

A polynomial-based Monte Carlo approach for estimating long-term collision probabilities

Andrea Zollo^{a,*}, Cristina Parigini^b, Roberto Armellin^b, Juan Félix San Juan Díaz^c, Annarita Trombetta^a, Ralph Kahle^a

^a German Space Operations Centre (GSOC), German Aerospace Centre DLR, Münchener Str. 20, 82234, Weßling, Germany

^b Te Pūnaha Atea - Space Institute, University of Auckland, Private Bag 92019, 1142, Auckland, New Zealand

^c Scientific Computation Research Institute, University of La Rioja, Madre de Dios 53, La Rioja, 26006, Logroño, Spain

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ABSTRACT

This paper introduces a versatile approach for computing the risk of collision specifically tailored for scenarios featuring low relative encounter velocities, but with potential applicability across a wide range of situations. The technique employs Differential Algebra (DA) to express the non-linear dynamical flow of the initial distribution in the primary-secondary objects relative motion through high-order Taylor polynomials. The entire initial uncertainty set is subdivided into subsets through Automatic Domain Splitting (ADS) techniques to control the accuracy of the Taylor expansions. The methodology samples the initial conditions of the relative state and evaluates the polynomial expansions for each sample while retaining their temporal dependency. The classical numerical integration of the initial statistics over the set of conditions for which a collision occurs is thus reduced to an evaluation of mono-dimensional time polynomials. Specifically, samples reaching a relative distance below a critical value are identified along with the time at which this occurs. The approach is tested against a Monte Carlo (MC) simulation for various literature test cases, yielding accurate results and a consistent gain in computational time. Specifically, the time reduction can reach up to 99.9 %, depending on the dynamics and the number of samples employed in the analysis, while the estimated final Probability of Collision (P_c), for long-term scenarios, deviates from the reference by no more than 0.018 %.

1. Introduction

With the continuous advancement of space technology, the frequency of spacecraft launches has increased steadily, leading to growing congestion in Earth's orbital environment. Over recent decades, this congestion has become increasingly hazardous to spacecraft operations, primarily due to the rising accumulation of space debris. Projections indicate that this trend will intensify, particularly with the deployment of large-scale satellite constellations, which will further saturate the most frequently used orbital regimes. Consequently, spacecraft operators are confronted with a growing number of close approach alerts and are required to perform an increasing number of Collision Avoidance Maneuvers (CAMs) to safeguard mission integrity [1,2].

A key aspect of managing these events is the assessment of their criticality, which mainly relies on evaluating the Probability of Collision (P_c) between the objects involved [3]. In the literature and collision

avoidance practice, these are referred to as the primary (typically the controlled asset of interest) and the secondary object (the one it encounters).

The methods for computing collision risk have been tailored for different conjunction types. Close approaches between satellites are in fact typically classified as either short-term or long-term encounters [4]. Short-term encounters involve objects with significantly different orbits, resulting in encounter velocities reaching several kilometers per second near the point of closest approach. These encounters last only a few seconds at most. Throughout the encounter, the relative velocity vector remains constant in both intensity and direction, leading to a straight-line relative trajectory. Moreover, the relative velocity uncertainty is deemed negligible in comparison to its pronounced mean. Consequently, the position error combined ellipsoid remains stable throughout the encounter since the positional uncertainties of the objects can be defined by two uncorrelated constant covariance matrices.

* Corresponding author.

E-mail addresses: andrea.zollo@dlr.de (A. Zollo), cristina.parigini@auckland.ac.nz (C. Parigini), roberto.armellin@auckland.ac.nz (R. Armellin), juanfelix.sanjuan@unirioja.es (J.F.S.J. Díaz), annarita.trombetta@dlr.de (A. Trombetta), ralph.kahle@dlr.de (R. Kahle).

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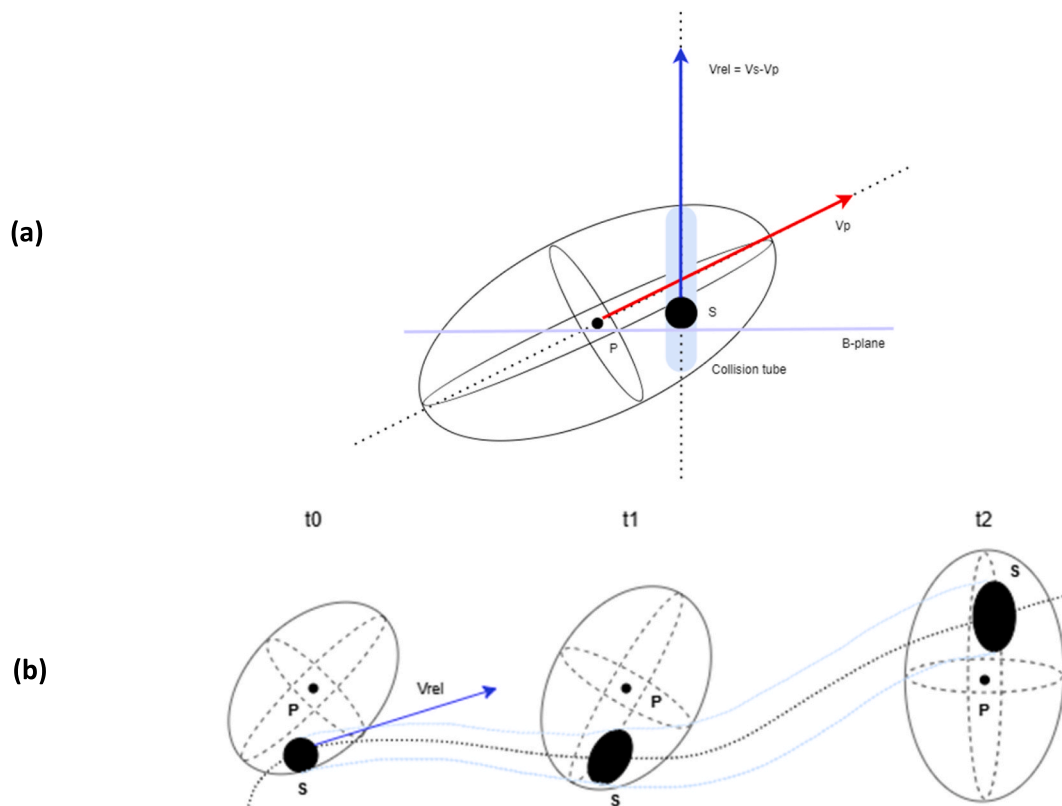


Fig. 1. representation of short-term (a) and long-term (b) encounter scenarios. By convention, the combined covariance position ellipsoid is centered on the primary object (P), while a sphere of radius equal to the Hard-Body Radius (HBR) is centered on the secondary object (S), accounting for the dimensions of both objects. In long-term encounters, the topological points on the hard-body sphere's surface at t_0 evolve into an ellipsoid as the combined covariance rotates and deforms over time.

Various methods are available in the literature to compute the collision probability for the short-term case. The problem has been, in fact, extensively studied by Foster [5], Patera [6,7], Alfriend et al. [8], Afano [9], Chan [10,11], and more recently by Serra et al. [12].

Typically, as illustrated in Fig. 1-(a), the collision risk is calculated by integrating the Probability Density Function (PDF) of the combined positional uncertainty over the volume defined by the combined hard-body sphere [6], as it moves along the relative trajectory. Given that this trajectory is rectilinear, the swept volume can be approximated as an infinite cylinder aligned with the direction of the relative velocity. This allows the 3D integral to be reduced to a 2D one, since the integration along the direction of relative velocity yields unity. As a result, the PDF is usually evaluated on a plane perpendicular to the relative velocity vector, commonly referred to as the B-plane.

The other type of encounter occurs between two satellites traveling along neighboring orbits, such as between two GEO satellites at adjacent longitude positions or, more generally, during close satellite operations, including rendezvous, formation, and cluster flights. It is worth noting that similar geometries can also occur naturally, albeit less frequently. These close approaches are characterized by a low relative velocity, typically on the order of a few meters per second. In such scenarios, the two objects remain in close proximity for an extended duration, approximately on the order of the orbital period. As shown in Fig. 1-(b), unlike in short-term close approaches, the relative velocity vector is not constant; it evolves over time in both direction and magnitude. For that, the relative trajectory usually bends and becomes non-linear. The uncertainty in the relative state can no longer be assumed constant; it also changes over time rotating and deforming. Additionally, the uncertainty associated with the relative velocity must be accounted for and cannot be neglected anymore. This results in a time-dependent, evolving combined covariance matrix during the encounter, and the collision tube can

exhibit complex, highly non-linear geometries that are very difficult to integrate.

The collision risk for long-term encounters has not been as thoroughly investigated as it has been for short-term encounters. A first category of methods, as in Refs. [13–15], tries to solve the bending tendency of the collision tube. The key concept is to discretize the collision tube into small subsections and to consider that for each segment the assumptions of a short-term encounter still hold. A better characterization of the collision volume is outlined in the works of Chan [16,17], where the swept-volume of the hard-body is described as an envelope of ellipsoids. However, it is important to note that these methods do not account for velocity uncertainties in the formulation of the problem. Coppola [18], on the other hand, presents a comprehensive mathematical framework that, for the first time, incorporates velocity uncertainties into the formulation. This marks one of the most extensive efforts to address the collision probability problem in a general manner. In this case, the intricate integration volume is continually mapped over time through the dynamic evolution of the initial conditions on the 3D surface of the hard-body sphere. However, the assumptions made do not accommodate for multiple encounters within the analysis timeframe, making the method not suitable for complex intersections of the integration volume.

As highlighted in Ref. [19], there is a notable absence in the literature of a general method capable of simultaneously characterizing the swept volume, especially when its shape is intricate, and computing the subsequent integral of the full relative state vector gaussian PDF over such a volume. In their work, it is suggested to approximate the swept volume using a Polynomial Superlevel Set (PSS) followed by a Monte Carlo integration to calculate the P_c .

