Small scale reconstruction in three-dimensional Kolmogorov flows using four-dimensional variational data assimilation

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Significant insights in computational fluid dynamics are obtained in recent years by adopting the data assimilation methods developed in the meteorology community. We apply the four dimensional variational method to reconstruct the small scales of three-dimensional turbulent velocity fields with a moderate Reynolds number, given a time sequence of measurement data on a coarse set of grid points. The problem presents new challenges because the evolution of the flow is dominated by the nonlinear vortex stretching and the energy cascade process, which are absent from two dimensional flows. The results show that, reconstruction is successful when the resolution of the measurement data, given in terms of the wavenumber, is at the order of the threshold value \( k_c = 0.2\eta_K \) where \( \eta_K \) is the Kolmogorov length scale of the flow. When the data are available over a period of one large eddy turn-over time scale, the filtered enstrophy and other small scale quantities are reconstructed with a 30\% or smaller normalized point-wise error, and a 90\% point-wise correlation. The spectral correlation between the reconstructed and target fields is higher than 80\% for all wavenumbers. Minimum volume enclosing ellipsoids (MVEEs) and MVEE trees are introduced to quantitatively compare the geometry of non-local structures. Results show that, for the majority samples, errors in the locations and the sizes of the reconstructed structures are within 15\%, and those in the orientations is within 15\(^\circ\). Overall, for this flow, satisfactory reconstruction of the scales two or more octaves smaller is possible if data at large scales are available for at least one large eddy turn-over time. In comparison, a direct substitution scheme results in three times bigger point-wise discrepancy in enstrophy. The spectral difference between the reconstructed and target velocity fields is more than ten times higher than what is obtained with the four dimensional variational method. The results show that further investigation is warranted to verify the efficacy of the method in flows with higher Reynolds numbers.

1. Introduction

Data assimilation (DA) uses available experimental or observational data to improve computational model predictions. It has had a long history in meteorology, in particular numerical weather prediction (NWP) research (see, e.g., Courtier \etal(1993) and Kalnay (2003) for reviews). Many DA methods have been developed. Four dimensional variational (4DVAR) methods and ensemble Kalman filter (EnKF) are among the most popular
methods (Kalnay 2003; Kalman 1960; Evensen 2009). 4DVAR are based on optimal control (Lions 1971), where a constrained optimization problem is solved. Spatial as well as temporal data are assimilated into the model by minimizing the difference between the model prediction and the data. In recent applications, time sequences of three dimensional spatial data have been assimilated, hence the name 4DVAR. EnKF is a sequential method where the error of the prediction is estimated when the prediction (known as the a priori estimate (Brown & Hwang 2012) or the background (Kalnay 2003)) is made. The observational data are assimilated by combining the a priori estimate with the measurement data, resulting in an updated prediction called the a posteriori estimate or the analysis. The weight for the measurement is chosen to minimize the error of the a posteriori estimate.

In recent years, these two methods have been applied to fluid dynamic problems (Hayase 2015). Gronskis et al. (2013) applies the variational method to reconstruct the inflow and initial conditions for two-dimensional (2D) mixing layers and wake flows behind a cylinder. Mons et al. (2016) considers 2D wake flows, where a comprehensive comparison between different DA schemes is presented. Kato & Obayashi (2013), Kato et al. (2015) and Li et al. (2017) use the EnKF method to improve Reynolds-Averaged Navier-Stokes (RANS) based prediction of turbulent flows. Meldi & Poux (2017) applies EnKF to large eddy simulations (LES) and detached eddy simulation (DES). Similar problems are also investigated with variational methods (Foures et al. 2014). In D’adamo et al. (2007), Artana et al. (2012), and Protas et al. (2013), variational methods are coupled with reduced order models based on Galerkin truncation. Variational methods are also used in state estimation in the context of flow control (Bewley & Protas 2004; Chevalier et al. 2006; Colburn et al. 2011), to extrapolate experimental data in a dynamically consistent way (Heitz et al. 2010; Combes et al. 2015) and to obtain optimal sensor placements (Mons et al. 2017). Mons et al. (2014) applies a variational method to investigate decaying isotropic turbulence, where the eddy-damped quasi-normal Markovian model is used.

The aforementioned research has demonstrated the potential of DA in turbulent simulations as well as in understanding the physics of turbulent flows. As having been demonstrated in NWP research, DA is unique in its ability to improve our modelling or prediction of instantaneous flow fields, in addition to their statistics. However, few studies have explored this aspect in the simulation of 3D turbulent fields. (see, e.g., the list of recent works on DA tabulated in Mons et al. (2017)). In Yoshida et al. (2005), the ability of a DA scheme to recover the instantaneous small scales in a 3D isotropic turbulent field is investigated. In this study, the data, given as Fourier modes, directly replace model predictions given by the NSEs at every time step. It is found that the small scale instantaneous velocity field can be asymptotically recovered exactly when Fourier modes with wavenumber up to a threshold value approximately equal \( k_c = 0.2\eta^{-1} \) are provided, where \( \eta_k \) is the Kolmogorov length scale. When the amount of data decreases towards the threshold, the time needed to recover the small scales tends to infinity (hence an infinitely long time sequence of measurement data is needed). The problem is investigated in Lalescu et al. (2013) from the perspective of chaos synchronization and a similar conclusion is reached. The DA method used in these works can be termed ‘direct substitution’ (DS). As far as we know, the ability of 4DVAR or EnKF to reconstruct the small scales has not been investigated. In this paper, we present an analysis based on the 4DVAR method. We consider a Kolmogorov flow in a 3D periodic box. It is assumed that a time sequence of velocity data are given on a set of grid points. 4DVAR is employed to reconstruct the initial velocity field such that the velocity at later times matches given measurement data. The time sequence of velocity fields computed from the initial
field are compared with the ‘true’ velocity fields (the target fields). The objective is to ascertain how well the instantaneous small scale velocity fields can be reconstructed with 4DVAR for a given set of parameters. The 3D turbulence in a periodic box is the simplest turbulent flow where the nonlinear inter-scale interaction plays the dominant role in the dynamics. The vortex stretching process, e.g., is absent in 2D flows. Therefore, although the mathematics is similar for 2D and 3D problems, the latter does present significant new challenges.

Using the 4DVAR method to reconstruct 3D fully developed (i.e., statistically stationary) turbulent field is the first contribution of this investigation. To evaluate the reconstruction, it is important to quantify the instantaneous difference between the reconstructed and the target velocity fields. Much information can be learned from point-wise correlations and/or the statistics of point-wise difference. However, these statistics are not best suited to capture the geometry of non-local structures populating the small scales of turbulent fields (such as the vortex filaments). This deficiency becomes a significant obstacle when the distribution of the non-local structures becomes one of our main interests.

The morphology of non-local structures have long been described in qualitative terms, assisted by visualization. Great efforts have been made in recent years to develop methods for quantitative description. Bermejo-Moreno & Pullin (2008) uses the probability density function (PDF) of the curvatures on the surface of a structure as its signature. Yang & Pullin (2011) describes the structures in a channel flow in terms of curvelets and angular spectra. Indices named shapefinders, defined in terms of the Minkowski functionals, are used in Leung et al. (2012) to classify the structures. These methods provide very detailed quantitative descriptions of the structures. However, they focus on the intrinsic geometry; the information about locations, orientations and sizes of the structures sometimes is missing, which happens to be important when we compare the geometry in two different fields. Therefore, as the second contribution of this investigation, we propose to use minimum volume enclosing ellipsoids (MVEEs) to describe the geometry of a nonlocal structure, and use MVEE trees where the structures are highly non-convex. MVEE is used widely in areas such as statistical estimate, cluster analysis and image processing (Todds 2016). Its application in turbulence research, however, has not been reported. The results demonstrate that MVEEs and MVEE trees are useful tool for the analysis of the non-local geometry in turbulence.

The paper is organized as follows. In Section 2 the 4DVAR formulation of the problem is introduced; the description of the small scales of homogeneous turbulence is reviewed, and then the calculation of MVEEs and MVEE tree is explained. In Section 3 the simulations and the results are presented and analyzed. The conclusions are summarized in Section 4.

2. The governing equations and the description of small scales of turbulent velocity fields

2.1. The problem setup and the optimality system

Let $B$ denote the three dimensional periodic box $[0,2\pi]^3$, and $v(x,t)$ be a time sequence of velocity fields in $B$ for $t \in [0,T]$ with $T > 0$. $T$ is called the optimization horizon hereafter. It is assumed that only part of $v(x,t)$ is known from measurement, and this part is denoted by $Fv$, where $F$ is a filter to be defined later. $F$ plays the same role as the measurement operator used in the meteorology community. The question is, from given $Fv(x,t)$, how to reconstruct the full field $v(x,t)$ for $t \in [0,T]$. 
To do so, an underlying ‘model’ for the data \( \mathbf{v}(x,t) \) has to be assumed. We assume \( \mathbf{v}(x,t) \) is the solution of the Navier-Stokes equations (NSEs), from some unknown initial condition (IC). Let \( \mathbf{u}(x,t) \) be a solution of the NSEs in \( B \). If one finds a velocity field \( \varphi(x) \) such that, if \( \mathbf{u}(x,0) = \varphi(x) \) then \( \mathbf{u}(x,t) \) agrees with \( \mathbf{v}(x,t) \) for \( (x,t) \in B \times [0,T] \), then \( \mathbf{u}(x,t) \) is a reconstruction of \( \mathbf{v}(x,t) \). It is unlikely to find \( \varphi(x) \) such that \( \mathbf{u}(x,t) = \mathbf{v}(x,t) \) exactly. In 4DVAR, one attempts to approximate \( \varphi(x) \) by the solution of a constrained optimization problem. The detail is described in what follows.

We define the inner product between two scalar or vector fields \( \mathbf{a}(x,t) \) and \( \mathbf{b}(x,t) \) as

\[
\langle \mathbf{a}, \mathbf{b} \rangle \equiv \frac{1}{(2\pi)^3} \int_{x \in B} \mathbf{a} \cdot \mathbf{b} \, dx. \tag{2.1}
\]

The cost function \( J \) for the optimization problem is defined as

\[
J = \frac{1}{2} \int_{0}^{T} \langle \mathcal{F} \mathbf{u} - \mathcal{F} \mathbf{v}, \mathcal{F} \mathbf{u} - \mathcal{F} \mathbf{v} \rangle dt. \tag{2.2}
\]

Qualitatively speaking, \( J \) is the total difference between the measurements based on \( \mathbf{u} \) and \( \mathbf{v} \) over the space-time domain \( B \times [0,T] \). The objective of the optimization problem is to find \( \varphi(x) \) such that \( J \) is minimized while \( \mathbf{u}(x,t) \) satisfies

\[
\mathbf{N}(\mathbf{u}, p) \equiv -\partial_t \mathbf{u} - \mathbf{u} \cdot \nabla \mathbf{u} - \nabla p + \nu \nabla^2 \mathbf{u} + \mathbf{f} = 0, \quad \nabla \cdot \mathbf{u} = 0, \quad \mathbf{u}(x,0) = \varphi(x), \tag{2.3}
\]

where \( p \) is the pressure, \( \nu \) is the kinematic viscosity and \( \mathbf{f}(x,t) \) is the forcing term. The flow has been assumed to be incompressible.

