

Evolution of topology in axi-symmetric and 3-d viscous flows

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Abstract

Topological methods are used to establish global and to extract local structure properties of vector fields in axi-symmetric and 3-d flows as function of time. The notion of topological skeleton is applied to the interpretation of vector fields generated numerically by the Navier-Stokes equations. The flows considered are swirling jets with super-critical swirl numbers that show low Reynolds number turbulence in the break-up region.

1 Introduction

A numerical study is conducted to investigate the temporal development of flow structures connected with the vortex breakdown in swirling free jets. There is experimental evidence (see Billant *et al.* [2]) that swirling jets exhibit a rich variety of flow phenomena. Simplified model equations assuming steady and rotationally symmetric motion above a conical stream surface show bi- and multi-stability (see [3], [4], [5], [6]) of several flow forms such as hysteresis and negative thrust phenomena. Direct numerical simulations at moderate Reynolds numbers allow relaxation of the model restrictions. The results show the development of the conical jet layer and its break up into three-dimensional vortex structures for moderate Reynolds numbers. The detection of these structures requires sophisticated tools, that are applicable to two and three dimensional, steady and unsteady flows. It is essential for the interpretation of these fields to reduce their properties to the bare minimum, hence the notion of the topological skeleton, introduced by Fomenko (see [7], [8]) for the classification of integrable Hamiltonian systems, is applied to the present case. The paper presents the theoretical aspects of vector field topology and numerical methods for their computation.

The organisation of the paper is as follows. First, the numerical method for the solution of the incompressible Navier-Stokes equations is briefly outlined. Then basic definitions and theorems for the topology of steady vector fields are discussed followed by the consideration of the topological skeleton and its temporal evolution. Numerical methods for the computation of the evolution of the vector field topology are developed and results for the swirling jet flows are presented.

2 Navier-Stokes solver

A hybrid spectral finite-difference method (developed in [9]) is used to solve the Navier-Stokes equations in cylindrical coordinates. The method will be outlined briefly, the details including the verification of its accuracy and convergence properties are available in [9] and [10]. The method takes advantage of the fact that all dependent variables φ are periodic with respect to azimuthal variable θ , hence is the discrete Fourier transform \mathcal{F}_N (Fourier collocation method, Canuto *et al.*, [11]) applicable

$$\tilde{\varphi}(k) = \mathcal{F}_N(\varphi) \equiv \frac{1}{N} \sum_{j=0}^{N-1} \varphi(\theta_j) \exp(-ik\theta_j), \quad k = -N/2, \dots, N/2 - 1$$

$$\varphi(\theta_j) = \mathcal{F}_N^{-1}(\tilde{\varphi}) \equiv \sum_{k=-N/2}^{N/2-1} \tilde{\varphi}(k) \exp(ik\theta_j), \quad j = 0, \dots, N-1 \quad (1)$$

where \mathcal{F}_N^{-1} denotes the inverse transform and the $\theta_j = 2\pi j/N$ are the discrete collocation points.

The dependent variables (see [9]) in the present formulation are the complex-valued Fourier modes (k denotes the discrete azimuthal wavenumber) of:

- The azimuthal velocity modes $\tilde{v}_\theta(r, k, z, t)$ for $0 \leq k \leq \frac{N}{2}$,
- the azimuthal vorticity modes $\tilde{\Omega}_\theta(r, k, z, t)$ for $0 \leq k \leq \frac{N}{2}$,
- the streamfunction modes $\tilde{\Psi}(r, k, z, t)$ for $0 \leq k \leq \frac{N}{2}$,
- the pressure modes $\tilde{p}(r, k, z, t)$ for $1 \leq k \leq \frac{N}{2}$.

The azimuthal momentum balance provides the equation for the velocity modes $\tilde{v}_\theta(r, k, z, t)$

$$\begin{aligned} \frac{\partial \tilde{v}_\theta}{\partial t}(k) + \mathcal{F}_N[v_r \frac{\partial v_\theta}{\partial r} + v_z \frac{\partial v_\theta}{\partial z} + \frac{v_\theta}{r} (\frac{\partial v_\theta}{\partial \theta} + v_r)] = -i \frac{k}{r\rho} \tilde{p}(k) \\ + \frac{1}{Re} [\frac{\partial}{\partial r} (\frac{1}{r} \frac{\partial}{\partial r} (r \tilde{v}_\theta(k))) - \frac{k^2}{r^2} \tilde{v}_\theta(k) + \frac{\partial^2 \tilde{v}_\theta}{\partial z^2}(k) + 2i \frac{k}{r^2} \tilde{v}_r(k)] \end{aligned} \quad (2)$$

where the convective terms are evaluated using FFT, Re denotes the Reynolds number. Application of the divergence operation to the momentum balances produces the equation for the pressure modes $\tilde{p}(r, k, z, t)$, which emerges in the form

$$\begin{aligned} \frac{1}{r} \frac{\partial}{\partial r} (r \frac{\partial \tilde{p}}{\partial r}) - \frac{k^2}{r^2} \tilde{p} + \frac{\partial^2 \tilde{p}}{\partial z^2} = \\ -\rho \mathcal{F}_N [(\frac{\partial v_r}{\partial r})^2 + \frac{1}{r^2} (\frac{\partial v_\theta}{\partial \theta} + v_r)^2 + (\frac{\partial v_z}{\partial z})^2 + \frac{2}{r} \frac{\partial v_\theta}{\partial r} (\frac{\partial v_r}{\partial \theta} - v_\theta) + 2 \frac{\partial v_r}{\partial z} \frac{\partial v_z}{\partial r} + \frac{2}{r} \frac{\partial v_z}{\partial \theta} \frac{\partial v_\theta}{\partial z}] \end{aligned} \quad (3)$$

The transport equation for the azimuthal vorticity component $\tilde{\Omega}_\theta$ follows from its definition and the momentum balances. It appears in the form

$$\begin{aligned} \frac{\partial \tilde{\Omega}_\theta}{\partial t} = \frac{\partial}{\partial r} [\mathcal{F}_N(v_\theta \Omega_r) - \mathcal{F}_N(v_r \Omega_\theta)] + \frac{\partial}{\partial z} [\mathcal{F}_N(v_\theta \Omega_z) - \mathcal{F}_N(v_z \Omega_\theta)] \\ + \frac{1}{Re} [\frac{\partial}{\partial r} (\frac{1}{r} \frac{\partial}{\partial r} (r \tilde{\Omega}_\theta(k))) - \frac{k^2}{r^2} \tilde{\Omega}_\theta(k) + \frac{\partial^2 \tilde{\Omega}_\theta}{\partial z^2}(k) + 2i \frac{k}{r^2} \tilde{\Omega}_r(k)] \end{aligned} \quad (4)$$

It is shown in [9] that complex-valued streamfunctions $\tilde{\Psi}(r, k, z, t)$ exist for each wavenumber k as a consequence of the Fourier transformed mass balance, which can be recast as the divergence of a 2-d vector being zero. This results in Poisson-Helmholtz type equations for the complex-valued $\tilde{\Psi}$

$$\frac{\partial}{\partial r} \left(\frac{1}{r} \frac{\partial}{\partial r} (r \tilde{\Psi}) \right) + \frac{\partial^2 \tilde{\Psi}}{\partial z^2} = \tilde{\Omega}_\theta + ik \frac{\partial \langle \tilde{v}_\theta \rangle}{\partial z} \quad (5)$$

The velocity components can be recovered from

$$-r \tilde{v}_z = \frac{\partial}{\partial r} (r \tilde{\Psi}), \quad \tilde{v}_r = \frac{\partial \tilde{\Psi}}{\partial z} - ik \langle \tilde{v}_\theta \rangle \quad (6)$$

once \tilde{v}_θ and the streamfunctions $\tilde{\Psi}$ have been computed. The angular brackets are essential for the streamfunction formalism (5), (6), they are defined by

$$\langle \Phi \rangle(r, k, z, t) \equiv \frac{1}{r} \int_0^r dr' \Phi(r', k, z, t) \quad (7)$$

The presence of singular coefficients in (2) to (5) requires careful analysis of the pdes for $r \rightarrow 0$ ([12], [13]). The results (details in [9]) can be summarized as follows:

(I) Scalar modes $\tilde{\Phi}(r, k, z, t)$ and axial vector components satisfy $O(\tilde{\Phi}(r, k, z, t)) = r^k$ and are symmetric with respect to r for k even and skew-symmetric for k odd.

(II) Radial and azimuthal vector components satisfy the kinematic conditions

$$\tilde{v}_r(0, k, z, t) + ik\tilde{v}_\theta(0, k, z, t) = 0$$

and both modes are symmetric with respect to r for $k = 1$, and

$$O(\tilde{v}_r(r, k, z, t)) = O(\tilde{v}_\theta(r, k, z, t)) = r$$

for $k = 0$ and

$$O(\tilde{v}_r(r, k, z, t)) = O(\tilde{v}_\theta(r, k, z, t)) = r^{k-1}$$

for $k > 1$, and they are skew-symmetric with respect to r for k even and symmetric for k odd. The complex-valued streamfunctions can be shown to obey

$$O(\tilde{\Psi}(k)) = r^{k+1}$$

and are skew-symmetric with respect to r for k even, symmetric for k odd.

