Learning Similarity Metrics for Numerical Simulations

Georg Kohl 1  Kiwon Um 1  Nils Thuerey 1

Abstract
We propose a neural network-based approach that computes a stable and generalizing metric (LSiM), to compare field data from a variety of numerical simulation sources. Our method employs a Siamese network architecture that is motivated by the mathematical properties of a metric. We leverage a controllable data generation setup with partial differential equation (PDE) solvers to create increasingly different outputs from a reference simulation in a controlled environment. A central component of our learned metric is a specialized loss function that introduces knowledge about the correlation between single data samples into the training process. To demonstrate that the proposed approach outperforms existing simple metrics for vector spaces and other learned, image-based metrics, we evaluate the different methods on a large range of test data. Additionally, we analyze benefits for generalization and the impact of an adjustable training data difficulty. The robustness of LSiM is demonstrated via an evaluation on three real-world data sets.

1. Introduction
Evaluating computational tasks for complex data sets is a fundamental problem in all computational disciplines. Regular vector space metrics, such as the $L^2$ distance, were shown to be very unreliable (Wang et al., 2004; Zhang et al., 2018), and the advent of deep learning techniques with convolutional neural networks (CNNs) made it possible to more reliably evaluate complex data domains such as natural images, texts (Benajiba et al., 2018), or speech (Wang et al., 2018). Our central aim is to demonstrate the usefulness of CNN-based evaluations in the context of numerical simulations. These simulations are the basis for a wide range of applications ranging from blood flow simulations to aircraft design. Specifically, we propose a novel learned simulation metric (LSiM) that allows for a reliable similarity evaluation of simulation data.

Potential applications of such a metric arise in all areas where numerical simulations are performed, or similar data is gathered from observations. For example, accurate evaluations of existing and new simulation methods with respect to a known ground truth solution (Oberkampf et al., 2004) can be performed more reliably than with a regular vector norm. Another good example is weather data for which complex transport processes and chemical reactions make in-place comparisons with common metrics unreliable (Jolliffe & Stephenson, 2012). Likewise, the long-standing, open questions of turbulence (Moin & Mahesh, 1998; Lin et al., 1998) can benefit from improved methods for measuring the similarity and differences in data sets and observations.

In this work, we focus on field data, i.e., dense grids of scalar values, similar to images, which were generated with known partial differential equations (PDEs) in order to ensure the availability of ground truth solutions. While we focus on 2D data in the following to make comparisons with existing techniques from imaging applications possible, our approach naturally extends to higher dimensions. Every sample of this 2D data can be regarded a high dimensional vector, so metrics on the corresponding vector space are applicable to evaluate similarities. These metrics, in the following denoted as shallow metrics, are typically simple, element-wise functions such as $L^1$ or $L^2$ distances. Their inherent problem is that they cannot compare structures on different scales or contextual information.

Many practical problems require solutions over time and need a vast number of non-linear operations that often result in substantial changes of the solutions even for small changes of the inputs. Hence, despite being based on known, continuous formulations, these systems can be seen as chaotic. We illustrate this behavior in Fig. 1, where two smoke flows are compared to a reference simulation. A single simulation parameter was varied for these examples, and a visual inspection shows that smoke plume (a) is more similar to the reference. This matches the data generation process: version (a) has a significantly smaller parameter change than (b), as shown in the inset graph on the right. LSiM robustly predicts the ground truth distances while the $L^2$ metric labels plume (b) as more similar. In our work, we focus on retrieving the relative distances of simulated data sets. Thus, we do not aim for retrieving the absolute parameter change but a relative distance that preserves ordering.
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Figure 1. Example of field data from a fluid simulation of hot smoke with normalized distances for different metrics. Our method (LSiM, green) approximates the ground truth distances (GT, gray) determined by the data generation method best, i.e., version (a) is closer to the ground truth data than (b). An $L^2$ metric (red) erroneously yields a reversed ordering.

with respect to this parameter.

Using existing image metrics based on CNNs for this problem is not optimal either: natural images only cover a small fraction of the space of possible 2D data, and numerical simulation outputs are located in a fundamentally different data manifold within this space. Hence, there are crucial aspects that cannot be captured by purely learning from photographs. Furthermore, we have full control over the data generation process for simulation data. As a result, we can create arbitrary amounts of training data with gradual changes and a ground truth ordering. With this data, we can learn a metric that is not only able to directly extract and use features but also encodes interactions between them. The central contributions of our work are as follows:

- We propose a Siamese network architecture with feature map normalization, which is able to learn a metric that generalizes well to unseen simulations methods.
- We propose a novel loss function that combines a correlation loss term with a mean squared error to improve the accuracy of the learned metric.
- In addition, we show how a data generation approach for numerical simulations can be employed to train networks with general and robust feature extractors for metric calculations.