The optimal solution for the problem will be found with the adjoint method. We introduce the Lagrangian functional:

\[
L(\mathbf{u}, \xi, \sigma, \lambda, \varphi) = J(\mathbf{u}) - \int_{0}^{T} \langle \xi, \mathbf{N} \rangle dt - \int_{0}^{T} \langle \sigma, \nabla \cdot \mathbf{u} \rangle dt - \langle \lambda, \mathbf{u}(x,0) - \varphi(x) \rangle, \tag{2.4}
\]

where \( \xi(x,t), \sigma(x,t), \) and \( \lambda(x) \) are the adjoint variables corresponding to the NSEs, the continuity equation, and the IC. \( \xi \) and \( \sigma \) are also called the adjoint velocity and the adjoint pressure. The minima of \( J \) are found at the stationary points for the Lagrangian where the functional derivatives of \( L \) with respect to \( \mathbf{u}, \xi, \sigma, \lambda, \) and \( \varphi \) are zero. These conditions give rise to equations for the adjoint variables (i.e. the adjoint equations) in addition to Eq. (2.3). The equations for \( \xi \) read:

\[
-\partial_t \xi - \mathbf{u} \cdot \nabla \xi + \nabla \mathbf{u} \cdot \xi + \nabla \sigma - \nu \nabla^2 \xi - \mathbf{F} = 0, \quad \nabla \cdot \xi = 0, \quad \tag{2.5}
\]

with the end-point condition:

\[
\xi(x,T) = 0. \tag{2.6}
\]

The forcing term \( \mathbf{F} \) in Eq. (2.5) is given by

\[
\mathbf{F}(x,t) \equiv -\frac{\partial J}{\partial \mathbf{u}} = -\mathcal{F}^+ \mathcal{F}[\mathbf{u}(x,t) - \mathbf{v}(x,t)] \tag{2.7}
\]

where \( \mathcal{F}^+ \) is the adjoint operator of \( \mathcal{F} \). The forcing term \( \mathbf{f}(x,t) \) in Eq. (2.3) has been assumed to be independent of \( \mathbf{u} \) and, as a consequence, does not have a counter part in the adjoint equations. When \( \mathbf{u}, p, \xi \) and \( \sigma \) satisfy Eqs. (2.3) and (2.5-2.7), the gradient of \( J \) with respect to \( \varphi \), denoted by \( DJ/D\varphi \), is given by

\[
\frac{DJ}{D\varphi} = \frac{\partial L}{\partial \varphi} = -\xi(x,0). \tag{2.8}
\]

Eqs. (2.3), (2.5-2.7) and the condition \( DJ/D\varphi = 0 \) constitute the optimality system of
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the optimization problem. The solution of the optimization problem is a solution of the optimality system.

The solution method of the optimality system will be explained later. The optimal solution for \( \varphi(x) \) will be denoted as \( \varphi^u(x) \). The solution \( u(x,t) \) of Eq. (2.3) with \( \varphi(x) = \varphi^u(x) \) is the reconstruction of \( v(x,t) \) from \( \mathcal{F}v(x,t) \). In this investigation, a sequence of known fully developed turbulent velocity fields in the 3D periodic box \( B \) is used as \( v(x,t) \). \( \mathcal{F} \) is chosen as a cutoff filter in the Fourier space with a cutoff wavenumber \( k_m \), such that only the low wavenumber Fourier modes are used in the optimization problem. \( k_m \) is an indicator of the spatial resolution of the measurement data. As a result, the cost function \( J \) can be written as

\[
J = \frac{1}{2} \int_{0}^{T} \int_{k \leq k_m} (\hat{u}(k,t) - \hat{v}(k,t)) \cdot (\hat{u}^*(k,t) - \hat{v}^*(k,t)) dk,
\]

where \( k = |k| \), and \( \hat{v}(k,t) \) and \( \hat{u}(k,t) \) denote the Fourier modes of \( v(x,t) \) and \( u(x,t) \), respectively. Letting \( \hat{F}(k,t) \) be the Fourier transform of \( F(x,t) \), we obtain

\[
\hat{F}(k,t) = \begin{cases} 
- [\hat{u}(k,t) - \hat{v}(k,t)] & \text{for } k \leq k_m, \\
0 & \text{otherwise.}
\end{cases}
\]

2.2. The questions to be answered

We call \( u(x,t) \) and \( v(x,t) \) the ‘reconstructed’ and the ‘target’ fields, respectively. The target field at \( t = 0 \), \( v(x,0) \), will also be denoted by notation \( \varphi^v(x) \), corresponding to its reconstruction \( u(x,0) \equiv \varphi^u(x) \). Since \( u(x,t) \) is the solution of the NSEs with \( \varphi^u(x) \) as the IC, it is expected \( \hat{u}(k,t) \approx \hat{v}(k,t) \) when \( k \leq k_m \). The interesting question is how closely \( \hat{u}(k,t) \) matches \( \hat{v}(k,t) \) when \( k > k_m \); in other words, to what extent small scales in \( v(x,t) \) can be reconstructed.

As \( u(x,t) \) and \( v(x,t) \) are both solutions of the forced NSEs, one expects their statistics to be the same for \( t \rightarrow T \) when \( u \) has fully developed. Therefore, for \( t \rightarrow T \), we are interested in the point-wise, instantaneous difference or correlation between \( u \) and \( v \). However, good reconstruction at \( t \rightarrow T \) is possible only when the IC \( \varphi^u(x) \) captures to some extent the features of real turbulence. Therefore, we will also examine selected statistics of \( \varphi^u(x) \). The statistics provide information on how much can be reconstructed at \( t = 0 \) given the measurement data \( \mathcal{F}v(x,t) \), hence is also relevant to the observability of the system. Nevertheless, the full discussion of this aspect is left for the future.

The efficacy of 4DVAR can be assessed by comparison with other methods. As we will show later, the straightforward Lagrangian interpolation fails badly. In Yoshida et al. (2005), an assimilation scheme we call ‘direct substitution’ (DS) is used. The authors integrated the NSEs starting from a Gaussian random field. The low wavenumber modes with \( k \leq k_m \) in the solution \( u(x,t) \) are replaced by those in the target field \( v(x,t) \) at every time step. Yoshida et al. (2005) shows that, if \( k_m \geq 0.2/\eta^3 \), the small scales in \( u(x,t) \) approach those in \( v(x,t) \) asymptotically as \( t \rightarrow \infty \) (see also Lalescu et al. (2013)). In most cases we are investigating, the resolution \( k_m \) is slightly below this threshold (c.f., the parameters in Sect. 3.1), and the measurement data are available only for \( t \in [0,T] \) where \( T \) is at the order of one large eddy turn-over time scale. As a consequence, the DS scheme will only achieve partial reconstruction. The DS scheme will be compared with the 4DVAR method, to demonstrate the improvement provided by the latter.

2.3. Description of the small scales of turbulent velocity fields

To compare the small scales of the reconstructed field \( u \) and the target field \( v \), the filtering approach is used to separate different scales and the analysis is conducted mainly
in the physical space. The filtered velocity field is defined by

$$\tilde{u}(x, t) = \int G_\Delta(x - y)u(y, t)dy,$$  \hfill (2.11)

where $\tilde{u}$ denotes the filtered velocity and $G_\Delta(x)$ is a filter with length scale $\Delta$. $G_\Delta$ separates $u$ into $\tilde{u}$ and the subgrid-scale (SGS) velocity. The parameters used to describe the structures of $\tilde{u}$ and its interactions with the SGS scales are now summarized. In the analysis of the velocity field $u$, we always use the Gaussian filter \cite{Pope2000}.

The local structures of $\tilde{u}$ are described by the filtered velocity gradient $\tilde{A}_{ij} \equiv \partial_j \tilde{u}_i$, the filtered strain rate tensor $\tilde{s}_{ij} \equiv (\tilde{A}_{ij} + \tilde{A}_{ji})/2$, and the filtered vorticity $\tilde{\omega}_i \equiv \varepsilon_{ijk} \tilde{A}_{kj}$. For $\tilde{A}_{ij}$, much insight has been learned from its tensor invariants

$$Q \equiv -\frac{1}{2} \tilde{A}_{ij}\tilde{A}_{ji}, \quad R \equiv -\frac{1}{3} \tilde{A}_{ij}\tilde{A}_{jk}\tilde{A}_{ki}. \hfill (2.12)$$

One of the key features of turbulence is that the joint PDF of $R$ and $Q$ displays a skewed tear-drop shape, which captures the prevalence of straining motion in turbulent flows; see, e.g., \cite{Cantwell1992}.

The eigenvectors of $\tilde{s}_{ij}$ are denoted by $e_{\alpha}^s$, $e_{\beta}^s$ and $e_{\gamma}^s$, with corresponding eigenvalues $\lambda_{\alpha}^s \geq \lambda_{\beta}^s \geq \lambda_{\gamma}^s$. $\lambda_{\alpha}^s + \lambda_{\beta}^s + \lambda_{\gamma}^s = 0$ due to the incompressibility of the filtered velocity field. The enstrophy of the resolved velocity field, defined by $\tilde{\omega}^2 \equiv \tilde{\omega}_i\tilde{\omega}_i$, is a measure of the magnitude of the resolved vorticity. The main source of growth for $\tilde{\omega}^2$ is the vortex stretching term $P_\omega \equiv \tilde{s}_{ij}\tilde{\omega}_i\tilde{\omega}_j$. Evidently,

$$P_\omega = \tilde{\omega}^2(\lambda_{\alpha}^s \cos^2 \theta_{\alpha}^s + \lambda_{\beta}^s \cos^2 \theta_{\beta}^s + \lambda_{\gamma}^s \cos^2 \theta_{\gamma}^s), \hfill (2.13)$$

where $\theta_{\alpha}^s$ is the angle between $\tilde{\omega}$ and $e_{\alpha}^s$ with $\theta_{\beta}^s$ and $\theta_{\gamma}^s$ defined in a similar way. Eq. (2.13) shows that $P_\omega$ are determined by the eigenvalues of $\tilde{s}_{ij}$ as well as the alignment between $\tilde{\omega}_i$ and the eigenvectors of $\tilde{s}_{ij}$. The eigenvectors of $\tilde{s}_{ij}$ form an orthogonal coordinate frame. The orientation of $\tilde{\omega}_i$ can be described by the polar and azimuthal angles relative to $e_{\alpha}^s$ in this frame, and they are denoted by $\theta_{sp}^\omega$ and $\phi_{sp}^\omega$. Fig. 1 gives an illustration of the angles. In turbulence, by examining the joint PDF of $\cos \theta_{sp}^\omega$ and $\phi_{sp}^\omega$ or the PDFs of $\theta_i^s$ ($i = \alpha, \beta, \gamma$), it has been shown that $\tilde{\omega}_i$ prefers to align with $e_{\alpha}^s$.

The dynamics of $\tilde{u}_i$ is governed by the filtered NSEs, which is unclosed due to the SGS stress tensor $\tau_{ij} = \tilde{u}_i\tilde{u}_j - \tilde{u}_i\tilde{u}_j$ (see, e.g., \cite{MeneveauKatz2000}). $\tau_{ij}$ represents the nonlinear interactions between the resolved and the SGS scales. An important parameter related to $\tau_{ij}$ is the SGS energy dissipation rate

$$\Pi_\Delta = -\tau_{ij}^d\tilde{s}_{ij} \hfill (2.14)$$

where $\tau_{ij}^d \equiv \tau_{ij} - \delta_{ij}\tau_{kk}/3$ is the deviatoric part of $\tau_{ij}$. $\Pi_\Delta$ is the turbulent kinetic energy flux cascading from resolved scales to the SGS scales. Correctly reproducing the statistics of $\Pi_\Delta$ is one of the main objectives in SGS modelling. As a consequence, the statistics of $\tau_{ij}^d$ and $\Pi_\Delta$ and their correlations with $\tilde{s}_{ij}$, $\tilde{\omega}_i$, $P_\omega$ and $\tilde{A}_{ij}$ have been extensively documented; see, e.g., \cite{MeneveauKatz2000} and \cite{Sagaut2002} for reviews on SGS modelling and large eddy simulations.

