Minimal box size for fractal dimension estimation

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Abstract: We extend Kenkel's model for determining the minimal allowable box size s^* to be used in computing the box counting dimension of a self-similar geometric fractal. This minimal size s^* is defined in terms of a specified parameter ε which is the deviation of a computed slope from the box counting dimension. We derive an exact implicit equation for s^* for any ε . We solve the equation using binary search, compare our results to Kenkel's, and illustrate how s^* varies with ε . A listing of the Python code for the binary search is provided. We also derive a closed form estimate for s^* having the same functional form as Kenkel's empirically obtained expression.

Introduction

One of the most widely used fractal dimensions is the box counting dimension d_B . Suppose we wish to estimate d_B from N points, e.g., pixels in a 2-dimensional image. We cover the N points with boxes of side length s. If the object is 2-dimensional, the boxes are squares; if the object is 3-dimensional, the boxes are cubes; in general for an E-dimensional object, the boxes are E-dimensional hypercubes. Let B(s) be the number of boxes containing at least one of the N points. When s is sufficiently large, all points lie in one box, so B(s)= 1. As s decreases, B(s) increases or remains constant. For all sufficiently small box sizes, each non-empty box contains a single point, so B(s) = N.

The box counting dimension d_B of a geometric object is defined as (Mandelbrot 1983)

$$d_{B} = -\lim_{s \to 0} \frac{\log B(s)}{\log s}$$

There are examples for which d_B does not exist (Falconer 2003) but in this paper we assume d_B exists. In practice, d_B is computed by evaluating B(s) for a set of J values of s, which we denote by s_j , j = 1, 2, ..., J. Then, typically, a line is fitted (often using linear regression) to the J pairs (-log s_j , log $B(s_j)$), and the slope of this line is the estimate of d_B . Assume the s_j values are ordered so that $s_1 < s_2 < ... < s_J$. We want $s_J - s_1$ as large as possible, so as to minimize the error in estimating d_B from the J pairs (log s_j , log $B(s_j)$). However, we cannot make s_1 arbitrarily small, since then, as noted above, B(s) approaches N, and the slope of the (log s, log B(s)) curve approaches 0. The study of the minimal and maximal box sizes has received a great deal of attention; see, e.g., the large number of references in Kenkel (2013).

A recent paper by Kenkel (2013) introduces a simple but very useful probabilistic model to obtain a relationship between *N* and s_1 . The model estimates the expected value of B(s) as a function of *s* and d_B . Then s_1 is chosen to be the value at which the "local slope" of the ($-\log s, \log B(s)$) curve deviates from d_B by no more than 0.001, where this accuracy was chosen since estimates of d_B are often expressed to three or four decimal places. Using 28 simulations in which *s* was decreased until the local slope is within 0.001 of d_B , Kenkel obtained the following approximation for the minimal usable box size:

$$s_1 \approx \left(\frac{N}{10}\right)^{-1/d_B} \ . \tag{1}$$

Since (1) was obtained numerically, and only for the accuracy 0.001, this result provides no guidance on how s_1 varies as a function of the accuracy. Indeed, in many applications, e.g., in the analysis of neurons (Karperien 2013), d_B is only estimated to two decimal places. Letting ε be the desired accuracy, in this paper we generalize Kenkel's result to compute s_1 for arbitrary ε . We show that s_1 can be computed, for any ε , by standard one-dimensional search techniques such as binary search.

The original model

Following Kenkel (2013), consider first the 1-dimensional case. If we randomly select *N* points on [0, 1] then the probability that a given box (i.e., interval) of size *s* does not contain any of these points is $(1-s)^N$. The probability that a given box of size *s* contains at least one of *N* randomly selected points is $1 - (1-s)^N$. Hence the expected number B(s) of nonempty boxes of size *s* is given by $B(s) = [1 - (1-s)^N]/s$.