2. Related Work

One of the earliest methods to go beyond using simple metrics based on $L^p$-norms for natural images was the structural similarity index (Wang et al., 2004). Despite improvements, this method can still be considered a shallow metric. Over the years multiple large databases for human evaluations of natural images were presented, for instance CSIQ (Larson & Chandler, 2010), TID2013 (Ponomarenko et al., 2015), and CID:IQ (Liu et al., 2014). With this data and the discovery that CNNs can create very powerful feature extractors that are able to recognize patterns and structures, deep feature maps quickly became established as means for evaluation (Amirshahi et al., 2016; Kang et al., 2014; Kim & Lee, 2017). Recently, these methods were improved by predicting the distribution of human evaluations instead of directly learning distance values (Prashnani et al., 2018; Talebi & Milanfar, 2018b). Zhang et al. compared different architecture and levels of supervision, and showed that metrics can be interpreted as a transfer learning approach, by applying a linear weighting to the feature maps of any network architecture to form the image metric $LPIPS v0.1$. Typical use cases of these image-based CNN metrics are computer vision tasks like detail enhancement (Talebi & Milanfar, 2018a), style transfer, and super-resolution (Johnson et al., 2016). Generative adversarial networks also leverage CNN-based losses by training a discriminator network in parallel to the generation task (Dosovitskiy & Brox, 2016).

Siamese network architectures are known to work well for a variety of comparison tasks such as audio (Zhang & Duan, 2017), satellite images (He et al., 2019) or the similarity of interior product designs (Bell & Bala, 2015). Furthermore, they yield robust object trackers (Bertinetto et al., 2016), algorithms for image patch matching (Hanif, 2019), and for descriptors for fluid flow synthesis (Chu & Thuerey, 2017). Inspired by these works we use a similar Siamese neural network architecture for our metric learning task. In contrast to other work on self-supervised learning that utilizes spatial or temporal changes to learn meaningful representations (Agrawal et al., 2015; Wang & Gupta, 2015), our method does not rely on tracked keypoints in the data.

While correlation terms have been used for learning joint representations by maximizing correlation of projected views (Chandar et al., 2016), and are popular for style transfer applications via the Gram matrix (Ruder et al., 2016), they were not used for learning distance metrics. As we demonstrate below, they can yield significant improvements in terms of the inferred distances.

Similarity metrics for numerical simulations are a topic of ongoing investigation. A variety of specialized metrics have been proposed to overcome the limitations of $L^p$-norms, such as the displacement and amplitude score from the area
of weather forecasting (Keil & Craig, 2009), and permutation based metrics for energy consumption forecasting (Haben et al., 2014). Turbulent flows, on the other hand, are often evaluated in terms of aggregated frequency spectra (Pitsch, 2006). Crowd-sourced evaluations based on the human visual system were also proposed to evaluate simulation methods for physics-based animation (Um et al., 2017), and for comparing non-oscillatory discretization schemes (Um et al., 2019). These results indicate that visual evaluations in the context of field data are possible and robust, but they require extensive (and potentially expensive) user studies. Additionally, our method naturally extends to higher dimensions, while human evaluations inherently rely on projections with at most two spatial and one time dimension.

3. Constructing a CNN-based Metric

In the following, we explain our considerations when employing CNNs as evaluation metrics. For a comparison that corresponds to our intuitive understanding of distances, an underlying metric has to obey certain criteria. More precisely, a function \( m : \mathbb{I} \times \mathbb{I} \rightarrow [0, \infty) \) is a metric on its input space \( \mathbb{I} \) if it satisfies the following properties \( \forall x, y, z \in \mathbb{I} \):

\[
\begin{align*}
    m(x, y) &\geq 0 \quad \text{non-negativity} \quad (1) \\
    m(x, y) &= m(y, x) \quad \text{symmetry} \quad (2) \\
    m(x, y) &\leq m(x, z) + m(z, y) \quad \text{triangle ineq.} \quad (3) \\
    m(x, y) = 0 &\iff x = y \quad \text{identity of indisc.} \quad (4)
\end{align*}
\]

The properties (1) and (2) are crucial as distances should be symmetric and have a clear lower bound. Eq. (3) ensures that direct distances cannot be longer than a detour. Property (4), on the other hand, is not really useful for discrete operations as approximation errors and floating point operations can easily lead to a distance of zero for slightly different inputs. Hence, we focus on a relaxed, more meaningful definition \( m(x, x) = 0 \), which leads to a so-called pseudometric. It allows for a distance of zero for different inputs but has to be able to spot identical inputs.