The conditions at the axis and the entrance, exit and outer boundaries are either Dirichlet or Neumann type or a combination of Dirichlet and Neumann conditions. Specifically, the axis conditions cannot be chosen arbitrarily but follows from the symmetry and smoothness properties at $r = 0$; the Fourier modes satisfy either homogenous Dirichlet or homogeneous Neumann conditions. At entrance and outer boundaries Dirichlet conditions are prescribed and convective boundary conditions are used at the exit.

The cross-sectional domain $(r, z) \in [0, R] \times [0, L_z]$ is rectangular, it may be semi-infinite ($R = \infty$). It is mapped onto the unit quadrangle using separable and analytic functions. It is discretized with respect to radial and axial directions with uniform spacing in the image domain $[0, 1] \times [0, 1]$. Either central differences are used for first and second derivatives with explicit filters (Kennedy & Carpenter [14]) applied to the increments of the transport equations or upwind-biased differences (Li[15]) for the convective terms and central differences for the non-convective terms. The filter order is chosen such that it does not degrade the accuracy of the finite-difference schemes, no filters are applied for the upwind-biased schemes. The numerical integration required in (7) is done using a fourth order accurate difference method.

The time integrator is an explicit 4th-order state space Runge-Kutta method that requires minimal storage (see [9]). The Poisson/Helmholtz equations (2) and (5) are solved using LU-decomposition (LINPACK, see Dongarra et al.[16]) combined with deferred corrections to reduce the bandwidth of the coefficient matrix.

3 Topology of Steady Vector Fields

The study of the topology requires clear definitions and numerical methods for their computation. We define in the following the basic notions of vector field topology relevant for the present purpose. The description is similar to Fomenko et al. [7], [8], but the results in our examples are fundamentally different since our fields do not allow a Hamiltonian function. We start with some simple terms.

First, we clarify two definitions widely used in fluid mechanics: Streamlines and pathlines. Given an unsteady vector field $\mathbf{v}(\mathbf{x}, t)$ we compute the streamlines $\Phi(t^*)$ as solutions of the initial value problem

$$\frac{d\Phi}{dt^*} = \mathbf{v}(\Phi(t^*), t) \quad (8)$$

with initial conditions determining the desired streamline. Streamlines are computed for the fixed time t (instantaneous integral curves or snapshots) and the picture generated by them changes with time if the vector field is unsteady. On the other hand are the pathlines the solutions of

$$\frac{d\Phi}{dt} = \mathbf{v}(\Phi(t), t) \quad (9)$$

plus initial conditions and it follows that pathlines are different from streamlines if the vector field is unsteady. The pathlines are the vector lines in the mathematical literature. We consider in

the present chapter steady vector fields and instant snapshots of unsteady fields. The vector lines are streamlines and the line parameter t is time if the field is steady, but for unsteady fields it is isomorphic to arclength.

Definition 3.1 A planar vector field is a map

$$\begin{aligned} v : D &\rightarrow \mathbb{R}^2, \\ x &\mapsto v(x). \end{aligned} \tag{10}$$

Definition 3.2 An integral curve through a point $x \in \mathbb{R}^2$ of a vector field $v : D \rightarrow \mathbb{R}^2$ is a map

$$\alpha_x : \mathbb{R} \supset I \rightarrow D, \tag{11}$$

where

$$\alpha_x(0) = x_0, \tag{12}$$

$$\dot{\alpha}_x(t) = v(\alpha_x(t)), \quad \forall t \in I. \tag{13}$$

Concerning the theorem on existence and uniqueness of integral curves, the *Lipschitz condition* has to be satisfied:

Definition 3.3 Let $U \subset \mathbb{R}^2$ be an open subset. A continuous vector field $v : \mathbb{R}^2 \rightarrow \mathbb{R}^2$ satisfies a **Lipschitz condition on U** provided there exists a real number $K > 0$ such that

$$\|Dv(x) - Dv(y)\| \leq K\|x - y\| \tag{14}$$

holds for all $x, y \in U$ and all $t \in J$ where Dv denotes the differential of the vector field v . The constant K is called **Lipschitz constant**.

This allows to formulate the existence and uniqueness theorem:

Theorem 3.4 (Existence and uniqueness of integral curves) Let $v : D \rightarrow \mathbb{R}^2$ be a vector field satisfying the Lipschitz condition on any open neighborhood around point $x \in \mathbb{R}^2$. Then there exists one and only one integral curve through any $x_0 \in \mathbb{R}^2$.

Proof: See [31, pp. 66–68].

We also need some terms for the different directions of the vector field at the boundary.

Definition 3.5 Let $D \subset \mathbb{R}^2$ be a compact domain of a Lipschitz-continuous vector field $v : D \rightarrow \mathbb{R}^2$. Let $d \in \partial D$ be a point on the boundary. We define the following three entities:

- (1) The point d is called an **outflow point** if every integral curve through d ends at d and there exists an integral curve through d that does not contain another point of ∂D . The set of all outflow points is called **outflow set**.
- (2) The point d is called a **inflow point** if every integral curve through d starts at d and there exists an integral curve through d that does not contain another point of ∂D . The set of all inflow points is called **inflow set**.
- (3) The point d is called a **boundary flow point** if there exists an integral curve $\alpha_d : (-\epsilon, \epsilon) \rightarrow D$, $\epsilon > 0$, through d lying completely inside ∂D . The set of all boundary flow points is called **boundary flow set**.
- (4) The point d is called **boundary saddle (switch point)**, if it does not belong to the other three types.

If a vector field satisfies the Lipschitz condition for an open neighborhood of every point, integral curves can be extended over the whole time line if they do not reach the boundary at an outflow point.

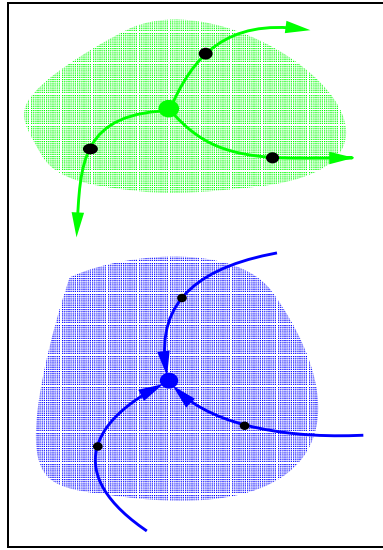


Figure 1: Around a source, every integral curve starts at the source; around a sink, every integral curve ends at the sink.

Lemma 3.6 *Let $v : D \rightarrow \mathbb{R}^2$ be Lipschitz continuous around each point. Then an integral curve can be defined up to $+\infty$ if it does not end at an outflow point. It can also be defined from $-\infty$ if it does not start at an inflow point.*

Since we want to keep the following notations simple, we define an integral curve at an inflow point as that point for the whole time before it enters the domain. Further, at an outflow point, we extend an integral curve by this point for all the time after it has left the domain. In this way, all integral curves are defined for all $t \in \mathbb{R}$. Topology is really concerned with the asymptotic behavior of integral curves, so we look at their asymptotic begin and end.

Definition 3.7 *Let $v : D \rightarrow \mathbb{R}^2$ be a Lipschitz-continuous vector field and $c : \mathbb{R} \rightarrow \mathbb{R}^2$ an integral curve. The set*

$$\{a \in \mathbb{R}^2 \mid \exists (t_n)_{n=0}^{\infty} \subset \mathbb{R}, t_n \rightarrow \infty, \lim_{n \rightarrow \infty} c(t_n) \rightarrow a\} \quad (15)$$

is called ω -limit set of c . The set

$$\{a \in \mathbb{R}^2 \mid \exists (t_n)_{n=0}^{\infty} \subset \mathbb{R}, t_n \rightarrow -\infty, \lim_{n \rightarrow \infty} c(t_n) \rightarrow a\} \quad (16)$$

is called α -limit set of c .

The most common types of limit sets are critical points.

Definition 3.8 *A critical point of $v : D \rightarrow \mathbb{R}^2$ is a point $p \in D$ with $v(p) = 0$.*

Two cases are of interest: (1) sinks and (2) sources.

Definition 3.9 *Let $v : D \rightarrow \mathbb{R}^2$ be a Lipschitz-continuous vector field, $z \in D$ a critical point. If there is a neighborhood $U \subset D$ of z such that all integral curves in U have a α -limit set consisting only of z , then z is called a **source**. If there is a neighborhood $U \subset D$ of z such that all integral curves in U have a ω -limit set consisting only of z , then z is called a **sink**. If there is a neighborhood $U \subset D$ of z such that there is a finite number of integral curves that start or end at z and all other integral curves leave U in both directions of time, z is called a **saddle**.*

Figure 1 shows a source in the upper part. All integral curves around the critical point start at this critical point. In the lower part, a sink is shown. Every integral curve in the neighborhood ends at the sink. Another type of limit sets are closed integral curves (cycles).