Considering next the 2-dimensional case, if we randomly select *N* points on the unit square $[0, 1] \times [0, 1]$ then the probability that a given box of size *s* does not contain any of these points is $(1 - s^2)^N$. The probability that a given box of size *s*

contains at least one of *N* randomly selected points is $1 - (1 - s^2)^N$. Hence the expected number B(s) of nonempty boxes of size *s* is given by $B(s) = [1 - (1 - s^2)^N]/s^2$.

Finally, consider the case where the *N* points are sampled from a self-similar fractal set with box counting dimension d_B . For example, *N* might be the total number of pixels in an image of a real-world fractal, such as a fern leaf. The expected number B(s) of nonempty boxes of size *s* is given by (Kenkel 2013)

$$B(s) = \frac{1 - (1 - s^{d_B})^N}{s^{d_B}}.$$
(2)

This equality is the basis for the analysis of the remainder of this paper.

From (2), Kenkel numerically computes the "local slope" of the (log *s*, log *B*(*s*)) curve, where the "local slope" *m*(*s*) is defined by $m(s) \equiv (\log B(s + \delta) - \log B(s))/(\log (s + \delta) - \log s)$ for some small increment δ . Then, s_1 is the value for which $m(s)+d_B = 0.001$ and empirically s_1 is found to be well approximated using (1). From (1) we obtain $s_1^{dB} = 10/N$. Substituting this in (2), we have

$$B(s_1) = \frac{1 - \left(1 - (10/N)\right)^N}{10/N}$$

which yields, for large *N*, the estimate $B(s_1) = N/10$. This means no box size *s* should be used, in the estimation of d_B , for which B(s) > N/10 (Kenkel 2013).

The generalized model

In this section, we generalize the original model by determining the minimal box size when the constant 0.001 described in Section 2 is replaced by a positive parameter ε . Rather than work with a local slope of the (log *s*, log *B*(*s*)) curve, we work with an actual derivative, which can be obtained in closed form. Using the chain rule for derivatives, we have

$$\frac{d \log B(s)}{d \log s} = \left(\frac{d \log B(s)}{d s}\right) \left(\frac{d s}{d \log s}\right) =$$

$$\left(\frac{d \log B(s)}{d s}\right) \left(\frac{d \log s}{d s}\right)^{-1} = \left(\frac{d \log B(s)}{d s}\right) s.$$
(3)

From (2), taking the derivative of $\log B(s)$ we have

$$\frac{d\,\log B(s)}{d\,s} = \tag{4}$$

$$\frac{d_B N s^{-1} (1 - s^{d_B})^{N-1} - d_B s^{-d_B - 1} (1 - (1 - s^{d_B})^N)}{(1 - (1 - s^{d_B})^N) s^{-d_B}}.$$

Combining (3) and (4) yields

$$\frac{d \log B(s)}{d \log s} =$$

$$\frac{d_B N (1 - s^{d_B})^{N-1} - d_B s^{-d_B} (1 - (1 - s^{d_B})^N)}{(1 - (1 - s^{d_B})^N) s^{-d_B}}.$$
(5)

Denoting the right hand side of (5) by F(s), our task is to find the value s^* such that $F(s^*) = -d_B + \varepsilon$. For $s < s^*$ we have $F(s) > -d_B + \varepsilon$, so such box sizes should not be used in the estimation of d_B . Thus s^* is the minimal usable box size. As a first step towards calculating s^* , define

$$\alpha \equiv (1 - s^{dB})^{N-1} . \tag{6}$$

While α is actually a function of *s*, for notational simplicity we write α rather than $\alpha(s)$. The equation $F(s) = -d_B + \varepsilon$ can now be rewritten as

$$\frac{d_B N \alpha - d_B s^{-d_B} \left(1 - \alpha (1 - s^{d_B})\right)}{\left(1 - \alpha (1 - s^{d_B})\right) s^{-d_B}} + d_B = \epsilon \,. \tag{7}$$