We realize these requirements for a pseudometric with an architecture that follows popular perceptual metrics such as LPIPS: The activations of a CNN are compared in latent space, accumulated with a set of weights, and the resulting per-feature distances are aggregated to produce a final distance value. Fig. 2 gives a visual overview of this process.

3.1. Base Network

The sole purpose of the base network is to extract feature maps from both inputs. The Siamese architecture implies that the weights of the base network are shared for both inputs, meaning all feature maps are comparable. In addition, this ensures the identity of indiscernibles because, for identical feature maps, a distance of zero is guaranteed by the following operations. In the last years, a variety of powerful CNN-based feature extraction architectures were proposed. We experimented with various networks, such as AlexNet (Krizhevsky et al., 2017), VGG (Simonyan & Zisserman, 2015), SqueezeNet (Iandola et al., 2016), and a fluid flow prediction network (Thuerey et al., 2018). In all cases, only the feature extracting layers are used, and the remaining layers, which are responsible for the original task (e.g., classification), are discarded. Building on this previous work, we consider three variants of the networks below: using the original pre-trained weights, fine-tuning them, or re-training the full networks from scratch. In contrast to typical CNN tasks where only the result of the final output layer is further processed, we make use of the full range of extracted features across the layers of a CNN (see Fig. 2). This implies a slightly different goal: while early features should be general enough to allow for extracting more complex features in deeper layers, this is not their sole purpose. Rather, features in earlier layers of the network can directly participate in the final distance calculation and can yield important cues. As we will demonstrate below, we achieved the best performance for our data sets using a custom architecture with five layers, similar to a reduced AlexNet, that was trained from scratch (see App. B.1).

![Figure 2](image-url)

*Figure 2.* Overview of the proposed distance computation for a simplified base network that contains three layers with four feature maps each in this example. The output shape for every operation is illustrated below the transitions in orange and white. Bold operations are learned, i.e., contain weights influenced by the training process.
3.2. Feature Map Normalization

The goal of normalizing the feature maps is to transform the extracted features of each layer, which typically have very different orders of magnitude, into comparable ranges. While this task could potentially be performed by the learned weights, we found the normalization to yield improved performance in general (see App. B.2). Zhang et al. proposed a length unit normalization using a division by the Euclidean norm in channel dimension to only measure the angle between the latent space vectors with a cosine distance. Instead, we suggest to interpret all possible feature maps as a normal distribution and to normalize them to a standard normal distribution. This is achieved via a preprocessing step using the full training data set: we subtract the mean of the feature maps and then divide by their standard deviation in channel dimension for each layer. As a result, we can measure angles for the latent space vectors and compare their magnitude in the global length distribution.

3.3. Latent Space Differences

Combining the latent spaces representations $\tilde{x}$, $\tilde{y}$ that consist of all extracted features from the two inputs $x$, $y$ lies at the core of the metric computation. Here, the most obvious approach to employ an element-wise difference, i.e., $\tilde{x}_i - \tilde{y}_i$, is not advisable as this would directly violate the metric properties above. Instead, non-negativity and symmetry can be ensured via $|\tilde{x} - \tilde{y}|$ or $(\tilde{x} - \tilde{y})^2$. We found that both work equally well in practice. Considering the importance of comparing the extracted features, the simple operations used for comparing the features do not seem optimal. Rather, one can imagine that improvements in terms of comparing one set of feature activations could lead to overall improvements for derived metrics. Hence, we experimented with replacing these operations with a pre-trained CNN-based metric for each feature map. This creates a recursive process or "meta-metric", that reformulates the initial problem of learning input similarities in terms of learning feature space similarities. However, as detailed in App. B.3, we have not found this recursive approach to yield any substantial improvements. This implies that once a large enough number of expressive features is available for comparison, the in-place difference of each feature is sufficient to compare two inputs. In the following, we compute the feature difference maps (Fig. 2, in yellow) via $(\tilde{x} - \tilde{y})^2$.

3.4. Aggregations

The subsequent aggregation operations (Fig. 2, in green) are applied to the difference maps to compress the contained per feature differences along the different dimensions into a single distance value. The aggregation operations only need to preserve the metric properties already established via the latent space difference. To aggregate the difference maps along the channel dimension, we found the weighted average proposed by Zhang et al. to work very well. Thus, we use one learnable weight to control the importance of a feature. The weight is a multiplier for the corresponding difference map before summation along the channel dimension. To preserve non-negativity and the triangle inequality, the weights are clamped to be non-negative. A negative weight would mean that a larger difference in this feature produces a smaller overall distance, which is not helpful. For spatial and layer aggregation, functions like a summation or averaging are sufficient and generally interchangeable. We tested more intricate aggregation functions, e.g., by learning a spatial average or determining layer importance weights dynamically from the inputs. When the base network is fixed and the metric only has very few trainable weights, this did improve the overall performance. But, with a fully trained base network, the feature extraction seems to automatically adopt these aspects making a more complicated aggregation unnecessary.

