
Square Packing as Algorithmic Art

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Flow of Topics

- Introduction to Algorithmic Art
- Packing Algorithms
- Square Tiling
 - Clustering and segmentation
 - Distance Transform
 - Square Fitting
- Results
- Conclusion

Objectives :

- Design a square packing algorithm for irregular bounded region.
- Create tiled representation of any coloured image using the packing algorithm with monochromatic square tiles of different fixed denomination sets.

Introduction to Algorithmic Art

- Algorithmic Art: Aesthetically appealing design considered as algorithmic art, if the features of the design are only determined by an algorithm without or negligible human interference.
- Historically manual algorithms or mathematical techniques are used to create art. For example, *Islamic geometric patterns*, *Italian Renaissance paintings* uses linear perspective and proportion.
- First computer-generated algorithmic art were created by Georg Nees, Frieder Nake, A. Michael Noll, Manfred Mohr and Vera Molnár in the early 1960s using computer driven plotter.
- With increasing accessibility of computer more mathematically intensive complex form of art works are explored by algorithmic artist, also called algorist.
- At SIGGRAPH in 1995, a panel called titled "Art and Algorithms", established identity of such algorists.
- Some popular form of visual algorithmic art are fractal art, cellular automata based art, genetic or evolutionary art.

Packing Problem

- Packing is covering a bounded region with non-overlapping elementary shapes .
- Packing problems are *optimization problems* ,where optimization criterion is fraction of area covered of the total bounded region, known as *packing density ratio*.
- Perfect Packing is *NP-complete problem*. Perfect packing algorithms are known some simple cases where container regions are regular .
- An example of perfect packing problem is tessalation, i.e perfect tiling of Euclidean plane .
- Some practical packing problem examples are stock cutting, VLSI chip design, advertisement placement, scheduling.

Square Tiling Algorithm

- Input: A coloured image, A set of denominations of square size
- Output: A set of squares each with size, position, axis alignment and colour information
- Objectives :
 - Packing Density Ratio should be high .
 - Number of square tiles used should be less.
 - The tiled image should capture significant features of the actual image
- Steps:
 - Clustering and Segmentation
 - Distance Transform
 - Square Fitting

Clustering and Segmentation

- Objective: Group the image pixels based on their colour and position similarity so that a single coloured tile may be fitted within cluster.
- Clustering is done by meanshift ,a discontinuity preserving clustering algorithm.
- Clustering is applied on a $d=p+2$ dimensional space, $p=3$ for colour images, on a joint spatial-range domain.
- Segmentation is done by separating two clusters with a line in between.
- To remove noisy spurs morphological cleaning and skeletonization is applied on the segmentation line image.

Meanshift

- This algorithm takes a radially symmetric kernel $K(x) = c_{k,d} k(\|x\|^2)$ where $c_{k,d}$ is a normalization function and $k(x)$ is monotonically decreasing function.
- Normal distribution is used as a kernel function.
- For each pixel kernel density is obtained as $f(x) = \frac{1}{nh^d} \sum_{i=1}^n K(\frac{x-x_i}{h})$
- The modes of the density function are located at the zeros of the gradient function $\nabla f(x) = 0$
- $\nabla f(x) = \frac{2c_{k,d}}{nh^{d+2}} \left[\sum_{i=1}^n g\left(\left\|\frac{x-x_i}{h}\right\|^2\right) \right] \left[\frac{\sum_{i=1}^n x_i g\left(\left\|\frac{x-x_i}{h}\right\|^2\right)}{\sum_{i=1}^n g\left(\left\|\frac{x-x_i}{h}\right\|^2\right)} - x \right]$, where $g(x) = -k'(x)$
- The second term $m_{h,g}(x) = \frac{\sum_{i=1}^n x_i g\left(\left\|\frac{x-x_i}{h}\right\|^2\right)}{\sum_{i=1}^n g\left(\left\|\frac{x-x_i}{h}\right\|^2\right)} - x$, is the mean shift
- The mean shift procedure, obtained by successive
 - computation of the mean shift vector $m_h(x_t)$,
 - translation of the window $x_{t+1} = x_t + m_h(x_t)$
- The converging is slow as the gradient decays. Hence a threshold parameter is kept.