$\Pi_\Delta$ is determined by the relative alignment between the eigenvectors of $\tau_{ij}^d$ and $\tilde{s}_{ij}$ as well as their eigenvalues. Specific preferential alignment has also been observed in turbulent flows \cite{Tao2002}. The eigenvectors of $-\tau_{ij}^d$ are denoted by $e_{\alpha}^d$, $e_{\beta}^d$, and $e_{\gamma}^d$, with corresponding eigenvalues $\lambda_{\alpha}^d \geq \lambda_{\beta}^d \geq \lambda_{\gamma}^d$ where $\lambda_{\alpha}^d + \lambda_{\beta}^d + \lambda_{\gamma}^d = 0$ by definition. The vorticity $\tilde{\omega}_i$ also displays preferential alignment with $-\tau_{ij}^d$ \cite{Horiuti2003}, specifically $\tilde{\omega}_i$ tends to be perpendicular to $e_{\alpha}^d$ while aligned with $e_{\beta}^d$.
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Figure 1. Definitions of the polar and azimuthal angles used to describe the orientation of $\tilde{\omega}_i$ in the eigen-frame of $\tilde{s}_{ij}$. Angles defined in a similar way for other vectors and eigen-frames are also used in the paper.

Figure 2. Instantaneous distributions of $\tilde{\omega} \equiv (\tilde{\omega}_i \tilde{\omega}_i)^{1/2}$ on a plane perpendicular to the $z$ axis. Left: from a target field; right: from the corresponding reconstructed field. $X$ and $Y$ are the labels of the grid points in the $x$ and $y$ directions, respectively.

2.4. Description of the non-local structures

The filtered velocity gradient $\tilde{A}_{ij}$ and related quantities describe the local structure of the filtered velocity field in an infinitesimal neighborhood of a spatial location. Non-local structures of finite sizes are also of great interests. The most well-known example of such structures is that of the vortex filaments that are characterized by strong enstrophy. The ability of the reconstructed velocity field to capture the locations, the dimensions and the orientations of these structures is not always clear from the point-wise statistics. To illustrate this point, Fig. 2 plots selected contours of $\tilde{\omega} \equiv (\tilde{\omega}_i \tilde{\omega}_i)^{1/2}$ in one target field and the corresponding reconstructed field. Detailed comparison is given later. However, it is already clear that the locations, orientation and sizes of the contours display remarkable similarity, especially in high $\tilde{\omega}$ regions. Usual statistics, including joint correlations between multiple points, do not always capture this similarity. Additional diagnostics are needed.

Note that, although an intuitive description of the structures visualized by the contours is straightforward, the precise description of their shapes, sizes and even locations is difficult, and actually there are no unique definitions for these concepts (also see discussions in, e.g., Bermejo-Moreno & Pullin (2008); Yang & Pullin (2011); Leung et al. (2012)). As explained in Section 1, we use the minimum volume enclosing ellipsoids (MVEEs) of a structure to define and describe its geometry. By definition, of all the
ellipsoids that contain a structure, the MVEE is the one with the smallest volume. The location, dimensions, and orientation of an MVEE are well-defined, and are given by its center, lengths of the three axes, and the orientations of the axes. We thus define the location, dimensions, and orientation of a structure using those of its MVEE. The structures in the reconstructed and the target fields are compared on this basis. The comparison includes four steps of calculations, which are explained below and illustrated in Fig. 3 using a 2D slice.

In the first step, the set $\mathcal{P}$ of the grid points where certain threshold condition is satisfied are extracted. $\mathcal{P}$ usually consists of a number of disjoint regions. The ‘structures’ we will be discussing refer to this type of regions; each structure corresponds to a disjoint region (c.f., the middle panel of Fig. 3).

In the second step, individual structures in $\mathcal{P}$ are extracted and distinguished from others. The programmatic realization of this task is non-trivial despite its innocuous appearance. It is accomplished by recasting it as a clustering problem, and using a density based clustering algorithm called DBSCAN [Ester et al. 1996; Schubert et al. 2017]. In this algorithm, the ‘neighbors’ of a point $p$ in $\mathcal{P}$ are first identified, where a neighbor is defined as a point whose distance to $p$ is less than a given value $\epsilon_c$. A point with fewer than $n_c$ neighbors is considered ‘noise’. A cluster is defined according to the following rule: the neighbors of a non-noise point $p$ are in the same cluster to which point $p$ belong. Algorithmically, the points in the cluster are identified by recursively applying this rule until all the points identified are noise. The same process is repeated for other non-noise points if they do not belong to any clusters that have been found.

For more details of the DBSCAN algorithm, the readers are referred to Ester et al. (1996). The outcome of applying the algorithm is that all the points in a structure are found and stored separately from those in other structures, as illustrated in the middle panel of Fig. 3. We normalize the distance in such a way that the distance between two consecutive grid points is 1. As a consequence, $\epsilon_c = D^{1/2}$ and $n_c = D + 1$ are used, where $D$ is the dimension of the embedding space (i.e. $D = 2$ for structures on a 2D slice whereas $D = 3$ for those in the 3D space).

In the third step, the MVEE for each structure is calculated. An ellipsoid is defined by a symmetric positive definite matrix $E$, which specifies the orientations and lengths of the axes, and a vector $c$, which specifies its center. Its volume is proportional to the determinant of $E^{1/2}$, $\det E^{1/2}$. Let $P$ be the set of points $p_i$ ($i = 1, ..., N$) in the
Figure 4. Calculation of sub-MVEEs and the MVEE tree. Left: one of the vortical structures shown in Fig. 3, which is enclosed by the level 0 sub-MVEE and is split into two groups along the dotted line. The red dots are the grid points in the structure. Middle: the level 1 sub-MVEE for the upper half. Right: the level 1 sub-MVEE for the lower half. $X$ and $Y$ are the labels of the grid points in the $x$ and $y$ directions, respectively.

The MVEE is given by the optimal $E$ and $c$ that minimize $\det E^{1/2}$, subject to the constraints

$$
(p^T_i - c^T)E^{-1}(p_i - c) \leq 1, \quad \forall p_i \in P. 
$$

The constraints ensure $p_i$ is inside the ellipsoid. This minimization problem can be solved by the Khachiyan algorithm (Khachiyan 1996; Todds 2016). We use the MATLAB implementation by Moshtagh (2009).

In the fourth (and last) step, the correspondence between a structure in the target field and its reconstruction in the reconstructed field is established. The above three steps are applied to both the reconstructed and the target fields. If the reconstructed field mimics the target field perfectly, one would then obtain two groups of MVEEs, with a one-to-one correspondence between the members. The correspondence can be trivially established. In reality, however, this naive method breaks down because the difference between the two fields sometimes is big enough to destroy the trivial one-to-one correspondence. The following procedure thus has been used, where, in essence, the structures are identified by their locations and sizes. The structures in the target field are first arranged into a list in the descending order according to their sizes, where the size of a structure is defined as the number of grid points in the structure. Let $S_T$ be a structure in the list. The structure in the reconstructed field whose distance to $S_T$ is the minimum is then identified, where the distance between two structures is defined as the shortest distance from one to the other. Let $S_O$ be this structure. If $S_O$ is not unique, the one with the largest size is chosen. This $S_O$ is taken as the reconstruction of $S_T$. $S_T$ and $S_O$ are called the matching structures. The above procedure is repeated for each structure in the target field, starting from the one largest in size, until all matching structures are identified.

The matching structures and corresponding MVEEs obtained after these four steps are then compared and statistics are calculated, including relative displacement, alignment, and relative sizes, among others. The analysis is applied to several different physical quantities, including high strain rate structures for which the magnitude of $\tilde{s}_{ij}$ is large, vortical structures with strong $\tilde{\omega}$, and structures with high SGS energy dissipation rate $\Pi_\Delta$ or negative $\Pi_\Delta$. In all cases, the analysis is performed directly on 3D structures.
MVEEs provide a good description of the enclosed structure when the latter is convex. A more complete description is desirable, however, if the structure is non-convex, such as the second structure from the top in the right panel of Fig. 3. In this case, we propose to use what we call the MVEE trees. The construction of MVEE trees is illustrated with Fig. 4. It is supposed that the MVEE for the structure has been found. In the context of MVEE trees, this MVEE is called the level 0 (L0) MVEE. The L0 MVEE allows one to split the grid points in the structure into two groups, using the symmetric plane of the MVEE perpendicular to its major axis (see the left panel of Fig. 4). The MVEEs for the two parts can then be found. These are called the L1 sub-MVEEs. The splitting can be repeated for each structure in this level, and those in each new level. The procedure thus forms a tree-like hierarchy of MVEEs, hence the name MVEE trees.

The MVEE trees are used to analyze strongly non-convex structures. For this purpose, the degree of non-convexity of a structure is defined as the ratio of the volume of the structure to the volume of its convex hull (Boyd & Vandenberghe 2004). The ratio is denoted by $R_V$. Smaller $R_V$ indicates stronger non-convexity. Only structures with small $R_V$ are selected when MVEE trees are calculated.

Moment of inertia ellipsoids (MoIE) have also been used in the past to characterize non-local structures. However, MoIEs do not capture the extremal points in a structure, hence tends to miss the details of highly non-convex structures. In comparison, MVEE and MVEE trees are better in this regard.

3. Simulations and results

3.1. Parameters and the computation of the target flow fields

The target fields $v(x,t)$ are obtained by numerically integrating the NSEs with a fully dealiased pseudo-spectral code in a $[0,2\pi]^3$ box with periodic boundary conditions. A second order explicit improved Euler method based on the trapezoidal rule is used in time-stepping. $v(x,t)$ is made statistically stationary by a constant forcing term $f$, where

$$f = (0, f_a \cos(k_f x_1), 0)$$

with $f_a = 0.15$ and $k_f = 1$ (these and the following parameters are all given in code units). Therefore, the flow is a 3D Kolmogorov flow with a sinusoidal mean velocity profile (Borue & Orszag 1996, Kang & Meneveau 2005). The Kolmogorov flow has been chosen because the forcing term for this flow is particularly simple.

In all cases $128^3$ grid points have been used. The viscosity is $\nu = 0.006$. Time step is $\delta t = 0.00575$. The time and length scales of $v$ are estimated from $f_a$, $k_f$, $\nu$, and the root-mean-square (RMS) velocity $v_{\text{rms}}$ which is found numerically to be about 0.65. The large eddy turnover time scale is thus $\tau_L \equiv v_{\text{rms}}/f_a \approx 4.35$. The energy dissipation rate $\epsilon$ is calculated numerically from the energy spectrum and $\nu$, which gives $\epsilon \approx 0.08$. The Kolmogorov time scale is $\tau_K \equiv (\nu/\epsilon)^{1/2} \approx 0.28$. The Kolmogorov length scale is $\eta_K \equiv (\nu^3/\epsilon)^{1/4} \approx 0.04$. Therefore, $\delta x/\eta_K \approx 1.25$ where $\delta x$ is the grid size of the simulations. The Reynolds number based on the Taylor micro-scale is $Re_{\lambda} \approx 75$, whereas the Reynolds number based on the forcing length scale is $Re_f \approx 340$.

3.2. The solution of the optimality system

The optimality system (Eq. (2.3) and Eqs. (2.5-2.8)) is solved iteratively. Given an initial guess for $\varphi$, the NSEs (Eq. (2.3)) are numerically integrated from $t = 0$ to $t = T$ to find $u(x,t)$. The cost $J$ is calculated from $u$. Unless $J$ is already smaller than a given tolerance, $\varphi$ has to be updated to reduce $J$. To do so, Eq. (2.5) is integrated backward in
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The time sequence of the forward solution \( u(x, t) \) is used in the backward integration. The solution for \( \xi \) provides \( DJ/D\varphi \) according to Eq. (2.8), which is then used to update \( \varphi \). The updating step uses the nonlinear conjugate gradient method using the Polak-Ribière formula (Nocedal & Wright 1999). The Brent’s method in the Numerical Recipe (Press et al. 1992) is used in the line search step.

