on the peening pattern denote different peening regimes. b) The target shape generated by application of the random pattern. The plate is 6 mm thick, and the Poisson's ratio equals 0.36. The deflections are at the original scale. c) The free pattern computed using the inverse problem resolution algorithm (see [Sushitskii et al., 2022b](#)). d) The clustered peening pattern computed by application of Algorithm 1 to the free pattern. e) The filtered clustered algorithm obtained by application of Algorithm 2 to the clustered pattern. We solved the forward problem for the patterns c), d) and e) and computed the dimensionless error Ω for each case. In the presented case, the filtered clustered pattern reproduces the original random pattern better than the free pattern, so it leads to a lower dimensionless error Ω .

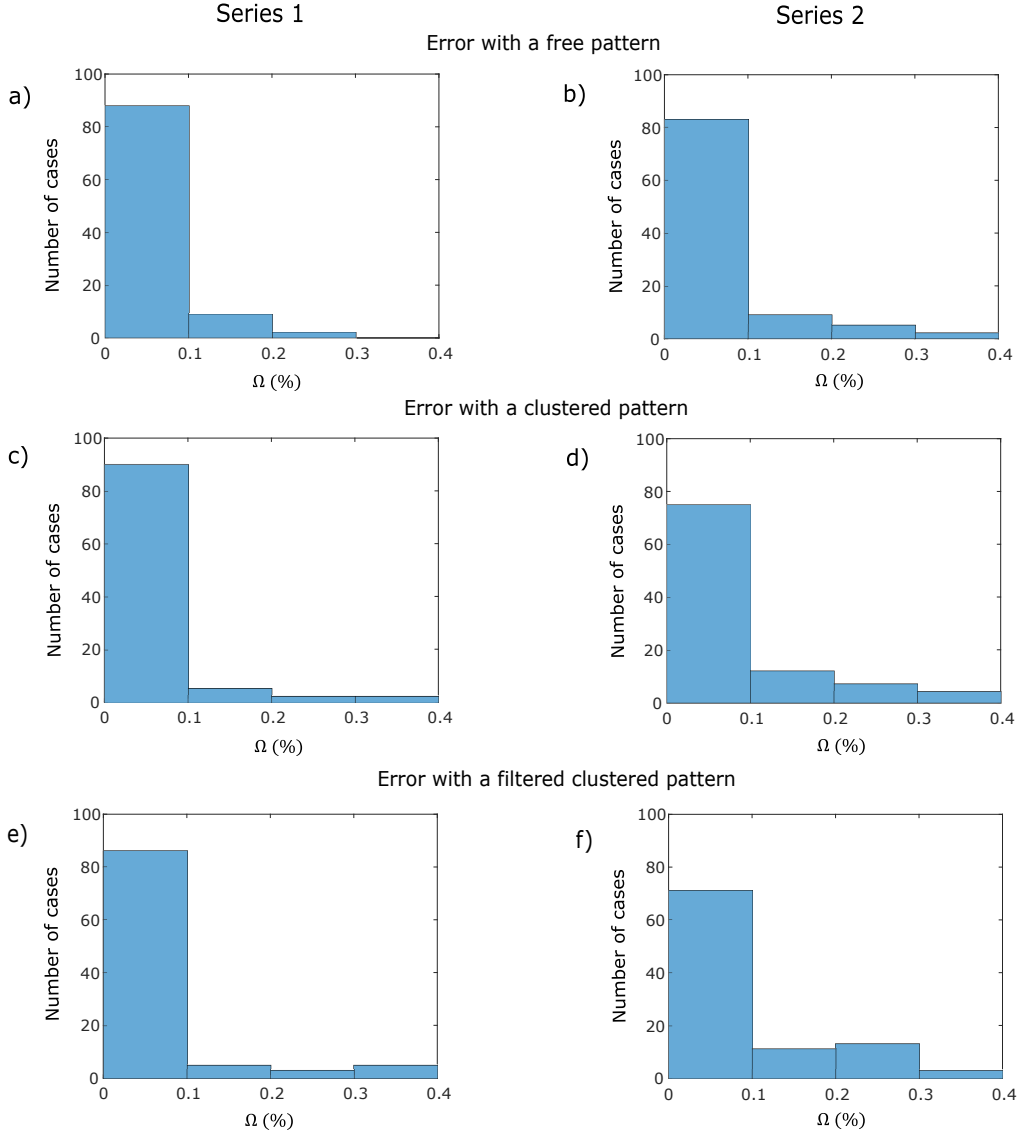


Figure 6. Histograms plotting the dimensionless error Ω between the target shapes and the final shapes computed during the validation of the clustering and filtering algorithms. We considered 200 test cases that were divided in two series of 100 cases each. In each case, we solved the inverse problem, clustered the peening pattern and applied the filtering algorithm. One peening regime was available for clustering in series 1, and four regimes were available in series 2. We simulated the application of the peening patterns at each stage and computed the dimensionless error Ω . Figures (a) and (b) plot the simulated Ω provided by the unclustered free patterns in series 1 and 2 respectively. Figures (c) and (d) trace the simulated Ω provided by the clustered patterns, and Figures (e) and (f) show Ω provided by the filtered clustered patterns. The filtered clustered patterns induce a slightly larger Ω than the free patterns in general, but Ω does not exceed 0.4% for all the types of patterns.