Mean Shift : Results with different threshold

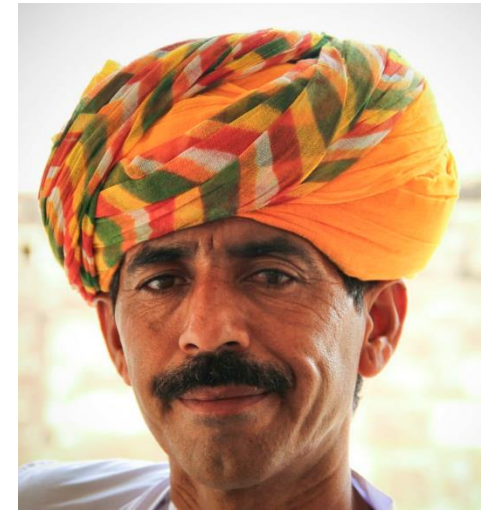
- Mean shift convergence rate decays as every pixel moves closer to its cluster mean . Hence total convergence takes too much time .
- Number of clusters increase as the threshold decreased .



Thr =0.2
Clusters=20

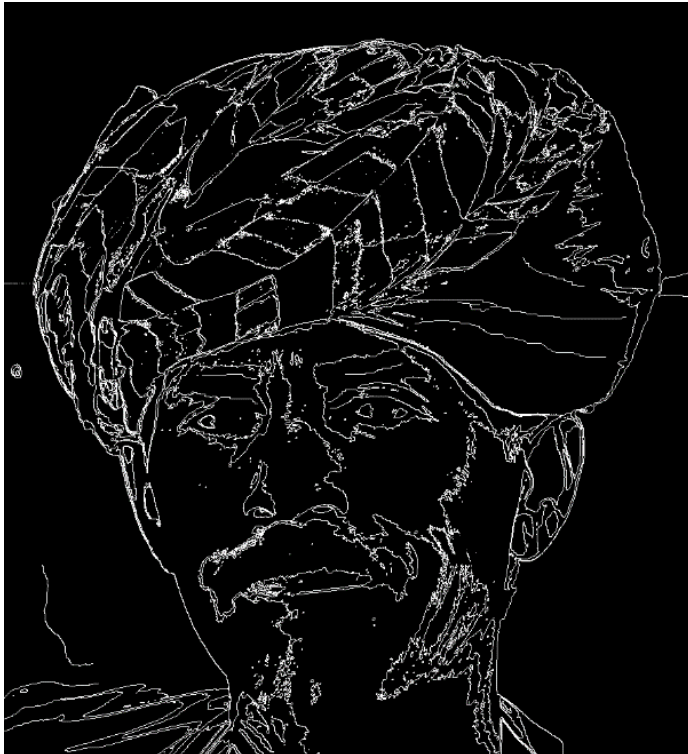


Thr =0.1
Clusters=55

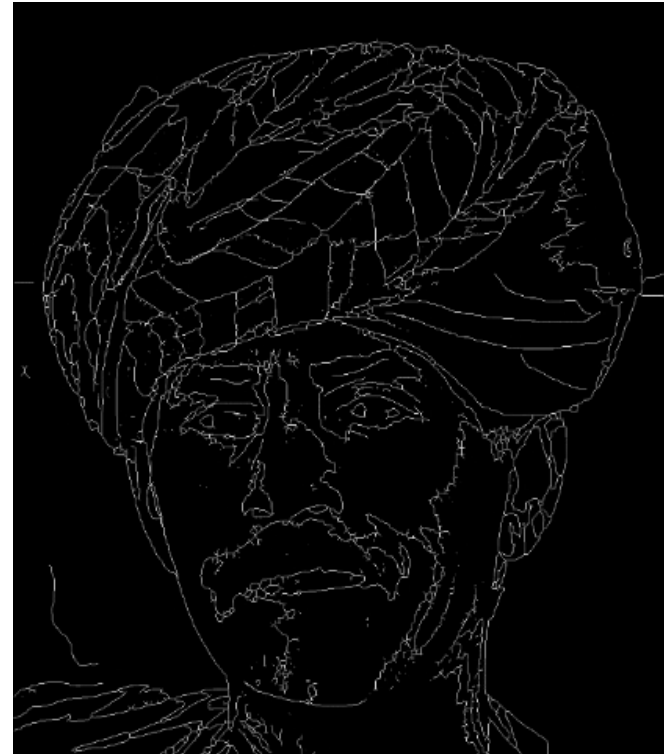


Segmentation and Morphological cleaning:

- The segmentation process gives a lot of noisy spurs, to reduce those morphological cleaning and skeletonization are done.



