









# Supplement: Uncertainty Visualization of Critical Points of 2D Scalar Fields for Parametric and Nonparametric Probabilistic Models

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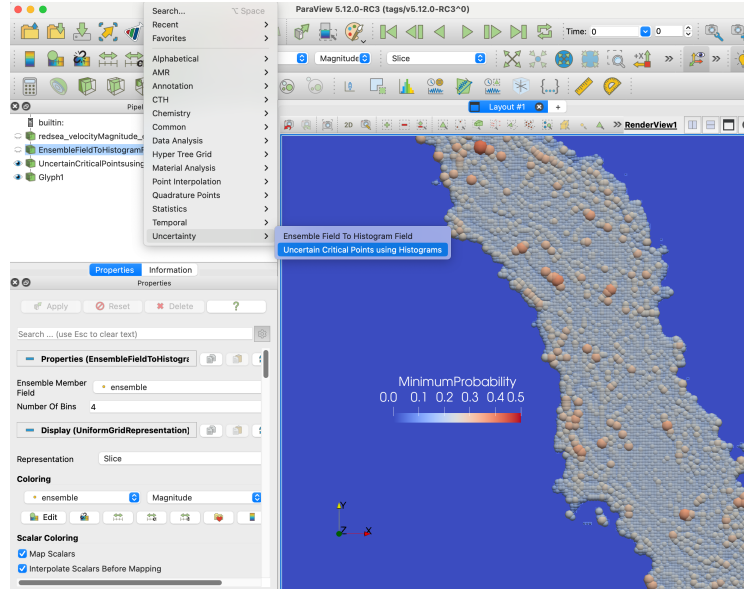


Fig. 1: Demonstration of our critical point uncertainty filter inside ParaView [2] to provide near-real-time results for the Red Sea ensemble dataset [6] with computations performed using VTK-m [5] parallel code on a GPU backend.

**Abstract**—This paper presents a novel end-to-end framework for closed-form computation and visualization of critical point uncertainty in 2D uncertain scalar fields. Critical points are fundamental topological descriptors used in the visualization and analysis of scalar fields. The uncertainty inherent in data (e.g., observational and experimental data, approximations in simulations, and compression), however, creates uncertainty regarding critical point positions. Uncertainty in critical point positions, therefore, cannot be ignored, given their impact on downstream data analysis tasks. In this work, we study uncertainty in critical points as a function of uncertainty in data modeled with probability distributions. Although Monte Carlo (MC) sampling techniques have been used in prior studies to quantify critical point uncertainty, they are often expensive and are infrequently used in production-quality visualization software. We, therefore, propose a new end-to-end framework to address these challenges that comprises a threefold contribution. First, we derive the critical point uncertainty in closed form, which is more accurate and efficient than the conventional MC sampling methods. Specifically, we provide the closed-form and semianalytical (a mix of closed-form and MC methods) solutions for parametric (e.g., uniform, Epanechnikov) and nonparametric models (e.g., histograms) with finite support. Second, we accelerate critical point probability computations using a parallel implementation with the VTK-m library, which is platform portable. Finally, we demonstrate the integration of our implementation with the ParaView software system to demonstrate near-real-time results for real datasets.

**Index Terms**—Topology, uncertainty, critical points, probabilistic analysis

## 1 INTRODUCTION

We divide this supplement into four parts. First, we present the detailed algorithms and illustrations for critical point probability computation

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for the four-pixel neighborhood case (Sec. 4.2 of the main paper). Second, we show quantitative and qualitative correctness of our algorithms for the uniform, Epanechnikov, and histogram (closed-form and semi-analytical) models through comparisons with the Monte Carlo (MC) solutions. Third, we present quantitative evaluation for the climate dataset [4] used in the main paper. Finally, we show seamless integration of our parallel VTK-m [5] implementation inside ParaView [2].

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## 2 CRITICAL POINT PROBABILITY COMPUTATION FOR FOUR-PIXEL NEIGHBORHOOD

### 2.0.1 Local Minimum Probability

The probability of point  $p$  being a local minimum,  $\Pr(p = l_{min})$ , can be computed by integrating the joint probability  $\Pr_{joint}$  over its support where random variable  $X_1$  is simultaneously smaller than all neighboring random variables (i.e.,  $X_2, \dots, X_5$ ). Mathematically, the local minimum probability computation can be represented as follows:

$$\begin{aligned} \Pr(p = l_{min}) &= \Pr[(X_1 < X_2) \text{ and } (X_1 < X_3) \text{ and } (X_1 < X_4) \text{ and } (X_1 < X_5)] \\ &= \int_{x_1=a_1}^{x_1=b_{min}} \int_{x_2=\max(x_1, a_2)}^{x_2=b_2} \dots \int_{x_5=\max(x_1, a_5)}^{x_5=b_5} (\text{Pdf}_{joint}) dx, \quad (1) \end{aligned}$$

where  $b_{min} = \min(b_1, b_2, b_3, b_4, b_5)$

Equation (1) represents the core integration formula for the computation of local minimum probability at a domain position  $p$ . We now explain the integral limits in Eq. (1) and our proposed piecewise integration algorithm to compute the formula in Eq. (1).

Limits  $a_1$  and  $b_{min}$  of the outer integral in Eq. (1): The outer integral of Eq. (1) indicates the portion of data range of a random variable  $X_1$  (i.e.,  $[a_1, b_1]$ ) that can result in point  $p$  being a local minimum. In particular, the portion  $[a_1, b_{min}]$  (with  $a_1 < b_{min}$ ) of random variable  $X_1$  can result in point  $p$  being a local minimum, where  $b_{min}$  denotes the minimum among  $b_1, \dots, b_5$ . In contrast, the data range  $[b_{min}, b_1]$  for  $b_{min} \neq b_1$  cannot result in a point  $p$  as a local minimum ( $l_{min}$ ) because it will be always greater than one of the random variables  $X_2, X_3, X_4$ , and  $X_5$  depending on if  $b_{min} = b_2, b_{min} = b_3, b_{min} = b_4$ , or  $b_{min} = b_5$  respectively. Mathematically, for any value  $x_1 \geq b_{min}$ ,  $\Pr(p = l_{min}) = 0$ . In the case  $b_{min} < a_1$ , then  $\Pr(p = l_{min}) = 0$  because at least one random variable among  $X_i$  with  $i \in \{2, 3, 4, 5\}$  will be always smaller than  $X_1$ .

Limits  $\max(x_1, a_i)$  and  $b_i$  of the inner integrals in Eq. (1): The four inner integrals in Eq. (1) integrate the joint distribution  $\text{Pdf}_{joint}$  over its support where random variables  $X_2, \dots, X_5$  are simultaneously greater than  $x_1 \in [a_1, b_{min}]$  in the outer integral. The inner integral lower limits are  $\max(x_1, a_i)$  for  $i \in \{2, 3, 4, 5\}$ . The maximum is taken because the support of a random variable  $X_i$  is restricted to  $[a_i, b_i]$ . Thus, for  $x_1 < a_i$  in any inner integral, the entire support  $[a_i, b_i]$  with  $i \in \{2, 3, 4, 5\}$  will always be greater than  $x_1$ , and the inner integral does not depend on the value of  $x_1$ . In contrast, for  $x_1 > a_i$ , the inner integration depends on the value of  $x_1$  because  $x_1$  assumes values in the support of distributions. It is guaranteed that the upper limit of inner integrals in Eq. (1) is greater than their respective lower limit, i.e.,  $b_i \geq \max(x_1, a_i)$ , for two reasons. First, for any random variable  $X_i$ , we assume  $a_i < b_i$ . Second, the maximum value of  $x_1$  is equal to  $b_{min} = \min(b_1, b_2, b_3, b_4, b_5)$  based on the outer integral (see the previous paragraph), and it cannot exceed the upper limits  $b_i$  with  $i \in \{2, 3, 4, 5\}$  in inner integrals. Depending upon if the  $\max(x_1, a_i)$  is equal to  $x_1$  or  $a_i$ , the integral in Eq. (1) can be simplified and computed differently, which necessitates the evaluation of the integral in Eq. (1) in a piecewise manner, as described next.