Definition 3.10 Let $v : D \rightarrow \mathbb{R}^2$ be a Lipschitz continuous vector field. A closed streamline $c : \mathbb{R} \rightarrow D$ of v is a streamline with a $t_0 \in \mathbb{R}$ with

$$c(t + nt_0) = c(t) \forall n \in \mathbb{N}. \quad (17)$$

(It is assumed that c does not stay at one point all the time, i.e. $c(t)$ is not a critical point.)

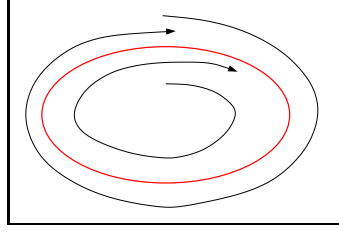


Figure 2: A limit cycle may attract streamlines in its neighborhood.

Figure 2 shows a typical example. Such a cycle is called structurally stable if, after small changes, the vector field still contains a closed streamline. Such stability arguments allow to concentrate on critical points, boundary regions and closed cycles as possible α - and ω -limit sets [29],[30]. For our analysis of a vector field, we have to determine the asymptotic behavior of all its integral curves. This is done by analyzing the union of all integral curves starting or ending at the same critical point, same cycle, or the same inflow/outflow region.

Definition 3.11 Let $v : D \rightarrow \mathbb{R}^2$ be a Lipschitz-continuous vector field and $M \subset D$. The union of all integral curves of v that converge to M for $t \rightarrow -\infty$ is called α -basin of M , denoted by $B_\alpha(M)$. The union of all integral curves of v that converge to M for $t \rightarrow \infty$ is called ω -basin of M , denoted by $B_\omega(M)$.

For sinks and sources, one can prove the following lemma:

Lemma 3.12 Let a be a critical point of a Lipschitz continuous vector field $v : D \rightarrow \mathbb{R}^2$. If a is a source then $B_\alpha(a)$ is an open subset of D . If a is a sink then $B_\omega(a)$ is an open subset of D .

Proof: See [30, pp. 181-182].

A similar result holds for cycles, i.e. a structurally stable cycle in 2D can be either a source and its α -basin is an open subset of the domain or it is a sink and its ω -basin is an open subset. A short comment on dimension prepares us for the final theorem of the section.

Definition 3.13 If a subset $M \subset \mathbb{R}^2$ allows a description as a pure n -dimensional manifold, we define the dimension of M as n .

Remark 3.14 We need the term dimension only for basins. The previous definition means that an open basin in D is a 2-manifold, a basin consisting of a finite number of streamlines is a 1-manifold and a basin consisting of a finite number of points is a 0-manifold.

Since we assume that we have only structurally stable limit sets, we obtain a decomposition of the domain into disjunct α -basins and a decomposition into disjunct ω -basins if we regard the connected component of the inflow set and the outflow set also as limit sets.

Theorem 3.15 Let $v : D \rightarrow \mathbb{R}^2$ be a Lipschitz continuous vector field with only structurally stable limit sets $L_i \subset D, i \in I$. Let $A_i \subset D$ denote the α -basins of the limit sets and $\Omega_i \subset D$ denote the ω -basins. Then:

$$D = \bigcup_{i \in I} A_i \quad (18)$$

$$D = \bigcup_{i \in I} \Omega_i \quad (19)$$

As topology of the vector field we define the set of limit sets L_i and the connected components of the intersections $T_{ij} = A_i \cap \Omega_j$.

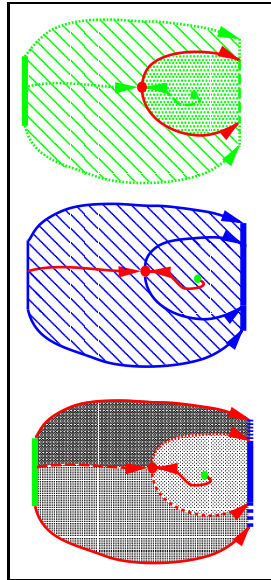


Figure 3: A simple vector field is first partitioned into α -basins and then into ω -basins. The resulting different intersections of α - and ω -basins are shown in the bottom image. All α -limit sets and α -basins are colored green, all ω -limit sets and ω -basins are colored blue and the separatrices and saddles are colored red. The intersections are shown in different black patterns.

Figure 3 shows the decomposition of a simple academic example flow into α -basins, ω -basins, and their intersections. Since the basins of sources, sinks, cycles, inflow, and outflow regions are all open subsets, the topology can be described by the one-dimensional basin intersections that form the boundaries of the open basin intersections. Because they separate the open basins, they are called **separatrices**. Since the boundary of an open set does not belong to the open set, the separatrices must start or end at saddles or boundary saddles. This fact is used to calculate the topology, i.e. we compute only the separatrices and obtain the other basin intersections as the resulting faces in the domain. All limit sets form the **topological skeleton** S_v together with the separatrices.

4 Evolution of Topology

We turn now to the analysis of unsteady vector fields. Our perspective is to look at the topology of the instantaneous integral curves (streamlines) and track their continuous evolution over time. The focus lies on the movement and substantial change of the limit sets. We do not consider the inflow and outflow regions here, but concentrate on critical points and the cycles. Again, the work is guided by results in the theory of dynamical systems. The main point is the search for structural changes of the vector field. As working definition, we use

Definition 4.1 *Let $v : D \times J \rightarrow \mathbb{R}^2$, $J \subset \mathbb{R}$ interval, be a Lipschitz continuous vector field. A time $s \in J$ is called **bifurcation time** if the topological skeleton $S(t)$ at $t \in (s - \delta, s)$ is different from $S(t)$ at $t \in (s, s + \delta)$, where δ is so small that the Skeleton $S(t)$ does not change for $t \in (s - \delta, s)$ or for $t \in (s, s + \delta)$.*

4.1 Local Bifurcations

There are bifurcations that involve only the vicinity of a critical point. They are called **local bifurcations**. We have two such types: static fold bifurcation and Hopf bifurcation [28].

Static Fold Bifurcation At the beginning, there are a saddle point and a sink, that are linked by a separatrix. If the attraction relation between both critical points along the separatrix weakens

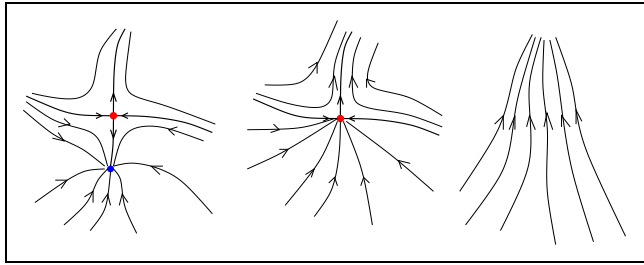


Figure 4: Pairwise annihilation

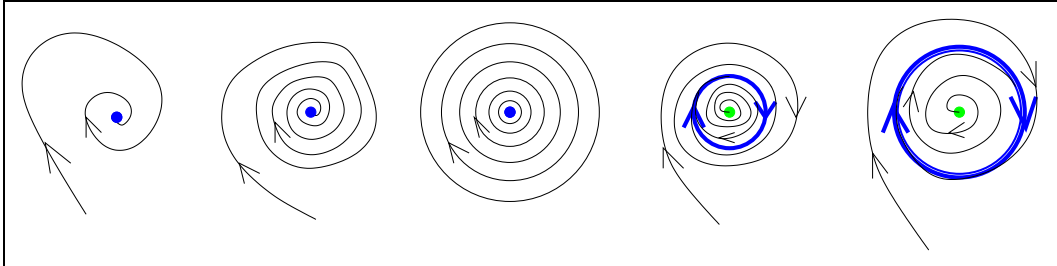


Figure 5: Hopf bifurcation

(i.e. the real positive eigenvalue of the Jacobian at the saddle point, corresponding to the considered separatrix, decreases and the negative real part of one of the Jacobian's eigenvalues at the sink increases), both critical points become closer and closer until they meet. At this point, an unstable critical point appears. As time goes on, this unstable structure vanishes: The new stable structure contains no critical point: This is a pairwise annihilation or static fold catastrophe (see figure 4). Inverting the direction of time, we get a pairwise creation. Similar transitions are obtained by inverting the direction of the flow (the sink becomes a source).