Segmentation Outline



Morphologically Cleaned and
skeletonized Segmentation Outline

Distance Transform:

- The distance transform map labels each pixel of a image with the nearest distance to the nearest boundary pixel or in our case segmentation outline.

- Distance Transform Metrics:

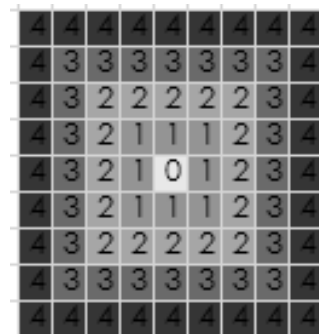
(a) *chessboard or 8 connected distance* , $d_8((i,j),(h,k)) = \max(|i-h|, |j-k|)$,

(b) *cityblock or 4 connected distance*, $d_4((i,j),(h,k)) = |i-h| + |j-k|$,

(c) *euclidean distance*, calculated as $d_e((i,j),(h,k)) = \sqrt{(i-h)^2 + (j-k)^2}$

- Distance Map contours:

d_8 mapping



d_4 mapping



d_e mapping



- This observation helps us hypothesize that d_8 metric distance calculation fits axes aligned squares better and likewise d_4 metric fits 45° rotated squares better and euclidean metric fits square of any alignment.

Distance Transform Methods:

- Distance Transform methods are of two basic types:

- (a) Parallel and (b) Sequential

- In parallel algorithms the mapping is done in a single pass starting from boundary points

$$S_0 \rightarrow L'(S_0) = S_1 \rightarrow L'(S_1) = S_2 \rightarrow \dots \rightarrow L'(S_{n-1}) = S_n = L(S).$$

- In sequential algorithms the mapping is done in multiple passes. In the n^{th} pass $L_n(i,j)$ is calculated from the previous mapping $L_{n-1}(i-1,j)$, $L_{n-1}(i,j-1)$ etc.
- Parallel algorithm is efficient in calculating *8 connected distance* and 4 connected distance and sequential distance mapping is suitable for euclidean distance calculation.
- Per Erik Daniellsson's 8 SED distance mapping is an efficient euclidean distance transform having error bound of 0.076.
- Parallel mapping have more memory requirement but takes one scan of the image compared to multiple scans in sequential mapping, hence faster.
- Both are $O(n)$ algorithm n being number of pixels.

Algorithm 1: 8 Sequential Euclidean Distance Transform

The picture L is a two-dimensional array with the elements

$$L(i,j) \quad 0 \leq i \leq M-1, 0 \leq j \leq N-1.$$

Each element is a *two-element vector*

$$\bar{L}(i,j) = (L_i, L_j), \quad L_i, L_j \text{ being positive integers,}$$

$$L(i,j) = (L_i(i,j), L_j(i,j)).$$

The size of a vector $L(i,j)$ is defined by

$$|L(i,j)| = \sqrt{L_i^2 + L_j^2}$$

8SED:

Initially:

$$L(i,j) = (0, 0) \text{ for } (i,j) \in S$$

$$L(i,j) = (Z, Z) \text{ for } (i,j) \in S'$$

S here denotes the background and S' the segmentation outline

First picture scan:

for $j = 1, 2, \dots, N-1$,

for $i = 0, 1, 2, \dots, M-1$

$$\bar{L}(i,j) = \min(\bar{L}(i,j), \bar{L}(i-1,j-1)+(1,1), \bar{L}(i,j-1)+(0,1), \bar{L}(i+1,j-1)+(1,1))$$

for $i = 1, 2, \dots, M-1$

$$\bar{L}(i,j) = \min(\bar{L}(i,j), \bar{L}(i-1,j)+(1,0))$$

for $i = M-2, M-3, \dots, 1, 0$

$$\bar{L}(i,j) = \min(\bar{L}(i,j), \bar{L}(i+1,j)+(1,0))$$

Second picture scan:

for $j = N-2, N-3, \dots, 1, 0$

for $i = 0, 1, 2, \dots, M-1$

$$\bar{L}(i,j) = \min(\bar{L}(i,j), \bar{L}(i-1,j+1)+(1,1), \bar{L}(i,j+1)+(0,1), \bar{L}(i+1,j+1)+(1,1))$$

for $i = 1, 2, \dots, M-1$

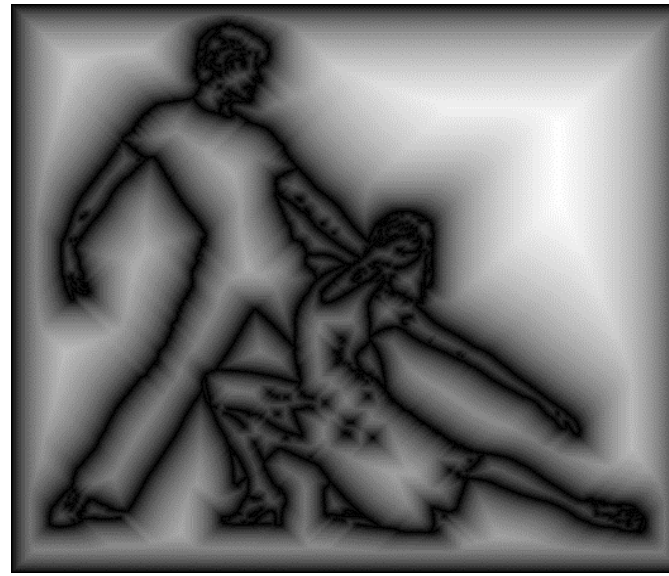
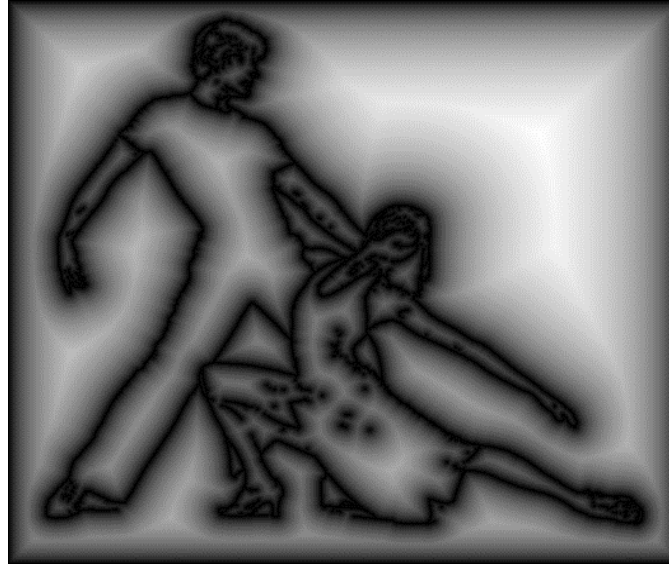
$$\bar{L}(i,j) = \min(\bar{L}(i,j), \bar{L}(i-1,j)+(1,0))$$

for $i = M-2, \dots, 1, 0$

$$\bar{L}(i,j) = \min(\bar{L}(i,j), \bar{L}(i+1,j)+(1,0))$$

The euclidean distance of any point may be calculated as $\sqrt{L_i^2 + L_j^2}$

Distance Mapping Image:



*top left- Segmented
Image, top right- d_8
distance transform,
bottom left- d_4
distance transform,
bottom right-
Euclidean distance
transform*