Piecewise simplification of Eq. (1): The core integration formula in Eq. (1) can be simplified differently for different subsets of the range of the outer integral (i.e.,  $[a_1, b_{min}]$ ). We refer to each subset as a piece  $P$ . For a piece  $P$ , where  $x_1 < a_i, \forall i \in \{2, 3, 4, 5\}$ , the inner integrals in Eq. (1) attain the range  $x_i \in [a_i, b_i], \forall i \in \{2, 3, 4, 5\}$ . In other words, the inner integrals do not depend on  $x_1$ . Thus, the integration over the entire support of random variables  $X_i, \forall i \in \{2, 3, 4, 5\}$ , simplifies Eq. (1) to the integration over a marginal distribution of  $X_1$  for the piece  $P$ , i.e.,  $\int_{x_1 \in P} \text{Pdf}_{X_1}(x_1) dx_1$ .

For a piece  $P$ , where  $x_1 \in [a_2, b_2]$  and  $x_1 < a_i, \forall i \in \{3, 4, 5\}$ , the inner integrals in Eq. (1) attain the range  $x_2 \in [x_1, b_2]$  and  $x_i \in [a_i, b_i], \forall i \in \{3, 4, 5\}$ . In this case, only the first inner integral related to the range of random variable  $X_2$  depends upon  $x_1$ . Thus, Eq. (1) simplifies as the integration over the joint distribution of  $X_1$  and  $X_2$  for the piece

**Function** *computeLocalMinimumProbability()*

**Input:** Intervals  $I = [[a_1, b_1], \dots, [a_5, b_5]]$

**Output:**  $\Pr(p = l_{min})$

$b_{min} = \min(b_1, \dots, b_5)$

**if**  $b_{min} < a_1$  **then**

    | return 0

**else**

    /\* Task1: Break the outer integral in Eq. (1) into pieces  $P_i$  \*/

$I_{sorted} = \text{sort}(I)$  based on start points  $a_i$

$id_{a1} =$  Index of the interval in  $I_{sorted}$  that starts with  $a_1$

$id_{bMin} =$

    Index of the interval in  $I_{sorted}$  that contains  $b_{min}$

    /\* Task2 and Task 3: Compute piece integrals \*/

$localMinimumProbability =$

$computePieceIntegrals(id_{a1}, id_{bMin}, I_{sorted}, b_{min})$

    return  $localMinimumProbability$

**end**

**Algorithm 1: Computation of local minimum probability**

$P$ , i.e.,  $\int_{x_1 \in P} \text{Pdf}_{X_1}(x_1) \text{Pdf}_{X_2}(x_2) dx_2 dx_1$ . In summary, various pieces of the outer integral in Eq. (1) can be simplified differently based on if the inner integrals depend on  $x_1$  or not.

Algorithm for computing local minimum probability: The computation of the core integration formula in Eq. (1) depends on the ordering of start points  $a_i$  with  $i \in \{1, 2, 3, 4, 5\}$  and  $b_{min}$ . Thus, there are  $6! = 720$  permutations of  $a_i$  and  $b_{min}$ . We, therefore, devise an efficient algorithm that computes the piecewise integrals on the fly depending on the observed permutation of  $a_i$  and  $b_{min}$  without needing to go through all permutations.

We now describe our Algorithm 1 for computation of local minimum probability at a domain position  $p$  (i.e.,  $\Pr[p = l_{min}]$ ). The algorithm comprises the three main tasks. **Task 1: Determination of pieces  $P_i$  of the outer integral in Eq. (1) needed for performing piecewise integration.** Initially, we compute  $b_{min}$ . If  $b_{min} < a_1$ , then there are no pieces or  $\Pr(p = l_{min}) = 0$ . If  $b_{min} > a_1$ , then we sort intervals representing uncertain data ranges (i.e.,  $[a_i, b_i]$ ) based on their start points  $a_i$  and keep them in the array named  $I_{sorted}$ . We note the index of the interval corresponding to  $[a_1, b_1]$  in  $I_{sorted}$  (referred to as  $id_{a1}$ ) and the index of interval in  $I_{sorted}$  from the end that contains  $b_{min}$  (referred to as  $id_{b_{min}}$ ), as  $a_1$  and  $b_{min}$  constitute limits of the outer integral in Eq. (1). Any start points  $a_i$  contained in the range of indices  $id_{a1}$  and  $id_{b_{min}}$  determine the pieces  $P_i$  for integration. This process generalizes to any ordering of  $a_i$  to determine the pieces  $p_i$  of the outer integral in Eq. (1).

Next, we compute the integration for piece  $P_1$  denoted as  $I_{P_1}$ . **Task 2: Integration over piece  $P_1$  of the outer integral range  $[a_1, b_{min}]$ .** The integration for piece  $P_1$  depends on the intervals that started before  $a_1$  because the inner integral in Eq. (1) depends on  $x_1$  for a random variable  $X_i$  (with  $i \in 2, 3, 4, 5$ ) started before  $a_1$ , as  $\max(x_1, a_i)$  is equal to  $x_1$ . The Task 2 in Algorithm 2, therefore, corresponds to finding the intervals that started before  $id_{a1}$  (by looking up the  $I_{sorted}$  array), which also determines the upper limits of inner integrals for piece  $P_1$  (denoted by the array  $upLimits$  in Algorithm 2) depending on observed order of intervals. In particular, the  $lowLimit$  and  $upLimits[1]$  in Task 2 of Algorithm 2 store the limits of the outer integral for piece  $P_1$ . The remaining entries in the  $upLimits$  array are used to store the upper limits of inner integrals based on the order they are observed.

The computation of the integral in Eq. (1) for piece  $P_1$  (as well as any arbitrary piece  $P_i$ ) simplifies to one of the five types of integration formulae, which we call integration templates. The integration templates are presented in Algorithm 3. Depending on the number of upper limits set (denoted by a variable  $h_i$  in Algorithm 3), the integral template varies. If there is no overlap with a piece, then the integral in Eq. (1) simplifies to the integration of the probability distribution of random variable  $X_1$  over a piece, which corresponds to setting  $h_1$  and the usage of Template 1 in Algorithm 3. If only one random variable is over-

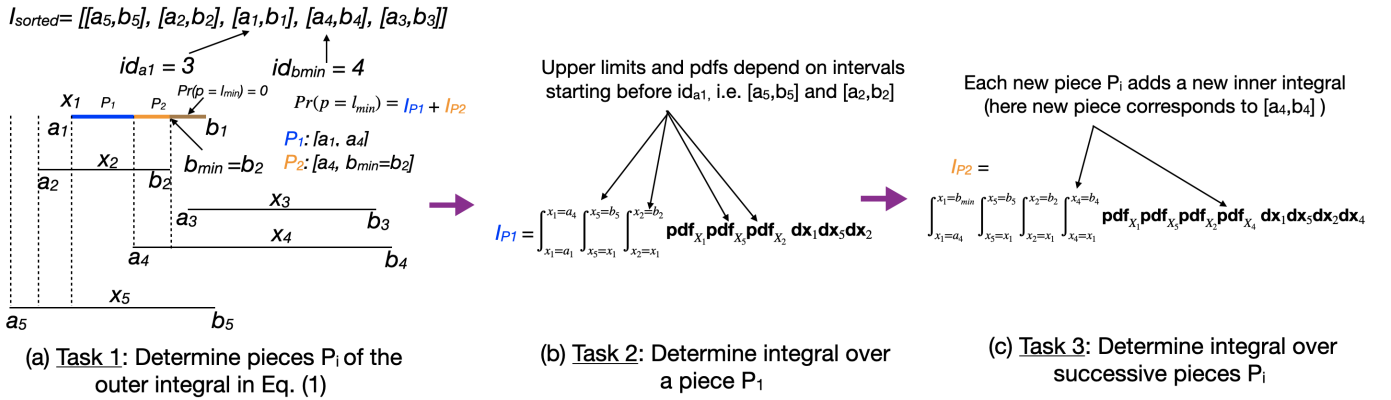


Fig. 2: Tasks summarizing our algorithm for computing local minimum probability for uncertain data.