Dividing both sides by d_B yields

$$\frac{N\alpha - s^{-d_B} \left(1 - \alpha (1 - s^{d_B})\right)}{\left(1 - \alpha (1 - s^{d_B})\right) s^{-d_B}} = \frac{\epsilon}{d_B} - 1.$$
(8)

Define $v \equiv (\varepsilon/d_B) - 1$. Multiplying the numerator and denominator of the left hand side of (8) by s^{d_B} yields $N\alpha s^{d_B} - 1 + \alpha - \alpha s^{d_B} = v(1-\alpha + \alpha s^{d_B})$, which simplifies to $\alpha s^{d_B}(N-1-v)=(v+1)$ (1- α). Recalling the definition of v, more algebra yields

$$\frac{Nd_B}{\epsilon} - 1 = \left(\frac{1-\alpha}{\alpha}\right)s^{-d_B}.$$
(9)

Finally, from (9) and the definition (6) of α , we obtain an implicit equation for *s**:

$$\left(\left(1-s^{d_B}\right)^{1-N}-1\right)s^{-d_B} = \epsilon^{-1}Nd_B - 1.$$
 (10)

Equation (10) is exact; no approximations were made. The parameter ε appears only in the right hand side, which is independent of *s*.

We compute a solution s^* of (10) using binary search over [0,1]; a code listing is provided in the Appendix. The binary



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Figure 2. Minimal usable box size for N = 1000 and for three values of ε , for $1 \le d_B \le 2$.

10 8 error 6 % 4 2 0 1 1.2 1.4 1.6 1.8 2 box counting dimension 0.1 0.09 • eps = 0.1 eps = 0.001 eps = 0.010.08 0.07 x 0.06 0.05 0.04 0.03 0.02 0.01 0 1 1.2 1.4 1.6 1.8 2 box counting dimension

search is halted when the width of the interval containing s^* is less than 1.0×10^{-7} , which takes 24 iterations. For N = 1000 and $\varepsilon = 0.001$, the percent error $100(s_1 - s^*)/s^*$, where s_1 is the estimate defined by (1), is plotted in Figure 1 as a function of d_B for $1 \le d_B \le 2$. The error in the approximation (1) is highest (about 10%) when $d_B = 1$.

Since, as noted in Section 1, in many applications d_B is estimated to only two decimal places, Figure 2 compares, for N=1000, the value s* for $\varepsilon=0.1$, $\varepsilon=0.01$, and $\varepsilon=0.001$, all for the same range $1 \le d_B \le 2$.

Approximating the minimal usable box size

We now derive a closed form approximation to s^* . Defining $x = s^{dB}$, we rewrite (10) as

$$x^{-1}\left(\left(1-x\right)^{1-N}-1\right) = \epsilon^{-1}Nd_B - 1.$$
 (11)

We have x < 1 since s < 1 and $d_B > 0$. Ignoring terms of degree four and higher, the Taylor series expansion of $(1-x)^{1-N}$ yields $(1-x)^{1-N} \approx 1-(1-N)x + (1-N)(-N)x^2/2 - (1-N)(-N)$ $(-N-1)x^3/6$. For large N we have $(1-x)^{1-N} \approx 1 + Nx + (N^2/2)x^2 + (N^3/6)x^3$. Substituting this in (11) yields

$$x^{-1}\left(\left(1-x\right)^{1-N}-1\right)$$

$$\begin{split} &\approx \quad x^{-1} \Big(1 + N x + (N^2/2) x^2 + (N^3/6) x^3 - 1 \Big) \\ &= \quad N + (N^2/2) x + (N^3/6) x^2 \\ &= \epsilon^{-1} N d_B - 1 \approx \epsilon^{-1} N d_B \,. \end{split}$$