Square Fitting

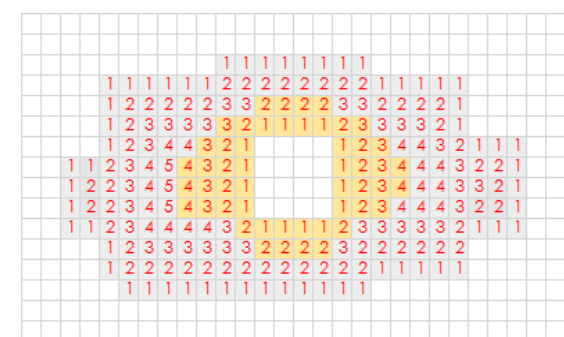
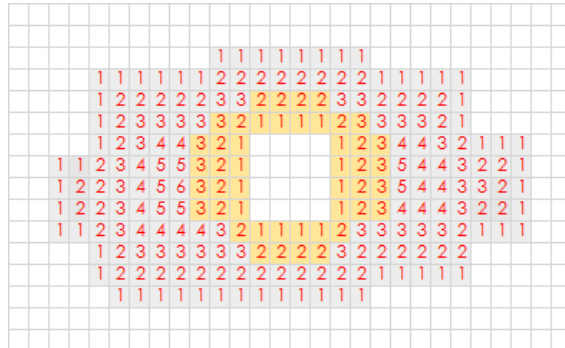
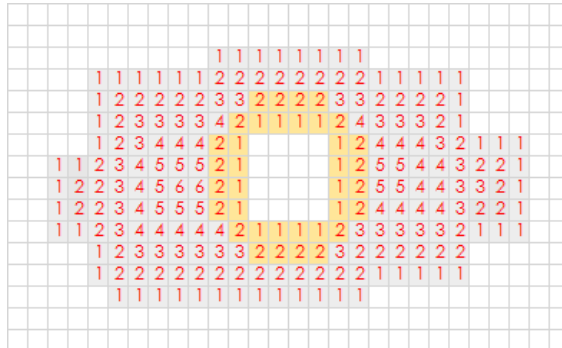
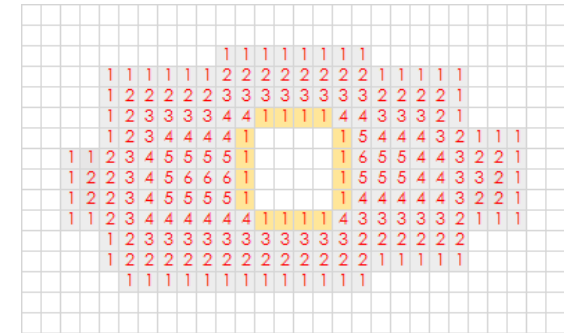
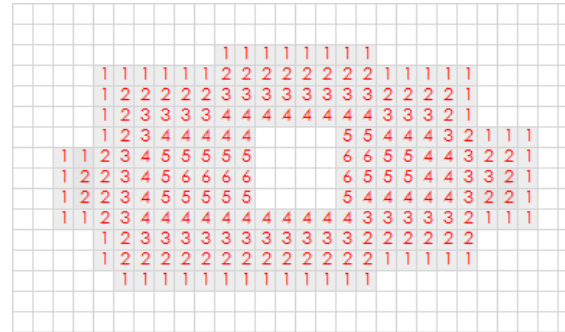
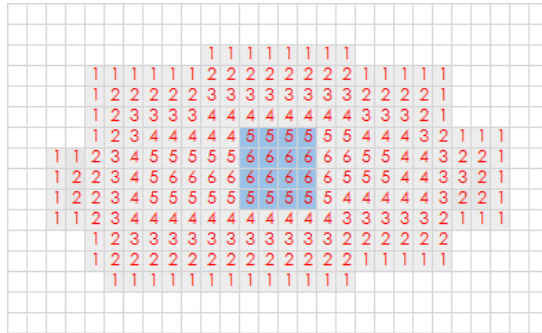
- The distance map gives an estimate of biggest square fitting possible at any pixel.
- Distance mapping value at pixel (i,j) denoted by d and largest fitting square size r_l different relations in different metrics.

$$r_l(i,j) = d_4(i,j) \text{ or } r_l = d_e(i,j) \text{ but } r_l(i,j) = \sqrt{2} \times d_8(i,j)$$

- Based on distance mapping, different values to alignment angle α is given. 0° in d_4 , 45° in d_8 and it is chosen out of a set of angles $(0^\circ, 15^\circ, 30^\circ, 45^\circ, 60^\circ, 75^\circ)$ in case of d_e .
- In case of d_e the alignment angle of square may be randomly assigned or based on *best fit angle calculation* heuristic.
- Overlapping square with segmentation outline or already drawn square are discarded.
- The distance recalculation after fitting a square is done locally .
- This distance recalculation may be easily done with parallel distance mapping starting with the new square boundary points.