### Function `computePieceIntegrals()`

```

Input:  $id_{a_1}, id_{b_{min}}, I_{sorted}, b_{min}$ 
Output: Sum of integrals over pieces  $P_i$ 

/* Initialize the upper limits of outer and
   inner integrals for a piece */
upLimits = [None, None, None, None, None]
totalIntegral = 0

/* Task2: Compute integral for piece  $P_1$  */
/* Determine the upper limit of piece  $P_1$  */
lowLimit =  $a_1$ 
if  $id_{a_1} < id_{b_{min}}$  then
  |  $upLimits[1] = I_{sorted}[id_{a_1} + 1][1]$ 
if  $id_{a_1} == id_{b_{min}}$  then
  |  $upLimits[1] = b_{min}$ 
/* Find intervals started before  $a_1$  and set
   respective upper limits for inner
   integrals */
for  $k = 1$  to  $(id_{a_1} - 1)$  do
  |  $upLimits[k + 1] = I_{sorted}[k][2]$ 
end
 $I_{P_1} =$ 
   $integralTemplate(lowLimit, upLimits[1], upLimits[2],$ 
     $upLimits[3], upLimits[4], upLimits[5], I_{sorted})$ 
 $totalIntegral += I_{P_1}$ 

/* Task3: Compute integral for successive
   pieces */
for  $k = (id_{a_1} + 1)$  to  $id_{b_{min}}$  do
  /* Determine piece limits */
   $lowLimit = I_{sorted}[k][1]$ 
  if  $k < id_{b_{min}}$  then
    |  $upLimits[1] = I_{sorted}[k + 1][1]$ 
  else
    |  $upLimits[1] = b_{min}$ 
  /* Determine the upper limit of new
   interval */
   $upLimits[k] = I_{sorted}[k][2]$ 
   $I_{P_{next}} =$ 
     $integralTemplate(lowLimit, upLimits[1], upLimits[2],$ 
       $upLimits[3], upLimits[4], upLimits[5], I_{sorted})$ 
   $totalIntegral += I_{P_{next}}$ 
end
return  $totalIntegral$ 

```

### Algorithm 2: Computation of the integral for piece $P_1$ and successive pieces $P_{next}$

lapping with a piece, then the integral in Eq. (1) simplifies to integral over the joint distribution of  $X_1$  and a random variable corresponding to overlapping interval. One overlapping interval corresponds to setting

only  $h_1$  and  $h_2$  and the usage of Template 2 in Algorithm 3. Note that the parameters  $\theta_i$  in Algorithm 3 capture the probability distribution functions to consider in templates by looking up the  $I_{sorted}$  array. The probability distributions to consider essentially depend on the ordering of intervals  $[a_i, b_i]$  and are captured on the fly by parameters  $\theta_i$ .

Having determined the integration for piece  $P_1$ , we compute integration for successive pieces (denoted as  $I_{P_{next}}$  in Algorithm 2). **Task 3: Integration over piece  $P_i$  with  $i > 1$ .** Essentially, each new start point  $a_i$  with  $i \in \{2, 3, 4, 5\}$  observed between the outer integral limits  $a_1$  and  $b_{min}$  of Eq. (1) creates a new piece. Generally speaking, each new start point  $a_i$  results in different simplification of Eq. (1) because  $\max(x_1, a_i)$  in Eq. (1) becomes equal to  $x_1$  at each new start point. Thus, encountering a new start point adds one inner integral in a simplified form compared to the piece before encountering a new start point. Finally, integration of all pieces (i.e.  $I_{P_1}$  and  $I_{P_{next}}$  in Algorithm 2) is summed to compute the local minimum probability at a point  $p$ , i.e.,  $Pr(p = l_{min})$ .

**Illustration of local minimum probability computation:** Fig. 2 illustrates our method for computing the local minimum probability for a domain point  $p$ . As shown for the example in Fig. 2,  $a_5 < a_2 < a_1 < a_4 < b_2 < a_3 < b_1 < b_5 < b_3 < b_4$ . Initially, we determine the range of a random variable  $X_1$  that can result in point  $p$  being a local minimum. As shown in Fig. 2a, each value in the range  $[x_1 = a_1, x_1 = (b_{min} = b_2)]$  has a nonzero probability of being simultaneously smaller than the neighboring random variables (i.e.,  $X_2, \dots, X_5$ ). In contrast, the range  $[x_1 = (b_{min} = b_2), x_1 = b_1]$  is always greater than the random variable  $X_2$ , and therefore, cannot result in point  $p$  as a local minimum.

In Task 1, we determine the pieces  $P_i$  of the range  $[x_1 = a_1, x_1 = (b_{min} = b_2)]$ . As depicted in Fig. 2, the array  $I_{sorted}$  has intervals ordered by  $a_i$ , where  $a_5 < a_2 < a_1 < a_4 < a_3$ . For this  $I_{sorted}$ ,  $id_{a_1} = 3$  and  $id_{b_{min}} = 4$ . Since  $a_4$  is a start point in interval indexed by  $id_{b_{min}}$ , it divides the outer integral range  $[a_1, b_{min}]$  in Eq. (1) into two pieces (depicted in blue and orange in Fig. 2).

In Task 2, we determine the simplification of the formula in Eq. (1) for piece  $P_1$ . The simplification for piece  $P_1 = [a_1, a_4]$  in Fig. 2 (denoted by blue) depends on the number of intervals that started before  $a_1$ . As observed in Fig. 2, the intervals  $[a_5, b_5]$  and  $[a_2, b_2]$  start before  $a_1$ . Since in piece  $P_1$ ,  $x_1 < a_3$  and  $x_1 < a_4$ , the formula in Eq. (1) integrates random variables  $X_3$  and  $X_4$  over their entire support and simplifies to the integration over the joint distribution of random variables  $X_1, X_5$ , and  $X_2$  shown in Fig. 2b.

In Task 3, we determine the simplification of the formula in Eq. (1) for successive pieces formed by each new start point  $a_i \in [a_1, b_{min}]$ . In Fig. 2, the start point  $a_4 \in [a_1, b_{min}]$  results in a new piece  $P_2$  shown in orange. All inner integrals for piece  $P_2$  stay the same as piece  $P_1$  except for one newly added inner integral with limits  $[x_1, b_4]$ , as shown in Fig. 2c, because  $x_1 = \max(x_1, a_4)$  for piece  $P_2$ , unlike the piece  $P_1$  in which  $a_4 = \max(x_1, a_4)$ . Thus, we make such updates to inner integrals for each new piece corresponding to a new start point  $a_i \in [a_1, b_{min}]$ .

**Time complexity:** Algorithm 1 initially sorts intervals based on start points  $a_i$  with  $i \in a_1 \dots a_5$  and  $b_{min}$  to determine pieces for inte-