So far, the most comprehensive methodologies capable of simultaneously accounting for uncertainties in both position and velocity, while

handling complex encounter geometries, are Monte Carlo (MC) techniques [20,21]. These methods offer a straightforward problem formulation and an intuitive approach to computing the P_c , without requiring an explicit mathematical characterization of the integration volume. Although MC simulations are often regarded as the standard for validating new P_c estimation methods due to their high accuracy, their primary drawback remains the significant computational cost. To address this, several studies have focused on speeding up the MC approach through techniques such as subset simulation [22] and line sampling [23] which use stepwise procedures based on Markov Chains to reduce the number of samples required for a given confidence level.

Rather than relying on a traditional MC approach, this work employs Differential Algebra (DA) [24] to efficiently characterize the collective behavior of subsets of sampled initial conditions. This approach builds upon the work of Morselli et al. [25], who first applied DA to estimate the P_c by expressing both the Time of Closest Approach (TCA) and the relative distance as Taylor expansions of the initial condition uncertainties, and then sampling these initial conditions to compute the P_c through MC evaluation of the DA representation of the miss distance. While their approach is elegant and influential, it is specifically tailored to short term encounters and the resulting P_c computation remains limited to a local analysis centered around a single TCA. Our study consolidates and extends recent developments by the authors [26], with the goal of adapting and generalizing the methodology in Ref. [25] for long-term encounter scenarios, including cases in which multiple close approaches may occur within the screening interval. In details, we propose to compute the dynamical evolution of the initial conditions as a patched 7D continuum, where each patch is represented by a high-order Taylor expansion in both time and the initial conditions at t_0 . The integration accuracy of the dynamical flow is controlled by the Automatic Domain Splitting (ADS) algorithm [27], which adaptively splits the initial domain of the combined covariance to ensure precision of the Taylor approximation. Subsequently, the PDF, defined in the initial relative state space, is sampled to generate specific realizations. Each patch is evaluated accordingly, resulting in a time-dependent Taylor expansion of the miss distance. Collision probability is computed as the fraction of samples for which the miss distance falls below a predefined threshold, determined by identifying the real roots of the corresponding Taylor polynomials.

On the one hand, compared with research approaches that diverge from MC-based formulations, the proposed methodology relaxes restrictive assumptions such as the neglect of velocity uncertainty and it is able to robustly manage complex intersections of the collision volume that arise from multiple close approaches within the computation interval. Moreover, it can accommodate any form of initial uncertainty without requiring the assumption of a Gaussian distribution. On the other hand, the methodology offers an inherent advantage over standard MC simulations: instead of performing a classical sample-by-sample propagation to check for collisions, it relies on a polynomial representation of the covariance evolution over time. By evaluating one-dimensional time polynomials, we achieve a substantial reduction in computational cost while also offering a flexible framework naturally suited for parallel implementation. This efficiency gain is achieved through a reformulation of the problem enabled by DA, and it remains fully compatible with common MC acceleration techniques such as [22, 23], since the sampling MC component remains an essential step within our algorithm.

The proposed methodology forms part of the authors' broader effort to extend and enhance the capabilities of the DLR German Space Operations Center's (GSOC) Collision Avoidance System (CAS) [28], preparing it for the challenges posed by an increasingly congested orbital environment. In particular, it is designed with direct applicability to GEO assets under their control, which may at times experience long-term encounters with neighboring satellites.

The paper is structured as follows: Section 2 introduces the mathematical formulation commonly used to compute the P_c with a MC

approach. Section 3 presents the core methodology, detailing how DA and ADS are employed to model the dynamical evolution of initial conditions, and how root-finding in the resulting polynomial expressions is used to determine collision events. Section 4 provides benchmark test cases from the literature, along with a comparison of computational performance against an equivalent standard MC simulation. Sub-section 4.3 applies the proposed approach to a real-world long-term conjunction scenario, offering practical guidance for operational implementation. Finally, Section 5 reports the conclusions and outlines directions for future work.

2. Problem description

Building on the derivation in Ref. [18], the statistical event for which a collision occurs is introduced as follows: given the initial distribution of the state for two space resident objects at time t_0 , the Hard-Body Radius $HBR = R_1 + R_2$ [6], where R_1 and R_2 are the radii of spheres whose diameters correspond to the maximum dimensions of the primary and secondary objects, and a maximum period of interest T , a collision between two objects is deemed to occur if there exists a time t , within the interval $I = [t_0, t_0 + T]$, such that the norm of the relative distance vector $\mathbf{d}(t)$, is less than or equal to HBR . To assess the likelihood of this event occurring, and consequently characterize the P_c in a comprehensive and general manner, we introduce the relative state vector of the two objects engaged in the encounter, $\mathbf{x}(t)$, which is a function of time. This vector is defined as

$$\mathbf{x}(t) = \begin{pmatrix} \mathbf{d}(t) \\ \mathbf{v}_{rel}(t) \end{pmatrix} \quad (1)$$

in which \mathbf{v}_{rel} is the relative velocity of the secondary object with respect to the primary. It is important to emphasize that $\mathbf{x}(t)$ is defined as the difference, expressed in an inertial reference frame, between the individual states of the two objects involved in a close approach denoted as $\mathbf{x}_p(t)$ and $\mathbf{x}_s(t)$, respectively. These two quantities represent six-dimensional general multivariate random variables, which may follow arbitrary distributions. However, within the scope of this study, they are assumed to be statistically independent, implying that their cross-covariance is zero:

$$COV(\mathbf{x}_p, \mathbf{x}_s) = COV(\mathbf{x}_s, \mathbf{x}_p) = \mathbf{0} \quad (2)$$

This assumption is widely adopted in the literature and is not overly restrictive, as $\mathbf{x}_p(t)$ and $\mathbf{x}_s(t)$ are typically obtained through separate orbit determination processes. These processes are often independent, relying on distinct dynamical models, measurement data, and associated uncertainties.

For notation simplicity, at the initial time $t_0 = 0$, the relative state vector is denoted as $\mathbf{x}(t_0) = \mathbf{x}_0$. Defining the PDF of the relative state vector at t_0 as $\rho_0(\mathbf{x}_0, t_0)$, we can, without loss of generality, define P_c as:

$$P_c = \Pr(\mathbf{x}_0 \in \mathbf{V}) = \int_{\mathbf{V}} \rho_0(\mathbf{x}_0, t_0) d\mathbf{x}_0 \quad (3)$$

where $\mathbf{V} \subseteq \mathbb{R}^6$ represents the initial set for which a collision occurs at some future time t . For long-term encounters, the time evolution of the set \mathbf{V} represents the same integration volume modelled in Refs. [16,17, 19]. The set \mathbf{V} can be interpreted as a sub-region of the multi-dimensional space \mathbb{R}^6 wherein every realization of the random vector \mathbf{x}_0 inevitably leads to the violation of the condition $\|\mathbf{d}(t)\| \leq HBR$ at a future time t . In mathematics:

$$\mathbf{V} = \{\mathbf{x}_0 \in \mathbb{R}^6 : \exists t \in I : \|\mathbf{d}(t)\| - HBR \leq 0\} \quad (4)$$

By definition of PDF, the 6D integral of the initial statistic over the set \mathbf{V} gives, in fact, the likelihood that $\mathbf{x}_0 \in \mathbf{V}$. Notably, the integrand in Eq. (3) represents the PDF of an arbitrary distribution. Hence, the methodology outlined in the following section is capable of handling the

dynamical evolution of any statistical distribution, provided that its form is known a priori at t_0 . Although the approach is not limited to any specific distribution, for the purposes of the present analysis, $\rho_0(\mathbf{x}_0, t_0)$ is modelled as Gaussian. This choice is justified by the fact that, at least at t_0 , the relative state \mathbf{x}_0 typically results from an orbit determination process, for which the Gaussian assumption is generally considered reasonable. Thus:

$$\rho_0(\mathbf{x}_0, t_0) = \frac{e^{\left(-\frac{1}{2}(\mathbf{x}_0 - \boldsymbol{\mu}_0)^T \mathbf{P}_0^{-1} (\mathbf{x}_0 - \boldsymbol{\mu}_0)\right)}}{\sqrt{(2\pi)^6} \sqrt{\|\mathbf{P}_0\|}} \quad (5)$$

where $\boldsymbol{\mu}_0$ and \mathbf{P}_0 represent respectively the mean and the covariance matrix of \mathbf{x}_0 .

One way of computing the integral in Eq. (3) is via a MC-based method. In fact, in such cases, the initial conditions \mathbf{x}_0 are sampled and trajectories over the time interval $[t_0, t_0 + T]$ are computed according to some dynamical model that propagates the relative state from time t_0 to T . The dynamics are usually expressed as an Ordinary Differential Equation (ODE) of the form:

$$\begin{cases} \dot{\mathbf{x}}(t) = \mathbf{f}(\mathbf{x}(t), \mathbf{u}(t), t) \\ \mathbf{x}(t_0) = \mathbf{x}_0 \end{cases} \quad (6)$$

where the vector $\mathbf{u}(t)$ represents an eventually modelled maneuver in the relative dynamics.

Even when the initial conditions are modelled as Gaussian, their propagation through the dynamics described in Eq. (6), which, in the context of long-term encounters, are nonlinear, generally leads to a loss of Gaussian properties in the state distribution at future times t . Unlike approaches such as [18], the proposed methodology does not require the assumption that the statistical distribution remains Gaussian throughout the time interval I .

Finally, to compute the P_c , each sample trajectory is analyzed to verify if, at some future time, a collision occurs. If one hit is recorded for a specific sample, it means that it originally belonged to the set \mathbf{V} . The probability is then computed by evaluating the ratio between the number of samples that produced a hit over the total number of samples.