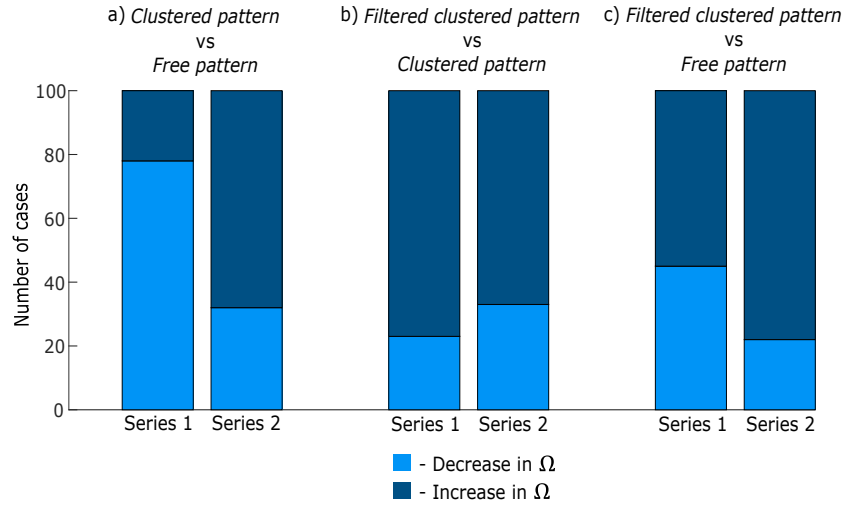


Figure 7. Influence of the clustering and the filtering algorithms on the dimensionless error Ω . The clustered patterns were obtained by application of the clustering algorithm to the free patterns, and the filtered clustered patterns were obtained by application of the filtering algorithm to the clustered patterns. Hence, the bars a) represent the influence of the clustering algorithm, the bars b) represent the influence of the filtering algorithm, and the bars c) represent the influence of the successive application of the two algorithms. The first series of tests implied one peening regime during generation of the training examples and clustering, while the second series implied four peening regimes. Both series consisted of 100 test cases. The results show that the influence of the successive application of the two algorithms is majorly negative.

466 neighboring elements that were assigned to different clusters was more signif-
 467 icant. Also, the four computed peening regimes did not exactly correspond
 468 to the original regimes.

469 The filtered clustered pattern increased Ω with respect to the clustered pat-
 470 tern for 77 cases in series 1 and for 67 cases in series 2, which is shown in
 471 Figure 7 b). The reason for this increase in most of the cases is that the
 472 filtering algorithm often enlarges the segments assigned to wrong clusters
 473 instead of eliminating them.

474 All in all, the successive application of the clustering and the filtering algo-
 475 rithms decreased Ω with respect to the free pattern for 67 cases out of 200
 476 (see Figure 7 c). The increase in Ω for 55 cases in series 1 and for 78 cases
 477 in series 2 is explained by the fact that the segmentation strategy does not
 478 account for Ω , so its output depends only on the input pattern.