gration (Task 1), which is a constant time operation. Task 2 and Task 3 in Algorithm 2 comprise a single loop, which runs a maximum of five times corresponding to five entries of a sorted interval array  $I_{sorted}$ . Each loop computes the integral template (Algorithm 3) on the fly in constant time. The algorithm is, therefore, linear time complexity with the number of input intervals and extremely efficient.

```

Function integralTemplates()
  Input: Integral limits  $l_1, h_1 = None, h_2 = None, h_3 = None, h_4 = None, h_5 = None$ 
  Input:  $I_{sorted}$  denoting parametric distribution ranges sorted by start points  $a_i$ .
  Output: Integral over piece  $[l_1, h_1]$ 
  /* Determine pdfs parameters  $\theta$  based on intervals sorted based on start points  $a_i$  except for interval  $[a_1, b_1]$  */
   $k = 1$ 
  for  $i = 1$  to 5 do
    if ( $i == id_{a_1}$ ) then
      continue
    else
       $\theta_k.a = I_{sorted}[i][1], \theta_k.b = I_{sorted}[i][2]$ 
       $k = k + 1$ 
    end
  /* Template 1 */
  if  $h_1 \neq None$  and others = None then
     $I = \int_{l_1}^{h_1} pdf_{x,a_1,b_1}$ 
  /* Template 2 */
  else if  $h_1 \neq None$  and  $h_2 \neq None$  and others = None then
     $I = \int_{l_1}^{h_1} \int_{x_1}^{h_2} pdf_{x,a_1,b_1} pdf_{x,\theta_1}$ 
  /* Template 3 */
  else if  $h_1 \neq None$  and  $h_2 \neq None$  and  $h_3 \neq None$  and others = None then
     $I = \int_{l_1}^{h_1} \int_{x_1}^{h_2} \int_{x_1}^{h_3} pdf_{x,a_1,b_1} pdf_{x,\theta_1} pdf_{x,\theta_2}$ 
  /* Template 4 */
  else if  $h_1 \neq None$  and  $h_2 \neq None$  and  $h_3 \neq None$  and  $h_4 \neq None$  and others = None then
     $I = \int_{l_1}^{h_1} \int_{x_1}^{h_2} \int_{x_1}^{h_3} \int_{x_1}^{h_4} pdf_{x,a_1,b_1} pdf_{x,\theta_1} pdf_{x,\theta_2} pdf_{x,\theta_3}$ 
  /* Template 5 */
  else if  $h_1, \dots, h_5 \neq None$  then
     $I = \int_{l_1}^{h_1} \int_{x_1}^{h_2} \int_{x_1}^{h_3} \int_{x_1}^{h_4} \int_{x_1}^{h_5} pdf_{x,a_1,b_1} pdf_{x,\theta_1} pdf_{x,\theta_2} pdf_{x,\theta_3} pdf_{x,\theta_4}$ 
  return  $I$ 
end

```

Algorithm 3: Integral templates for arbitrary piece  $P_i$

## 2.0.2 Local Maximum Probability

Having derived a probabilistic framework for computation of local minimum probability (Sec. 2.0.1), the computation of the local maximum probability  $Pr(p = l_{max})$  at each grid vertex  $p$  is fairly straightforward. Computation of the local maximum probability corresponds to computing  $Pr([(X_1 > X_2) \text{ and } (X_1 > X_3) \text{ and } (X_1 > X_4) \text{ and } (X_1 > X_5)])$ , which is equivalent to computing  $Pr([(X_1 < -X_2) \text{ and } (X_1 < -X_3) \text{ and } (X_1 < -X_4) \text{ and } (X_1 < -X_5)])$ . This negation format is equivalent to Eq. (1). Thus, we negate the intervals for random variables  $X_1, \dots, X_5$  to create new random variables  $X'_1 = -X_1, \dots, X'_5 = -X_5$ . We then apply our proposed local minimum probability computation algorithm (Algorithm 1) to these new random variables  $X'_i$  for computing the local maximum probability in closed form.

## 2.0.3 Saddle Probability

Here, we provide a detailed description of the saddle point probability computation algorithm. As explained in the main paper, we derive only our closed-form computations and algorithm for the term  $t_1 = Pr[(X_1 < X_2) \text{ and } (X_1 > X_3) \text{ and } (X_1 < X_4) \text{ and } (X_1 > X_5)]$  in Eq. 4 of the main paper.

## Function *computeSaddleProbability()*

**Input:** Intervals  $I = [[a_1, b_1], \dots, [a_5, b_5]]$ . Without loss of generality,  $a_2 < a_4$  and  $b_3 < b_5$ . So if  $a_2 > a_4$  or  $b_3 > b_5$  in the original data, then we swap the two intervals. Swapping these intervals do not change the probability computation in Eq.2 of the main paper.

**Output:**  $Pr(p = l_s)$

$a_{max}^{alt} = \max(a_1, a_3, a_5)$

$b_{min}^{alt} = \min(b_1, b_2, b_4)$

**if**  $b_{min}^{alt} \leq a_{max}^{alt}$  **then**  
 | **return** 0

**else**

/\* Task1: Break the outer integral in Eq. 2 of the main paper into pieces  $P_i$  \*/

$sortedInterestPoints = sort([a_{max}^{alt}, a_2, a_4, b_3, b_5, b_{min}^{alt}])$

$id_{a_{max}^{alt}} = \text{Index of } a_{max}^{alt} \text{ in } sortedInterestPoints$

$id_{b_{min}^{alt}} = \text{Index of } b_{min}^{alt} \text{ in } sortedInterestPoints$

/\* Task2 and Task 3: Compute piece integrals \*/

$saddleProbability =$

$computePieceIntegrals(id_{a_{max}^{alt}}, id_{b_{min}^{alt}}, sortedInterestPoints, I)$

**return**  $saddleProbability$

**end**

Algorithm 4: Computation of saddle probability

$$\begin{aligned}
 t_1 &= Pr[(X_1 < X_2) \text{ and } (X_1 > X_3) \text{ and } (X_1 < X_4) \text{ and } (X_1 > X_5)] \\
 &= \int_{x_1=a_{max}^{alt}}^{x_1=b_{min}^{alt}} \int_{x_2=\max(x_1, a_2)}^{x_2=b_2} \int_{x_3=\min(x_1, b_3)}^{x_3=a_3} \int_{x_4=\max(x_1, a_4)}^{x_4=b_4} \int_{x_5=\min(x_1, b_5)}^{x_5=a_5} \dots \\
 &(\text{Pdf}_{joint}) dx,
 \end{aligned}$$

where  $a_{max}^{alt} = \max(a_1, a_3, a_5)$ ,  $b_{min}^{alt} = \min(b_1, b_2, b_4)$

(2)

We now explain the integral limits in Eq. (2) and the piecewise integration algorithm to compute the formula in Eq. (2), similar to our explanations for the local minimum probability computation in Sec. 2.0.1.

Limits  $a_{max}^{alt}$  and  $b_{min}^{alt}$  of the outer integral in Eq. (2): First, we identify the portion of data range of a random variable  $X_1$  (i.e.,  $[a_1, b_1]$ ) that can result in point  $p$  being a saddle. The portion  $[a_{max}^{alt}, b_{min}^{alt}]$  of outer integral in Eq. (2) indicates the data values that can result in point  $p$  being saddle, where  $a_{max}^{alt}$  denotes the maximum value among  $a_1, a_3$ , and  $a_5$ ,  $b_{min}^{alt}$  denotes the minimum value among  $b_1, b_2$ , and  $b_4$ , and  $a_{max}^{alt} < b_{min}^{alt}$ . The superscript *alt* represents alternate neighboring random variables of a random variable  $X_1$  under consideration. The data portion  $[a_1, a_{max}^{alt}]$  of random variable  $X_1$  with  $a_1 < a_{max}^{alt}$  will always be smaller than either  $X_3$  or  $X_5$  depending on if  $a_{max}^{alt}$  is equal to  $a_3$  or  $a_5$ , respectively. Thus, the range  $[a_1, a_{max}^{alt}]$  cannot result in point  $p$  as a saddle. Similarly, the data portion  $[b_{min}^{alt}, b_1]$  of random variable  $X_1$  with  $b_{min}^{alt} < b_1$  will always be greater than either  $X_2$  or  $X_4$  depending on if  $b_{min}^{alt}$  is equal to  $b_2$  or  $b_4$ , respectively. Thus, the range  $[b_{min}^{alt}, b_1]$  cannot result in point  $p$  as a saddle. To the contrary, each point in the range  $[x_1 = a_{max}^{alt}, x_1 = b_{min}^{alt}]$  has a nonzero probability of simultaneously being smaller than the random variables  $X_2$  and  $X_4$  and greater than the random variables  $X_3$  and  $X_5$ , and therefore, represents a valid range for point  $p$  being a saddle.