3. Methodology

3.1. DA to express the dynamical evolution of \mathbf{x}_0

As outlined in the preceding section, our methodology employs DA techniques to express the dynamical evolution of the initial relative conditions. DA provides a computational framework that enables the treatment of functions as n th order Taylor polynomial expansions within a computer environment, rather than handling them solely as floating-point values. This framework holds considerable potency as it allows to extract more information on a function rather than its mere raw values [24]. Within this context, the time t can be expressed as a DA variable τ and scaled with respect to the maximum time of interest T , such that $\tau \in [-1, 1]$:

$$\tau = \frac{2(t - t_0)}{T} - 1 \quad (7)$$

Before introducing a vector of DA variables that corresponds to the variation of the initial relative state vector from its mean at t_0 , we perform a rotation to the space in which $\mathbf{x}(t)$ is defined. This last may be, in general, either an inertial reference frame or a frame co-moving with the target along its trajectory. Such frames are commonly employed to express relative dynamics. Regardless of the chosen initial frame, which can be arbitrary, a transformation is applied to align the state space with the principal axes of the initial combined covariance matrix \mathbf{P}_0 . Specifically, the transformation is based on the eigen-decomposition of \mathbf{P}_0 , such that

$$\mathbf{P}_0 = \mathbf{Q} \boldsymbol{\Lambda} \mathbf{Q}^T, \quad (8)$$

where $\mathbf{Q} \in \mathbb{R}^{6 \times 6}$ is an orthogonal matrix, whose columns are the eigenvectors of \mathbf{P}_0 , and $\boldsymbol{\Lambda} \in \mathbb{R}^{6 \times 6}$ is a diagonal matrix containing the corresponding eigenvalues. This decomposition enables the mapping of a vector $\mathbf{m} \in \mathbb{R}^6$ from the original reference frame into an equivalent vector $\mathbf{m}' \in \mathbb{R}^6$, expressed in a coordinate system aligned with the principal axes of the uncertainty ellipsoid defined by \mathbf{P}_0 , according to:

$$\mathbf{m}' = \mathbf{Q}^T (\mathbf{m} - \boldsymbol{\mu}_0) \quad (9)$$

From this point onward, all references to the relative state vector and its DA representation will be made in the transformed coordinate system. To avoid overly pedantic notation, we deliberately omit the prime symbol (\cdot).

At this point, consistently with the DA variable τ , we introduce a vector of DA variables, $\delta \mathbf{x}_0$, representing the deviation of the initial relative state vector from $\boldsymbol{\mu}_0$. Each component is normalized by the maximum expected variation, $\Delta \mathbf{x}_0$, such that it is defined within the interval $[-1, 1]$. This is

$$\delta \mathbf{x}_0 = \frac{\mathbf{x}_0 - \boldsymbol{\mu}_0}{\Delta \mathbf{x}_0} \quad (9)$$

The relative state at a given scaled instant τ , can now be computed via integration of the dynamics described in Eq. (6) and expressed in the DA framework as:

$$\mathbf{x} = \mathcal{T}_{\mathbf{x}}(\tau, \delta \mathbf{x}_0) \quad (10)$$

That is a vector of high order polynomials that are functions of the deviations of the scaled time τ and the initial normalized relative statistics, $\delta \mathbf{x}_0$. The Taylor map $\mathcal{T}_{\mathbf{x}}$ establishes a relationship between the perturbed initial state vector and the corresponding state vector at a specified time within I , utilizing the dynamical model defined in (6). This mapping from the initial set to the final one bears conceptual similarity to the mathematical notion of a manifold well described in Ref. [29]. Within the context of this research, the employed dynamical model may be arbitrarily complex and may include highly nonlinear relative motion. The only requirement is that it must be described by an analytical law that establishes a functional relationship between the state at time t and the initial conditions at t_0 .

Utilizing DA in this context offers several advantages. Firstly, it enables the representation of an infinite set solely through its Taylor expansion coefficients. This preserves a specific analytical structure in contrast to a mere point-wise set representation. Secondly, and perhaps most significantly, it allows the propagation of entire sets through a function using straightforward DA arithmetic operations. Unlike a standard MC simulation, where the ODE flow of equation (6) is integrated for each sample, in this scenario, only a single integration is required. The resulting DA expansion represents the outcome of propagating all points from the initial domain through the ODE in (6).

3.2. ADS to control integration accuracy

The challenging part of this approach arises from the nonlinear dynamics involved usually in long-term encounters. When the Taylor map needs to approximate a strongly non-linear function, the convergence of the ODE expansion across the domain becomes inaccurate. Consequently, the DA map, which is a local representation of the function, poorly represents the actual evolution of the whole domain, even though the description is accurate in the vicinity of the center of the expansion.

To address this issue, we employ the technique proposed in Ref. [27]. The ADS algorithm identifies instances where the ODE flow expansion over the initial conditions no longer accurately describes the dynamics.

Once such a scenario is detected, the domain of the original polynomial expansion is divided along one of the expansion variables into two domains, each half the size of the original. By re-expanding the

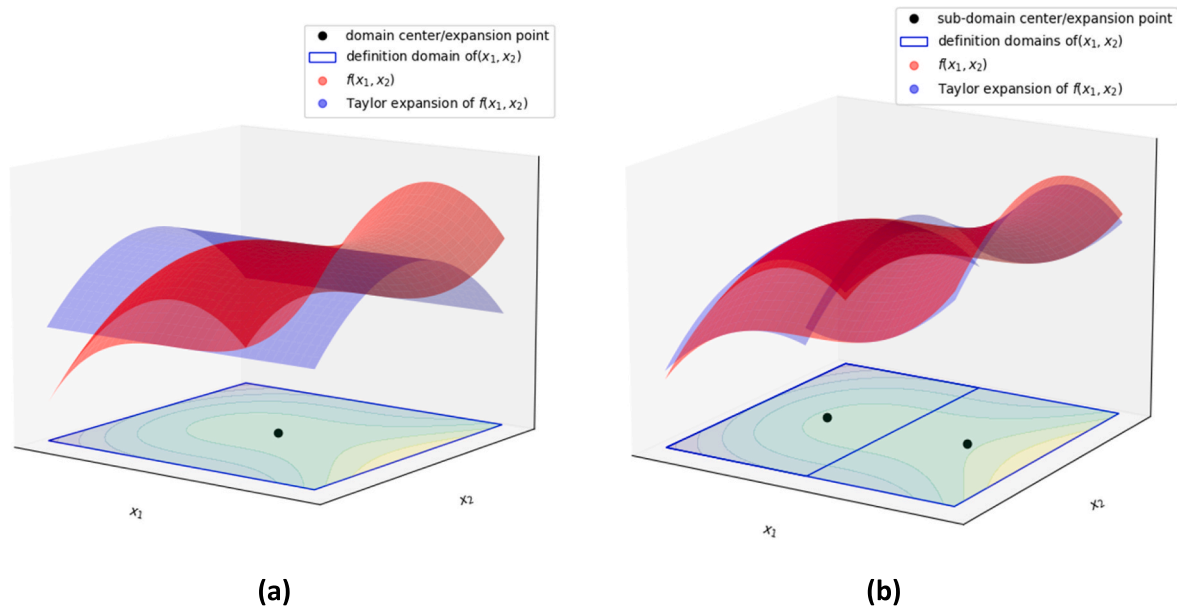


Fig. 2. ADS algorithm illustration. (a) Taylor expansion of $f(x_1, x_2)$ around initial domain's midpoint. (b) Taylor expansions of $f(x_1, x_2)$ recalculated around the new domain centers.

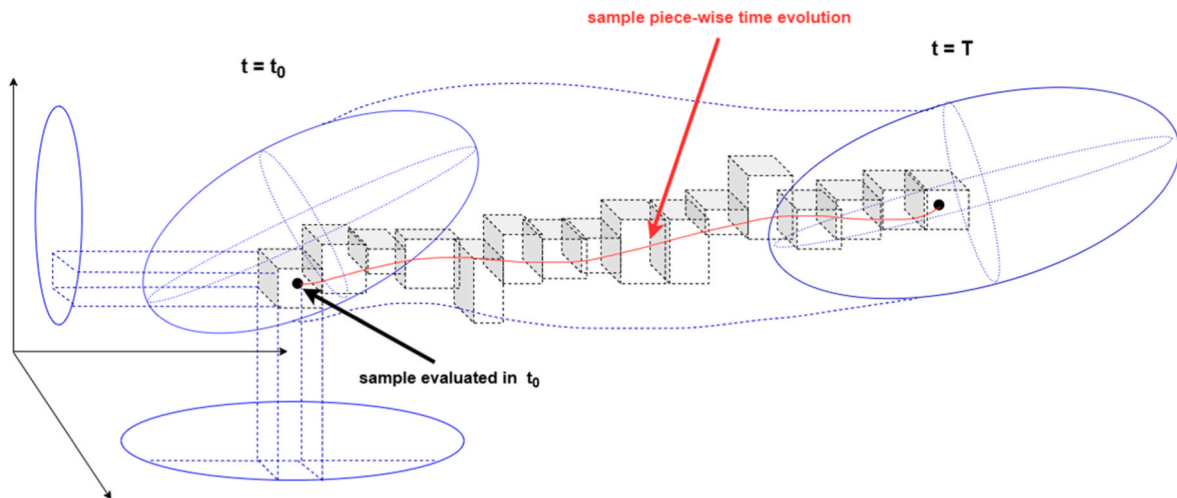


Fig. 3. Dynamical flow evolution of the initial conditions x_0 . For representation purposes, only spatial coordinates are shown.

polynomials around the new center points, two separate polynomial expansions are generated.

To aid the reader's understanding and provide a visual interpretation of the concept, this process is illustrated in Fig. 2 -(a) and (b). In the visualization, a 2D function $f(x_1, x_2)$ is depicted alongside the defined domain of the variables (x_1, x_2) . Additionally, Taylor expansion centered around the domain's midpoint is displayed to approximate $f(x_1, x_2)$. The accuracy of the approximation is high near the center but diminishes towards the domain edges. To ensure accuracy, the algorithm iteratively splits the initial domain into two segments whenever the Taylor series representation diverges from the actual function by a user-defined margin ϵ . Subsequently, the expansions are recalculated around the new centers, and this process continues until all expansions accurately represent the function within the specified threshold ϵ .