Local Distance recalculation

- After each square fitting, the distance map changes.
- This may be easily handled by parallel distance mapping (d_4 or d_8) taking the square boundary points as the initial set of points in effectively $O(1)$ time complexity.
- The stopping criterion is newly calculated distances being equal to previous distances



Algorithm 2 : Square Packing

Input: Image I, Image M, Set D

Output: Set S

begin

 for each d in D

$\alpha \leftarrow \text{assign_angle}()$;

$r \leftarrow k * d$

$\{\text{sq_boundary}, \text{sq_internal}\} \leftarrow \text{precalculate_square}(r, \alpha)$

 while M(d) is not null

$P \leftarrow M(d).\text{get_first}()$;

 bool f=false;

 for p in sq_boundary

 if (M(P+p) is 0)

 f = true;

 end_if

 end_for

 if(f is true or variation_abv_threshold(I,sq_internal,P,threshold))

 continue;

 end_if

 bool f1=draw_square(P,sq_internal);

 if(f1)

$S \leftarrow S \cup \text{Sq}(P, I, r, \alpha)$;

$M(\{\text{sq_internal}\}) \leftarrow 0$;

 distance_recalculation(P,sq_boundary,M);

 end_if

 end_while

 end_for

end

Time complexity : Square Fitting

- The algorithm is run sequentially for each of the denomination in the set D and for each denomination $d \in D$ there is a $O(1)$ computation involved in `precalculate_square()` function.
- The inside while loop draws square for a particular denomination d_i and maximum $\alpha_i n / d_i^2$ squares are drawn.

n = number of pixels in image α_i = fraction of area covered by squares of denomination d_i .

- Drawing a square of denomination d_i takes $O(d_i^2)$ and an additional $O(1)$ for `distance_recalculation()`.
- Hence overall complexity formula is given as,

$$\begin{aligned} & O(|D|) \cdot O(1) + \sum_i \alpha_i n / d_i^2 (O(d_i^2) + O(1)) \\ &= O(|D|) + \sum_i \alpha_i n \leq O(|D|) + O(n) \text{ , as } \sum_i \alpha_i \leq 1 \end{aligned}$$

- In case of squares discarded due to failing variance check it adds an additional $O(\sum_i k_i d_i^2)$ time complexity where k_i number squares of size d_i are discarded.

Results: Comparison of three distance metric

Image Fig. No.	Size	DM	N_r	D	N_s	Thr	ρ	Cpu Time
Dog (a)	800x600	d_4	17	D1	4210	∞^*	0.8497	8 m16s
Dog (b)	800x600	d_8	17	D1	3900	∞	0.9204	4 m07s
Dog (c)	800x600	d_e	17	D1	6537	∞	0.9020	4 m19s



(a)



(b)



(c)

Results: Comparison of algorithm with or without variance check:

Image Fig. No.	Size	DM	N_r	D	N_s	Thr	ρ	Cpu Time
Salsa (a)	997x1132	d_8	31	D1	5566	∞	0.9517	4 m 6 s
Salsa (b)	997x1132	d_8	31	D1	6176	50	0.9126	9 m 59s



(a)

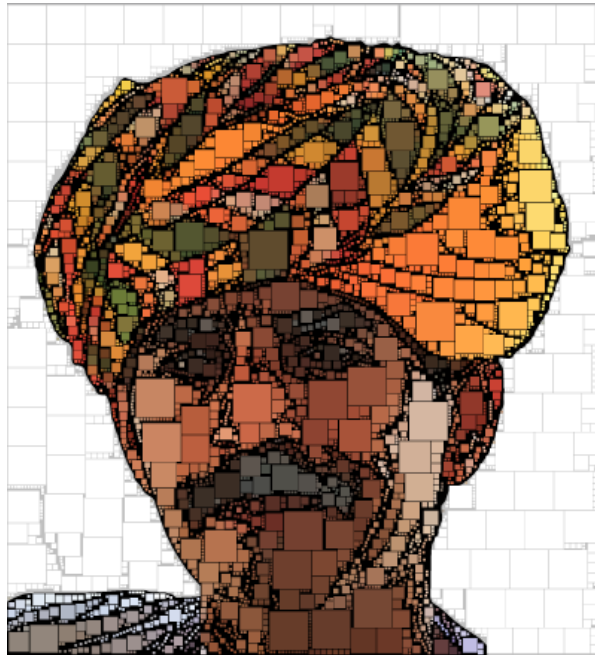


(b)