Hopf Bifurcation. We start with a sink with spiraling behavior in its neighborhood. If the attracting effect of this sink weakens, the number of streamline rotations around this critical point increases (the convergence “slows down”). This corresponds mathematically to an increasing negative real part of the complex conjugate eigenvalues of the Jacobian matrix at the critical point. When the real part is zero, we get a center point, which is an unstable structure: A bifurcation has occurred. If the real part increases further, a new stable structure appears which consists of an attracting closed orbit typically moving away from the critical point. The critical point is transformed into a source. So, a sink has changed into a source with the emission of a cycle: This is a Hopf bifurcation. An illustration of this evolution is given in figure 5. Inverting the direction of time, we get the transition from a cycle containing a source into a sink over an instantaneous center. Similar transitions are obtained by inverting the direction of the flow (that is by replacing sources by sinks and vice versa).

4.2 Global Bifurcations

There are also three important global bifurcations in 2D, i.e. bifurcations that do not take place in an arbitrary small neighborhood of a point. The three types are periodic fold bifurcation, periodic blue sky in 2D bifurcation, and basin bifurcation. We describe them in detail.

Periodic Fold Bifurcation. Figure 6 shows one structurally stable cycle enclosing another one in the left picture. In this situation without critical point in the ring between two cycles, one cycle is a source and the other cycle is a sink. If the two cycles drift towards each other, they eventually merge into an unstable cycle at the bifurcation time as can be seen in the middle of the figure. After this, both cycles have disappeared as in the right picture. Of course, a reversion of time is also possible, where two cycles around at least one critical point are created out of nothing.

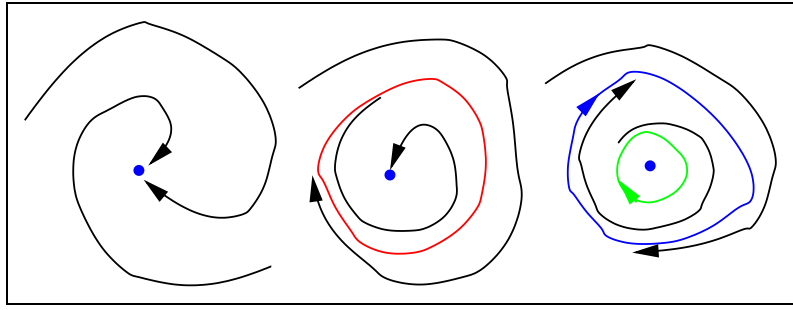


Figure 6: Periodic fold bifurcation

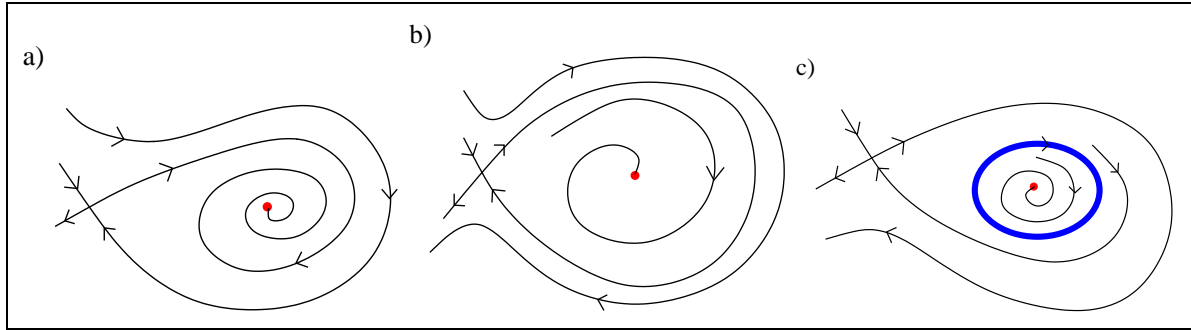


Figure 7: Periodic Blue Sky in 2D

Periodic Blue Sky in 2D. In this type of bifurcation there are two different types of critical points involved: a saddle and a sink. Figure 7a shows the situation. As the attracting effect of the sink gets weaker and weaker, we see a homoclinic connection after some time where the saddle is connected to itself as shown in figure 7b. This results in a bifurcation: when this configuration breaks up again we find a limit cycle which simply appears out of the blue. The reason for the occurrence of the closed streamline is that the sink is totally unaffected by the whole event. Since there is an outflow to the critical point inside and to the saddle there must be a critical point or a closed streamline in this region according to the Poincaré-Bendixson theorem [29]. Because of the fact that there are only the two critical points a closed streamline emerged. This configuration is shown in figure 7c. Other bifurcations of the same type can be constructed by replacing the sink with a source or by inverting the direction of time.

Basin Bifurcation The remaining bifurcation that plays a role in our analysis does not involve critical points or cycles but only separatrices. It is called basin bifurcation, since only the basins and therefore the connectivity of the limit sets changes. It is characterized by a heteroclinic connection between two saddles at the bifurcation time. Figure 8 shows this event. The separatrices starting at the two saddles come closer and closer until they coincide at the bifurcation time. After this, the connection between the limit sets has changed — we have other basins than before. The evolution of topology can now be defined.

Definition 4.2 Let $v : D \times J \rightarrow \mathbb{R}^2$, $J = [s_0, s_1] \subset \mathbb{R}$ an interval, be a Lipschitz continuous vector field. Let the vector field $v_t : D \rightarrow \mathbb{R}^2$, $v_t(x) = v(x, t)$ contain only structurally stable limit sets for all $t \in J$ except at a finite number of bifurcation times $t_1, \dots, t_i, \dots, t_m \in J$. The **evolution of topology** of v is the topology at s_0 , the maps of the movement of all limit sets, the maps of the movement of the separatrices, and the type and location of all bifurcations.

We assume that our unsteady vector field is structurally stable, so that there takes place only one bifurcation at a bifurcation time. Especially, we do not consider combinations of bifurcations that interfere with each other. The evolution can be described by the movement of the limit sets, the movement of the separatrices and the bifurcations. The information about the bifurcations includes type, time, location, and the involved limit sets as well as separatrices.

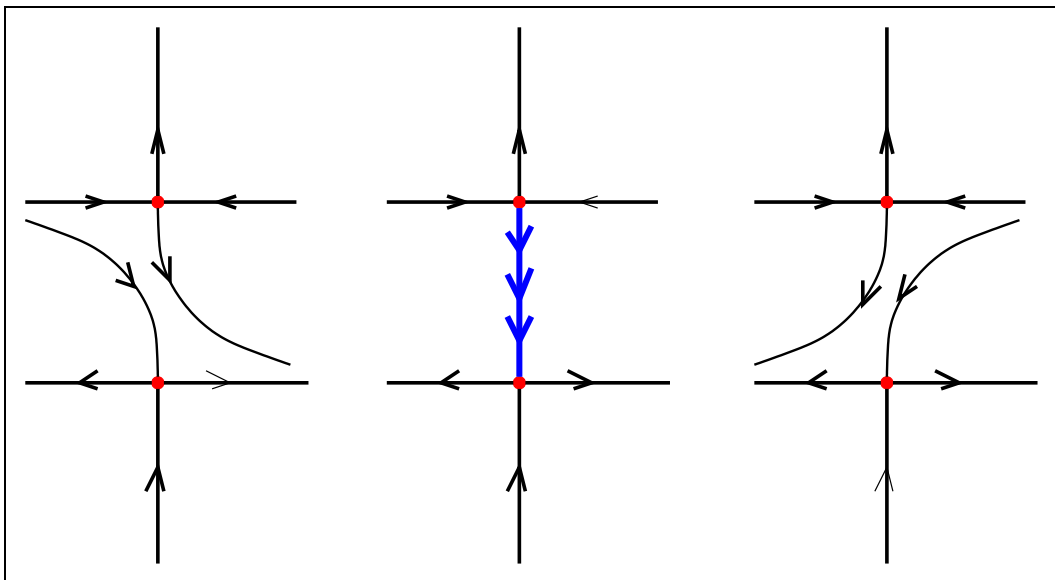


Figure 8: Basin bifurcation

5 Computation of Topology

For the topology, we need the limit sets and the basin boundaries as defined in section 3. Since we leave the boundary regions aside (for a clear analysis, see [33]), we are left with critical points, cycles, and the separatrices originating at saddles.

5.1 Critical Points

A critical point is a zero of the field, so we have to do a zero search. This task depends heavily on the used interpolation scheme. In our case, we obtain a cartesian grid in a plane cut along the rotation axis of the cylindric domain from the numerical method. Since we want to keep the computation effort, especially in the next section, at a manageable level, we triangulate the grid and use linear interpolation in each triangle. For the critical points, we can now solve a simple linear equation in each triangle. If the resulting zero is in the triangle, we have a critical point. Its type is easily found by calculating the (complex) eigenvalues of the Jacobian:

- Two negative real parts correspond to a sink.
- Two positive real parts correspond to a source.
- Mixed real parts correspond to a saddle.

A zero eigenvalue means a whole critical line of zeros in the triangle which we have excluded by our assumption of structurally stable vector fields. (We have also not found this case in practice.) Figure 9 shows the different cases. An analysis of critical points at the vertices of the grid is a little bit more difficult, but it can be done, see [34].