In a similar fashion, in our case the initial 7D domain, defined by the variables τ and δx_0 is split into different sub-domains. The dynamical evolution of the initial condition assumes the shape of a patched 7D continuum, mathematically defined as a manifold object [30]. Fig. 3 attempts to give a visual representation of this last, considering only the

position components of the relative state vector and the time. To each patch at a given time corresponds a Taylor expansion, function of τ and of δx_0 , that approximates locally the dynamical flow. Once a single integration has been performed and the evolution of the initial condition is approximated by patched polynomials, our methodology proceeds to calculate the function d^2 within the DA framework:

$$d^2 = \mathcal{F}_{d^2}(\tau, \delta x_0) \quad (11)$$

Here, d^2 is a high order polynomial representation of the relative distance squared expressed as function of time and initial state. It is again piece-wise defined, and its definition interval depends on the ADS accuracy control algorithm.

Subsequently, the initial set x_0 is sampled. Each realization X_{0i} of the random vector is linked to its respective initial patch and evaluated only in space and velocity. This evaluation reduces the dimensions of the d^2 polynomials, resulting in a set of one-dimensional Taylor expansions depending solely on time:

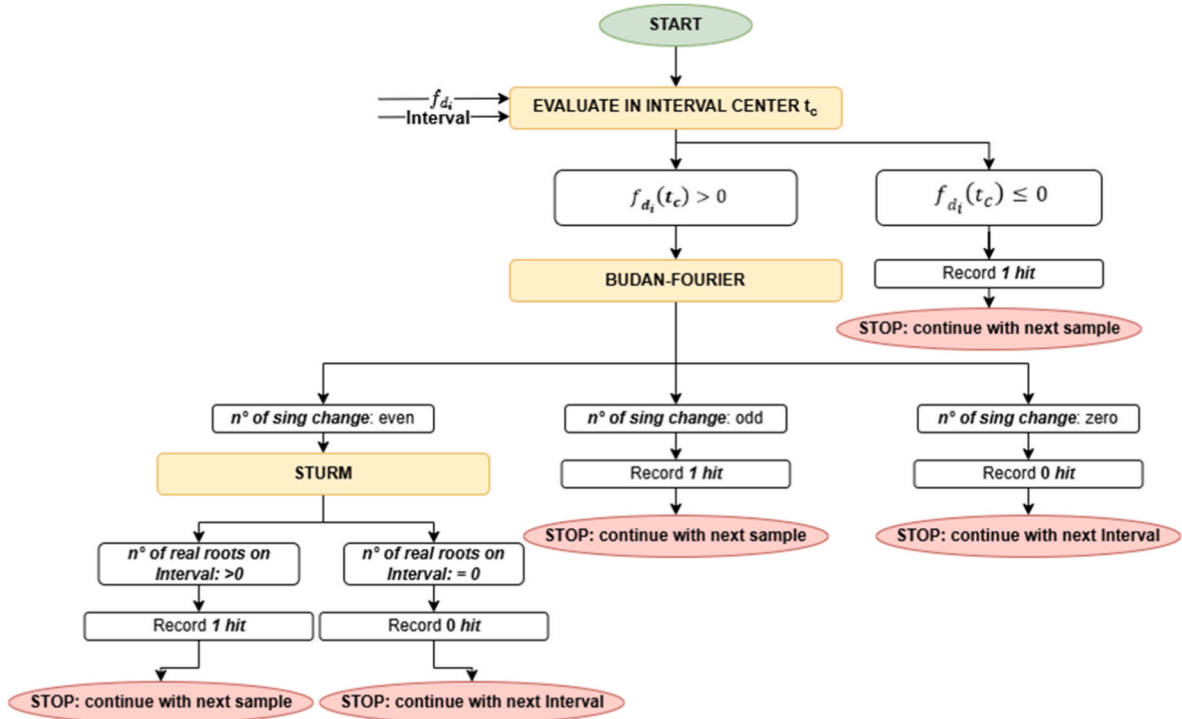


Fig. 4. Roots-finding algorithm to individualize the zeros of high order mono-dimensional time polynomials.

$$d_i^2 = \mathcal{T}_{d^2}(\tau, \delta \mathbf{x}_0 = \mathbf{X}_{0i}) = \mathcal{T}_{d_i^2}(\tau), \quad (12)$$

This expansion holds significant importance as it is analyzed by the algorithm to determine if a collision occurs for a specific sample. Specifically, this process is reduced to the task of locating the roots of

$$f_{d_i} = d_i^2 - HBR^2 = \mathcal{T}_{f_{d_i}}(\tau) \quad (13)$$

within the defined bounds of the Taylor expansion. This process is again illustrated in Fig. 3, where a sample is evaluated within the split initial set, resulting in a subset of one-dimensional polynomials that approximate the time evolution of the trajectory highlighted in red.

3.3. Finding the roots of high order one-dimensional time polynomials

At this stage, the task of determining whether a given sample trajectory results in a collision, and thus contributes to the cumulation of P_c , is reformulated as a problem of finding the roots of high-order univariate polynomials that depend solely on time. This is accomplished through the application of a series of analytical theorems, following a methodology analogous to that employed in a different context in Ref. [31]. The root-finding algorithm is detailed in Fig. 4. After evaluating the sample and computing the DA expression in Eq. (13), we iterate through all the polynomials that define a sample trajectory. For each Taylor expansion and its corresponding definition interval, we initially verify if its center falls into the negative range to rule out the possibility of finding no roots due to the segment trajectory already being below the collision threshold. Subsequently, we determine the number of sign changes of the polynomial coefficients using the Budan-Fourier theorem [32]. This theorem considers the number of roots by examining the sequence of coefficient sign variations in the polynomial. Specifically, if the number of sign changes is odd, it indicates the presence of at least one real root within the polynomial's interval. In such cases, the algorithm registers a hit and proceeds to analyze the next sample. If there are no sign changes (i.e., the count is zero), we conclude that the polynomial has no real root in the interval. Thus, the algorithm proceeds to analyze the adjacent Taylor expansion

as time progresses along the trajectory.

The situation differs when the number of sign changes is even. In such instances, the theorem does not provide conclusive results because the number of roots can be a multiple of an even number, potentially including zero. Therefore, the workflow further investigates using the Sturm algorithm [33]. This algorithm is a robust root isolation method that precisely determines the number of roots of a high-order polynomial within an interval by recursively performing Euclidean divisions to construct a sequence of polynomials. The sign variations in this sequence are analyzed to ascertain the number of roots. As before, if there are no roots, the algorithm proceeds to analyze the neighboring polynomials. However, if at least one root is found, a hit is recorded, and the polynomial approximation of the subsequent trajectory is studied.

The procedure described above is executed for each sample within a loop over the ADS-defined intervals. To improve the efficiency of the algorithm, the intervals are sorted in ascending order of their initial time. Before testing whether a given interval covers the sample, the algorithm verifies that its initial time is not earlier than the final time of the previously analyzed interval; if it is, the interval is skipped without further checks.

3.4. Monte Carlo tool for validation

To verify our methodology and measure the efficiency of our approach, we compare the results obtained with those generated by a standard, in-house MC simulation. To this end, the initial distribution is sampled in an analogous manner and each sample is propagated from t_0 to T . A grid of equally time-spaced points in miss distance is created, followed by interpolation where the real roots of a localized cubic polynomial are extracted. To elaborate, curve fitting is executed using a technique called parabolic blending [34,35], where a set of four equally spaced points is utilized to construct a third-order polynomial by merging two quadratic polynomials generated from the initial three points and the last three points. The minimum of the fitted curve is then determined by extracting the roots of the polynomial's first derivative. The MC process then assesses whether a collision occurs for a given sample by checking if the relative distance at any point within the

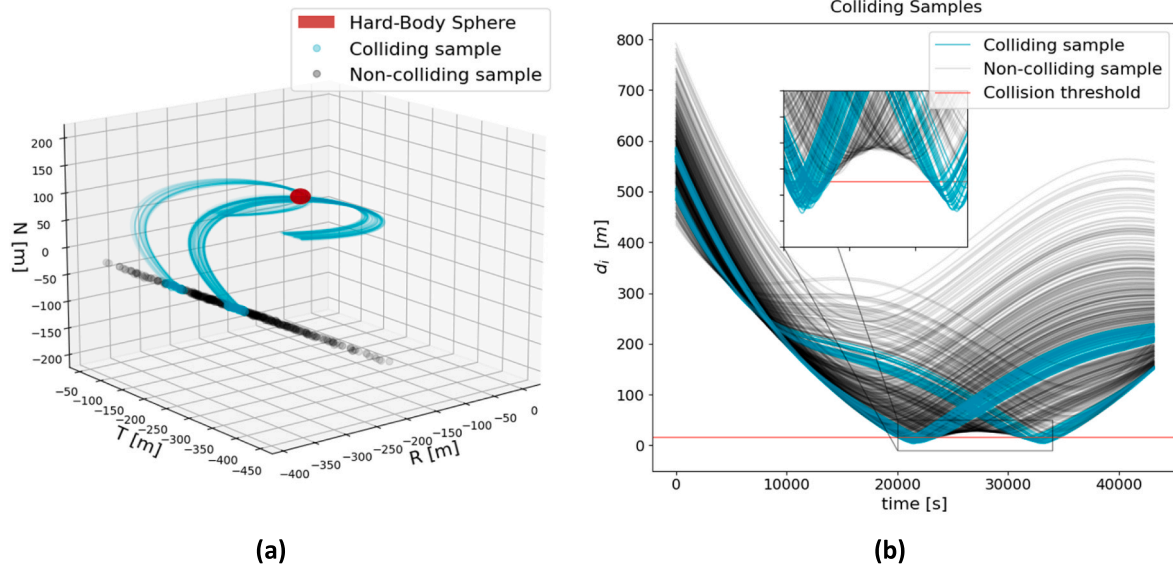


Fig. 6. (a) Samples of the combined covariance at t_0 , with colliding samples and their trajectories shown in cyan for test case #1 (b) Evolution of the relative distance d_i (in magnitude) for each sample, with trajectories that violate the collision threshold highlighted in cyan. (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)