Limits  $\max(x_1, a_i)$  and  $\min(x_1, b_i)$  of the inner integrals in Eq. (2): Inner integrals in Eq. (2) integrate the joint distribution Pdf<sub>joint</sub> over its support where random variables  $X_2$  and  $X_4$  are simultaneously greater and  $X_3$  and  $X_5$  are simultaneously smaller than  $x_1 \in [a_{max}^{alt}, b_{min}^{alt}]$  in the outer integral. The inner integral lower limit

**Function** *computePieceIntegrals()*

```

Input:  $id_{a_{max}^{alt}}, id_{b_{min}^{alt}}, sortedInterestPoints, I$ 
Output: Sum of integrals over pieces  $P_i$ 

/* Task2: Compute integral for piece  $P_1$  */
/* Lower (l) and upper (h) limits of piece  $P_1$  */
l = sortedInterestPoints[id_{a_{max}^{alt}}]
h = sortedInterestPoints[id_{a_{max}^{alt}} + 1]
totalIntegral = 0
/* Initialize the limits of inner integrals
for piece  $P_1$  assuming  $a_2, a_4$  starting after
 $a_{max}^{alt}$  and  $a_3, a_5$  are starting before  $a_{max}^{alt}$  */
tempd3 = a_3
tempd5 = a_5
tempu2 = None
tempu4 = None
/* Update the initialized limits depending on
which of interest points  $a_2, a_4, b_3, b_5$  lie
before  $a_{max}^{alt}$ . */
for k = 1 to (id_{a_{max}^{alt}} - 1) do
  if sortedInterestPoints[k] == a_2 then
    tempu2 = I[2][2]; // i.e., b_2
  if sortedInterestPoints[k] == a_4 then
    tempu4 = I[4][2]; // i.e., b_4
  if sortedInterestPoints[k] == b_3 then
    tempd3 = None
  if sortedInterestPoints[k] == b_5 then
    tempd5 = None
end
I_{P_1} = integralTemplate(l, h, l3 = tempd3, l5 =
tempd5, h2 = tempu2, h4 = tempu4, I)
totalIntegral += I_{P_1}

/* Task3: Compute integral for successive
pieces */
for k = (id_{a_{max}^{alt}} + 1) to (id_{b_{min}^{alt}}) do
  /* Determine piece limits */
  l = sortedInterestPoints[k]
  h = sortedInterestPoints[k + 1]
  /* Update the lower or upper limit for
each new interest point */
  if sortedInterestPoints[k] == a_2 then
    tempu2 = I[2][2]; // i.e., b_2
  if sortedInterestPoints[k] == a_4 then
    tempu4 = I[4][2]; // i.e., b_4
  if sortedInterestPoints[k] == b_3 then
    tempd3 = None
  if sortedInterestPoints[k] == b_5 then
    tempd5 = None
  I_{P_{next}} = integralTemplate(l, h, l3 = tempd3, l5 =
tempd5, h2 = tempu2, h4 = tempu4, I)
  totalIntegral += I_{P_{next}}
end
return totalIntegral

```

end

**Algorithm 5: Computation of integral for piece  $P_1$  and successive pieces  $P_{next}$** 

corresponds to  $\max(x_1, a_i)$  for  $i \in \{2, 4\}$  and upper limit corresponds to  $\min(x_1, b_i)$  for  $i \in \{3, 5\}$ . These integral limits are derived using a reasoning similar as in the case of local minimum probability described Sec. 2.0.1. The maximum or minimum is taken because the support of random variable  $X_i$  is restricted to  $[a_i, b_i]$ . In particular, for  $x_1 < a_i$  in inner integral with  $i \in \{2, 4\}$ , the support  $[a_i, b_i]$  will always be greater than  $x_1$  and the inner integral does not depend on the value of  $x_1$ . Similarly, for  $x_1 > b_i$  in any inner integral with  $i \in \{3, 5\}$ , the support  $[a_i, b_i]$  will always be smaller than  $x_1$ , and the inner integral does not depend on the value of  $x_1$ . In contrast, when  $x_1$  assumes values in the support

**Function** *integralTemplates()*

```

Input: Integral limits
l, h, l3 = None, l5 = None, h2 = None, h4 = None
Output: Integral over piece  $[l, h]$ 
/* Template 1 */
if l3 ≠ None and others = None then
  I = ∫_l^h ∫_{l3}^{x_1} pdf_{x,a_1,b_1} pdf_{x,a_3,b_3}
/* Template 2 */
else if l5 ≠ None and others = None then
  I = ∫_l^h ∫_{l5}^{x_1} pdf_{x,a_1,b_1} pdf_{x,a_5,b_5}
/* Template 3 */
else if h2 ≠ None and others = None then
  I = ∫_l^h ∫_{x_1}^{h2} pdf_{x,a_1,b_1} pdf_{x,a_2,b_2}
/* Template 4 */
else if h4 ≠ None and others = None then
  I = ∫_l^h ∫_{x_1}^{h4} pdf_{x,a_1,b_1} pdf_{x,a_4,b_4}
/* Template 5 */
else if l3 ≠ None and l5 ≠ None and others = None then
  I = ∫_l^h ∫_{l3}^{x_1} ∫_{l5}^{x_1} pdf_{x,a_1,b_1} pdf_{x,a_3,b_3} pdf_{x,a_5,b_5}
/* Template 6 */
else if l3 ≠ None and h2 ≠ None and others = None then
  I = ∫_l^h ∫_{l3}^{x_1} ∫_{x_1}^{h2} pdf_{x,a_1,b_1} pdf_{x,a_3,b_3} pdf_{x,a_2,b_2}
/* Template 7 */
else if l3 ≠ None and h4 ≠ None and others = None then
  I = ∫_l^h ∫_{l3}^{x_1} ∫_{x_1}^{h4} pdf_{x,a_1,b_1} pdf_{x,a_3,b_3} pdf_{x,a_4,b_4}
/* Template 8 */
else if l5 ≠ None and h2 ≠ None and others = None then
  I = ∫_l^h ∫_{l5}^{x_1} ∫_{x_1}^{h2} pdf_{x,a_1,b_1} pdf_{x,a_5,b_5} pdf_{x,a_2,b_2}
/* Template 9 */
else if l5 ≠ None and h4 ≠ None and others = None then
  I = ∫_l^h ∫_{l5}^{x_1} ∫_{x_1}^{h4} pdf_{x,a_1,b_1} pdf_{x,a_5,b_5} pdf_{x,a_4,b_4}
/* Template 10 */
else if h2 ≠ None and h4 ≠ None and others = None then
  I = ∫_l^h ∫_{x_1}^{h2} ∫_{x_1}^{h4} pdf_{x,a_1,b_1} pdf_{x,a_2,b_2} pdf_{x,a_4,b_4}
/* Template 11 */
else if l3 ≠ None and l5 ≠ None and h2 ≠ None and
h4 = None then
  I =
∫_l^h ∫_{l3}^{x_1} ∫_{l5}^{x_1} ∫_{x_1}^{h2} pdf_{x,a_1,b_1} pdf_{x,a_3,b_3} pdf_{x,a_5,b_5} pdf_{x,a_2,b_2}
/* Template 12 */
else if l3 ≠ None and l5 ≠ None and h4 ≠ None and
h2 = None then
  I =
∫_l^h ∫_{l3}^{x_1} ∫_{l5}^{x_1} ∫_{x_1}^{h4} pdf_{x,a_1,b_1} pdf_{x,a_3,b_3} pdf_{x,a_5,b_5} pdf_{x,a_4,b_4}
/* Template 13 */
else if l3 ≠ None and h2 ≠ None and h4 ≠ None and
l5 = None then
  I =
∫_l^h ∫_{l3}^{x_1} ∫_{x_1}^{h2} ∫_{x_1}^{h4} pdf_{x,a_1,b_1} pdf_{x,a_3,b_3} pdf_{x,a_2,b_2} pdf_{x,a_4,b_4}
/* Template 14 */
else if l5 ≠ None and h2 ≠ None and h4 ≠ None and
l3 = None then
  I =
∫_l^h ∫_{l5}^{x_1} ∫_{x_1}^{h2} ∫_{x_1}^{h4} pdf_{x,a_1,b_1} pdf_{x,a_5,b_5} pdf_{x,a_2,b_2} pdf_{x,a_4,b_4}
/* Template 15 */
else if l3 ≠ None and l5 ≠ None and h2 ≠ None and
h4 ≠ None then
  I = ∫_l^h ∫_{l3}^{x_1} ∫_{l5}^{x_1} ∫_{x_1}^{h2} ∫_{x_1}^{h4} pdf_{x,a_1,b_1} pdf_{x,a_3,b_3}
pdf_{x,a_5,b_5} pdf_{x,a_2,b_2} pdf_{x,a_4,b_4}
return I