Additional details for the steps above are given in App. A, where we also show that the proposed Siamese architecture by construction qualifies as a pseudometric.

4. Data Generation and Training

Similarity data sets for natural images typically rely on changing already existing images with distortions, noise, or other operations and assigning ground truth distances according to the strength of the operation. Since we can control the data creation process for numerical simulations directly, we can generate large amounts of simulation data with increasing dissimilarities by altering the parameters used for the simulations. As a result, the data contains more information about the nature of the problem, i.e., which changes of the data distribution should lead to increased distances, than by applying modifications as a post-process.

4.1. Data Generation

Given a set of model equations, e.g., a PDE from fluid dynamics, typical solution methods consist of a solver that, given a set of boundary conditions, computes discrete approximations of the necessary differential operators. The discretized operators and the boundary conditions typically contain problem dependent parameters, which we collectively denote with $p_0, p_1, \ldots, p_i, \ldots$. in the following. We only consider time dependent problems, and our solvers start with initial conditions at $t_0$ to compute a series of time steps $t_1, t_2, \ldots$ until a target point in time ($t_\text{t}$) is reached. At that point, we obtain a reference output field $o_0$ from one of the PDE variables, e.g., a velocity.

For data generation, we incrementally change a single parameter $p_i$ in $n$ steps $\Delta_i, 2 \cdot \Delta_i, \ldots, n \cdot \Delta_i$ to create a series...
of \( n \) outputs \( o_1, o_2, \ldots, o_n \). We consider a series obtained in this way to be increasingly different from \( o_0 \). To create natural variations of the resulting data distributions, we add Gaussian noise fields with zero mean and adjustable variance to an appropriate simulation field such as a velocity. This noise allows us to generate a large number of varied data samples for a single simulation parameter \( p_i \). In addition, it is similar in nature to numerical errors introduced by discretization schemes. Thus, these perturbations enlarge the space covered by the training data, and we found that training networks with suitable noise levels improves robustness as we will demonstrate below. The process for data generation is summarized in Fig. 3.

As PDEs can model extremely complex and chaotic behaviour, there is no guarantee that the outputs always exhibit increasing dissimilarity with the increasing parameter change. This behaviour is what makes the task of similarity assessment so challenging. Even if the solutions are essentially chaotic, their behaviour is not arbitrary but rather governed by the rules of the underlying PDE. For our data set, we choose the following range of representative PDEs:

- We include a pure Advection-Diffusion model (AD) and Burger’s equation (BE), which introduces a viscosity term.
- Furthermore, we use the full Navier-Stokes equations (NSE), which introduce a conservation of mass constraint. When combined with a deterministic solver and a suitable parameter step size, all these PDEs exhibit chaotic behaviour at small scales and the medium to large scale characteristics of the solutions shift smoothly with increasing changes of the parameters \( p_i \). The noise \( n \) amplifies the chaotic behaviour to larger scales to create an environment with a controlled amount of perturbations. This lets the network learn about the nature of the chaotic behaviour of PDEs without overwhelming it with data where patterns are not observable anymore. The latter can easily happen when \( \Delta \) or \( n \) grows too large and produce essentially random outputs. Instead, we specifically target solutions that are difficult to evaluate in terms of a shallow metric. We choose the smallest \( \Delta \) and \( n \) such that the ordering of several random output samples with respect to their \( L^2 \) difference drops below a correlation value of 0.8.

### 4.2. Training

For training, the 2D scalar fields from the simulations were individually normalized and augmented with random color maps, flips, rotations, and cropping to obtain an input size of \( 224 \times 224 \). Afterwards, each input is standardized to a standard normal distribution by subtracting the mean and dividing by the standard deviation (pre-computed from all training data). Unless noted otherwise, networks were trained with a batch size of 1 for 40 epochs with the Adam optimizer using a learning rate of \( 10^{-5} \) that was reduced to \( 5 \cdot 10^{-6} \) after 15 epochs. To evaluate the trained networks on validation and test inputs, only a bilinear resizing and the standardization step is applied.