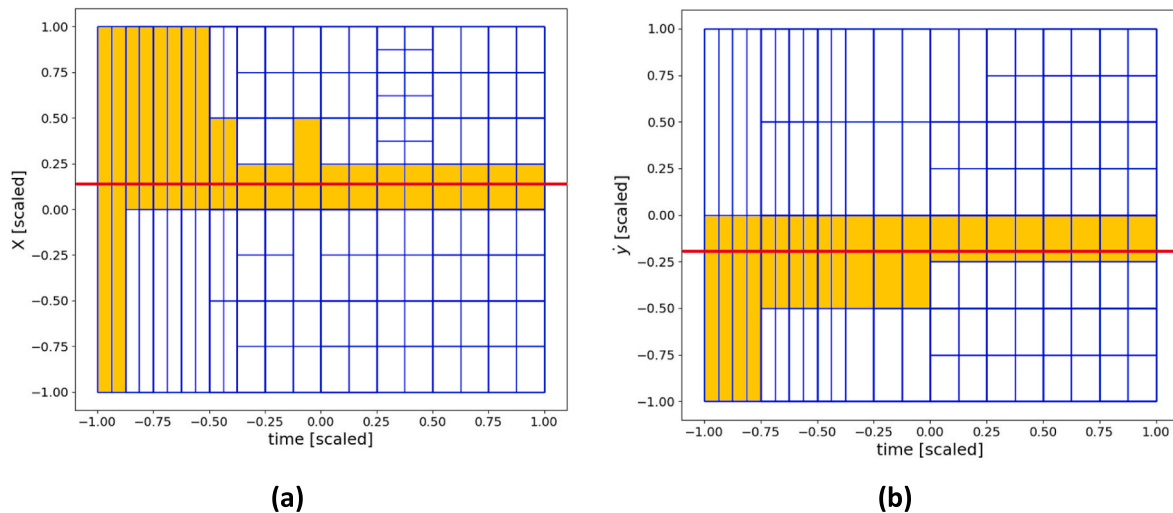


Fig. 7. Subdomains time evolution of relative state (a) first-component and (b) fifth-component. In orange all the subdomains associated to a given sample trajectory. (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)

the interested reader.

Regardless of the dynamics employed to consistently derive the conditions at t_0 , the workflow outlined in Fig. 5(a) is followed in all cases. First, Eq. (6) is integrated using the same dynamics adopted to generate the initial conditions, followed by the sampling of the initial statistic. The minimum number of samples required for statistical significance is then determined using the same statistical bounding criteria as presented in Ref. [20].

In the following discussion, we provide a detailed explanation of the methodology, offering insights into its operation and presenting plots corresponding to the first test cases based on Clohessy–Wiltshire dynamics. An overview of the complete set of tests conducted with different dynamical models is then summarized in Table 2.

4.1. Test case #1 (Clohessy–Wiltshire)

The first test case considered involves two satellites in GEO having a

non-rectilinear encounter. Fig. 6(a) illustrates the combined positional covariance sampled at the initial time. The cyan points represent a subset of samples, denoted as \mathbf{V} in Eqs. (3) and (4), for which a collision occurs within the timeframe of analysis. As depicted in Fig. 6(b), the test case was deliberately designed so that even the mean of the initial distribution results in a collision, leading to a notably high final P_c reference value of $2.1783\text{E-}01$. This elevated P_c level necessitates fewer than 16,000 Monte Carlo runs for the results to attain statistical significance with 5 % accuracy and 95 % confidence.

As described in section 3, the evolution of the initial condition generates a 7D manifold, with the initial sub-domains established by the ADS routine. Fig. 7 illustrates how the domain in the first and fifth components of the initial relative state and the time is split. Each sub-domain defines the range of these variables for which a single Taylor expansion can represent the relative state vector with the required accuracy. The red line represents a particular determination of the initial relative state component, based on which all the associated sub-domains

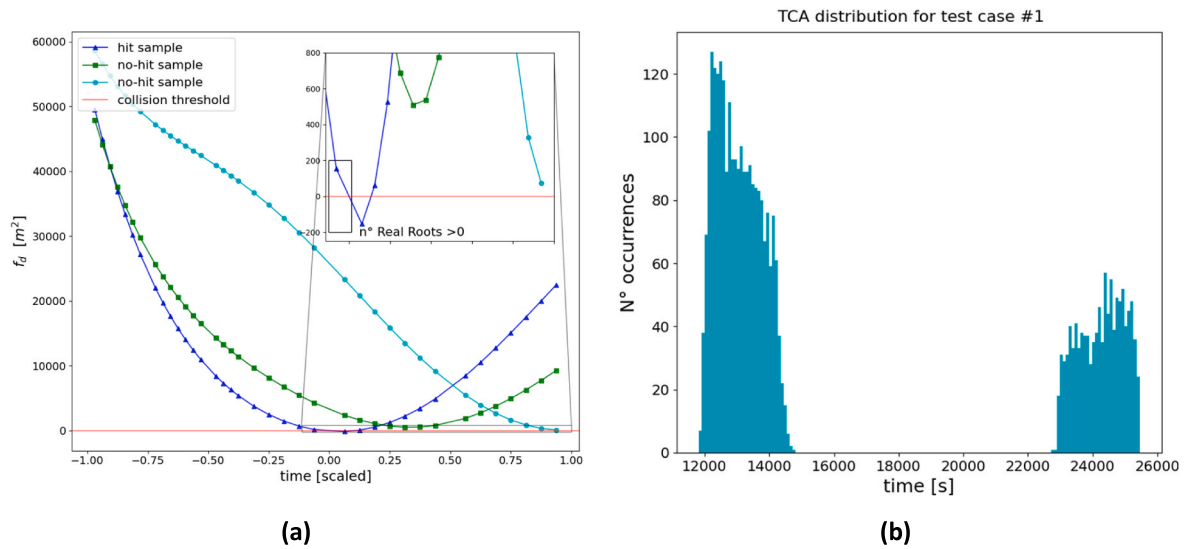


Fig. 8. (a) Taylor expansions of f_d evaluated at the centre of each time sub-domain for two “hit” samples and one “no-hit” sample. (b) TCA distribution for test case #1.

are selected.

The evaluation of all these polynomials for a given sample of the initial relative state allows computing the DA expansion outlined in Eq. (13), which is a function only of the time. This is illustrated in Fig. 8-(a), where the Taylor expansions of f_d are evaluated at the center of each sub-domain in time for three different samples. Two samples do not result in a collision (cyan and green curves), and one does lead to a collision (blue). The zoomed view in the figure highlights that the polynomial identifying the sample as colliding is not the one with its center below the zero line, but rather the one enclosed within the rectangle.

The cumulative collision probability determined through the DA polynomial evaluation method stabilizes at a value of $2.1783e-01$,

mirroring the result obtained from our in-house Monte Carlo simulation. Over the analysis period, the initial conditions evolve in a manner that, on average, leads to an accumulation of P_c at two distinct times. This test case illustrates the method's ability to address multiple conjunctions and, consequently, multiple TCAs within the analysis time-frame. Fig. 8-(b) depicts the TCA distribution in the form of a histogram, clearly indicating that among the samples resulting in a collision, a subset hits around 12,000 s after the start of the simulation. Successively, the P_c stops accumulating after about 2000 s, and then increases again approximately 9500 s later. The computational time is compared to that of the in-house Monte Carlo simulation with an integration time-step of 5 s. As shown in Table 2, approximately 16,000 iterations are completed in about $2.62E-01$ s on an Intel(R) Core (TM) i9-14900K CPU

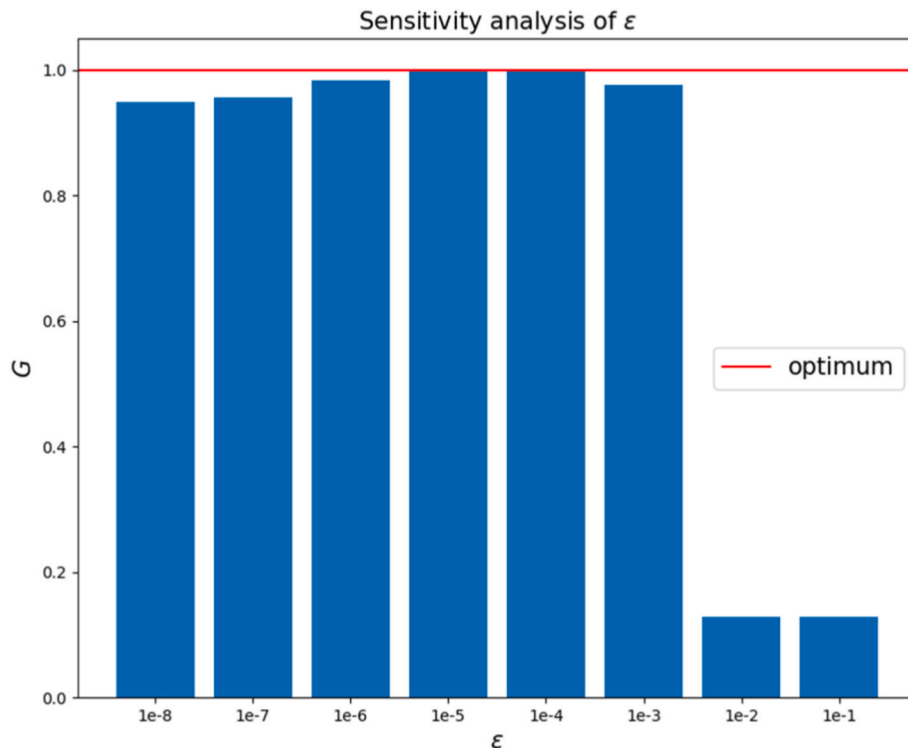


Fig. 9. Sensitivity analysis of the splitting tolerance ϵ

Table 1
Description of test cases.