The above several steps constitute the main loop of the solution algorithm. Due to the chaotic nature of the solution of the NSEs, our experience is that the iterations may become difficult to converge when \( T \) is large. We thus solve the optimality system first with a small \( T \), using the iterations outlined above. The optimal \( \varphi \) for this problem is used as the initial guess for the problem with a larger \( T \). The procedure is repeated until the intended optimization horizon is reached. As the computational time is usually very long on the computer we used (see below for more information), we save the optimal solutions for the intermediate \( T \) values on the hard disk, which safeguards against unexpected interruptions of the computation. Therefore, this implementation is also useful from a practical point of view. The choices for the initial small \( T \) and the increment in \( T \) are mostly based on trial and error. An initial \( T = 0.25\tau_L \approx 4\tau_K \) with an increment equal 0.025\( \tau_L \approx 0.4\tau_K \) has been used in most cases, but we expect other similar choices will also work.

We now explain briefly the solution of the individual equations in the main loop. \( u(x, t) \) is solved from the NSEs using the same numerical methods and parameters as those for \( v(x, t) \). The initial guess for \( u(x, 0) \equiv \varphi \) is usually a divergence free Gaussian random velocity field, with the Fourier modes with \( k \leq k_m \) replaced by those in the target data \( \varphi^v(x) \equiv v(x, 0) \) (c.f., Section 2.1). Our tests show that the initial guess has no effects on the statistics of optimal solutions. It is possible to use an initial guess with the energy spectrum matching the initial target field. However, our tests showed that it did not improve the results nor speed up convergence. A possible reason is that matching the energy spectrum imposes only a weak constraint on the control \( \varphi \) since the latter has a very large number of components.

As will be explained below, we run several related cases with different parameters. Sometimes the optimal solution \( \varphi^u \) for a solved case is then used as the initial guess for \( \varphi \) in a related case. For example, the optimal solution obtained with a larger tolerance has been used as the initial guess for a case with a smaller tolerance; the one with a smaller \( T \) used as the initial guess for a case with a larger \( T \).

The adjoint equations (Eq. (2.5)) are solved with the same numerical schemes used for \( v \) and \( u \). The divergence free condition for \( \xi \) is used to eliminate the adjoint pressure \( \sigma \), in the same way as \( p \) is eliminated from the NSEs when they are solved numerically. However, Eq. (2.5) has to be integrated backwards in time, as it is supplemented with the end-point condition in Eq. (2.6). The time sequence of \( u(x, t) \ (t \in [0, T]) \) is needed for the solution of \( \xi \). Therefore, \( u(x, t) \) has to be saved in the forward integration of the NSEs. In our computation, typically the available RAM is not large enough to hold the whole time sequence. We thus use checkpointing, where \( u(x, t) \) is saved on the hard disk at selected t’s (the checkpoints) in the forward integration. The sequence of \( u(x, t) \) between two checkpoints, which will be short enough to store in the RAM, are computed (using the data on a checkpoint as the initial condition) when the backward integration of \( \xi \) reaches the time interval between the two checkpoints. Checkpointing thus incurs several passes of forward integrations in one single backward integration. But it avoids having to read data at every time step from the hard disk, which typically is a time consuming operation.

The technique can be optimized by choosing the checkpoints judiciously although we do
Table 1. The parameters for the cases being investigated. $N_R$ is the number of realizations. Case C1 is the baseline case with the parameters highlighted in bold.

| Case | $T/\tau_L$ | $k_m$ | $10^4 \epsilon_{\text{tot}}$ | $N_R$ |
|------|------------|-------|-----------------------------|-------|
| C1   | 1          | 4     | 2 5                         | 1 2   |
| C2   | 1          | 4     | 1.5 2                       | 2     |
| C3   | 0.5        | 2     | 2 2                         | 2     |
| C4   | 0.5        | 2     | 1 2                         | 2     |
| C5   | 0.5        | 2     | 0.5 2                       | 2     |
| C6   | 0.5        | 4     | 2 5                         | 5     |
| C7   | 0.5        | 4     | 1 2                         | 2     |
| C8   | 0.5        | 4     | 0.5 2                       | 2     |
| C9   | 0.5        | 8     | 2 2                         | 2     |
| C10  | 0.5        | 8     | 1 2                         | 2     |
| C11  | 0.5        | 8     | 0.5 2                       | 2     |

not attempt to do so in this investigation. More details about checkpointing can be found in, e.g., [Wang et al.] (2009).

3.3. The choices of computational parameters

We assume that the data $\mathcal{F}v$ is available continuously, so that the forcing term in Eq. (2.7) is applied at each time step. We assume the optimal solution is found when $J/J(u = 0)$ is less than a tolerance $\epsilon_{\text{tot}}$, where, according to Eq. (2.2), $J(u = 0)$ is the total kinetic energy of the measurement data $\mathcal{F}v(x, t)$ ($0 \leq t \leq T$). Due to our limited computational resources, this investigation has used only $128^3$ grid points. However, several values of $T$, $k_m$ and $\epsilon_{\text{tot}}$ are examined, and up to five realizations are run for each set of parameters. The target data $v(x, t)$ in different realizations are different segments of the same time series of DNS data.

The parameters are summarized in Table 1. We consider $T$ up to one large eddy turnover time scale $\tau_L$ and resolutions $k_m = 2, 4$ and 8. However, for $T = \tau_L$, only cases with $k_m = 4$ are reported for the following reasons. Firstly, for $(T, k_m) = (\tau_L, 2)$, we failed to find convergent results (see the discussion of the results below for further comments). Secondly, for $(T, k_m) = (\tau_L, 8)$, it turns out that the optimal solutions are the same as those with $(T, k_m) = (0.5\tau_L, 8)$ with $\epsilon_{\text{tot}} = 0.5 \times 10^{-4}$ (i.e., the cases with a shorter $T$ but more stringent tolerances). Therefore, the results for these two $k_m$’s have been omitted.

For $(T, k_m) = (\tau_L, 4)$, optimal solutions are found for $\epsilon_{\text{tot}} = 2 \times 10^{-4}$ in five realizations (case C1). In each of these realizations, it took around 15 days to find the optimal solution, using up to 48G RAM and 4 cores on an Intel Ivybridge or Westmere based CPU, with the optimal solutions from Case 6 as the initial guesses. In two of the realizations, the optimal solutions are further improved by additional iterations to reduce $J/J(u = 0)$ below $1.5 \times 10^{-4}$ (case C2). The improvement required about 15 more days of computer time. Given the significant computational cost, we did not attempt to reduce the cost function further. On the other hand, for $T = 0.5\tau_L$, optimal solutions were obtained for $\epsilon_{\text{tot}} = 10^{-4}$ and $0.5 \times 10^{-4}$ in two realizations for all three $k_m$ values. The cost decreases rapidly initially, but further improvement becomes very slow. Fig. 5 plot the change of $J/J(u = 0)$ in the iterations for a run in case C6, which depicts this behavior. The inset zooms into the curve to show the change at later iterations. Abrupt jumps in $J$ are observed when $T$ is extended. In this run, the optimal solution for $T = 0.5\tau_L$ is found after 124 iterations. In each iteration, around 5 forward integrations are performed in order to find the conjugate gradient direction. Taking backward integration into account, $700 \sim 800$ passes of integration are used, although we note that part of these integrations are over shorter optimization horizons.
We consider Case C1 (shown in bold in Table 1) the baseline case. The comparison between this and other cases will illustrate the effects of the resolution of the measurement data $k_m$, the optimization horizon $T$ and the tolerance $\epsilon_{\text{tot}}$. However, the majority of results is presented only for the baseline case. The statistics are averaged over all the realizations in each case.

3.4. The energy spectra and spectral correlation for different computational parameters

The spectral distributions of $u$ and $v$ obtained with different computational parameters are compared in this subsection. The purpose is to give an overview of the quality of the reconstruction. More detailed investigation, focusing on the baseline case, will be presented later. Symbols $u$ and $v$ in the subscripts or superscripts are used to denote results from $u$ (the reconstruction) and $v$ (the target), respectively.

Fig. 6 shows the energy spectra for $\phi_u$ and $\phi_v$, which are plotted with solid and empty symbols, respectively. The figure shows that perfect match between $\phi_u$ and $\phi_v$ is obtained for $k \leq k_m$, as expected. Fig. 6(a) and (b) show that, for $k_m = 4$, $\phi_u$ under-predicts the spectrum for intermediate $k$'s and over-predicts it for large $k$'s. Nevertheless, the spectra of $\phi_u$ broadly follow those of $\phi_v$. The agreement does not change much with $T$ and $\epsilon_{\text{tot}}$ for the ranges of the parameters we are considering. Note that it is non-trivial to observe the fairly good agreement. The dashed line in Fig. 6(a) plots the energy spectrum for the initial guess for $\phi_u(x)$, which shows that the small scales in $\phi_u$ are not present in the initial guess and are generated by the data assimilation process.

The resolution of the measurement data $k_m$ has significant effects on $E_u(k)$, as is demonstrated in Fig. 6(c). $E_u(k)$ for $k_m = 2$ essentially fails to capture the features of $E_v(k)$. On the other hand, the spectrum for $k_m = 8$ shows similar level of discrepancy as for $k_m = 4$ at the high wavenumber end, which is somewhat surprising. Reducing the tolerance to $\epsilon_{\text{tot}} = 0.5 \times 10^{-4}$ does not improve the agreement, as shown in Fig. 6(d).

Since both $u(x,t)$ and $v(x,t)$ are solutions of the forced NSEs, their statistics are expected to be the same for $t \rightarrow T$. It is interesting, however, to investigate the instantaneous point-wise difference between $u(x,t)$ and $v(x,t)$ at a later time $t$ for $t \in [0,T]$. Two quantities are used to quantify the difference. The first one is the energy spectrum for the velocity difference $\delta u(x,t) = u(x,t) - v(x,t)$, denoted by $E_D(k)$, and called spectral difference. Obviously, a smaller $E_D(k)$ indicates better match between $u(x,t)$ and $v(x,t)$, hence a better reconstruction. The second one is the spectral
Figure 6. The energy spectra for $\varphi^u(x)$ (solid symbols) and $\varphi^v(x)$ (empty symbols of same shapes). (a) with different optimization horizon $T$ (the dashed line shows the energy spectrum of the initial guess for $\varphi^u(x)$ in one of the realizations); (b) with different tolerance $e_{\text{tot}}$; (c) with different measurement resolution $k_m$; (d) with different tolerance for $k_m = 8$.

correlation (i.e., normalized co-spectrum) between $u$ and $v$, defined by

$$\hat{\rho}_{uv}(k) = \frac{\int \langle \hat{u}^*(k,t) \cdot \hat{v}(k,t) \rangle dS_k}{2 \sqrt{E_u(k)E_v(k)}}.$$  \hspace{1cm} (3.2)

The pointed brackets in the numerator denotes ensemble average whereas the integral is a surface integral over the sphere $|k| = k$. By definition, $0 \leq \hat{\rho}_{uv} \leq 1$, and a larger $\hat{\rho}_{uv}$ indicates better point-wise agreement between $u$ and $v$. Note that $E_D(k)$ and $\hat{\rho}_{uv}(k)$ both depend on time $t$, although the dependence is not given explicitly in the expressions.

Fig. 7 compares $E_D(k)$ and $\hat{\rho}_{uv}(k)$ from different cases in various ways. Fig. 7(a) and (b) show that, in Case C1 and C2, the point-wise agreement between $u$ and $v$ improves over time within the optimization horizon. $\hat{\rho}_{uv}$ reaches 0.8 or higher at $t = \tau_L$ for all wavenumbers, whereas $E_D(k)/E_v(k)$ can be reduced down to 30% even for $k = 40$. Note that the measurement data contain approximately $8^3/2 = 256$ Fourier modes, constituting less than 0.1% of the Fourier modes in the whole flow field.