### 5. Correlation Loss Function

The central goal of our networks is to identify relative differences of input pairs produced via numerical simulations. Thus, instead of employing a loss that forces the network to only infer given labels or distance values, we train our networks to infer the ordering of a given sequence of varying inputs \( o_1, \ldots, o_n \). We propose to use the Pearson correlation coefficient (see Pearson, 1920), which yields a value in \([-1, 1]\) that measures the linear relationship between two distributions. A value of 1 implies that a linear equation describes their relationship perfectly. We compute this coefficient for a full series of outputs such that the network can learn to extract features that arrange this data series in the correct ordering. Each training sample of our network consists of every possible pair of inputs of the sequence \( o_1, \ldots, o_n \) and the corresponding ground truth distance distribution \( c \in [0,1]^{0.5n(n-1)} \) representing the parameter change from the data generation. For a distance prediction \( d \in [0,\infty)^{0.5n(n-1)} \) of our network for one sample, we compute the loss with:

\[
L(c, d) = \lambda_1 (c - d)^2 + \lambda_2 (1 - \frac{(c - \bar{c}) \cdot (d - \bar{d})}{\|c - \bar{c}\|_2 \|d - \bar{d}\|_2}) \quad (5)
\]

Here, the mean of a distance vector is denoted by \( \bar{c} \) and \( \bar{d} \) for ground truth and prediction, respectively. The first part of the loss is a regular MSE term, which minimizes the difference between predicted and actual distances. The second part is the Pearson correlation coefficient, which is inverted such that the optimization results in a maximization of the correlation. As this formulation depends on the length of the input sequence, the two terms are scaled to adjust
their relative influence with $\lambda_1$ and $\lambda_2$. For the training, we chose 10 variations for each reference simulation, i.e., $n = 11$. If $n$ should vary during training, the influence of both terms needs to be adjusted accordingly. We found that scaling both terms to a similar order of magnitude worked best in our experiments.

In Fig. 4, we investigate how the proposed loss function compares to other commonly used loss formulations for our full network and a pre-trained network. In addition to our full loss function, we consider a loss function that replaces the Pearson correlation with a simpler cross-correlation $(c \cdot d) / (\|c\|_2 \|d\|_2)$. We also include networks trained with only the MSE or only the correlation terms for each of the two variants.

Figure 4. Results on our test data of the proposed approach (LSIM) and a smaller model (AlexNet) for different loss functions.

As shown in Fig. 4, a simple MSE loss yields a low accuracy of less than 0.6. Using any correlation based loss function for the AlexNet metric (see Section 6.2) improves the results, but there is no major difference due to the limited number of only 1152 trainable weights. For LSIM, the proposed combination of MSE loss with the Pearson correlation performs significantly better than using cross-correlation or a variant without the MSE loss. Interestingly, combining cross correlation with MSE yields worse results than cross correlation alone. This is caused by the cross correlation term influencing absolute distance values, which potentially conflicts with the MSE term. For our loss, the Pearson correlation only handles the relative ordering while MSE deals with the absolute distances, leading to an improved correlation for the inferred distances.

6. Results

In the following, we will discuss how the data generation approach was employed to create a large range of training and test data from different PDEs. Afterwards, the proposed metric is compared to other metrics, and its robustness is evaluated with several external data sets.

6.1. Data Sets

We created four training (Smo, Liq, Adv and Bur) and two test data sets (LiqN and AdvD) with ten parameter steps for each reference simulation. Based on two 2D NSE solvers, the smoke and liquid simulation training sets (Smo and Liq) add noise to the velocity field and feature varied initial conditions such as fluid position or obstacle properties, in addition to variations of buoyancy and gravity forces. The two other training sets (Adv and Bur) are based on 1D solvers for AD and BE, concatenated over time to form a 2D result. In both cases, noise was injected into the velocity field, and the varied parameters are changes to the field initialization and forcing functions.

For the test data set, we substantially change the data distribution by injecting noise into the density instead of the velocity field for AD simulations to obtain the AdvD data set and by including background noise for the velocity field of a liquid simulation (LiqN). In addition, we employed three more test sets (Sha, Vid, and TID) created without PDE models to explore the generalization for data far from our training data setup. We include a shape data set that features multiple randomized moving rigid shapes, a video data set consisting of frames from random video footage, and the perceptual image data set TID2013 (Ponomarenko et al., 2015). Below, we additionally list a combined correlation score for all test sets (All) apart from TID, which is excluded due to its different structure.

Simulation examples for each data set are shown in Fig. 5 and generation details with additional samples can be found in App. D.

6.2. Performance Evaluation

To evaluate the performance of a metric on a data set, we first compute the distances from each reference simulation to all corresponding variations. Then, the predicted and the ground truth distance distributions over all samples are combined and compared using Spearman’s rank correlation coefficient (see Spearman, 1904). Like the Pearson correlation, it is a value in $[-1, 1]$ to compare distributions, but it measures the correlation between ranking variables, i.e., monotonic relationships.