Test case	Description
Case #2	It shares the same characteristics and relative motion as Case #1, involving two GEO satellites; however, it features a smaller HBR, resulting in a mean miss distance that does not intersect the combined collision sphere.
Case #3	It also considers a GEO encounter but represents a short-term conjunction, where the P_c accumulates instantaneously at TCA.
Case #4	It involves non-rectilinear relative motion between two GEO objects and is designed to illustrate how the shape and orientation of the relative covariance can lead to P_c accumulation occurring after TCA.
Case #5, #6, #7	Cases #5, #6, and #7 correspond to LEO encounters, with each case representing progressively more pronounced non-linear relative motion: from marginal non-linearity in Case #5 to strongly non-linear conditions in Case #7.
Case #8	It features a long-term encounter in MEO.
Case #9	Cases #9 and #10 involve encounters between satellites in highly eccentric orbits, with non-rectilinear relative motion and a mean miss distance at TCA that exceeds the HBR. Due to the high eccentricity of these orbits, only the Yamanaka-Ankersen model is used to analyze these two cases.
Case #10	

@ 5.5 GHz, compared to roughly 14 s for the MC approach, resulting in a gain in computational time of more than 5000 %.

Before extending the analysis to additional test cases and different dynamical models, we present a sensitivity study on the splitting tolerance ε for test case #1. In the previous section, ε was introduced as a user-defined parameter that controls how accurately the Taylor polynomials reproduce the dynamical evolution of the initial conditions. Our aim is to provide the reader with intuition and practical guidance on how to select an appropriate value for this parameter. Intuitively, a smaller value of ε requires a Taylor expansion of higher order to accurately follow the dynamics, which in turn leads to a finer subdivision of the initial domain into a larger number of subdomains. Conversely, a larger ε allows the use of fewer subdomains and a lower-order polynomial approximation, but at the cost of degrading the accuracy of the estimated final P_c . This degradation occurs because, when the polynomial representation does not sufficiently match the true trajectory evolution, some samples that should result in a collision may no longer

do so, and vice versa. Fig. 9 illustrates this trade-off through a parametric analysis of ε . For each value of the splitting tolerance, we compute the gain function G , defined as the product of the computational-time reduction (relative to MC) and the P_c accuracy percentage (relative to MC) normalized with respect to the highest accuracy achieved. The gain function G highlights the values of ε that offer the best balance between accuracy and computational efficiency. In particular, values of G close to 1 correspond to an optimal choice of ε , whereas values approaching 0 indicate a suboptimal configuration. As shown, values of ε between 1e-5 and 1e-4 provide the best balance between computational efficiency and estimation accuracy. Beyond 1e-3, the gain function G decreases noticeably, even though the corresponding error in P_c remains on the order of only 1–2 %. It is also worth noting that, even for very small values of ε , the reduction in computational time remains significant, as indicated by G remaining close to 1. For case #1 and the test cases presented in the following subsections, a reference value of 1e-4 has therefore been adopted.

4.2. Additional test cases: different dynamical models and computational time

Additional test cases were examined to assess the performance of the proposed method across a range of scenarios and with the dynamical models introduced in the section above. A brief description of each case is provided in Table 1 to give the reader an overview of its structure and purpose.

Table 2 presents a comprehensive comparison of the DA-based methodology against a standard MC approach across various dynamical models. For each test case, the table reports the number of samples used, the final P_c obtained, the percentage error in P_c relative to the MC result, the respective computational times, and the time required by the DA and ADS components to integrate Eq. (6). Test case #7 is included in the description above (Table 1) for completeness, but it is not further analyzed due to the elevated number of samples required and its equivalence to cases #5 and #6. Moreover, cases #9 and #10 are only applicable to the Yamanaka-Ankersen dynamics, since in both scenarios the primary follows a highly eccentric orbit.

The results show that the DA-based methodology achieves excellent

Table 2
Test cases overview for different scenarios and different relative dynamical models.

Case	n° Samples	DA Polynomials			Monte Carlo		Pc err [%]
		Pc	Cmp. Time [s]	ADS integration time [s]	Pc	Cmp. Time [s]	
Clohessey-Wiltshire							
# 1	1.57E+04	2.1783439490e-01	2.2803e-01s	3.699e-03	2.1783439490e-01	1.3612e+01	0.0000
# 2	6.52E+06	1.6044057777e-02	1.4247e+02	7.310e-03	1.6043137529e-02	5.2175e+03	0.0057
# 3	9.54E+05	1.0017956031e-01	4.4498e+01	1.3565e-02	9.9781236209e-02	7.2086e+02	0.3991
# 4	1.24E+06	7.4077658315e-02	1.6199e+01	3.1722e-03	7.4077658315e-02	1.1144e+03	0.0000
# 5	2.30E+06	4.9284041611e-02	2.3036e+02	3.8621e-02	4.9277954638e-02	2.1331e+03	0.0123
# 6	2.45E+07	3.2113120024e-02	4.4939e+03	7.7043e-02	3.2113120024e-02	2.3475e+04	0.0000
# 8	3.11E+06	3.5209770838e-02	9.8045e+01	7.0816e-03	3.5209770838e-02	3.0456e+03	0.0000
Yamanaka-Ankersen							
# 1	1.57E+04	2.1974522293e-01	7.7953e-01	7.8263e-02	2.1974522293e-01	2.4096e+02	0.0000
# 2	6.52E+06	1.5625208494e-02	1.5904e+02	3.1008e-02	1.5623828121e-02	9.4998e+04	0.0088
# 3	9.54E+05	1.0004958087e-01	7.8968e+01	4.9884e-02	9.9618761904e-02	1.3478e+04	0.4325
# 4	1.24E+06	7.3597818057e-02	1.9757e+01	4.5257e-02	7.3597818057e-02	1.8311e+04	0.0000
# 5	2.30E+06	4.9344170491e-02	3.5407e+02	1.6887e-01	4.9335474794e-02	3.5586e+04	0.0176
# 6	2.45E+07	3.2108784306e-02	5.1490e+03	3.0087e-01	3.2108784306e-02	4.4743e+05	0.0000
# 8	3.11E+06	3.5504638603e-02	8.7062e+01	4.8339e-02	3.5504638603e-02	4.6758e+04	0.0000
# 9	1.80E+05	3.6322070245e-01	1.5878e+01	2.2812e-01	3.6322070245e-01	2.5490e+03	0.0000
# 10	1.92E+05	3.6289961354e-01	1.6699e+01	2.1105e-01	3.6289961354e-01	2.6738e+03	0.0000
D'Amico ROEs (J2 only)							
# 1	1.57E+04	2.1229299363e-01	1.1358e+01	2.9052e-01	2.1229299363e-01	4.0226e+01	0.0000
# 2	6.52E+06	1.5595278886e-02	5.0212e+03	2.9113e-01	1.5595278886e-02	1.6352e+04	0.0000
# 3	9.54E+05	9.9111422782e-02	7.9797e+02	4.8696e-01	9.8611421210e-02	2.6991e+03	0.5070
# 4	1.24E+06	7.3588081261e-02	1.0374e+03	2.8894e-01	7.3588081261e-02	3.1196e+03	0.0000
# 5	2.30E+06	1.3047958428e-01	1.8123e+03	4.4447e+00	1.3046827989e-01	6.8514e+03	0.0087
# 6	2.45E+07	3.214569346e-02	4.4890e+04	4.9542e+00	3.214569346e-02	2.6569e+05	0.0000
# 8	3.11E+06	3.5636115989e-02	5.3679e+03	8.5806e-01	3.5635151356e-02	1.2422e+04	0.0027

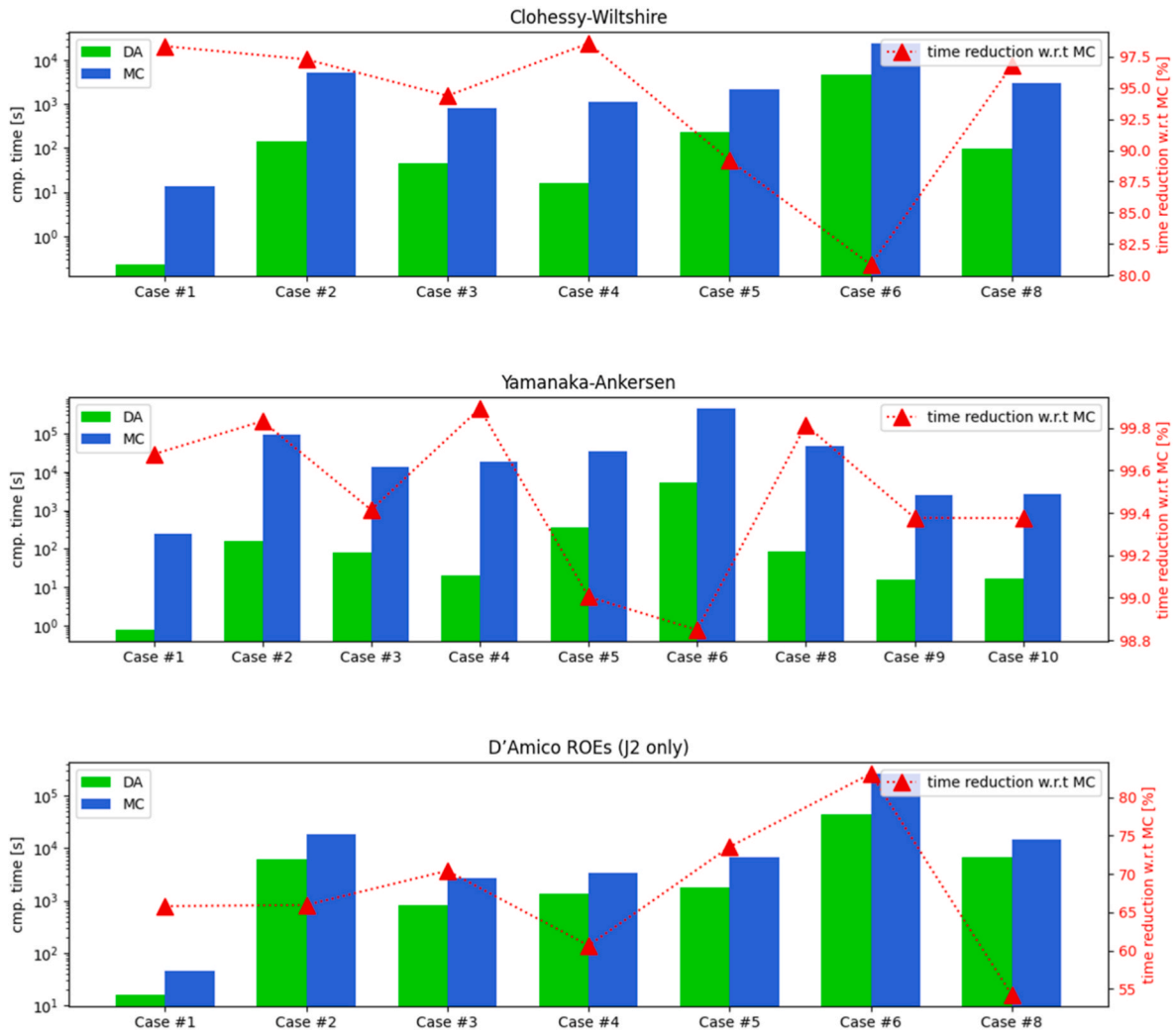


Fig. 10. Comparison of computational times for different test cases between the DA-based methodology and the corresponding MC simulation. The left axis shows the computational time in logarithmic scale, while the right axis shows the associated computational time reduction.