Fig. 7(a) and (b) also show that imposing a smaller tolerance $e_{\text{tot}}$ can lead to significantly better agreement at later times, even if it is only marginally smaller (i.e., $1.5 \times 10^{-4}$ versus $2 \times 10^{-4}$). On the other hand, the effect is quite limited at earlier times (e.g., at $t = 0$), where significant discrepancy between $u$ and $v$ is observed for both $e_{\text{tot}}$. This
Figure 7. (a) and (b): $E_D(k)/E_v(k)$ and $\hat{\rho}_{uv}(k)$ at different time $t$ for case C1 (solid symbols) and case C2 (empty symbols), with $(T, k_m) = (\tau_L, 4)$ and $\varepsilon_{tot} = 2 \times 10^{-4}$ or $1.5 \times 10^{-4}$. (c): $\hat{\rho}_{uv}(k)$ for case C1 where $T = \tau_L$ (solid symbols) and case C6 where $T = 0.5\tau_L$ (empty symbols). (d): $\hat{\rho}_{uv}(k)$ for different $\varepsilon_{tot}$ (cases C6, C7, C8; empty symbols) at $t = 0.5\tau_L$. Solid symbols: the result for case C1 as a comparison. (e): $\hat{\rho}_{uv}(k)$ at $t = 0.5\tau_L$ for different $k_m$ (cases C3, C6, C9). (f): $E_D(k)/E_v(k)$ at $t = \tau_L$ obtained from 4DVAR (case C1; triangles), from DS (squares), and from 5th-order Lagrange interpolation (circles).

observation is interesting, suggesting that the improvements at later times are due to effects that are not completely revealed by $E_D(k)$ and $\hat{\rho}_{uv}$ at earlier times.

The effects of optimization horizon $T$ are investigated in Fig. 7(c), where $\hat{\rho}_{uv}$ for $T = 0.5\tau_L$ and $T = \tau_L$ is shown with empty and filled symbols, respectively. The results
for $T = 0.5\tau_L$ are shown for $t = 0.25\tau_L$ (squares) and $0.5\tau_L$ (triangles). Those for $T = \tau_L$ are shown for the same times and $t = \tau_L$ too. It is observed that larger $T$ improves the point-wise agreement, and the improvement increases over time. The agreement, again, can be improved by imposing a smaller tolerance. Fig. 7(d) compares $\hat{\rho}_{uv}$ at $t = T$ for three $e_{tot}$’s. For $e_{tot} = 0.5 \times 10^{-4}$, the correlation can be improved to about the same level obtained with a longer horizon $T = \tau_L$. Nevertheless, in order to obtain satisfactory agreement over the whole range of wavenumbers, even smaller $e_{tot}$ or $T > 0.5\tau_L$ is needed.

The parameter $k_m$ affects the results strongly. Fig. 7(e) plots $\hat{\rho}_{uv}$ at $t = T$ for $T = 0.5\tau_L$ with $k_m = 2, 4, 8$. When $k_m = 2$, $u$ shows little correlation with $v$. For $k_m = 8$, almost perfect correlation is obtained. To understand these results, it is instructive to compare the 4DVAR method with the DS method (Yoshida et al. 2005; Lalescu et al. 2013) (c.f. Section 1), which is shown in Fig. 7(f). In Yoshida et al. (2005), the authors integrate the NSEs with a Gaussian random field as the IC. The low wavenumber modes with $k \leq k_m$ in the solution $u(x, t)$ are replaced by those in the target field $v(x, t)$ at each time step. They show that, if $k_m \geq k_c = 0.2\eta_k^{-1}$, the small scales in $u(x, t)$ approach those in $v(x, t)$ asymptotically as $t \to \infty$, assuming an infinite sequence of data are available. In our investigation, the threshold wavenumber is $k_c = 5$ so $k_m = 4$ is slightly below $k_c$, and the measurement data are available only for $t \in [0, T]$. As a consequence, the DS scheme will only achieve partial reconstruction. The reconstruction at $t = \tau_L$ using the DS scheme with $k_m = 4$ is compared with the 4DVAR method in Fig. 7(f). It shows that the 4DVAR method improves the reconstruction significantly. A more simplistic reconstruction, obtained with a 5th-order Lagrange interpolation, is also examined in Fig. 7(f). The reconstruction has essentially no correlation with the target field for $k > k_m$.

Fig. 7(e) and (f) together thus show that, for $k_m$ around or greater than $k_c$, 4DVAR can exploit a short time sequence of data to obtain better (for $k_m = 4$) or almost perfect (for $k_m = 8$) reconstruction. For $k_m$ much smaller than $k_c$ (e.g., for $k_m = 2$), 4DVAR will fail. These results suggest that the threshold $k_c$ found with the DS method is also valid for 4DVAR. Nevertheless, a more comprehensive investigation is needed to confirm this preliminary observation. Simulations with other $k_m$’s around $k_c$, as well as other Reynolds numbers, are needed. We leave this investigation for the future.

3.5. Statistics of the reconstructed fields at $t = 0$

In this subsection, we examine other dynamically important statistics of the reconstructed initial fields (i.e. $\varphi^u(x)$). We consider the cases with $k_m = 4$ only and focus mostly on the baseline case C1 where $T = \tau_L$, $k_m = 4$ and $e_{tot} = 2 \times 10^{-4}$. We conduct the analysis in the physical space, using the filtering approach where necessary (c.f. Section 2.3). The Gaussian filter is used, and the filter scale $\Delta$ is usually taken as $\Delta_m/2$ or $\Delta_m/4$ where $\Delta_m \equiv \pi/k_m$ is the equivalent filter scale associated with $k_m$.

The joint PDFs of normalized $Q$ and $R$ are plotted in Fig. 8. The contours from the target fields depict the well-known skewed tear-drop shape with the Vieillefosse tail extended in the fourth quadrant (Vieillefosse 1984; Cantwell 1992). Events in the second and the fourth quadrants occur with higher probabilities. These features are qualitatively reproduced by the contours of the reconstructed fields (dashed lines). However, the Vieillefosse tail is shorter, and the probabilities for large excursions are lower by a rather significant amount. The lowest contours for the reconstructions with larger $T$ and smaller $e_{tot}$ (the top and middle rows) are closer to those for the target, although the improvement is very small. The right column shows that the discrepancy is bigger when the filter scale is smaller.

The mean strain rate, enstrophy, and vortex stretching term, normalized by the Kolmogorov time scale $\tau_K$, are given in Fig. 9 as functions of the filter scale. Not
Figure 8. The joint PDF of $Q^* \equiv Q/(\bar{A}_{ij}\bar{A}_{ij})$ and $R^* \equiv R/(\bar{A}_{ij}\bar{A}_{ij})^{3/2}$. Left column: $\Delta = \Delta_m/2$; right column: $\Delta = \Delta_m/4$. Top row: case C1; middle row: case C2; bottom row: case C6. Grey scales: from $\varphi^v(x)$; dashed lines: from $\varphi^u(x)$. 
Figure 9. \langle \tilde{s}_{ij} \tilde{\omega}_i \tilde{\omega}_j \rangle \) (circles), \langle \tilde{s}_{ji} \tilde{s}_{ij} \rangle \) (squares), and \langle \tilde{\omega}_i \tilde{\omega}_i \rangle \) (diamonds) as functions of filter scale \(\Delta\) from case C1. Filled and empty symbols: from \(\varphi^v(x)\) and \(\varphi^u(x)\), respectively. The vertical line marks the scale where \(\Delta = \Delta_m\).

Figure 10. PDFs of \(| \cos \theta^s_i|\) with \(\theta^s_i\) being the angle between \(\tilde{\omega}_i\) and \(e^s_\alpha\) (diamonds), \(e^s_\beta\) (squares), and \(e^s_\gamma\) (circles). Filled symbols: \(\varphi^v(x)\); empty symbols: \(\varphi^u(x)\). From case C1 and \(\Delta = \Delta_m/2\).

Surprisingly, the differences between the results in the two fields are larger for smaller filter scales. For \(\langle \tilde{s}_{ij} \tilde{s}_{ij} \rangle\) and \(\langle \tilde{\omega}_i \tilde{\omega}_i \rangle\) the discrepancy is below 20% for all filter scales. The discrepancy for the stretching term however is much bigger. The latter can be partially explained by the mis-alignment between \(\tilde{\omega}_i\) and \(\tilde{s}_{ij}\) in \(\varphi^u\), which is shown in Fig. 10 with the PDFs for the cosines of the angles between \(\tilde{\omega}_i\) and the eigenvectors of \(\tilde{s}_{ij}\). The results for \(\varphi^u\) display qualitatively correct features, i.e., a preferred alignment between \(\tilde{\omega}_i\) and \(e^s_\beta\), and the tendency for \(\tilde{\omega}_i\) to be perpendicular to \(e^s_\gamma\). However, the peaks of the PDFs are lower to a significant degree. Therefore, the preferable alignment in \(\varphi^u\) is weaker, and this mis-alignment contributes to the observed discrepancy in the vortex stretching term.

The mean SGS energy dissipation \(\langle \Pi_\Delta \rangle\) is shown in the left panel of Fig. 11. The results from \(\varphi^u\) display the same trends as those from \(\varphi^v\). This result is a direct evidence that the
nonlinear interaction between different scales are partially reconstructed. At $\Delta = \Delta_m/2$ ($\Delta/\eta_K \approx 10$), the value calculated from $\varphi^u(x)$ is smaller by about 25%. The results for the two different $T$'s show only very slight difference. The PDFs for the normalized $\Pi_\Delta$ at $\Delta = \Delta_m/2$ (filled symbols) and $\Delta = \Delta_m/4$ (empty symbols), from $\varphi^v$ (circles) and $\varphi^u$ (diamonds) for case C1.

The conditionally averaged SGS energy dissipation $\langle \Pi_\Delta \rangle / \epsilon$ has also been calculated, where $Q^* \equiv Q / \langle A_{ij} A_{ij} \rangle$ and $R^* \equiv R / \langle A_{ij} A_{ij} \rangle^{3/2}$. Shown in Fig. 12 is $\langle \Pi_\Delta | Q^*, R^* \rangle$ weighted by the joint PDF $P(R^*, Q^*)$ and normalized by the mean energy dissipation rate $\epsilon$. The results from $\varphi^v$ (the left column) show that higher SGS dissipation is observed around the Vieillefosse tail where high straining motion happens. The behavior is reproduced qualitatively by $\varphi^u$, shown in the middle and on the right for $T = \tau_L$ and $0.5\tau_L$, respectively. Negative SGS dissipation, representing energy back-scattering, is found mainly in the first quadrant, a feature also captured by $\varphi^u$. However, overall, $\varphi^u$ tends to underestimate the magnitude of the weighted SGS energy dissipation. The results at $\Delta = \Delta_m/2$ are closer than those at $\Delta = \Delta_m/4$. There are some small improvements with $T = \tau_L$ compared with $T = 0.5\tau_L$.

To summarize, the results in this subsection demonstrate that $\varphi^u(x)$ captures the main statistical features of real turbulence with reasonable agreement. This is the basis for the reconstruction of the instantaneous velocity fields at later times. The discrepancy between $\varphi^v(x)$ and $\varphi^u(x)$, manifested by the several statistics we considered, can be reduced slightly by extending the optimization horizon from $T = 0.5\tau_L$ to $T = \tau_L$ or by using a smaller $e_{\text{tot}}$. Given the chaotic nature of the flow, these small differences might be the explanation for the significant improvement found at later time $t \rightarrow T$ shown in Section 3.4. This conjecture may be quantified by calculating the sensitivities of the statistics although this is beyond the scope of this paper.

3.6. Reconstruction of the instantaneous local structures for $t \rightarrow T$

For the reconstructed field $u(x, t)$ at later times, it is expected that its statistics would agree well with those in the target field $v(x, t)$, for the simple reason that both are the
stationary solutions of the NSEs. However, there is no a priori reason to assert that the $u(x,t)$ would have point-wise agreement with $v(x,t)$. The point-wise differences have been examined briefly using $E_D(k)$ and $\hat{\rho}_{uv}(k)$ in Section 3.4 to demonstrate the effects of the choices of computational parameters. In this subsection, the base line case C1 (c.f. Table 1) is examined in further detail. Note that time $t$ is now another parameter. We usually choose $t = T$. However, when comparing results with different $T$’s, other values of $t$ are also used.