```

end

**Algorithm 6: Integral templates for arbitrary piece  $P_i$** 

of a distribution, the inner integral then depends on  $x_1$ . It can also be verified that the lower limit of any inner integral does not exceed the

upper limit of their respective integral. For example,  $a_3 \leq \min(x_1, b_3)$  in all cases because our initial assumption is  $a_3 < b_3$  for the random variable  $X_3$  (see the Background and Problem Setting section of the main paper). Also,  $x_1$  in the outer integral of Eq. (2) attains values in the range  $[a_{max}^{alt} = \max(a_1, a_3, a_5), b_{min}^{alt} = \min(b_1, b_2, b_4)]$  (see the previous paragraph). Since the minimum value  $x_1$  is equal to  $\max(a_1, a_3, a_5)$ ,  $x_1 \geq a_3$  is guaranteed. Thus,  $a_3 \leq \min(x_1, b_3)$  is true in all cases. A similar reasoning is applicable to the other inner integrals too.

**Piecewise simplification of Eq. (2):** The saddle probability again needs to be computed in a piecewise manner for reasons similar to the case of local minimum probability computation described in Sec. 2.0.1. Essentially, at each start point  $a_i$  with  $i \in \{2, 4\}$  in Eq. (2), the value of  $\max(x_1, a_i)$  changes, which simplifies the integral computation differently. Similarly, for each end point  $b_i$  with  $i \in \{3, 5\}$  in Eq. (2), the value of  $\min(x_1, b_i)$  changes, which simplifies the integral computation differently. The simplified formula of a piecewise integral, therefore, depends on the ordering of the points  $a_2, a_4, b_3,$  and  $b_5$  with respect to the limits of the outer integral in Eq. (2), i.e.,  $a_{max}^{alt}$  and  $b_{min}^{alt}$ .

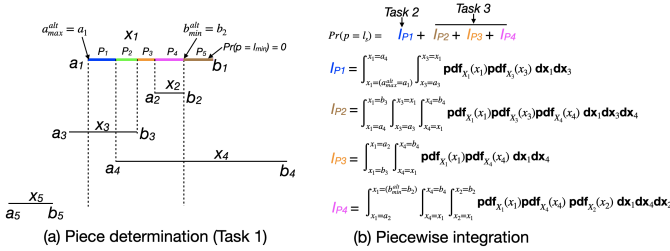


Fig. 3: Illustration of saddle probability computation. (a) Pieces are determined as a part of Task 1. Here, pieces are  $P_1 = [a_{max}^{alt} = a_1, a_4]$ ,  $P_2 = [a_4, b_3]$ ,  $P_3 = [b_3, a_2]$ , and  $P_4 = [a_2, b_{min}^{alt} = b_2]$ . (b) The saddle probability is computed by computing integral for the first piece  $P_1$  (Task 2) followed by integrals for successive pieces  $P_{next}$  (Task 3) and summing all piecewise integrals.

**Algorithm for computing saddle probability:** The computation of the formula in Eq. (2) depends on the ordering of points  $a_2, a_4, b_3,$  and  $b_5$  with respect to the limits of the outer integral in Eq. (2), i.e.,  $a_{max}^{alt}$  and  $b_{min}^{alt}$ . Because of these six quantities, there are again  $6! = 720$  permutations similar to the local minimum probability case. We, therefore, devise an efficient algorithm that can compute the integral in Eq. (2) without needing to go through all permutations. Our algorithm matches closely with the one for the local minimum probability computation. Our algorithm computes the simplified integrals per piece  $P_i$  on the fly depending on the observed permutation of points  $a_2, a_4, b_3, b_5, a_{max}^{alt} = \max(a_1, a_3, a_5)$ , and  $b_{min}^{alt} = \min(b_1, b_2, b_4)$ . If  $b_{min}^{alt} \leq a_{max}^{alt}$ , the saddle probability is 0. If  $a_{max}^{alt} < b_{min}^{alt}$ , we divide our algorithm into three tasks. In Task 1, we determine the pieces  $P_i$  of the outer integral in Eq. (2) for performing piecewise integration (see Task 1 in Algorithm 4). For that, we sort the intervals based on the ordering of  $a_2, a_4, b_3, b_5, a_{max}^{alt} = \max(a_1, a_3, a_5)$ , and  $b_{min}^{alt} = \min(b_1, b_2, b_4)$ . This operation is similar to the  $I_{sorted}$  array computation in the case of local minimum probability computation in Algorithm 1. We then find the indices of  $a_{max}^{alt}$  and  $b_{min}^{alt}$  in the sorted intervals and points in between, which determine the pieces  $P_i$ .

In Task 2, we compute the integration for the first piece  $P_1$  depending on which points in  $a_2, a_4, b_3,$  and  $b_4$  are starting before or after  $a_{max}^{alt}$ . The algorithm for integration for the first piece  $P_1$  is presented in Algorithm 5 as Task 2 (similar to the Task 2 in the case of local minimum probability computation in Algorithm 2). Finally, we compute the integration for successive pieces (denoted as  $I_{P_{next}}$  in Algorithm 5) depending on the order of points  $a_2, a_4, b_3,$  and  $b_4$  observed between  $a_{max}^{alt}$  and  $b_{min}^{alt}$ , which is similar to the Task 3 in Algorithm 2 for the local minimum probability computation. Again, we derive the integral templates (i.e., simplifications) similar to those in the case of local minimum probability computation, which are detailed in Algorithm 6.

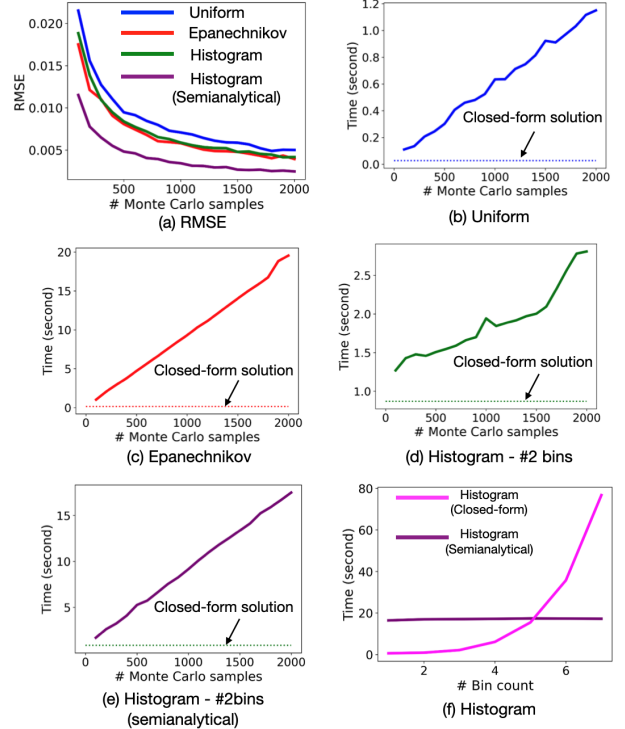


Fig. 4: Quantitative proof of correctness of our proposed closed-form computations and performance results. The RMSE between our closed-form/semianalytical solutions and the MC solution (image a) drops with the increase in the number of MC samples, which confirms the correctness of our algorithms. As depicted in images b-e, the closed-form computation (dotted line) provides significantly high performance compared to the MC sampling (solid curves). The closed-form histogram computation time exponentially grows with an increase in the number of bins, but the semianalytical solution time stays about constant (image f) for 1000 MC samples.

These integration templates compute the integral for each piece on the fly based on the observed piece limits.