agreement with the MC simulation across all dynamical regimes, consistently reproducing the P_c with high accuracy. In some cases, there is a negligible discrepancy in the final P_c with an error that never exceeds 0.0176 %. These minor deviations are attributed to an inherent limitation of the DA framework: it requires setting a threshold for the uncertainty in the initial deviation $\delta \mathbf{x}_0$ to construct the patched 7D manifold. In this study, the threshold has been conservatively set to five standard deviations of the initial uncertainty. In rare cases, especially when a large number of samples is involved, a few may fall outside this 5σ boundary at t_0 . Such samples are discarded and not included in the analysis, which can lead to a slight under-/or over-estimation of P_c compared to the complete MC reference. Nevertheless, the overall accuracy and efficiency of the DA approach remain robust across all test scenarios.

In case #3, across all dynamics, the error with respect to the reference never exceeds 0.5 %, as the test case essentially represents a 2D encounter. In such situations, the P_c accumulates almost instantaneously at TCA, requiring the algorithm to perform a large number of splits around that time to properly capture the dynamical evolution, with a corresponding reduction in the splitting tolerance ϵ . This leads to a loss in computational efficiency, highlighting the inevitable trade-off between runtime and P_c estimation accuracy in 2D cases. However, this limitation is not particularly critical, since for short-term encounters the use of a MC based approach is generally not recommended, since far more efficient methods exist to compute P_c with high accuracy.

In terms of computational speed, it is important to note that the DA integration consistently requires significantly less time, often by several orders of magnitude, than the evaluation of the individual samples, to the point where its contribution becomes practically negligible. This highlights a fundamental strength of the methodology: it captures the full evolution of the initial condition set through a single, efficient integration. For a visual comparison of computational times between the DA-based approach and the MC method, the reader is referred to Fig. 10. All computational times were obtained using the same hardware configuration described in the previous section.

The plot displays computational time on the left axis (in logarithmic scale) and the percentage time reduction in computational time relative to the corresponding MC simulation on the right axis. Overall, the DA-based approach outperforms the MC method across all test cases and dynamical models, with computational time reductions ranging from a minimum of approximately 55 % to nearly 99.9 %. Notably, the gap in computational efficiency between the DA-based method and the MC approach widens as the complexity of the underlying dynamics increases. While the MC method may benefit from slightly faster sample propagation in simpler models, such as Clohessy–Wiltshire or the D'Amico formulation, the DA-based approach becomes significantly more advantageous as the dynamics involve more complex transformations, as in the case of Yamanaka–Ankersen.

Table 3

State vectors and associated uncertainties of real test case scenario.

Real long-term conjunction scenario	
State vector and covariance at epoch time	
Primary object	Secondary object
X = 35037.184606 [km]	X = -35035.891661 [km]
Y = -23466.448768 [km]	Y = -23467.794741 [km]
Z = 15.784810 [km]	Z = 15.864584 [km]
X_DOT = 1.710187 [km/s]	X_DOT = 1.709784 [km/s]
Y_DOT = -2.554690 [km/s]	Y_DOT = -2.554927 [km/s]
Z_DOT = -0.001790 [km/s]	Z_DOT = 0.000755 [km/s]
CR_R = 1.341040E+03 [m**2]	CR_R = 1.353645E+05 [m**2]
CT_R = -1.285933E+02 [m**2]	CT_R = -3.104145E+05 [m**2]
CT_T = 2.221449E+04 [m**2]	CT_T = 2.118139E+07 [m**2]
CN_R = 1.062929E+04 [m**2]	CN_R = 5.530171E+02 [m**2]
CN_T = 2.613577E+02 [m**2]	CN_T = -2.019151E+03 [m**2]
CN_N = 8.435330E+04 [m**2]	CN_N = 4.600785E+04 [m**2]
CRDOT_R = 0.000000E+00 [m**2/s]	CRDOT_R = 1.620946E+01 [m**2/s]
CRDOT_T = 0.000000E+00 [m**2/s]	CRDOT_T = -1.582021E+03 [m**2/s]
CRDOT_N = 0.000000E+00 [m**2/s]	CRDOT_N = 1.073235E-01 [m**2/s]
CRDOT_RDOT = 0.000000E+00 [m**2/s**2]	CRDOT_RDOT = 1.189744E-01 [m**2/s**2]
CTDOT_R = 0.000000E+00 [m**2/s]	CTDOT_R = -9.649515E+00 [m**2/s]
CTDOT_T = 0.000000E+00 [m**2/s]	CTDOT_T = 1.190034E+01 [m**2/s]
CTDOT_N = 0.000000E+00 [m**2/s]	CTDOT_N = -3.914278E-02 [m**2/s]
CTDOT_RDOT = 0.000000E+00 [m**2/s**2]	CTDOT_RDOT = -3.817790E-04 [m**2/s**2]
CTDOT_TDOT = 0.000000E+00 [m**2/s**2]	CTDOT_TDOT = 6.930053E-04 [m**2/s**2]
CNDOT_R = 0.000000E+00 [m**2/s]	CNDOT_R = -2.373264E-02 [m**2/s]
CNDOT_T = 0.000000E+00 [m**2/s]	CNDOT_T = -4.966605E-01 [m**2/s]
CNDOT_N = 0.000000E+00 [m**2/s]	CNDOT_N = -3.501316E-01 [m**2/s]
CNDOT_RDOT = 0.000000E+00 [m**2/s**2]	CNDOT_RDOT = 4.222343E-05 [m**2/s**2]
CNDOT_TDOT = 0.000000E+00 [m**2/s**2]	CNDOT_TDOT = 1.972771E-06 [m**2/s**2]
CNDOT_NDOT = 0.000000E+00 [m**2/s**2]	CNDOT_NDOT = 9.341607E-05 [m**2/s**2]

4.3. Real test case

We now consider a real conjunction scenario, as the test cases presented in Ref. [20], while academically relevant, are not operationally realistic due to their extremely high P_c . The event involves a close approach between an asset in GEO and a secondary object in a very similar orbit, with a relative distance of 1.83 km and an HBR of 27.05 m. The encounter features a normal component of just 81 m and a relative velocity at TCA of approximately 3 m/s, classifying it as a long-term conjunction.

For this event, the TCA conditions are summarized in Table 3. Specifically, the state vectors are provided in an ECI frame, while the corresponding covariance matrices are expressed in the RTN frame for each object. Fig. 11-(a) illustrates the evolution of the relative distance between the two objects over a 2-day period centered on the TCA. This interval reveals multiple close approaches, indicated by local minima in the relative distance. In addition to the situation at TCA, two notable minima (shown in the plot as local minimum A and B) occur approximately 24 h before and after the closest approach. In the first one, the objects reach a separation of 10 km with a normal component of 178 m; in the second, the normal separation is 221 m. The instantaneous P_c , computed using the 2D methodology reported in Ref. [3], peaks at 1.42E-04 at TCA, while at the other minima, it is roughly an order of magnitude lower.

The methodology proposed in this work is tested on this scenario, and the results are compared to a MC simulation as done for the other test cases. Based on the Instantaneous P_c at TCA, the required number of samples has been estimated: approximately $3.0E+07$ are needed to achieve statistical relevance with 5 % accuracy and 95 % confidence. The resulting P_c evolution is given in Fig. 11-(b) and (c). As it can be seen the P_c accumulates first around the first minimum, reaching the value of

1.76E-06, then accumulates for a period of 20 min, bracketing TCA and then accumulates again at the third minimum, settling to a value of 1.4173E-04. All the samples that are producing a hit in the MC simulation are producing a hit also in the DA methodology reaching a 0.0 % estimation error while being approximately 87.8 times faster.

5. Conclusion and future work

We presented a general methodology for computing the P_c between two space objects, particularly suited for long-term encounters with relative velocities on the order of meters per second, yet in principle applicable to two-dimensional scenarios as well. The approach employs DA to capture the non-linear time evolution of the multivariate initial relative state vector. This last is modelled as a patched 7D continuum, where each patch is represented by a high-order Taylor expansion in both time and the initial conditions at t_0 . The integration accuracy of the dynamical flow is controlled by the ADS algorithm, which adaptively splits the initial domain of the combined covariance to ensure precision of the Taylor approximation. The initial PDF is then sampled and each patch is evaluated in a specific state-vector realization to definitively determine the time evolution of a given sample. The identification of collisions is reduced to finding the real roots of the DA approximation of the miss distance.

The proposed methodology retains all the advantages of a MC simulation while offering a substantial improvement in computational speed, owing to its DA foundation. In principle, it can be applied to any form of initial uncertainty, provided that the initial PDF is known a priori, without requiring the restrictive assumption of Gaussianity at t_0 or throughout the evolution of the initial conditions. Moreover, the approach can incorporate uncertainties in velocity and naturally accounts for multiple conjunctions occurring within the screening interval. Additionally, while this study has focused on three specific dynamical models, the technique is applicable to any form of relative or absolute dynamics, provided that an analytical formulation of the dynamics is available.