479 5. Discussion

480 In our experiments, the quantity of peening regimes N was fixed for each
 481 series of tests. However, in practice N is not always pre-determined. In a
 482 such case, the clustering algorithm can be launched several times for different
 483 N , and the best suitable N can be determined based on the minimal Ω . A
 484 larger N provides more flexibility in terms of target shapes but extends the
 485 time needed for application of the pattern in practice. Here, more practical
 486 considerations will need to be taken into account by engineers using the
 487 method such as what kind of treatments are available and how much time
 488 can be allotted to the treatment.

489 Both the clustering and the filtering algorithms can either increase or decrease
 490 the Ω , which introduces uncertainty in the solution. One potential way to
 491 mitigate this problem is to implement a black-box optimization algorithm,
 492 which would optimize the eigenstrains prescribed to each peening regime after
 493 the filtering in order to reduce Ω . However, this approach would significantly
 494 increase the computational burden.

495 In practice, the actual peening parameters can be related to the eigenstrains
 496 constituting the segmented pattern using experiments in combination with
 497 the eigenstrain simulations. Thus, for each couple $(\varepsilon^{(1)}, \varepsilon^{(2)})$, one can numer-
 498 ically simulate a uniform treatment of a dummy specimen, such as an Almen
 499 strip, and calculate the resulting shape of the specimen. Next, the peening

parameters can be adjusted in order to achieve the computed shape in practice. Furthermore, one can establish a continuous relationship between one of the peening parameters, such as the peening pressure, and the induced eigenstrains. However, a precise workflow for establishing this relationship is beyond the scope of this paper.

6. Conclusion

We developed a general strategy for the segmentation of the peening pattern, which makes every pattern applicable with shot peening equipment. This step is essential. Without this clustering algorithm, the continuous pattern that comes out of any shot peening optimization method cannot be applied directly. Our method allows discretizing the pattern into a finite number of achievable treatment segments. The segmentation strategy is an essential step towards a fully automated forming process, which is more cost-efficient and repeatable than the manual operation. The segmentation strategy computes the best suitable peening regimes, and the segments where they should be applied. It includes a clustering algorithm, which is inspired by the k-means method, and a filtering algorithm. The clustering algorithm is designed for the bilayer eigenstrain formulation, and any other formulation of the trough-thickness eigenstrain profile can be reduced to the bilayer. When applied consecutively, the clustering and the filtering algorithms remove from the pattern all eigenstrain variations that are not reproducible in practice. Hence, the strategy provides a ready-to use peening pattern, which is suitable for industrial reproduction using masking or a narrow peening stream.

The clustering and the filtering algorithms were validated numerically using the existing bilayer shell model. The validation showed that the clustering algorithm had a weak influence on the quality of the peening pattern, so that the original and the clustered patterns induced similar final shapes. Out of 200 tested cases, the error due to the segmentation, quantified with the normalised Hausdorff distance, did not attain more than 0.4% for any tested pattern. A clustered pattern involving one peening regime is less affected by the checkerboard problem than this involving multiple regimes, but a large number of regimes provides more flexibility in terms of target shapes. The filtering also has a slight but majorly negative influence on the final shape. Nevertheless, it efficiently deals with the checkerboard problem and rapidly

535 eliminates local clustering errors.

536 The described segmentation strategy can be applied as a post-processing
537 tool for eigenstrain-based inverse problem solvers that deal with laminated
538 shape-shifting structures. Such structures are present in the fields of Micro-
539 Electro-Mechanical Systems (MEMS) or 4D printed composites. Moreover,
540 metal plates shaped with laser peening or English wheel can also be virtually
541 represented as laminates and modeled using the eigenstrain approach. The
542 segmentation strategy eliminates the need for trial-and-error determination
543 of the best suitable eigenstrains and thus accelerates the inverse problem
544 resolution.

545 The advantage of the proposed segmentation strategy in comparison to the
546 existing inverse problem solvers is the automated determination of the opti-
547 mal peening regimes. Moreover, the filtering strategy ensures that the treated
548 segments contain more than one finite element, which was not yet encoun-
549 tered in the peen forming literature. In addition, the bilayer formulation
550 applies restriction on the cluster centroids in terms of the k-means method,
551 and this problem was overcome using the method of Lagrange multipliers.
552 Hence, the novel modified k-means method can be of interest for other types
553 of clustering problems not related to peen forming.