The top part of Tab. 1 shows the performance of the shallow metrics $L^2$ and SSIM as well as the LPIPS metric (Zhang et al., 2018) for all our data sets. The results clearly show that shallow metrics are not suitable to compare the samples in our data set and only achieve good correlation values on the TID2013 data set, which contains a large number of pixel-based image variations without contextual structures. The perceptual LPIPS metric performs better in general and outperforms our method on the image data sets Vid and TID. This is not surprising as LPIPS is specifically trained
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Figure 5. Samples from our data sets. For each subset the reference is on the left, and three variations in equal parameter steps follow. From left to right and top to bottom: Smo (density, velocity, and pressure), Adv (density), Liq (flags, velocity, and levelset), Bur (velocity), LiqN (velocity), AdvD (density), Sha and Vid. For such images. For the simulation data sets LiqN and Sha, however, it performs significantly worse than for the image content. The last row of Tab. 1 shows the results of our LSiM network with a very good performance across all data sets. Note that although it was not trained with any natural image content, it still performs well for the image test sets.

The middle block of Tab. 1 contains several interesting variants: AlexNetFrozen and AlexNetFrozen are small models, where the base network is the original AlexNet with pre-trained weights. AlexNetRandom contains purely random aggregation weights without training, whereas AlexNetFrozen only has trainable weights for the channel aggregation and therefore lacks the flexibility to fully adjust to the data distribution of the numerical simulations. The random model performs surprisingly well, pointing to powers of the underlying CNN architecture.

Recognizing that many PDEs include transport phenomena, we investigated optical flow (Horn & Schunck, 1981) as a means to compute motion from field data. For the Optical flow metric, we used FlowNet2 (Ilg et al., 2016) to bidirectionally compute the optical flow field between two inputs and aggregate it to a single distance value by summing all flow vector magnitudes. On the data sets Sha and Vid that are similar to the training data of FlowNet2, it performs relatively well, but in most other cases, it performs poorly. This shows that computing a simple warping from one input to the other is not enough for a stable metric although it seems like an intuitive solution. A more robust metric needs the knowledge of the underlying features and their changes to generalize better to new data.

For the Non-Siamese metric, we used a non-Siamese architecture that directly predicts the distance from both inputs to evaluate whether a Siamese architecture is really beneficial. For this purpose, we employed a modified version of AlexNet that reduces the weights of the feature extractor by 50% and of the remaining layers by 90%. As expected, this metric works great on the validation data but has huge problems with generalization. In addition, even simple metric properties like symmetry are no longer guaranteed because this architecture does not have the inherent constraints of

| Metric            | Validation data sets | Test data sets |
|-------------------|----------------------|----------------|
|                   | Smo | Liq | Adv | Bur | TID | LiqN | AdvD | Sha | Vid | All |
| $L^2$             | 0.67 | 0.80 | 0.72 | 0.59 | 0.83 | 0.72 | 0.58 | 0.50 | 0.77 | 0.56 |
| SSIM              | 0.67 | 0.75 | 0.75 | 0.68 | 0.80 | 0.25 | 0.70 | 0.35 | 0.70 | 0.48 |
| LPIPS v0.1        | 0.72 | 0.75 | 0.75 | 0.72 | 0.80 | 0.63 | 0.60 | 0.81 | 0.82 | 0.66 |
| AlexNetRandom     | 0.64 | 0.75 | 0.67 | 0.64 | 0.84 | 0.64 | 0.67 | 0.61 | 0.78 | 0.62 |
| AlexNetFrozen     | 0.67 | 0.70 | 0.68 | 0.70 | 0.79 | 0.40 | 0.64 | 0.84 | 0.81 | 0.62 |
| Optical flow      | 0.62 | 0.57 | 0.36 | 0.37 | 0.55 | 0.49 | 0.28 | 0.61 | 0.75 | 0.48 |
| Non-Siamese       | 0.72 | 0.82 | 0.76 | 0.69 | 0.29 | 0.73 | 0.62 | 0.60 | 0.72 | 0.63 |
| Skip from scratch | 0.65 | 0.84 | 0.74 | 0.67 | 0.80 | 0.79 | 0.59 | 0.83 | 0.79 | 0.70 |
| LSiM noiseless    | 0.64 | 0.83 | 0.74 | 0.60 | 0.80 | 0.81 | 0.58 | 0.84 | 0.76 | 0.69 |
| LSiM strong noise | 0.63 | 0.82 | 0.71 | 0.61 | 0.80 | 0.78 | 0.50 | 0.80 | 0.77 | 0.65 |
| LSiM (ours)       | 0.68 | 0.82 | 0.76 | 0.70 | 0.78 | 0.80 | 0.61 | 0.85 | 0.76 | 0.71 |
the Siamese setup. Finally, we experimented with multiple fully trained base networks. As re-training existing feature extractors only provided small improvements, we used a custom base network with skip connections for the $\text{L} \text{S} \text{i} \text{M}$ metric. Its results already come close to the proposed approach on most data sets.