The technique has been extensively validated against benchmark test cases from the literature. In all scenarios, the estimated P_c shows excellent agreement with the corresponding MC simulations used for validation. The percentage reductions in computational time are consistently significant, ranging from approximately 55 % to nearly 99.9 %, with the gap in efficiency between the DA-based method and the MC approach increasing as the complexity of the underlying dynamics grows.

This work also presents a real-case scenario involving an encounter between two GEO asset involved in a long-term conjunction. The proposed methodology successfully captures multiple conjunction events and the corresponding P_c accumulation occurring in different intervals within the computation window.

In future work, we plan to enhance the fidelity of the dynamics by extending the presented models to include additional orbital perturbations, particularly those relevant to the GEO environment.

While the reduction in computational time is evident, a key objective remains the adaptation of the methodology for operational use at GSOC. Having successfully passed an extensive validation phase, the next goal is its full integration into the CAS, enabling the handling of long-term conjunctions across multiple missions simultaneously. In doing so, the system will be capable of providing outputs that function as reliable decision-support tools for flight dynamics engineers in real-world encounter scenarios, ultimately contributing to the safe operation of more than 40 satellites.

CRedit authorship contribution statement

Andrea Zollo: Writing – review & editing, Writing – original draft, Visualization, Validation, Software, Methodology, Investigation, Formal analysis, Data curation, Conceptualization. **Cristina Parigini:** Writing –

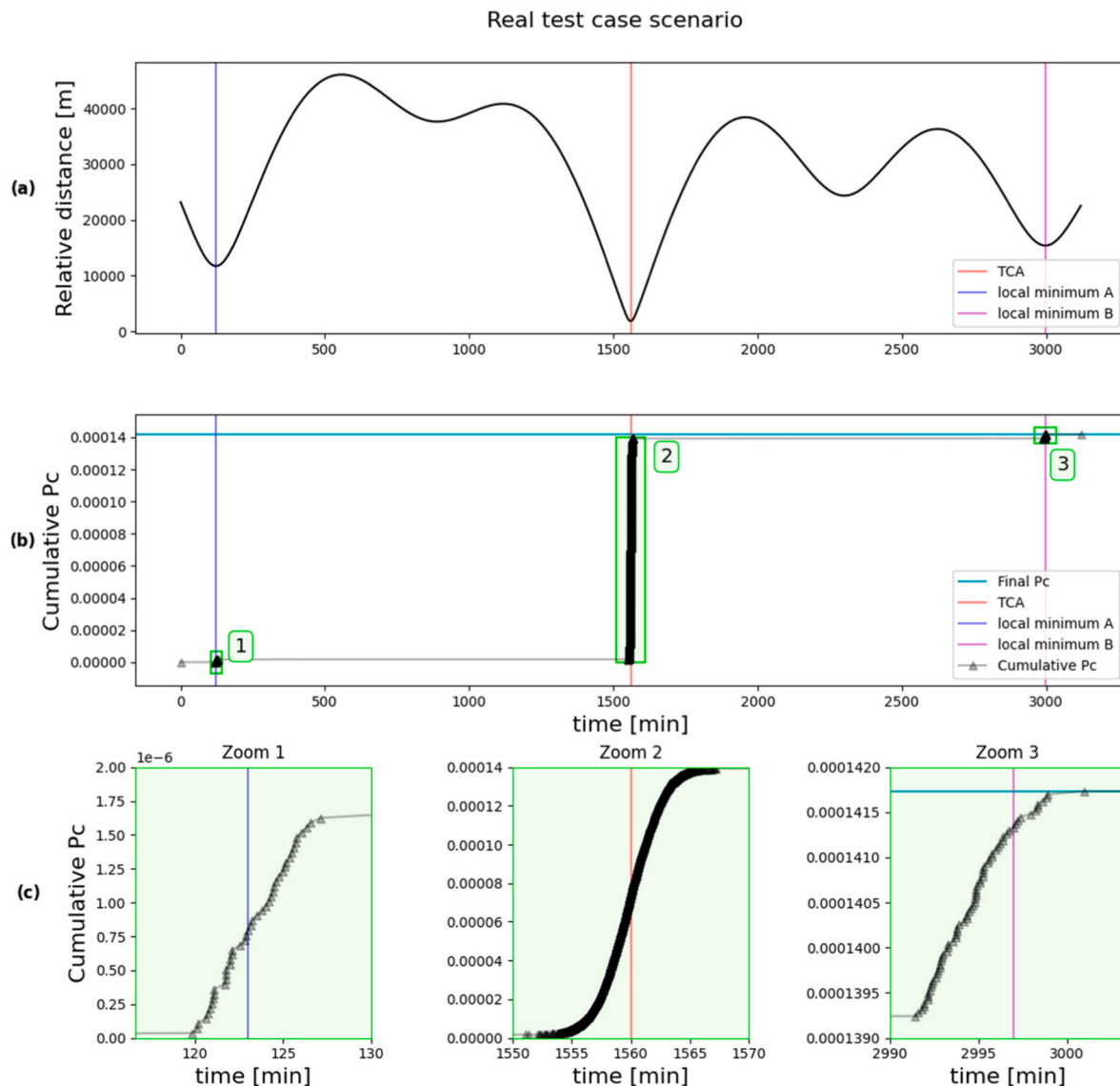


Fig. 11. (a) Evolution of relative distance for real test case. (b) Evolution of P_c . (c) Zoomed view highlighting regions of P_c accumulation.

review & editing, Supervision, Software, Methodology, Investigation, Conceptualization. **Roberto Armellini**: Writing – review & editing, Supervision, Methodology, Investigation, Conceptualization. **Juan Félix San Juan Díaz**: Writing – review & editing, Validation, Supervision, Software, Investigation, Formal analysis. **Annarita Trombetta**: Writing – review & editing, Software, Investigation. **Ralph Kahle**: Writing – review & editing, Supervision, Project administration, Funding acquisition.

Disclosure statement

Artificial intelligence tools have been utilized to enhance the grammar, spelling and overall readability of this manuscript.

Declaration of competing interest

We wish to confirm that there are no known conflicts of interest associated with this publication and there has been no significant financial support for this work that could have influenced its outcome.

We confirm that the manuscript has been read and approved by all named authors and that there are no other persons who satisfied the criteria for authorship but are not listed.

We further confirm that the order of authors listed in the manuscript has been approved by all of us.

We confirm that we have given due consideration to the protection of intellectual property associated with this work and that there are no impediments to publication, including the timing of publication, with respect to intellectual property.

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Appendix

Nomenclature

List of abbreviations	
ADS	Automatic Domain Splitting
CAM	Collision Avoidance Maneuver
CAS	Collision Avoidance System
DA	Differential Algebra
GEO	Geostationary Orbit
GSOC	German Space Operations Center
HBR	Hard Body Radius
LEO	Low Earth Orbit
MC	Monte-Carlo
ODE	Ordinary Differential Equation
P_c	Probability of Collision
PDF	Probability Density Function
PSS	Polynomial Superlevel Set
ROEs	Relative Orbital Elements
STM	State Transition Matrix
TCA	Time of Closest Approach
List of symbols	
t_0	Initial time
T	Maximum period of interest and end of simulation time
I	Screening interval in which the methodology computes the P_c
$\mathbf{d}(t)$	Relative distance vector between Primary and Secondary objects
R_1	Radius of a sphere whose diameter correspond to the maximum dimension of the primary object.
R_2	Radius of a sphere whose diameter correspond to the maximum dimension of the secondary object.
HBR	Radius of the combined sphere that accounts for both primary and secondary dimensionality
$\mathbf{x}(t)$	Relative state vector between Primary and Secondary objects at time t
\mathbf{x}_0	Relative state vector at initial time t_0
$\mathbf{v}_{rel}(t)$	relative velocity of the secondary object with respect to the primary
$\mathbf{x}_p(t)$	State vector of the primary object
$\mathbf{x}_s(t)$	State vector of the secondary object
$COV(\mathbf{x}_p, \mathbf{x}_s)$	Cross-covariance of the two random variables \mathbf{x}_p and \mathbf{x}_s
$\rho_0(\mathbf{x}_0, t_0)$	PDF of relative state vector at t_0
\mathbf{V}	Integration volume to compute the P_c
μ_0	Mean of the relative state distribution at t_0
\mathbf{P}_0	Covariance matrix of the relative state distribution at t_0
$\mathbf{u}(t)$	Input maneuver modelled in the relative dynamics
τ	Simulation time expressed as DA variable
\mathbf{Q}	Orthogonal matrix whose columns are the eigenvectors of \mathbf{P}_0
Λ	Diagonal matrix containing the eigenvalues of \mathbf{P}_0
\mathbf{m}	Given vector expressed in an initial reference frame that can either be inertial or co-moving with the primary object.
\mathbf{m}'	Vector \mathbf{m} transformed to a reference frame aligned with the principal axes of \mathbf{P}_0
$\delta\mathbf{x}_0$	Vector of DA variables representing the deviation of \mathbf{x}_0 from μ_0 .
$\Delta\mathbf{x}_0$	Vector containing maximum expected deviation of each component of \mathbf{x}_0
\mathbf{x}	Relative state at a given scaled instant τ expressed in the DA framework
\mathcal{T}_x	Taylor map that links the deviations $\delta\mathbf{x}_0$ at t_0 and \mathbf{x}
ε	Splitting tolerance
d^2	DA representation of the relative distance squared
\mathcal{T}_{d^2}	Taylor map that links the deviations $\delta\mathbf{x}_0$ at t_0 and d^2
\mathbf{X}_{0i}	Given realization of $\delta\mathbf{x}_0$
d_i^2	One-dimensional Taylor polynomial depending solely on time representing d^2 after sample evaluation
f_{d_i}	DA expression representing the difference between d_i^2 and HBR squared used to check for collisions
G	Gain function introduced to individualize the optimum of ε

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