554 One limitation of the filtering strategy is the dependence of the minimal size
555 of the treated segment on the size of elements. Hence, the smallest treated
556 segment can still be narrower than the shot stream in the case of a dense
557 mesh, which can be mitigated by narrowing the peening nozzle. In addition,
558 the proposed segmentation strategy is limited to isotropic eigenstrain models,
559 because the ability to vary the principal growth direction from one element
560 to another is not yet included in the algorithms.

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583 10. Author contribution

- 584 • **Vladislav Sushitskii (corresponding author):** Conceptualization,
585 Methodology, Software, Validation, Formal analysis, Investigation, Writ-
586 ing - Original Draft, Writing - Review & Editing.
- 587 • **Hong Yan Miao:** Resources, Writing - Review & Editing, Supervi-
588 sion.
- 589 • **Martin Lévesque:** Conceptualization, Methodology, Writing - Re-
590 view & Editing, Supervision, Project administration, Funding acquisi-
591 tion.
- 592 • **Frédéric Gosselin:** Conceptualization, Methodology, Validation, Re-
593 sources, Writing - Review & Editing, Supervision, Project administra-
594 tion, Funding acquisition.

595 11. Data availability statement

596 The datasets generated during and/or analysed during the current study
 597 are available from the corresponding author on reasonable request. The
 598 programming code that was used to generate the datasets is available at:
 599 https://github.com/lm2-poly/peen_forming_2022.

600 Appendix A. k-means clustering of the trilayer eigenstrain pat- 601 tern

602 Another common way to idealize the through-thickness eigenstrain profile
 603 is to represent it as a trilayer $\gamma_{tri}(z)$. The trilayer formulation implies the
 604 assignment of eigenstrain to the two outer layers of a variable thickness:

$$\gamma_{tri}(z) = \begin{cases} \varepsilon_*^t & \text{for } \left(\frac{h}{2} - h_*^t\right) < z < \frac{h}{2}, \\ 0 & \text{for } \left(h_*^b - \frac{h}{2}\right) < z < \left(\frac{h}{2} - h_*^t\right), \\ \varepsilon_*^b & \text{for } -\frac{h}{2} < z < \left(h_*^b - \frac{h}{2}\right), \end{cases} \quad (\text{A.1})$$

605 where ε_*^t and ε_*^b stand for the eigenstrains assigned to the top and bottom
 606 layers respectively, while h_*^t and h_*^b denote the thicknesses of these layers.
 607 Hence, in this formulation, the eigenstrain assigned to each finite element e
 608 is described with four variables $(\varepsilon_{*e}^t, h_{*e}^t, \varepsilon_{*e}^b, h_{*e}^b)$.

609 The range of peening regimes that can be applied from the top and bottom
 610 sides is the same, so the eigenstrains assigned to the top and bottom layers
 611 are clustered simultaneously. This means that each element generates two
 612 points on a coordinate plane with axes ε_* and h_* . The coordinates of these
 613 points are $(\varepsilon_{*e}^t, h_{*e}^t)$ and $(\varepsilon_{*e}^b, h_{*e}^b)$. In this case, the points can be clustered
 614 using the standard k-means algorithm (Lloyd, 1982), except for the fact that
 615 the zero centroid $(0,0)$ corresponding to the absence of treatment is fixed
 616 at the origin (Algorithm 3). The presence of this centroid suggests that N
 617 peening regimes actually induce $k = N + 1$ cluster centroids. The other
 618 N centroids are initialized using the *Maxmin* method, which proves itself as
 619 the most efficient initialization technique for the k-means algorithm (Fränti &
 620 Sieranoja, 2019). With this method, the centroids are initialized in sequence,
 621 and each subsequent centroid is placed at the location of the furthest point
 622 from the previous centroid. The first centroid in this sequence is the zero
 623 centroid.