3.6.1. Geometry of the local structures

The instantaneous difference in the geometrical structures of the reconstructed fields $u$ and the target fields $v$ is first investigated. The first quantity we examine is the ‘cross-alignment’ between the vorticity in $v$ ($\tilde{\omega}_v$) and the eigenvectors of the strain rate tensor in $u$ ($\tilde{s}_{ij}^u$). The alignment is measured by the joint PDF of $\cos \theta_{sp}$ and $\phi_{sp}$, where the angles define the orientation of $\tilde{\omega}_v$ in the eigen-frame of $\tilde{s}_{ij}^u$ (c.f., Fig. 1). The joint PDF is shown with gray scale contours in Fig. 13. The joint PDF for the usual alignment between $\tilde{\omega}_v$ and $\tilde{s}_{ij}^u$, both from $v$, is plotted in dashed lines as a comparison. The left column of Fig. 13 shows that the joint PDF for $\tilde{\omega}_v$ and $\tilde{s}_{ij}^u$ is very close to the one for

Figure 12. The normalized weighted conditional average of the SGS energy dissipation $\epsilon^{-1}P(R^*,Q^*)/\langle H \Delta \vert R^*,Q^* \rangle$. Left: from $\varphi^v$; middle: from $\varphi^u$ in case C1; right: from $\varphi^u$ in case C6. Top row: $\Delta = \Delta_m/2$; bottom row: $\Delta = \Delta_m/4$. 

stationary solutions of the NSEs. However, there is no a priori reason to assert that the $u(x,t)$ would have point-wise agreement with $v(x,t)$. The point-wise differences have been examined briefly using $E_D(k)$ and $\hat{\rho}_{uv}(k)$ in Section 3.4 to demonstrate the effects of the choices of computational parameters. In this subsection, the base line case C1 (c.f. Table 1) is examined in further detail. Note that time $t$ is now another parameter. We usually choose $t = T$. However, when comparing results with different $T$’s, other values of $t$ are also used.

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Figure 13. The joint PDFs of \((\cos \theta_{sp}, \phi_{sp})\) for angles between \(\tilde{\omega}^\gamma_i\) and \(\tilde{s}^u_{ij}\) (gray scales) and those between \(\tilde{\omega}^\gamma_i\) and \(\tilde{s}^u_{ij}\) (dashed lines). Top row: \(\Delta = \Delta_m/2\); bottom row: \(\Delta = \Delta_m/4\). Left: \(t = T\) for case C1; middle: \(t = 0.5T\) for case C1; right: \(t = T\) for case C6.

\(\tilde{\omega}^\gamma_i\) and \(\tilde{s}^u_{ij}\) for both \(\Delta = \Delta_m/2\) and \(\Delta_m/4\). Both joint PDFs display a high peak around \((0, 0)\), which shows the well-known preferred alignment between the vorticity vector with \(e^{\gamma}_\alpha\) in turbulence. The peak probability density for the cross alignment is only slightly lower. However, the fact that replacing \(\tilde{s}^u_{ij}\) by \(\tilde{s}^u_{ij}\) still yields a close joint PDF proves that the instantaneous orientations of the eigenvectors of \(\tilde{s}^u_i\) and \(\tilde{s}^u_{ij}\) are very close to each other. Same behaviours are observed for the results in the middle column, where the results at an earlier time \(t = T/2\) for the same \(T = \tau_L\) are shown. The joint PDFs display only slightly larger discrepancy. The result obtained with a shorter horizon \(T = 0.5\tau_L\), given in the right column, shows much larger discrepancy, as the highest level contour is much smaller even though the location of the peak is correctly found at \((0, 0)\).

Fig. 14 considers the cross alignment between \(\tilde{\omega}^\gamma_i\) and the eigenvectors of \(-\tau_{ij}^{d,u}\). The dashed lines show the joint PDF for the usual alignment in \(v\). The two peaks of the joint PDF depict the known feature in turbulence, where \(\tilde{\omega}_i\) tends to align with \(e^\tau_\alpha\) and \(e^\tau_\beta\). The cross alignment between \(\tilde{\omega}^\gamma_i\) and \(-\tau_{ij}^{d,u}\) qualitatively reproduces these features. The result at \(\Delta = \Delta_m/2\) show excellent quantitative agreement, although the discrepancy
Figure 14. The joint PDFs of \((\cos \theta_{sp}^\omega, \phi_{sp}^\omega)\) for angles between \(\tilde{\omega}_v^{\gamma}\) and \(-\tau_{ij}^{d,v}\) (gray scales) and those between \(\tilde{\omega}_i^{\gamma}\) and \(-\tau_{ij}^{d,v}\) (dashed lines), for \(\Delta = \Delta_m/2\) (left) and \(\Delta_m/4\) (right) with \(t = T\) in case C1.

Figure 15. The joint PDF of \((\cos \theta_{sp}^{s,i}, \phi_{sp}^{s,i})\) for angles made by \(e_{s,u}^i\) in the eigen-frame of \(\tilde{s}_{vij}\). Left: \(\Delta = \Delta_m/2\) and \(t = T\) from case C1; middle: \(\Delta = \Delta_m/4\) and \(t = T\) from case C1; right: \(\Delta = \Delta_m/2\) and \(t = T\) for case C6. Solid lines: \(i = \alpha\); dashed lines: \(i = \beta\); dotted lines: \(i = \gamma\). The contour levels are 0.3, 3, and 30.

becomes larger for \(\Delta = \Delta_m/4\). The comparison at \(t = 0.5\tau_L\) for \(T = 0.5\tau_L\) and \(\tau_L\) is not shown, because it simply confirms our expectation that the results with a smaller optimization horizon \(T\) display larger discrepancy.

The tensorial structures of \(\tilde{s}_{ij}^u\) and \(\tilde{s}_{ij}^v\) are compared using the orientation of the eigenvectors of the former in the eigen-frame of the latter. The alignment is given in the left panel of Fig. 15 for \(\Delta = \Delta_m/2\) and \(T = \tau_L\), in terms of the joint PDFs of \((\cos \theta_{sp}^{s,i}, \phi_{sp}^{s,i})\), where \(\theta_{sp}^{s,i}\) and \(\phi_{sp}^{s,i}\) are the angles made by \(e_{s,u}^i\) in the eigen-frame of \(\tilde{s}_{ij}^v\) (c.f., Fig. 2). According to the definitions of the angles, perfect alignment between \(e_{s,v}^i\) and \(e_{s,u}^i\) corresponds to \(\cos \theta_{sp}^{s,\alpha} = 1\) and an undetermined \(\phi_{sp}^{s,\alpha}\). The solid lines in Fig. 15
show a sharp ridge of peak values at $\cos \theta^{\alpha,\beta} = 1$, which indicates that $\mathbf{e}^{\alpha,\beta}_{\text{sp}}$ and $\mathbf{e}^{\alpha,\beta}_{\text{sp}}$ are strongly aligned. Perfect alignment between $\mathbf{e}^{\beta,\gamma}_{\text{sp}}$ and $\mathbf{e}^{\beta,\gamma}_{\text{sp}}$ corresponds to a peak at $(0,0)$, whereas the perfect alignment between $\mathbf{e}^{\gamma,\alpha}_{\text{sp}}$ and $\mathbf{e}^{\gamma,\alpha}_{\text{sp}}$ corresponds to a peak at $(0,\pi/2)$. The joint PDFs in these two cases (shown with dashed and dotted lines, respectively) indeed display high peak probabilities around these points. Therefore, these joint PDFs demonstrate excellent agreement between the orientations of $\mathbf{s}^u_{ij}$ and $\mathbf{s}^v_{ij}$. The results at $\Delta = \Delta_m/4$ given in the middle shows that strong alignment persists to smaller filter scales. The joint PDFs obtained with $T = 0.5\tau_L$, shown on the right, have much weaker peaks, indicating a larger discrepancy in the orientation of the two tensors.

Fig. 16 plots the same statistics as those in Fig. 15 but for the SGS stress tensors $-\tau^d_{ij}$ and $-\tau^e_{ij}$. The observations made from Fig. 15 are also observed in Fig. 16 except that the alignment shown in the latter is in general somewhat weaker.

### 3.6.2. Correlations

We will look into the covariance and the correlation coefficients in this subsection. For two random variables $X$ and $Y$, they are denoted by $C(X,Y)$ and $\rho(X,Y)$, respectively. By standard definitions,

$$ C(X,Y) \equiv \langle (X - \langle X \rangle)(Y - \langle Y \rangle) \rangle, \quad \rho(X,Y) \equiv \frac{C(X,Y)}{\sigma_X \sigma_Y}, $$

where $\sigma_X \equiv \sqrt{C(X,X)}$ is the RMS value of $X$ and $\sigma_Y$ is defined similarly. The physical quantities in $\mathbf{u}(\mathbf{x},t)$ and $\mathbf{v}(\mathbf{x},t)$ will be denoted with superscripts $\mathbf{u}$ and $\mathbf{v}$, respectively. The difference in $X$ is quantified by the RMS value of $X^u - X^v$, $\sigma_{X^u-X^v}$, which will be normalized by the RMS value of $X$ in the target fields $\sigma_{X^v}$.

The left panel in Fig. 17 shows $\rho(X^u, X^v)$ where $X \equiv \mathbf{s}_{ij}\mathbf{s}_{ij}$. The correlation deteriorates with decreasing filter scales, but improves with $t/T$ for $t$ within the optimization horizon. Excellent correlation across all scales is achieved at $t/T = 1$ for $T = \tau_L$. The difference in $X = \mathbf{s}_{ij}\mathbf{s}_{ij}$ is also examined in the right panel in terms of the RMS value of $X^u - X^v$ normalized by the RMS value of $X^v$. For $t = T = \tau_L$, the value can be reduced to about 27% at the Kolmogorov scale where $\Delta/\eta_K \approx 1$. For results from the DS method...
Figure 17. The correlation coefficient $\rho(X^u, X^v)$ (left) and the normalized RMS value of $X^u - X^v$ (right) as a function of $\Delta/\eta_K$, where $X = \tilde{s}_{ij}\tilde{s}_{ij}$. Solid symbols: from case C1; open symbols: from case C6. Circles: $t = 0$; upright triangles: $t = 0.5\tau_L$; left-pointing triangles: $t = \tau_L$ (for case C1 only). Right-pointing solid triangles: from the DS method at $t = \tau_L$ for $k_m = 4$. The vertical line marks the scale $\Delta = \Delta_m$.

Figure 18. Left: The normalized RMS values of the differences $X^u - X^v$ from case C1, where $X \equiv \tilde{s}_{ij}\tilde{s}_{ij}$ (circles), $\tilde{\omega}_i\tilde{\omega}_i$ (squares), $P_\omega(\equiv \tilde{s}_{ij}\tilde{\omega}_i\tilde{\omega}_j)$ (diamonds), and $\Pi_\Delta$ (triangles), respectively. Right: the correlation coefficients $\rho(X^u, X^v)$ from case C1. The vertical line marks the scale $\Delta = \Delta_m$.

(right-pointing solid triangles), the correlation $\rho$ at the smallest $\Delta/\eta_K$ is about 70% and the RMS difference is about 80%, indicating significantly higher discrepancy. This contrast is consistently observed in other statistics. Therefore, we will not present more results from the DS method.

$\sigma_{X^u-X^v}/\sigma_{X^v}$ and $\rho(X^u, X^v)$ for $X = \tilde{\omega}_i\tilde{\omega}_i$, $P_\omega$ and $\Pi_\Delta$ are plotted in Fig. 18, where the results for $\tilde{s}_{ij}\tilde{s}_{ij}$ are also given as comparison. For all quantities, almost perfect correlation is observed. The RMS difference for $\tilde{\omega}_i\tilde{\omega}_i$ is similar to $\tilde{s}_{ij}\tilde{s}_{ij}$, both reaching about 27% at the Kolmogorov scale where $\Delta/\eta_K \approx 1$. The discrepancy for $\Pi_\Delta$ and $P_\omega$ is somewhat larger with the normalized RMS difference found at about 35% at the smallest filter scale.