Dividing by N, we obtain $(N^2/6)x^2 + (N/2)x + (1 - \varepsilon^{-1}d_B) \approx 0$. Using $1 - \varepsilon^{-1}d_B \approx -\varepsilon^{-1}d_B$ we get

$$\begin{split} x &= \frac{-\frac{N}{2} \pm \sqrt{\frac{N^2}{4} - 4\frac{N^2}{6}(-\epsilon^{-1}d_B)}}{(2/6)N^2} \approx \frac{N\sqrt{\frac{2}{3}\epsilon^{-1}d_B}}{(1/3)N^2} \\ &= \frac{1}{N}\sqrt{\frac{6d_B}{\epsilon}} \,. \end{split}$$

By definition, $x = s^{d_B}$, so $s^{d_B} = (1/N)\sqrt{(6d_B/\varepsilon)}$. Solving for *s* yields an approximation \tilde{s} for the minimal usable box size:

$$\widetilde{s} = \left(N\sqrt{\frac{\epsilon}{6\,d_B}}\right)^{-1/d_B} \,. \tag{12}$$

The estimate \tilde{s} , which is a decreasing function of ε , looks identical to (1), except that the empirically derived constant 0.1 in (1) is replaced by $\sqrt{(\varepsilon/(6d_B))}$ in (12). However, \tilde{s} is not a good approximation to s^* . For example, with N = 1000, $d_B = 2$, and $\varepsilon = 0.001$, we have $s^* \approx 0.099$ while

 $\tilde{s} \approx 0.331$. The error arises from the third order Taylor series approximation, since for $x = (s^*)^2$ we have $(1-x)^{1-N} \approx 19700$ while $1 + Nx + (N^2/2)x^2 + (N^3/6)x^3 \approx 219$. Thus, while (12) yields the functional form obtained by Kenkel, in practice s^* should be computed using (10) and binary search.

Concluding remarks

We generalized Kenkel's model for finding the minimal usable box size s^* for computing the box counting dimension. For box sizes smaller than s^* , the slope of the (-log s, log B(s)) curve flattens out, and deviates by more than a specified accuracy ε from d_B . Whereas Kenkel considered only the choice $\varepsilon = 0.001$, we derived an exact implicit equation for s^* for any ε . Binary search was used to actually compute s^* , and a Python implementation is provided in the Appendix. We also used the implicit equation to derive a closed form approximation for s^* having the same functional form as Kenkel's empirically obtained expression.

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References

- Falconer, K. 2003. Fractal Geometry: Mathematical Foundations and Applications, 2nd edn. Wiley, Chichester.
- Karperien A., H. Ahammer and H.F. Jelinek. 2013. Quantitating the subtleties of microglial morphology with fractal analysis. *Frontiers in Cellular Neuroscience* 7: 3, doi: 10.3389/fncel.2013.00003.
- Kenkel, N.C. 2013. Sample size requirements for fractal dimension estimation. *Community Ecol.* 14: 144–152.
- Mandelbrot, B.B. 1983. *The Fractal Geometry of Nature*. W.H. Freeman, New York.

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Appendix: Binary search code

Python code used to implement the binary search method used in Section 3 to compute the *s** solving (10). The code performs binary search over the interval [0, 1] to compute s* to an accuracy of 10^{-7} for $\varepsilon = 0.001$ and for d_B from 1 to 2, in increments of 1/200. The variable Estimate is the estimate given by (1).

Binary Search	
1	import math
2	N = 1000.
3	epsilon = .001
4	for i in range(201):
5	d = 1.0 + float(i)/200.
6	RHS = -1.0 + N*d/epsilon
7	Estimate = (0.1*N)**(-1.0/d)
8	low = 10 * * (-8)
9	high = 1.0
10	while (low + $10**(-7) \le $ high):
11	s = (low + high)/2.0
12	x = s**d
13	LHS = $((1.0-x)**(1.0-N) -1.0)/x$
14	if LHS $<$ RHS:
15	low = s
16	else:
17	high = s
18	error = 100*(Estimate-s)/s
19	print " d ", d, " s ", s, " error ", error