Algorithm 3 The clustering algorithm in the trilayer case

```
1: for each finite element  $e$  do
2:   Initialize two points with coordinates  $(\varepsilon_{*e}^t, h_{*e}^t)$  and  $(\varepsilon_{*e}^b, h_{*e}^b)$ 
3: end for
4: Initialize the zero centroid at the origin
5: Initialize  $N$  centroids at random positions in coordinates  $(\varepsilon_*, h_*)$  using
   the Maxmin method
6: while at least one centroid changes its position do
7:   for each point do
8:     Compute the squared Euclidean distance to each centroid
9:     Assign the point to the cluster formed by the closest centroid
10:  end for
11:  for each cluster except for the zero cluster do
12:    Compute the mean of all points in the cluster
13:    Assign the mean as the new cluster centroid
14:  end for
15: end while
16: for each finite element  $e$  do
17:   Assign the top layer with a couple  $(\varepsilon_*^t, h_*^t)$  corresponding to the cen-
     troid of the cluster containing point  $(\varepsilon_{*e}^t, h_{*e}^t)$ 
18:   Assign the bottom layer with a couple  $(\varepsilon_*^b, h_*^b)$  corresponding to the
     centroid of the cluster containing point  $(\varepsilon_{*e}^b, h_{*e}^b)$ .
19: end for
```

624 **Appendix B. Additional test cases**

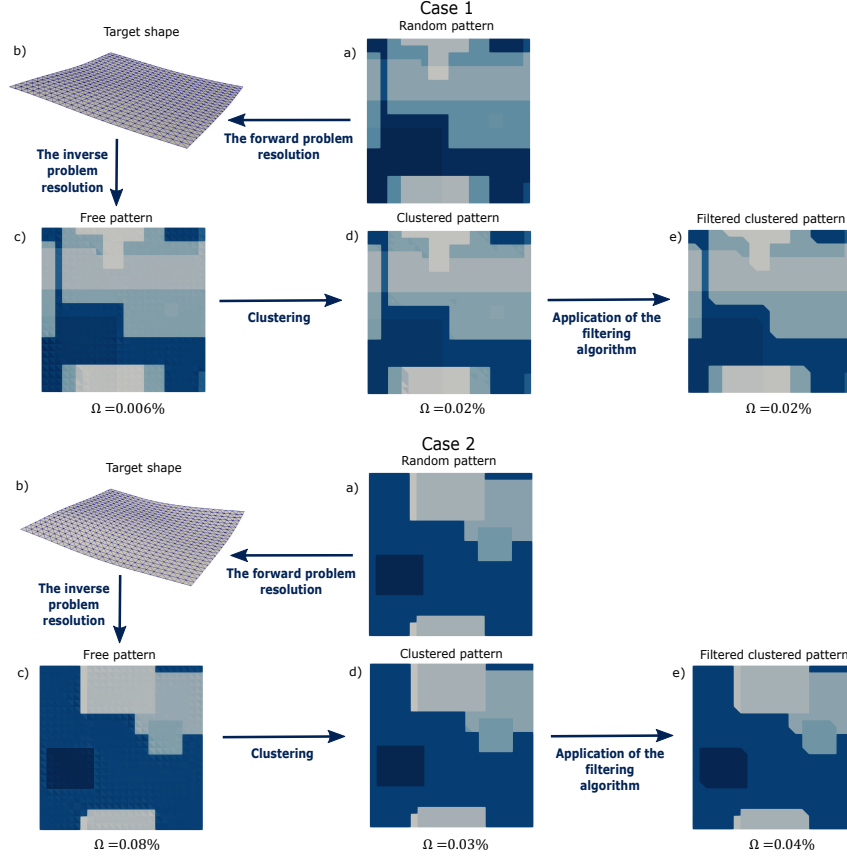


Figure B.8. Two test cases from series 2. a) The random peening pattern generated on a 1×1 m sized plate using the algorithm presented by [Sushitskii et al. \(2022b\)](#). Different colors on the peening pattern denote different peening regimes. b) The target shape generated by application of the random pattern. The deflections are at the original scale. c) The free pattern computed using the inverse problem resolution algorithm (see [Sushitskii et al., 2022b](#)). d) The clustered peening pattern computed by application of Algorithm 1 to the free pattern. e) The filtered clustered algorithm obtained by application of Algorithm 2 to the clustered pattern. We solved the forward problem for the patterns c), d) and e) and computed the dimensionless error Ω for each case. In case 1, the free pattern reproduces the original random pattern the best. This is due to several elements attributed to wrong clusters during clustering, so the dimensionless error Ω is higher for the clustered pattern. In case 2, the lowest Ω is provided by the clustered pattern, which almost exactly replicates the random pattern. Although the filtering algorithm makes the pattern fully reproducible, in case 2 the filtered clustered pattern is further from the random pattern than the clustered one.

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