**Illustration of saddle probability computation:** We illustrate the three tasks comprising the saddle probability computation in Fig. 3. For the example shown in Fig. 3,  $a_5 < b_5 < a_3 < a_1 < a_4 < b_3 < a_2 < b_2 < b_1 < b_4$ . The Task 1 includes determining the portion of random variable  $X_1$  that can result in point  $p$  being a saddle and determining pieces for integration. As observed in Fig. 3a, each point outside the range  $[x_1 = a_{max}^{alt}, x_1 = b_{min}^{alt}]$  has a zero probability of being a saddle. For example, in Fig. 3a, the range  $[x_1 = b_{min}^{alt}, b_1]$  (i.e.,  $[b_2, b_1]$  shown in brown) has a zero probability of being smaller than the random variable  $X_2$ , which violates the condition of a saddle in Eq. (2). Thus, only the data range  $[x_1 = a_{max}^{alt}, x_1 = b_{min}^{alt}]$  can result in point  $p$  being a saddle.

We then determine pieces of integration. For the example illustrated in Fig. 3a, the ordering of points of interest (points involved in the integration limits of Eq. (2)) that determines the piece limits is  $b_5 < a_{max}^{alt} = a_1 < a_4 < b_3 < a_2 < b_{min}^{alt} = b_2$ . Thus, there are four pieces  $P_1 = [a_{max}^{alt} = a_1, a_4]$ ,  $P_2 = [a_4, b_3]$ ,  $P_3 = [b_3, a_2]$ ,  $P_4 = [a_2, b_{min}^{alt} = b_2]$ , depicted by four colors in Fig. 3a. The piecewise integration process is again similar to the local minimum probability computation described in Sec. 2.0.1.

In Task 2, we compute the integration formula for piece  $P_1 = [a_1, a_4]$  denoted by a blue range in Fig. 3a. We determine how many intervals are overlapping with the piece  $P_1$  to compute integration over piece  $P_1$ , i.e.,  $I_{P_1}$ . The process of finding which intervals are overlapping with the piece  $P_1$  is similar to the one in the case of local minimum probability computation (the only difference is that the points of interest in the case of local minimum probability computation are  $a_1, \dots, a_5$  and  $b_{min}$ ). Since  $b_5 < (a_{max}^{alt} = a_1)$  and  $a_4, a_2 > a_1$  in Fig. 3a, the intervals of random variables  $X_5, X_4,$  and  $X_2$  do not overlap with the piece  $P_1$ .

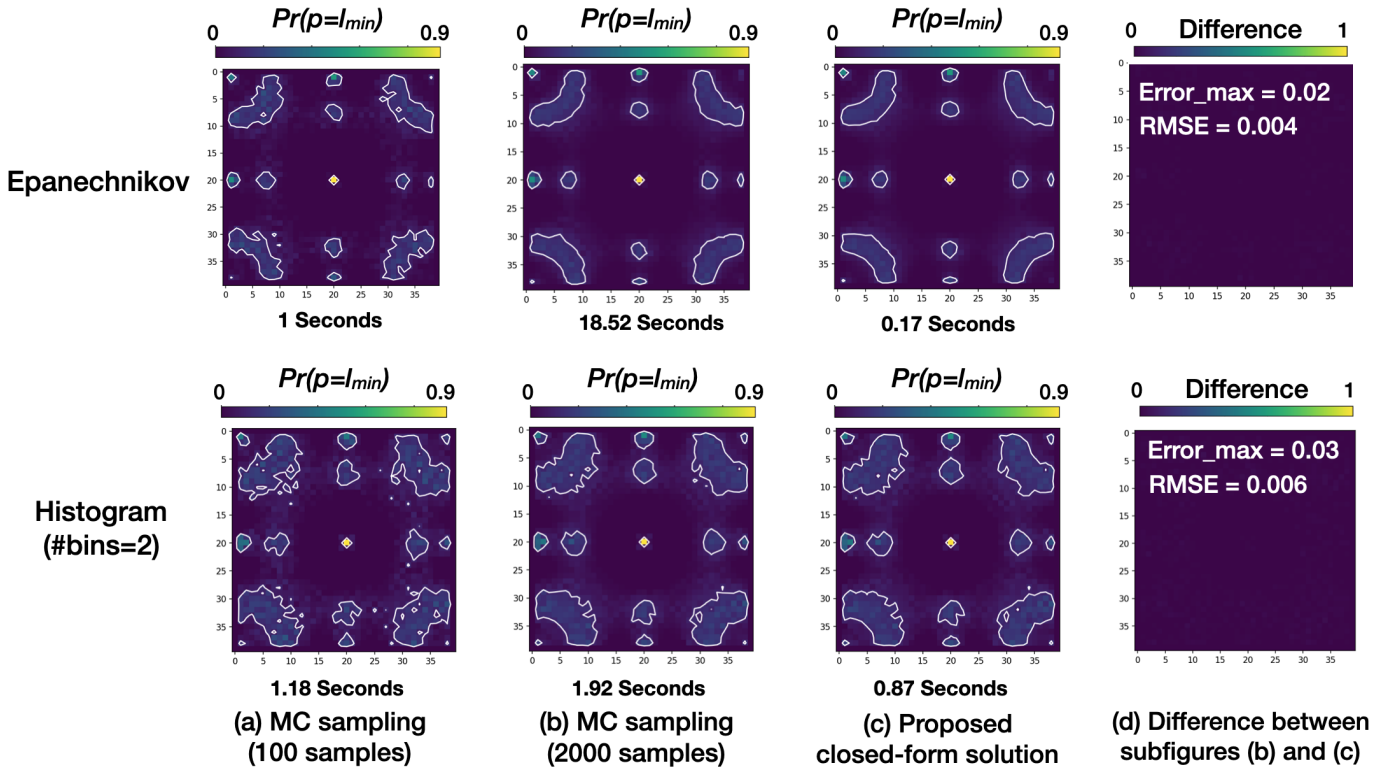


Fig. 5: Qualitative proof of correctness of our proposed closed-form computations (column c) for the Epanechnikov (top row) and Histogram models (bottom row) using the MC sampling approach (columns a and b) as the baseline. The solution obtained with 2000 MC samples converges to our closed-form computations (see the difference image in column d), thereby confirming the correctness of our methods and implementation. Our method increases the speed for both Epanechnikov and histogram with 2 bins. Note that the closed-form histogram computation, although accurate, exponentially grows with an increase in the number of bins.

Only the interval for random variable  $X_3$ , i.e.,  $[a_3, b_3]$  overlaps with the piece  $P_1$ . Thus, the core integration formula in Eq. (2) simplifies to a double integral over the marginal joint distribution of random variables  $X_1$  and  $X_3$ , as shown with the formula for  $I_{P_1}$  in Fig. 3b.

In Task 3, the integration is computed for successive pieces. In particular, for the pieces  $P_2$  (green),  $P_3$  (orange), and  $P_4$  (magenta) in Fig. 3a, the integral updates based on if a new piece results from a new start point (i.e.  $a_i$  with  $i \in \{2, 4\}$ ) or end of an interval (i.e.  $b_i$  with  $i \in \{3, 5\}$ ). For example, for the piece  $P_2$ , the start point  $a_4$  causes the interval for random variable  $X_4$ , i.e.,  $[a_4, b_4]$  to overlap with the piece  $P_2$  in addition to the interval for random  $X_3$ , i.e.,  $[a_3, b_3]$ . Thus, the core integration formula in Eq. (2) simplifies to a triple integral over the marginal joint distribution of random variables  $X_1$ ,  $X_3$ , and  $X_4$ , as shown with the formula for  $I_{P_2}$  in Fig. 3b. On the contrary, the end point  $b_3$  terminates the overlap of the interval  $[a_3, b_3]$  with the piece  $P_3$ . Thus, the core integration formula in Eq. (2) simplifies to a double integral over the marginal joint distribution of random variables  $X_1$  and  $X_4$ , as shown with the formula for  $I_{P_3}$  in Fig. 3b. The integration formula for the piece  $P_4$  updates in a similar manner because of a new start point  $a_2$ .