The last block in Tab. 1 shows variants of the proposed approach, trained with varied noise levels. This inherently changes the difficulty of the data. Hence, $\text{L} \text{S} \text{i} \text{M}_{\text{noiseless}}$ was trained with relatively simple data without perturbations, whereas $\text{L} \text{S} \text{i} \text{M}_{\text{strong noise}}$ was trained with strongly varying data. Both cases decrease the generalizing capabilities of the trained model on the test data. This indicates that the network needs to see a certain amount of variation at training time in order to become robust, but overly large changes hinder the learning of useful features (see App. B and C).

6.3. Evaluation on Real-World Data

To evaluate the generalizing capabilities of our trained metric, we turn to three representative and publicly available data sets of captured and simulated real-world phenomena, namely buoyant flows, turbulence, and weather. For the former, we make use of the ScalarFlow data set from Eckert et al., which consists of captured velocities of buoyant scalar transport flows. Additionally, we include data from the Johns Hopkins Turbulence Database (JHTDB) (Perlman et al., 2007), which represents direct numerical simulations of fully developed turbulence. As a third case, we use temperature and geopotential fields from the WeatherBench repository (Rasp et al., 2020), which contains global climate data on a Cartesian latitude-longitude grid of the earth.

For the evaluation, we extracted sequences of frames with fixed temporal and spatial intervals from each data set to obtain a ground truth ordering. We use six different interval spacings with 60-240 sequences each, yielding more than 360 evaluations for every data source (details in App. E). We then measure how well our metric recovers the original ordering in the presence of the complex changes of content, driven by the underlying physical processes. We employ the pre-trained $\text{L} \text{S} \text{i} \text{M}$ metric model that was generated as outlined in the previous sections. It recovers the ordering of all three cases with high accuracy yielding averaged correlations of $0.92 \pm 0.02$, $0.95 \pm 0.04$, and $0.92 \pm 0.08$ for ScalarFlow, JHTDB, and WeatherBench, respectively. Visual examples of the three data sets are shown on the left of Fig. 6 while the right side summarizes the robustness of the $\text{L} \text{S} \text{i} \text{M}$ evaluation.

7. Conclusion

We have presented the $\text{L} \text{S} \text{i} \text{M}$ metric to reliably and robustly compare outputs from numerical simulations. Our method significantly outperforms existing shallow metric functions and provides better results than other learned metrics. We demonstrated the usefulness of the correlation loss, showed the benefits of a controlled data generation environment, and highlighted the stability of the obtained metric for a range of real-world data sets.

Our trained $\text{L} \text{S} \text{i} \text{M}$ metric has the potential to impact a wide range of fields: from the fast and reliable accuracy assessment of new simulation methods, over robust optimizations of parameters for reconstructions of observations, to guiding generative models of physical systems. Furthermore, it will be highly interesting to evaluate other loss functions, e.g., mutual information (Bachman et al., 2019) or contrastive predictive coding (Hénaff et al., 2019). We also plan to evaluate our approach for an even larger collection of PDEs as well as for 3D and 4D data sets. Especially turbulent flows are a highly relevant and interesting area for future work on learned evaluation metrics.
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Appendix: Learning Similarity Metrics for Numerical Simulations

This supplemental document contains an analysis of the proposed metric design with respect to properties of metrics in general (App. A), and details to the used network architectures (App. B). Afterwards, material that deals with the data sets is provided. It contains examples and failure cases for each of the data domains and analyzes the impact of the data difficulty (App. C and D). Next, the evaluation on real-world data is described in more detail (App. E). Finally, we explore additional metric evaluations (App. F), and give an overview on the used notation (App. G).

The source code for using and evaluating the trained LSiM metric, and some examples from each data set are published alongside this submission. The full source code for retraining the model, the data generation scripts, and the entire data sets will be published in the near future.

A. Discussion of Metric Properties

To analyze if the proposed method qualifies as a metric, it is split in two functions $m_1 : I \rightarrow \mathbb{L}$ and $m_2 : \mathbb{L} \times \mathbb{L} \rightarrow [0, \infty)$ which operate on the input space $I$ and the latent space $\mathbb{L}$. Through flattening elements from the input or latent space into vectors, $I \simeq \mathbb{R}^a$ and $\mathbb{L} \simeq \mathbb{R}^b$ where $a$ and $b$ are the dimensions of the input data or all feature maps respectively, and both values have a similar order of magnitude. $m_1$ describes the non-linear function computed by the base network combined with the following normalization and returns a point in the latent space. $m_2$ uses two points in the latent space to compute a final distance value, so it includes the latent space difference and the aggregation along the spatial, layer, and channel dimensions. With the Siamese network architecture the resulting function for the entire approach is