5.2 Closed Streamlines

As mentioned before, a closed streamline $d : \mathbb{R} \rightarrow \mathbb{R}^2, t \mapsto d(t)$ is a streamline of a vector field v such that there is a $t_0 \in \mathbb{R}/\{0\}$ with $d(t + nt_0) = d(t) \forall n \in \mathbb{N}$ and d not constant. In this subsection we present an algorithm that detects whether an arbitrary streamline c converges to such a closed curve. This means that c has d as α - or ω -limit set depending on the orientation of integration. We do not assume any knowledge on the existence or location of the closed curve, so that the algorithm can detect closed streamlines. The principle of the algorithm works on any piecewise defined planar vector field where one can determine the topology inside the pieces. The

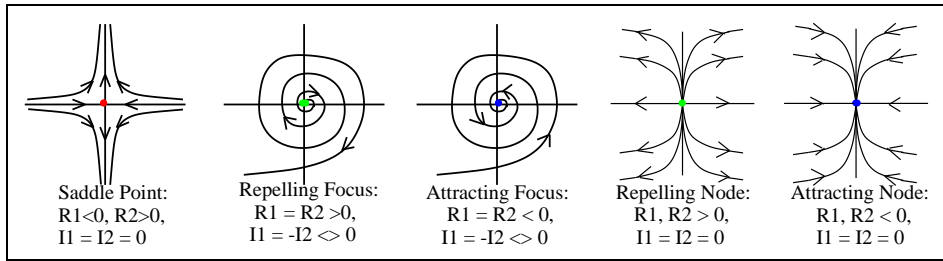


Figure 9: Common types of critical points.

basic idea of the algorithm is to determine a region of the vector field that is never left by a streamline. In case of a continuous vector field the Poincaré-Bendixson-Theorem ensures that this streamline approaches a closed streamline if no critical point exists in that region. A streamline approaching a closed streamline has to reenter the same cell. In this case we check if the cells were crossed by the streamline in the same order for the last two turns. This results in a *cell cycle* which identifies the above mentioned region. To examine if this cell cycle is left by the streamline we detect possible changes by checking the edges of the cells of the cell cycle. Therefore we identify points on each edge which we call *potential exits* where an outflow out of the cell cycle may occur in the vicinity. These points are identical with the vertices of the edge and points where the vector field is tangential to the edge.

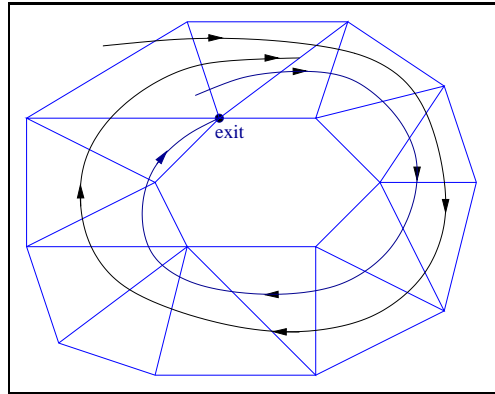


Figure 10: If a real exit can be reached, the streamline will leave the cell cycle.

We have to figure out if the actually investigated streamline will leave the cell cycle near such an exit. Therefore we integrate a streamline backwards from the potential exit to see if it leaves the cell cycle. If it does not leave after it crossed every cell of the cell cycle it converges to our streamline. We call this potential exit a *real exit* because the streamline will leave the cell cycle after a finite number of turns near that exit. Figure 10 displays an example for that case.

If the backward integrated streamline leaves the cell cycle, there will also be an entry point as shown in figure 11. A streamline starting at that point cannot be crossed by our actually investigated streamline. Consequently we cannot leave the cell cycle at this exit.

If there is no real exit for the streamline, we have proven that the streamline will never leave the cell cycle. If there is no critical point inside the cell cycle the Poincaré-Bendixson-Theorem ensures that there exists a closed streamline in our cell cycle and the integral curve tends toward it. The exact position of the closed streamline can be found using the Poincaré map.

Let us assume that we have a two-dimensional continuous vector field containing one closed streamline. Then we can choose a point P on the closed streamline and draw a *cross section* S which is a line segment nowhere parallel to the vector field. This line segment is then called a *Poincaré section*. If we start a streamline at an arbitrary point x on S and follow it until we cross the Poincaré section S again, we get another point $R(x)$ on S . This results in the *Poincaré map* R . Figure 12 illustrates the situation. The left part shows the Poincaré section with the closed

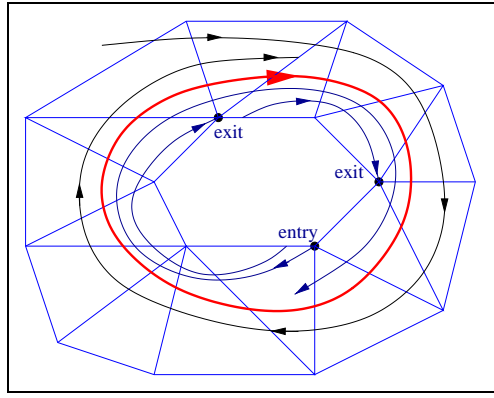


Figure 11: If no real exit can be reached, the streamline will approach a closed streamline.

streamline in the middle, drawn with a thicker line. The right part displays the Poincaré map itself.

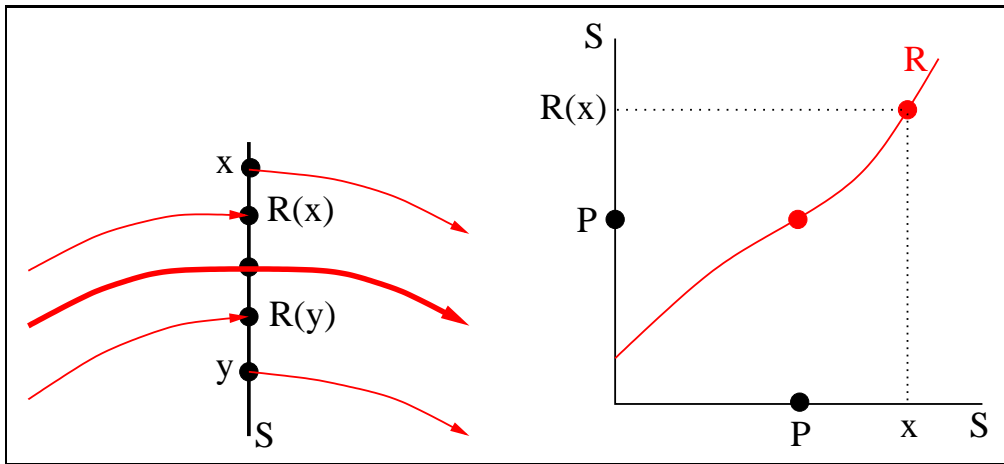


Figure 12: Poincaré section and Poincaré map.

R maps the point on the closed streamline onto itself because the closed streamline will intersect the Poincaré section always at the same point. Consequently, it is sufficient to find the fix point to get a point on the closed streamline. If we find a cell cycle we can use the edge where we detected the cell cycle for the first time as a Poincaré section. To find the fix point of the Poincaré map we do a binary search: first we divide the edge into two parts at the mid point of the edge. Then we check which part gets intersected by the streamline starting at the intersection point after one turn. This part is subdivided again and we start another streamline. This continues until we are close enough at the fix point of the Poincaré map. If we start a streamline at this point we get the whole closed streamline after we crossed every cell of the cell cycle. This method terminates because we proved in the previous step where we detected the cell cycle that we converge to a closed streamline.

5.3 Separatrices

In section 3, it was explained that the separatrices can be found at the saddle points. Since we have only linear saddles in our vector field, we can use the eigenvectors of the Jacobian to find the separatrices in the triangle containing the saddle, see figure 13. We extend the separatrices by using an adaptive Runge-Kutta-Fehlberg integration [32]. We check for reaching the boundary, running into a critical point, and approaching a cycle as explained above. This completes our

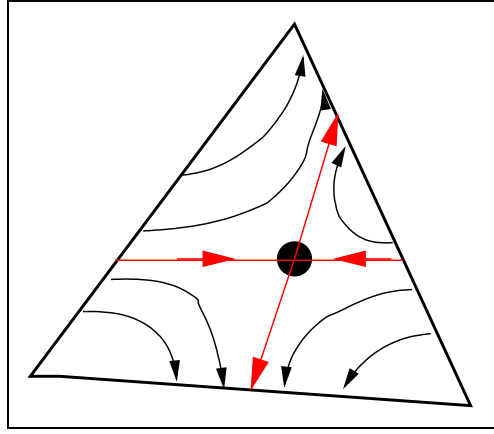


Figure 13: Separatrices for linear saddle.