3.6.3. Joint PDFs and conditional statistics

To examine the large fluctuations in the differences between the target and reconstructed parameters, we look into the joint PDFs and conditional statistics in this subsection. Fig. 19 plots the cumulative distribution functions for the normalized differences.
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Figure 19. The cumulative distribution functions for the normalized differences $|X^u - X^v|$ from case C1, where $X = \tilde{s}_{ij} \tilde{s}_{ij}$ (circles), $\tilde{\omega}_i \tilde{\omega}_i$ (squares), $P_\omega \equiv \tilde{s}_{ij} \tilde{\omega}_i \tilde{\omega}_j$ (diamonds), and $\Pi_{\Delta}$ (triangles). Filled symbols: $\Delta = \Delta_m/2$; empty symbols: $\Delta = \Delta_m/4$. The vertical line marks the 20% threshold.

Figure 20. Left: the averaged $\Pi_{\Delta}$ in $u(x, t)$ conditioned on $\Pi_{\Delta}$ in $v(x, t)$ (circles) and the joint PDF of the two (contours). The error bars are the conditional RMS values of $\Pi_{\Delta}^u/\langle \Pi_{\Delta}^v \rangle$. Right: the same for $P_\omega$. For case C1 with $\Delta = \Delta_m/4$ and $t = T$.

The results for both $\Delta = \Delta_m/2$ and $\Delta_m/4$ are given. The vertical line indicates an arbitrarily chosen 20% threshold. The probabilities at $\Delta = \Delta_m/4$ are slightly lower than those at $\Delta_m/2$, and those for $\tilde{s}_{ij} \tilde{s}_{ij}$ are slightly lower than those for the other quantities. The probability to observe a less than 20% difference ranges from 75% to 90% for different quantities.

The conditional and joint statistics, e.g., the average of $X^u$ conditioned on $X^v$ and the joint PDF of the two, are presented next. The left panel of Fig. 20 is the results for $X \equiv \Pi_{\Delta}$ with $\Delta = \Delta_m/4$. For a perfectly reconstructed field, the conditional average will fall on the diagonal (dotted line). Therefore, the figure shows that $\Pi_{\Delta}^u$ slightly underestimates the magnitude of $\Pi_{\Delta}^v$, and the underestimate increases slightly with the magnitude of $\Pi_{\Delta}^v$. Nevertheless, the agreement is very good. The joint PDF does concentrate around the diagonal, although there is significant scattering. The
scattering is indicated by the error bars, which represent \( \pm \sigma_\Pi u | \Pi v \Delta | \langle \Pi v \Delta \rangle \) with \( \sigma_\Pi u | \Pi v \Delta | \equiv \left[ \langle (\Pi u \Delta)^2 \rangle \langle (\Pi v \Delta)^2 \rangle \right]^{1/2} \) being the conditional RMS value of \( \Pi u \Delta \). The error bar increases mildly with the magnitude of \( \Pi v \Delta \). The right panel of Fig. 20 plots the same results for \( P_\omega \equiv \tilde{s}_{ij} \tilde{\omega}_i \tilde{\omega}_j \). The results for enstrophy and strain rate are shown in Fig. 21. Good agreement is also observed for these quantities. The error bars are slightly larger for \( P_\omega u \) and \( \tilde{\omega}_i u \tilde{\omega}_i u \), compared with the other two quantities.

### 3.7. Reconstruction of non-local structures

Fig. 2 has shown that the shape, orientation and locations of the strong vortices in the reconstructed field can be remarkably similar to those in the target field. We quantify the similarity of the vortical and other structures using MVEEs and MVEE trees in this subsection.

#### 3.7.1. Vortical structures

The calculation of MVEEs have been explained in Section 2.4. Let the directions of the axes of an MVEE be denoted by \( e_i \) (\( i = \alpha, \beta, \gamma \)), whereas the length of axis \( i \) be denoted by \( \ell_i \). The indices \( \alpha, \beta \) and \( \gamma \) denote the major, intermediate and the minor axes, respectively, so that \( \ell_\alpha \geq \ell_\beta \geq \ell_\gamma \). As before, the parameters in the target fields are indicated by superscript/subscript \( v \) whereas those in the reconstructed fields by \( u \).

We now consider strong vortical structures where \( \tilde{\omega} \geq 3 \langle \tilde{\omega}_i \tilde{\omega}_i \rangle^{1/2} \) with filter scale \( \Delta = \Delta_m / 2 \) and \( \Delta_m / 4 \). As it turns out, the number of isolated structures in a field may be too large to process. If this happens, only the largest \( N_s \equiv 10 \) structures in the target \( v \) and their matching structures in \( u \) are used. Only the data in the five realizations of case C1 are analyzed. There are thus in total \( N_e \equiv 5N_s = 50 \) pairs of matching MVEEs at each time step. As the number of samples is modest, medians instead of means are used to delineate the average behaviors, because the former is less sensitive to extraneous fluctuations. Additional details are given by percentiles and histograms.

Fig. 22 compares the sizes of the vortical structures in \( u \) and \( v \) in terms of \( R \), which is the median of the ratio between the numbers of grid points in the two matching vortices. There are \( N_s = 10 \) groups of vortices of different sizes. The thick line with filled circles shows \( R \) calculated over all \( N_s = 10 \) groups, whereas the thin lines with empty circles show \( R \) in each group separately. The figure shows that the vortices in \( u \) tend to be smaller initially, the median ratio being only about 10%. However, it increases quickly,
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Figure 22. The median $R$ of the ratio between the numbers of grid points in two matching structures (circles with thick line). The green thin lines show the medians in different size groups. Left: $\Delta = \Delta_m/2$; right: $\Delta = \Delta_m/4$. From case C1.

Figure 23. The normalized histograms of $\ell_u^i/\ell_v^i$ for $i = \alpha$ (left column), $i = \beta$ (middle column), and $i = \gamma$ (right column), with $\Delta = \Delta_m/2$ (top row) and $\Delta = \Delta_m/4$ (bottom row). From case C1 with $t = T$.

reaching 90% at $t/T \approx 0.5$ and approximately 1 at the end of the optimization horizon. The same behavior is observed at two filter scales. The thin lines show that there is significant statistical variation between different groups.

The sizes can also be measured with the lengths of the axes of the MVEEs. Fig. 23 plots the histograms of the ratios of the axes in $u$ to those in $v$ at $t/T = 1$ for case C1. The results are consistent with Fig. 22. The peaks of the histograms show that 80% of the axes of the MVEEs in $u$ fall within $\pm10\%$ of those in $v$.

The median and 80th percentile for $|\ell_u^i - \ell_v^i|/\ell_v^i$ (i.e., the relative error in the lengths of the axes) are shown in Fig. 24 for two filter scales. Both values decrease over time. At $t/T = 1$, the median is reduced down to somewhere between 5% and 10%, i.e., for about half of MVEEs, the relative error in the length of the axes is less than $5 \sim 10\%$. The result for the 80th percentile shows that, for 80% MVEEs the relative error is below $10 \sim 15\%$. The difference between different axes is very small.
Figure 24. The medians (empty symbols) and the 80th percentiles (solid symbols) of $\|\ell_i^u - \ell_i^v\|/\ell_i^v$ ($i = \alpha, \beta, \gamma$) for the MVEEs of matching vortices. Circles: $i = \alpha$; squares: $i = \beta$; diamonds: $i = \gamma$. Left: $\Delta = \Delta_m/2$; right: $\Delta = \Delta_m/4$. From case C1.

Figure 25. The histograms of $|\cos(e_i^v, e_i^u)|$ for $i = \alpha$ (top row), $i = \beta$ (middle row), and $i = \gamma$ (bottom row), at $t/T = 0$ (left column), $1/2$ (middle column), and $1$ (right column). From case C1 with $\Delta = \Delta_m/2$.

The alignment between $e_i^v$ and $e_i^u$ is examined in Fig. 25, where the histograms of $|\cos(e_i^v, e_i^u)|$ are plotted ($t/T = 0, 1/2, 1$ from left to right, respectively). The $i$th axes of the matching MVEEs would align perfectly if $|\cos(e_i^v, e_i^u)| = 1$. Initially, the correct alignment between the axes is visible, but the peaks are not strong. Perfect alignment
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develops quickly. The histograms at $t/T = 1/2$ and 1 basically concentrate around 1. In Fig. 26, the medians and the 20th percentiles of the cosines are plotted for two filter scales. In both cases, $|\cos(e^x_i, e^u_i)| \geq 0.95$ is observed at $t/T = 1$ for the 20th percentiles for all axes, which corresponding to angles smaller than 18°. Therefore, only 20% samples have mis-alignment angle bigger than 18°.
Figure 29. The median $R$ of the ratio between the numbers of grid points in two matching straining structures (circles with thick line). The green thin lines show the medians in different size groups. Left: for straining structures; middle: for structures with large positive $\Pi_\Delta$; right: for structures with large negative $\Pi_\Delta$. From case C1 with $\Delta = \Delta_m/4$.

Figure 30. The medians (empty symbols) and the 20th percentiles (solid symbols) of $|\cos(\mathbf{e}_v^i, \mathbf{e}_u^i)|$ for straining structures (left), structures with large positive $\Pi_\Delta$ (middle), and structures with large negative $\Pi_\Delta$ (right). Circles: $i = \alpha$; squares: $i = \beta$; diamonds: $i = \gamma$. From case C1 with $\Delta = \Delta_m/4$.

Figure 31. The medians (empty symbols) and the 80th percentiles (solid symbols) of $|\ell_v^i - \ell_u^i|/\ell_y^i$ for straining structures (left), structures with large positive $\Pi_\Delta$ (middle), and structures with large negative $\Pi_\Delta$ (right). Circles: $i = \alpha$; squares: $i = \beta$; diamonds: $i = \gamma$. From case C1 with $\Delta = \Delta_m/4$.

Fig. 27 and 28 compare the locations of the MVEEs. The quantity being examined is $d_{c,i}/\ell_y^i$ ($i = \alpha, \beta, \gamma$), where $d_{c,i} = |(\mathbf{c}_v^i - \mathbf{c}_u^i) \cdot \mathbf{e}_v^i|$ is the displacement between the centers of the MVEEs in the direction of $\mathbf{e}_v^i$. Thus, $d_{c,i}/\ell_y^i$ is the displacement relative to the lengths of the axes of the MVEE in $\mathbf{v}$. The median and the 80th percentile of $d_{c,i}/\ell_y^i$ are given in Fig. 27. The 80th percentile is quite large initially, but decreases quickly over time. For $\Delta = \Delta_m/2$, the relative displacement at $t/T = 1$ is less than 10% for the 80th percentiles, and less than 5% for the medians. For $\Delta = \Delta_m/4$, the displacement is slightly larger, but still is less than 15% for the 80th percentile.
3.7.2. Straining structures and SGS energy dissipation structures

The MVEEs are used in this subsection to investigate the strain rate and the SGS energy dissipation rate. As the results show behaviors similar to those for the vortical structures, only selected results are presented. For the strain rate, the structures with \((\bar{s}_{ij}\bar{s}_{ij})^{1/2} \geq 2.5(\bar{s}_{ij}\bar{s}_{ij})^{1/2}\) are considered. For \(\Pi_\Delta\), we examine both structures with large positive values and those with large negative values. The latter represents the regions where backscattering happens. These structures are defined by \(\Pi_\Delta \geq 10\langle \Pi_\Delta \rangle\) and \(\Pi_\Delta \leq -2.5\langle \Pi_\Delta \rangle\), and referred to as \(P^+\) and \(P^-\) structures, respectively.