### 3 VALIDATION AND PERFORMANCE OF THE PROPOSED ALGORITHMS

We show the quantitative results for the uniform, Epanechnikov, and histogram noise models and qualitative results for Epanechnikov and histogram noise models (similar to Fig. 6 in the main paper for the uniform noise) for the local minimum probability computations on the Ackley dataset [1] used in the main paper. Figure 4 shows the quantitative comparison of MC sampling solution and our closed-form computations using the proposed algorithms. For all uniform, Epanechnikov, and histogram (closed-form and semianalytical) noise models, the root mean squared error (RMSE) between the MC and closed-form

solution drops as the number of MC samples is increased, as observed in Fig. 4a. This convergence confirms the correctness of our closed-form computations and algorithms. In Fig. 4a, the semianalytical (purple curve) solution has a lower RMSE than the histogram with MC sampling (green curve) for all neighbors per grid point, thereby showing the higher accuracy of our semianalytical approach. Further, as seen from Fig. 4b-e, the closed-form solutions provide the highest performance (dotted line) compared to the MC method (solid curves). As observed in Fig. 4f, the closed-form histogram computation time exponentially grows with an increase in the number of bins, as described in the main paper. The semianalytical histogram solution (the purple curve in Fig. 4f), however, takes an about constant amount of time for a fixed number of MC samples because it does not depend on the bin count (see the main paper for details). Figure 5 visualizes the qualitative results similar to the Fig. 6 of the main paper, but for the Epanechnikov and histogram noise models. It shows that the MC solution converges to our closed-form solution as we increase the number of samples from 100 (Fig. 5a) to 2000 (Fig. 5b). This qualitative convergence can be seen through the difference image in Fig. 5d that exhibits negligible maximum error ( $Error_{max}$ ) and RMSE between Fig. 5c and Fig. 5b.

### 4 QUANTITATIVE EVALUATION FOR THE CLIMATE DATA

In Fig. 6, we present a quantitative evaluation for the climate data [4] used in the main paper. First, we computed the difference  $D$  between the number of critical points of the decompressed and original field as a function of probability threshold. In particular, at probability threshold  $t$ , we consider all critical points of decompressed data that have probability greater than  $t$ . Thus, for  $t = 0$ , all critical points of a decompressed field are considered. However, as we increase  $t$ , critical points with probability smaller than  $t$  get filtered out. For a probability threshold  $t = 0$ , the difference  $D$  is high, as all critical points of decompressed field are considered (which contain several

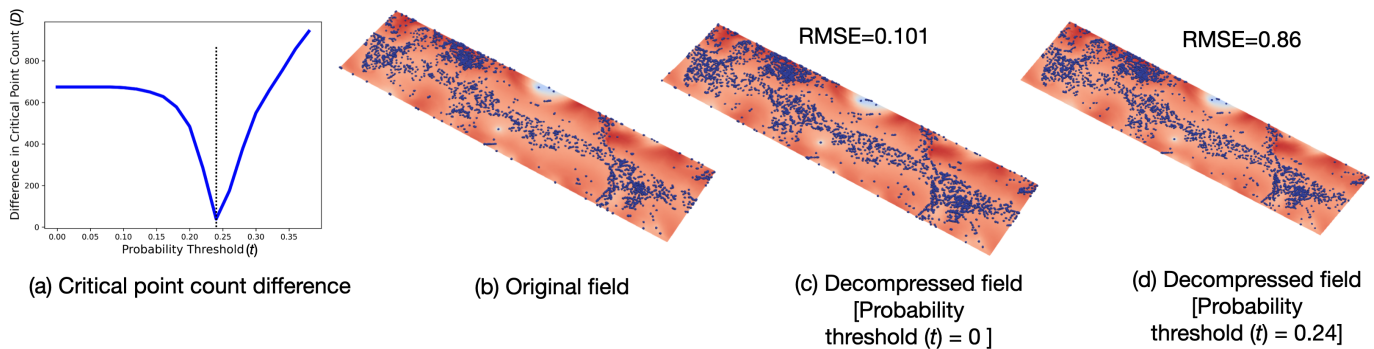


Fig. 6: Quantitative evaluation of the climate data. (a) Difference ( $D$ ) between the critical point count of original and decompressed fields plotted as a function of a probability threshold ( $t$ ). In particular, for a threshold  $t$ , all critical points with probability smaller than  $t$  are removed in the decompressed field. At threshold  $t = 0.24$ , the number of critical points of the original and decompressed fields is nearly same. (b) Critical points of the original field. (c) Critical points of the decompressed data without applying filtering ( $t = 0$ ). (d) Critical points of the decompressed data filtered by probability threshold of  $t = 0.24$ . The filtered critical points in (d) exhibit lower error (RMSE = 0.86) with respect to the nonfiltered critical points in (c) (RMSE = 0.101) with the original field in (a) as the reference, which is illustrative of the utility of probabilistic methods in identifying robust critical points.

new critical points from compression errors). As we start increasing the threshold, the difference  $D$  drops and reaches the minimum at a probability threshold  $t = 0.24$ , as observed in Fig. 6a. For probability threshold  $t > 0.24$ , the difference  $D$  again increases because the number of thresholded critical points in the decompressed field become smaller than the number of critical points in the original field. Thus, at  $D = 0.24$ , both the original and decompressed fields have about the same number of critical points.

We compute the root mean squared error (RMSE) to quantitatively evaluate probabilistic results. We consider the decompressed data without critical point filtering ( $t = 0$ ) in Fig. 6c and with critical point filtering ( $t = 0.24$ ) in Fig. 6d for evaluation. Initially, we assign 1 to each position in the original field where the critical point appears and 0 everywhere else to create a binary field. We generate similar binary fields denoting critical point positions for thresholds  $t = 0$  (decompressed data with all critical points) and  $t = 0.24$  (decompressed data with critical points that have probability greater than  $t = 0.24$ ). We then compute the RMSE between these binary image representations. The RMSE for nonfiltered critical points in Fig. 6c with respect to original data critical points is 0.101. This RMSE drops to 0.86 as we filter critical points based on probability in Fig. 6d, which indicates that removing low-probability critical points from decompressed data provides answer closer to the true critical points in the original field, thereby presenting the utility of probability computations for critical points.

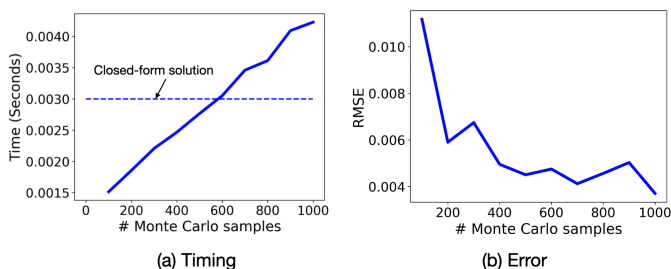


Fig. 7: Performance and accuracy results of our VTK-m implementation on the AMD GPU. (a) The performance reduces with an increase in the number of MC samples. (b) The RMSE between the MC solution and solution of our algorithms reduces with an increase in the number of MC samples, thereby confirming the correctness of our algorithms and parallel implementation.

The performance and accuracy results for the climate dataset using our VTK-m parallel implementation on the AMD GPU are shown in Fig. 7. The AMD GPU resource is courtesy of Oak Ridge National

Laboratory’s Frontier supercomputer [3]. The performance of the MC solutions decreases with an increase in the number of samples, as observed from Fig. 7a. The decrease in RMSE between the MC solution and our closed-form solution with an increase in the number of MC samples, as observed in Fig. 7b, confirms the correctness of our algorithms.

## 5 INTEGRATION OF VTK-M CODE WITH PARAVIEW

One of the main benefits offered by VTK-m [5] includes its easy integration with ParaView using a plugin approach, thereby making VTK-m filters accessible to a broader community. Figure 1 visualizes the integration of our VTK-m critical point uncertainty code with ParaView. The ParaView plugin is generated by wrapping a VTK-m class inside VTK class, which can then be integrated in ParaView. Using GPU as a backend, reliable uncertainty visualizations can be generated using the proposed nonparametric models (i.e., histograms) in near-real-time within ParaView. We provide a small video demonstration as another supplement to show the usage of our VTK-m critical point uncertainty filter inside ParaView.

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