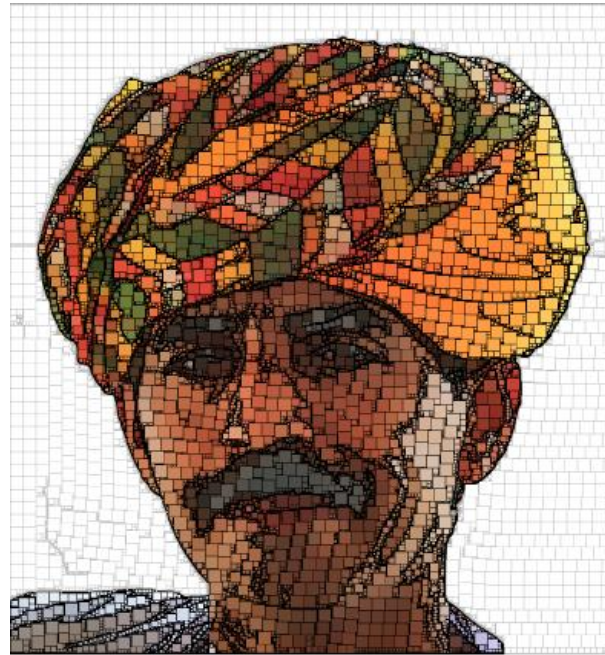
$D1=(100,80,60,40,30,25,20,15,12,10,8,5,3,2)$

Results: Comparison of algorithm different sets denomination:

Image Fig. No.	Size	DM	N_r	D	N_s	Thr	ρ	Cpu Time
Turban (a)	896x982	d_8	20	D2	5413	100	0.8775	10m16s
Turban (b)	896x982	d_8	20	D3	12082	100	0.8523	17m45s



(a)



(b)

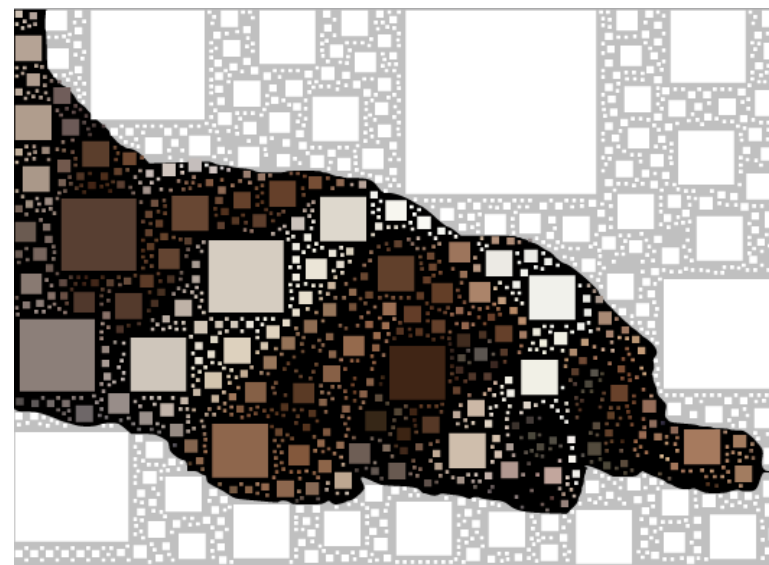
$D2=(30,25,20,16,12,10,5,3,2),$
 $D3=(10,8,7,5,3,2)$

Results: Comparison of d_4 and d_8 as local distance recalculation metric:

Image Fig. No.	Size	DM	N_r	D	N_s	Thr	ρ	Cpu Time
Dog (a)	800x600	d_8 (DR= d_4)	17	D1	3900	∞	0.9204	4 m07 s
Dog (a)	800x600	d_8 (DR= d_8)	17	D1	2153	∞	0.7104	20m15s



(a)



(b)

Conclusion:

- From the results we see that d_8 distance mapping gives the best packing and d_4 as local distance recalculation metric.
- The packing algorithm gives as high as 95% packing in some case .The segmentation boundary points and their adjacent pixels are mostly left.
- If the cluster points are completely separated then square fitting may be done in parallel for each of the segment reducing time requirement .

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Thank You