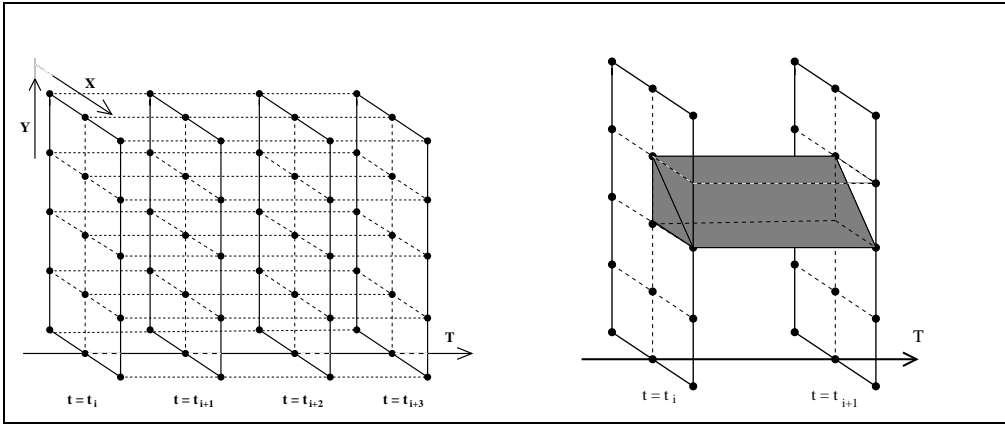


Figure 14: Space/Time grid

numerical analysis of the topology.

6 Computation of the Evolution

For the calculation of the evolution of the topology, we need an interpolation scheme over time. Since we want to keep the calculations simple, we do a linear interpolation between the stored time steps from the numerical simulation. The resulting grid is shown in figure 14.

Definition 6.1 Let $v_j : D \rightarrow \mathbb{R}^2$ be the given vector fields at time steps $t_0, \dots, t_i, \dots, t_m$. Then we define

$$\begin{aligned}
 v : D \times [t_0, t_m] &\rightarrow \mathbb{R}^2 \\
 (x, t) &\mapsto \frac{t_{i+1} - t}{t_{i+1} - t_i} v_i(x) + \frac{t - t_i}{t_{i+1} - t_i} \vec{a}_{i+1} v_{i+1}(x), \quad t \in [t_i, t_{i+1}].
 \end{aligned} \tag{20}$$

We can now use the results from section 5 to obtain all critical points, cycles, and separatrices at t_0 . We have to find now all the paths of critical points in all the prism cells, connect them and find the local bifurcations along the way.

6.1 Evolution of Critical Points

We locate and track critical points over time, using the interpolation scheme presented above. In particular, we seek the local bifurcations that may modify the topological structure: fold and Hopf bifurcations. To reduce the memory needs of our method, we do not process the whole grid at once: We consider only two consecutive time planes and handle the prism cells connecting them, forming a “time slice”. More precisely, we first determine for each prism cell the possible entry and exit positions of singular points and then reconnect these pieces of information to track every singularity from one time plane to the next.

Cellwise Analysis. The interpolant has been chosen so that it contains at most one critical point for a fixed value of t in a cell. This is due to the affine linear nature of its restriction to any time plane. If t moves from t_i to t_{i+1} , the singularity position describes a 3D curve. Yet, we are only interested in the curve sections that intersect the interior domain of the considered prism cell. A simple way to determine them is to consider the singularities lying on the side faces of the prism: Two triangular faces lying in $t = t_i$ and $t = t_{i+1}$ and 3 quadrilaterals connecting them. Finding the position of a singularity in each prism face requires the solution of a simple linear (resp. quadratic) system. If this is done, we sort the found (3D) positions in ascending time order and associate them pairwise. Indeed, since only one critical point can be present in a prism cell at a given time t , we know that a critical point must first leave the cell before it reenters it later. So, for each pair, we identify an entry and an exit position. We now need to pay attention to the singularity types to complete the topological information. The generic types of linear singularities are *source*, *sink* or *saddle* (omitting *center* points that correspond to a transition between source and sink). The transition from one type to another constitutes a bifurcation. Since a prism cell contains at most one critical point for a given t , we can assert that no fold bifurcation (involving two critical points) can occur inside it: *A fold bifurcation must occur at the cell boundary.* Consequently, we only seek Hopf bifurcations occurring between two consecutive entry and exit points. Practically, the determination of a singularity type is based upon the eigenvalues of the Jacobian at its position. Thus we compute the Jacobian matrix and its associated eigenvalues at each entry and exit point and check if they are the same. If this is the case, we can assert that no Hopf bifurcation has occurred since, otherwise, at least two bifurcations have taken place, which is impossible since our interpolant varies linearly over time. If the type has changed, a Hopf bifurcation is on the way. Because a Hopf bifurcation corresponds to a transition from a repelling to an attracting nature, the instantaneous nature of the intermediate singularity is a center and the trace of the Jacobian is zero at this point, which we use as criterion to determine the exact time location of the bifurcation. With the expression above, we obtain:

$$\text{tr}(J(t)) = \text{tr} \begin{pmatrix} a_{11}(t) & a_{12}(t) \\ a_{21}(t) & a_{22}(t) \end{pmatrix} = a_{11}(t) + a_{22}(t) = 0.$$

After straightforward calculus, we get finally

$$t = t_i - (t_{i+1} - t_i) \times \frac{a_{11}^i + a_{22}^i}{a_{11}^{i+1} - a_{11}^i + a_{22}^{i+1} - a_{22}^i}.$$

The corresponding position $(x(t), y(t), t)$ is the location of a Hopf bifurcation.

Tracking between the Prisms. At this stage, every cell knows the successive entry and exit positions of the critical point paths through its interior domain, as well as the possible presence of a Hopf bifurcation on the way. Yet, this information is scattered and must be reconnected to offer a global view of the topology evolution over time. A fundamental aspect of this task is to detect and identify the bifurcations that may take place on the faces. They are detected when connecting critical point path sections lying in neighbor cells as we show next.

We start in the time plane $\{t = t_i\}$. The restriction of the grid to this plane is a triangulation. In a boolean array, we indicate for each triangle if its corresponding prism must be further inspected or not. If it has to be inspected, we check the information collected during the cell analysis for a path section starting at an entry point located on the considered triangle face. This provides us

with an exit point at which the tracking must be proceeded. This exit point can either lie on a side (quadrilateral) face of the prism or on the triangle face lying in $\{t = t_{i+1}\}$. In the latter case, we signify in the boolean array of the next time plane $\{t = t_{i+1}\}$ that this triangle must be further inspected. If the path leaves the cell at $t^* < t_{i+1}$ (side face), then it enters a neighboring cell at the same time. Back to a 2D representation, we get in time plane $\{t = t^*\}$ a critical point lying on the common edge of both neighboring triangle cells. Due to the discontinuity of the Jacobian matrix through this edge, the critical point may have another type when considered from the other side. This simple argument explains the possible existence of a bifurcation in such cases. Two situations may actually occur.

- There is a simple crossing of a critical point from one cell to another cell where no critical point was present so far. In this case, the type may change but, a saddle remains a saddle. A source may become a sink and vice versa. As mentioned previously, the transition source/sink or sink/source is a so-called Hopf bifurcation. We easily detect it by checking if the type of a source (resp. a sink) path changes after passing the face.
- The second situation corresponds to a merging of two coexisting critical points in neighboring cells on their common face (common edge in 2D). Now, an important property of the piecewise affine linear interpolant over a triangulation is that two neighboring triangles cannot contain two critical points with the same type which implies that only a saddle and a source or a saddle and a sink can merge that way. This type of bifurcation has been considered in section 4: It is a static fold bifurcation or its inverse.

When we leave the current cell through a side face, and a possible bifurcation has been detected, we identify this position. We proceed with tracking the path by asking the neighbor cell for the next exit position. We get three possible answers.

1. The exit position lies in the time plane $\{t = t_{i+1}\}$: We have reached the next time plane. The corresponding triangle face is marked `true` in the boolean array corresponding to the next time plane.
2. The “exit” position lies in the time plane $\{t = t_i\}$: If we were tracking the current path forward so far, we have found a pairwise annihilation. In this case, the corresponding triangle is set to `false` in the boolean array of $\{t = t_i\}$, indicating that this cell has been processed now.
3. The exit position lies on a side face for $t = t^\#$. If $t^\# < t^*$ and we were tracking the current path forward so far, we have found a pairwise annihilation and proceed with backward tracking. If $t^\# > t^*$ and we were tracking the current path backward so far, we have found a pairwise creation and proceed with forward tracking. In this case, we are back to the current situation where one enters a cell through a side face.

The possible cases are illustrated in figure 15.

6.2 Tracking the Cycles

Since there must be a critical point inside each closed streamline we use the critical point path containing a Hopf bifurcation as a starting point for our streamline algorithm from subsection 5.2 which detects the closed streamline if it exists. Therefore we follow the critical point path in discrete steps in positive and negative directions starting at the bifurcation. After we have found the cell cycle containing the closed streamline we find the exact position using the Poincaré-map. Then we have to check if the closed streamline really surrounds the critical point. This is necessary because the streamline may have ran in another closed streamline in a totally different region of the flow. Obviously, closed streamlines surrounding the critical point occur only in one of the two temporal directions. This process continues until the cycles reach either another bifurcation which breaks them up or the border of the grid.