Fig. 29 shows the sizes of the reconstructed structures using the median ratio \(R\). The reconstructed structures tend to be quite small initially, but the discrepancy decreases over time quickly, similar to the results for vortical structures in Fig. 22. At \(t/T = 1\), all curves level off at about 1. The sizes for the \(P^-\) structures need longer time to improve.

The medians and selected percentiles for alignment cosine \(|\cos(e^v, e^u)|\), relative difference in axis length \(|\ell^u - \ell^v|/\ell^v\), and relative displacement \(d_c/\ell^v\) are plotted in Figs. 30 to 32. For both straining and \(P^+\) structures, excellent alignment is observed at \(t/T = 1\), with the medians and the 20th percentile being mostly above 0.9 (Fig. 30). The medians and 80th percentiles for \(|\ell^u - \ell^v|/\ell^v\) and \(d_c/\ell^v\) are found to be around 5% and 15%, respectively, at \(t/T = 1\), similar to those for the vortical structures (Fig. 31 and 32). For \(P^-\) structures, however, the results show stronger discrepancies between \(v\) and \(u\). For the alignment at \(t/T = 1\), the medians still reach above 0.9. However, the 20th percentile result for the intermediate axis is only about 0.7 (c.f. Fig. 30). The medians and 80th percentiles for \(|\ell^u - \ell^v|/\ell^v\) and \(d_c/\ell^v\) are found at 10% and 20%, respectively. These values are slightly higher than those for the other two structures (see Fig. 31 and 32).

3.7.3. MVEE Trees for vortical structures

The analyses so far have examined the overall sizes, orientations and locations of the non-local structures as defined by their MVEEs. For strongly non-convex structures, the MVEEs do not provide full descriptions. These structures are further examined with the MVEE trees in this sub-section (c.f. Section 2.4). Note that, if two MVEEs are significantly different (e.g., seriously mis-aligned), the difference between their sub-MVEEs will only be bigger. Therefore, this analysis is meaningful mainly for matching structures whose MVEEs are in good agreement; it provides a more detailed comparison for these structures. We will limit our discussion to vortical structures, as the results for other structures are similar, and only the structures at \(t = T\) for the base line case C1 are investigated. The analysis is applied to the non-convex vortical structures with \(R_V \leq 0.8\) where \(R_V\) is the ratio of the volume of a structure to the volume of its convex hull (c.f. Section 2.4). Five pairs of matching vortical structures are found in each realization, and 25 pairs in total from 5 realizations. Two splitting are used to find the sub-MVEEs (c.f. Section 2.4) so that 3 levels of sub-MVEEs are calculated: level 0, 1, and 2, which will be
Figure 33. Left: median of $R_V$, the ratio of the volume of a structure to the volume of its convex hull at different levels. Circle: for $u$; squares: for $v$. Right: the median of $\ell_\alpha/\ell_\gamma$ (solid symbols) and $\ell_\alpha/\ell_\beta$ (empty symbols) at different levels. Circle: for $u$; squares: for $v$. For strongly non-convex vortical structures only. For case C1 with $t/T = 1$ and $\Delta = \Delta_m/2$.

Figure 34. The percentiles for $|\cos(e_i^\alpha, e_i^u)|$ ($i = \alpha, \beta, \gamma$) for MVEEs of matching vortices and their sub-MVEEs. From bottom: the 10th, 20th, 30th, 40th, and 50th percentiles. The dashed lines highlight the 20th and 50th percentiles. From case C1 with $t/T = 1$ and $\Delta = \Delta_m/2$.

Figure 35. The percentiles for $d_{c,i}/\ell_i^\gamma$ ($i = \alpha, \beta, \gamma$) for MVEEs of matching vortices and their sub-MVEEs. From bottom: the 10th to 100th percentiles with increment of 10 percent. The dashed lines highlight the 50th and 80th percentiles. For case C1 with $t/T = 1$ and $\Delta = \Delta_m/2$. 
referred to as L0, L1, and L2 sub-MVEEs. L0 sub-MVEEs are the same as the MVEEs already presented.

The geometry of the sub-structures at different levels is different. Fig. 33 displays the difference in two aspects. The median of $R_V$ (left) shows that $R_V$ is larger at higher levels. Therefore, the sub-structures at higher levels are closer to their convex hulls. The right panel shows the median of the ratio of the lengths of the axes for the sub-MVEEs. The ratio is smaller for a higher level. The implication is that the sub-structures are rounder than their parent structures.

The alignment results for the sub-MVEEs are given in Fig. 34. The five curves in the figure (the spaces between the curves are filled with different shades of gray) are the 10th, 20th, 30th, 40th and 50th percentiles. The values for the L0 sub-MVEEs are close to those in Fig. 26, although they are not exactly the same since here only strongly non-convex structures are considered. For the major axes (shown in the left panel), the alignment becomes weaker at higher levels, since a given percentile increases as the level number increases. For example, the 20th percentile is nearly 1 at L0 and approximately 0.85 at L2. It means that 80% matching L0 sub-MVEEs align almost perfectly, while we are only able to say the major axes of 80% matching L2 sub-MVEEs are aligned to within 32°, corresponding to $|\cos(e_{i}^n, e_{i}^{\alpha})| \geq 0.85$. The 50th percentile for L2 is also nearly 1, indicating a near perfect alignment is still observed for more than a half matching sub-MVEEs. On the other hand, the 10th percentile drops quite significantly at L2. This shows that, although the MVEEs for these matching structures indicate very good agreement in their overall shapes, the compositional parts of these structures do differ quite substantially. This happens for about 10% of the matching structures. For the intermediate (middle panel) and minor axes (right panel), similar behaviours can be observed, although the variation with respect to the levels is somewhat more complicated. However, even at the highest level (L2), the 20th percentile is still almost 1 for the minor axes, and 0.8 for the intermediate axes. Thus, very good alignment is observed for more than 80% samples.

There are several competing factors that affect the alignment of the sub-MVEEs. The higher level sub-structures are smaller and are closer to being convex (c.f., the left panel of Fig. 33). As such, the orientations of the sub-MVEEs are less sensitive to the details of the shapes, so that it is easier to obtain better alignment. On the other hand, the misalignment has a cumulative effect. That is, the mis-alignment between the sub-MVEEs at a lower level naturally affects the alignment between those at higher levels. Furthermore, higher level structures tend to be rounder (c.f., the right panel of Fig. 33). This too increases the chance of mis-alignment at higher levels, because small perturbation to
a sphere could lead to ellipsoids with very different axis directions. The competition of these mechanisms may lead to the observed non-monotonic behavior. For the same reason, misalignment between the MVEEs does not necessarily imply large deviation in the shapes of the structures.

The relative displacement of the sub-MVEEs is presented in Fig. 35, which shows higher percentiles increase quicker with the level numbers (see, e.g., the 80th percentiles). This is likely due to the cumulative effect of displacement. The curves for smaller percentiles (which correspond to cases with small relative displacements) are rather flat. Therefore, if the matching MVEEs locate closely to each other, so do their sub-MVEEs. Quantitatively, the 50th percentiles at L2 are approximately 0.08, 0.07 and 0.06 for the major, intermediate and minor axes, respectively. Thus, in about 50% cases, the relative displacement is less than 8%, 7%, and 6% in the major, intermediate and minor axis directions, respectively. Fig. 36 plots the relative difference in the lengths of the axes. For this quantity, the relative differences at L2 and L0 are not much different; the median value is found around 5%. It is probably not surprising since there is no cumulative effect for the lengths of the axes.

To sum up, we observe that, if the matching MVEEs already are close to each other, their sub-MVEEs also show good agreement in orientation, sizes and locations. Therefore, the description provided by the MVEEs might already be enough in some applications.

4. Discussion and conclusions

The four dimensional variational (4DVAR) method has been used to reconstruct the small scales of a sequence of 3D Kolmogorov turbulent velocity fields with a moderate Reynolds number. Velocity on a coarse grid from a time sequence of DNS velocity fields is used as the measurement data. We focus on the reconstruction of instantaneous distributions of the small scales, and present detailed assessment at scales one and two octaves smaller than the measurement grid. In terms of Fourier modes, the number of modes contained down to these scales are eight and sixty-four times of those in the measurement grid, respectively. The filtered vorticity \( \tilde{\omega}_i \), filtered strain rate tensor \( \tilde{s}_{ij} \), the vortex stretching term \( \tilde{s}_{ij} \tilde{\omega}_i \tilde{\omega}_j \), and the subgrid-scale energy dissipation \( P_\Delta = -\tau_{ij} \tilde{s}_{ij} \) are used to characterize the small scales. In order to quantitatively assess the reconstruction of non-local structures, minimum volume enclosing ellipsoids (MVEE) and MVEE trees are introduced. A method to apply these objects in the analysis is proposed and applied to the 4DVAR data.

Different parameters are tested, including the resolution of the measurement data \( k_m \), the tolerance for the convergence test \( e_{\text{tot}} \), and the optimization horizon \( T \). \( k_m \) has the most significant effects on the reconstruction. In reference to the threshold value \( k_c = 0.2\eta^{-1} K \) found previously in the literature with a direct substitution method, the reconstruction failed when \( k_m \) is significantly smaller than \( k_c \). Accurate reconstruction is found for \( k_m \) slightly smaller than \( k_c \). Although more research is needed to prove that the threshold value is the same in 4DVAR, these results demonstrate that reconstruction from a finite time sequence of data using 4DVAR can be accomplished for \( k_m \) at or above the threshold value \( k_c \).

The reconstruction at the initial time \( t = 0 \) and at the end of the optimization horizon \( t = T \) is examined separately for the successful cases. Larger \( T \) and smaller \( e_{\text{tot}} \) improve the reconstruction at \( t = T \). For \( T \) equal the large eddy turnover time scale, accurate reconstruction at scales at least two octaves smaller than the measurement grids is obtained for a range of quantities. The main findings are summarized as follows, where approximate numbers are quoted to give an estimate of the efficacy of the method:
(i) The reconstruction achieves higher than 95% point-wise correlation for the filtered enstrophy and the other parameters. The spectral correlation for velocity is higher than 80% across the spectrum. The normalized RMS point-wise difference and the normalized spectral difference increase towards the small scales, and reaches around 30% at the Kolmogorov scale.

(ii) The non-local structures are reconstructed accurately. For most quantities, the misalignment is less than 15° for the majority of the samples, the sizes differing within ±10%, and the locations displaced by less than 15% of the lengths of the axes. The structures with strong negative SGS energy dissipation have slightly larger errors.

(iii) A direct substitution scheme provides much poorer reconstruction, where the RMS point-wise difference is about three times as big for the filtered enstrophy, and is almost ten times as big in terms of the spectral difference of the velocity fields. Meanwhile, Lagrangian interpolation is simply infeasible.

The reconstruction at \( t = 0 \) can capture the qualitative features of the small scales in turbulence, including qualitatively correct energy spectrum and mean SGS energy dissipation. However, significant discrepancy exists and the agreement is improved very little by tuning the optimization horizon or the tolerance. A plausible explanation is that our optimal solutions have not truly converged. Due to the high computational costs, we have not been able to further improve the solution by using even smaller \( e_{\text{tot}} \). Given the chaotic nature of the flow, it is challenging to obtain convergent results if \( T \) is significantly larger or \( e_{\text{tot}} \) is much smaller.

This study confirms the efficacy of the 4DVAR scheme, and demonstrates that MVEEs and MVEE trees can provide useful information about non-local structures for CFD with data assimilation when the prediction of instantaneous velocity field is one of the main interests. As the application of data assimilation in CFD is still in an early stage, many questions remain open. Most importantly, the Reynolds number for the flow considered here is relatively low. The natural next step is to investigate the quality of reconstruction when coarse-grained models (such as large eddy simulations) are used to model the data. This will allow us to investigate flows with higher Reynolds numbers. To obtain better converged results, a better understanding of the behavior of the adjoint fields is useful. These questions are the focus of our on-going research.

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