The left of figure 16 shows the result of this step, where we have found the closed streamlines at various timesteps. The closed streamlines are approximated by several line segments and the paths

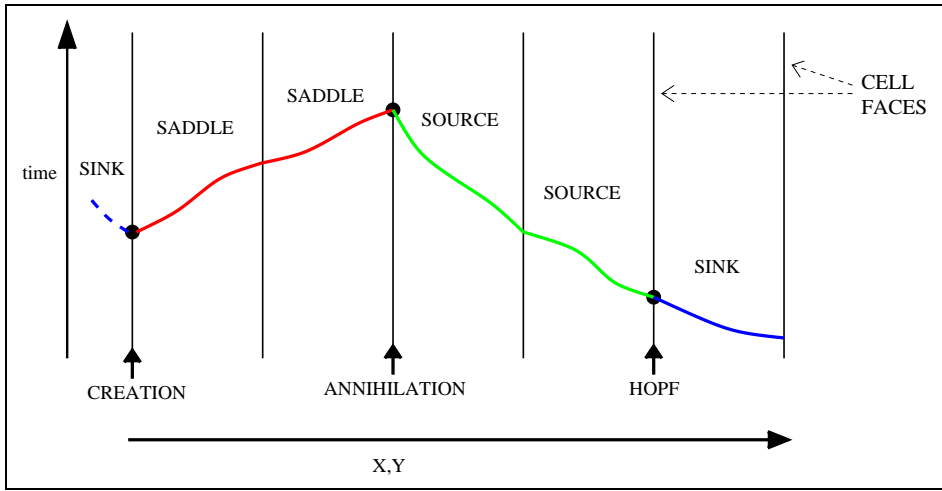


Figure 15: Possible cases when leaving a prism cell through a rectangular face

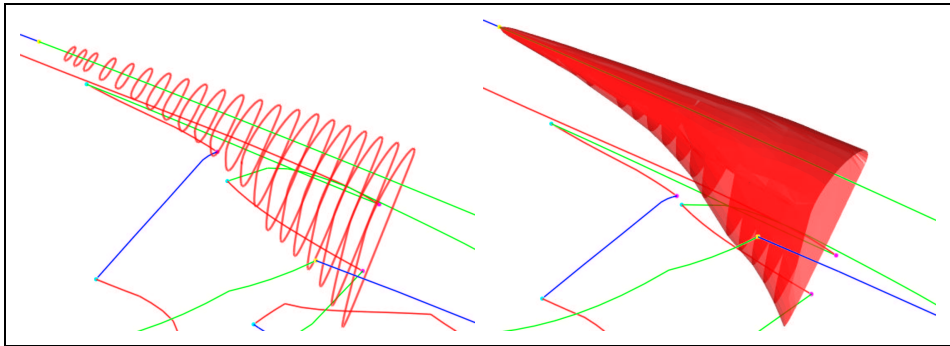


Figure 16: Closed streamlines visualized as wireframe and as a tube over time.

of the critical points are also shown using the same colors as in the original paper [35]. The Hopf bifurcation, where we started to detect the closed streamlines, is marked with a yellow sphere. In this example the life cycle of the closed streamline is started by a Hopf bifurcation and terminated by a Periodic Blue Sky in 2D bifurcation.

To visualize the evolution of closed streamlines, we construct tubes from the various closed streamlines similar to the pictures by Abraham and Shaw [28]. Therefore we construct surfaces consisting of triangles which connect the approximating line segments of the closed streamlines. The bifurcation point is connected to the tube using a parabolic surface approximated with triangles. The result is shown at the right of figure 16. We used an alpha value of 0.7 to get a better three dimensional impression and to prevent the tubes from hiding the paths of the critical points.

6.3 Separatrices

For the separatrices, we calculate each of them in smaller steps than the stored time steps from the simulation. We check for changes in the connectivity to find global bifurcations. It is possible to connect the instantaneous separatrices by surfaces visualizing their evolution, see [35]. Since these surfaces tend to cover the whole display, we do not show them in the final results. Right now, we do not have a method to find the basin bifurcations directly like the local bifurcations, so this is a good point for further research. During the calculation of the separatrices, we check for cycles all the time using the method from section 5. In this way, we hope to detect periodic fold bifurcations. If we find a cycle from one side, we go on with integral curve computation on an arbitrary point on the other side of the cycle to check for enclosed cycles. This allows to find all cycles for a structurally stable vector field. (Of course, we may miss some cycles due to our disregard of boundary saddles. Checks with random integral curves have not shown such missed

cycles up to now.)

7 Results and discussion

The experiments of Billant *et al.* [2] document the breakdown of swirling water jets for Reynolds numbers in the range $300 < Re < 1200$ and swirl numbers S in the range $[0, 1.42]$. The swirl number S is defined by [2]

$$S \equiv \frac{2v_\theta(R/2, z_0)}{v_z(0, z_0)} \quad (21)$$

where $z_0 = 0.4D$, $D = 2R$ being the nozzle diameter. The Reynolds number is

$$Re \equiv \frac{v_z(0, z_0)D}{\nu} \quad (22)$$

The simulation of the 3d viscous flow calculates the values on a cylindrical grid. From this grid, we only take the values on a halfplan with the axis of the cylinder as one side and the radius as the other side. This results in an cartesian grid due to the set up of the direct numerical simulation. In figure 17, one can see the topology of the instantaneous integral curves after 20 timesteps. The figure does not show the whole grid but most of it. It is not difficult to see that the whole action takes place in a small area in the lower right corner. One can see vortical motion in action and might ask for a closer look at the critical points and cycles. For a better analysis, it may be mentioned that the program sets very small values to zero at the vertices to avoid a large number of critical points in the empty zone of the grid that may occur from numerical errors. (If the values are very close to zero, the simulation method creates a large number of extremely weak critical points that do not have any physical evidence, but result in a very disturbing visualization.) In figure 18, we present a closer look at the topological structure of the flow. The flow does not contain a cycle right now, but one can see various blue sinks and green sources. All the curves are separatrices created by the red colored saddle points. The green part of the grid around the dominant structure in the middle is now used for an analysis of the topology evolution. We show this evolution in figure 19. The cone below shows the direction of time from time step 20 at the circle to time step 70 at the top. One can see clearly the paths of the saddles, sinks, and sources. All the small spheres stand for bifurcations. A static fold is shown by a purple sphere. A light blue sphere shows its inverse, i.e. a pairwise creation of a sink and a saddle or a source and a saddle. The yellow spheres denote Hopf bifurcations. As before, the transparent red surfaces show the evolution of cycles. They can either end/start at a Hopf bifurcation or at a blue sky in 2d bifurcation. As mentioned before, we have found no evidence for periodic fold bifurcations. Since we do not show the separatrices, one can also not see basin bifurcations.

We use the axisymmetric simulation to indicate some of the problems that we still have to solve. Figure 20 shows a large number of Hopf bifurcations along the trace of some of the sinks/sources. This is difficult for the analysis since it is nearly impossible to check for the cycles created (and destroyed) by these bifurcations. We present also a close-up in figure 21. Here, one can see that there are also sometimes larger number of static fold bifurcations that one would like to ignore since they are obviously not important for the behavior of the flow.

8 Conclusions

The flow visualisations of Billant *et al.*[2] showed the development of a conical shear layer in swirling jets. Using a simplified analytical model, it can be shown that this conical shear layer produces sheets of azimuthal vorticity of both signs. From the DNS data it can be inferred that these sheets emerge as conical shear layer and it is demonstrated to have a spatial structure similar to a conical jet with layers of vorticity of opposite sign. The break up of this conical jet produces helical vortex tubes. The structure of these tubes was shown to be essentially counter-rotating strands of vorticity. The cosine of the angle between velocity and vorticity is mostly negative, indicating opposite orientation of the vectors, but positive values are also present to a lesser extent. Hence, it must be concluded that tangled vortex lines are present within the tubes, which is also

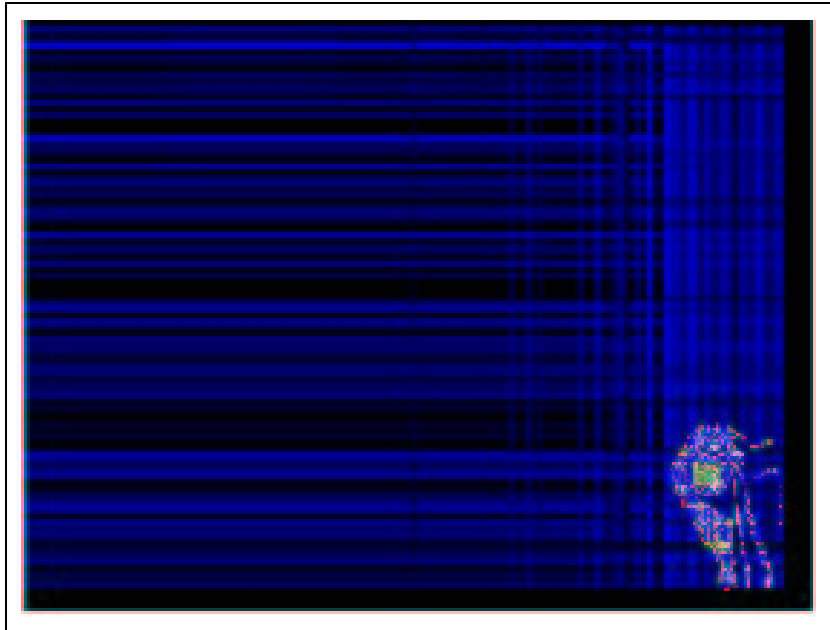


Figure 17: Topology of 3d viscous jet at time step 20.

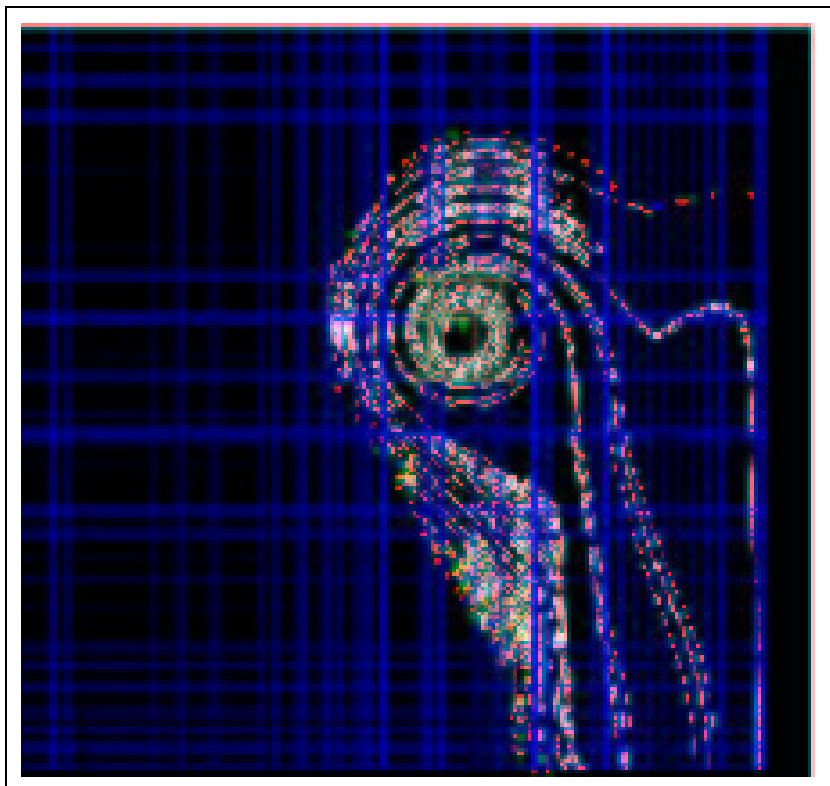


Figure 18: Interesting part of 3d viscous jet at time step 20. The green part of the grid is analyzed over time in the next figure.

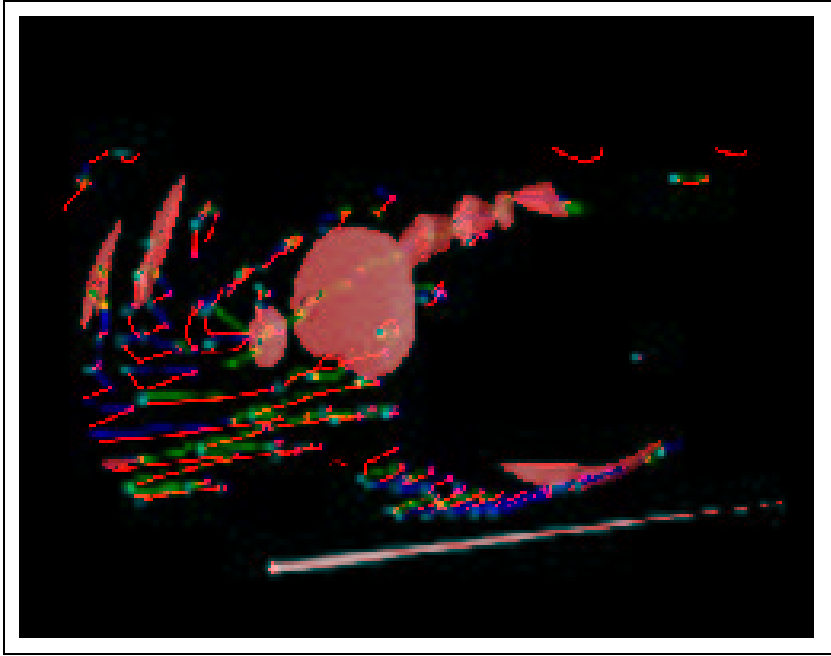


Figure 19: Evolution of the topology in the marked area over time.

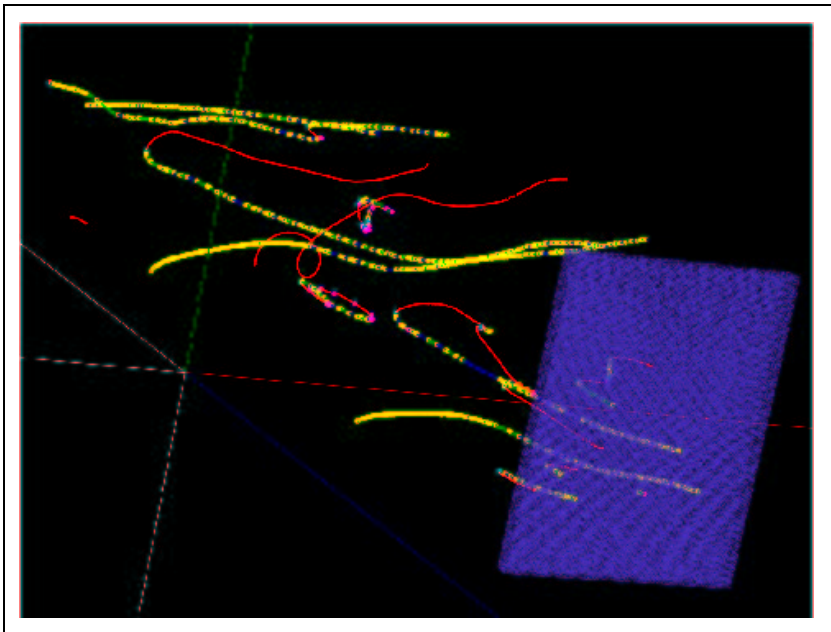


Figure 20: Cascade of Hopf bifurcations in the axisymmetric case.

indicated by the distribution of helicity density. The tubes are recognizable for about two jet diameters and become engulfed in the weakly turbulent flow in the core of the break-up region further downstream.

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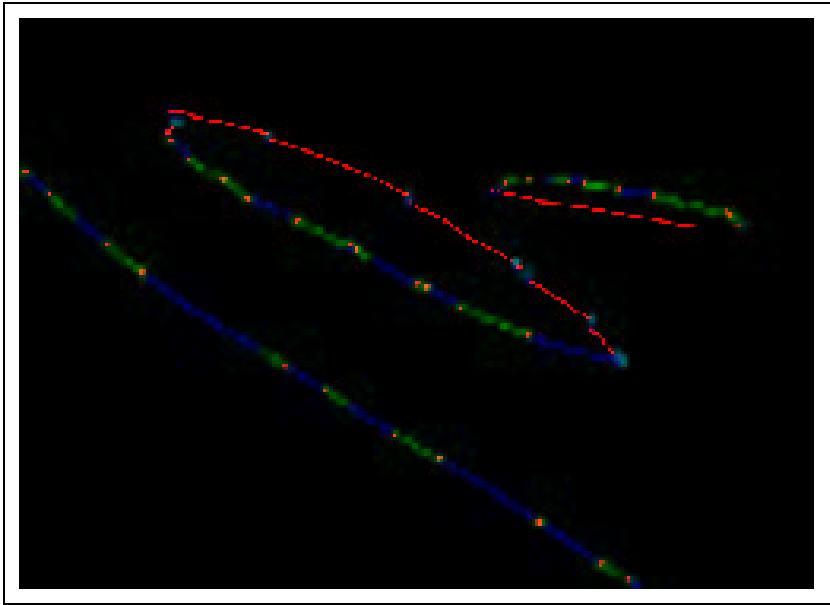


Figure 21: Zoom into some of the cascades.