$$m(x, y) = m_2(m_1(x), m_1(y)).$$

The identity of indiscernibles mainly depends on $m_1$ because even if $m_2$ itself guarantees this property, $m_1$ could still be non-injective, which means it can map different inputs to the same point in latent space $\tilde{x} = \tilde{y}$ for $x \neq y$. Due to the complicated nature of $m_1$ it is difficult to make accurate predictions about the injectivity of $m_1$. Each base network layer of $m_1$ recursively processes the result of the preceding layer with various feature extracting operations so intuitively, significant changes in the input should produce different feature map results in a layer. But very small changes in the input do lead to zero valued distances predicted by the CNN (i.e. an identical latent space for different inputs), meaning $m_1$ is in practice not injective. In an additional experiment, the proposed architecture was evaluated on about 3500 random inputs from all our data sets, where the CNN received one unchanged and one slightly modified input. The modification consisted of multiple pixel adjustments by one bit (on 8-bit color images) in random positions and channels. When adjusting only a single pixel in the $224 \times 224$ input, the CNN predicts a zero valued distance on about 23% of the inputs, but we never observed an input where seven or more changed pixels resulted in a distance of zero in all experiments.

In this context, the problem of numerical errors is important, because even two slightly different latent space representations could lead to a result that seems to be zero if the difference vanishes in the aggregation operations or is smaller than the floating point precision. On the other hand, an automated analysis to find points that have a different input but an identical latent space image is a challenging problem and left as future work.

The evaluation of the base network and the normalization is deterministic, and hence $\forall x : m_1(x) = m_1(x)$ holds. Further, we know that $m(x, x) = 0$ if $m_2$ guarantees that $\forall m_1(x) : m_2(m_1(x), m_1(x)) = 0$. Thus, the remaining properties non-negativity, symmetry, and the triangle inequality only depend on $m_2$, since for them the original inputs are not relevant, only their respective images in the latent space. The resulting structure with a relaxed identity of indiscernibles is called a pseudometric, where $\forall \tilde{x}, \tilde{y}, \tilde{z} \in \mathbb{L}$:

$$m_2(\tilde{x}, \tilde{y}) \geq 0$$
$$m_2(\tilde{x}, \tilde{y}) = m_2(\tilde{y}, \tilde{x})$$
$$m_2(\tilde{x}, \tilde{y}) \leq m_2(\tilde{x}, \tilde{z}) + m_2(\tilde{z}, \tilde{y})$$

Notice, that $m_2$ has to fulfill these properties with respect to the latent space and not the input space. If $m_2$ is carefully constructed the metric properties still apply, independently of the actual design of the base network or the feature map normalization.

A first observation concerning $m_2$ is that if all aggregations were sum operations and the element-wise latent space difference was the absolute value of a difference operation, $m_2$ would be equivalent to computing the $L^1$-norm of the difference vector in latent space.

$$m_2^{sum}(\tilde{x}, \tilde{y}) = \sum_{i=1}^{b} |\tilde{x}_i - \tilde{y}_i|$$
Similarly, adding a square operation to the element-wise
distance in the latent space and computing the square root
at the very end leads to the $L^2$-norm of the latent space
difference vector. In the same way, it is possible to use any
$L^p$-norm with the corresponding operations.

$$m_2^{\text{sum}}(\bar{x}, \bar{y}) = \left( \sum_{i=1}^{b} |\bar{x}_i - \bar{y}_i|^p \right)^{\frac{1}{p}}$$

In both cases, this forms the metric induced by the corre-
sponding norm which by definition has all desired properties
(6), (7), (8), and (9). If we change all aggregation meth-
ods to a weighted average operation, each term in the sum
is multiplied by a weight $w_i$. This is even possible with
learned weights, as they are constant at evaluation time, if
they are clamped to be positive as described above. Now, $w_i$
can be attributed to both inputs by distributivity, meaning
each input is element-wise multiplied with a constant vector
before applying the metric, which leaves the metric prop-
ties untouched. The reason is that it is possible to define
new vectors in the same space, equal to the scaled inputs.
This renaming trivially provides the correct properties.

$$m_2^{\text{weighted}}(\bar{x}, \bar{y}) = \sum_{i=1}^{b} w_i |\bar{x}_i - \bar{y}_i|$$

Accordingly, doing the same with the $L^p$-norm idea is possible,
each $w_i$ just needs a suitable adjustment before distribu-
tivity can be applied, keeping the metric properties once
again.

$$m_2^{\text{weighted}}(\bar{x}, \bar{y}) = \left( \sum_{i=1}^{b} w_i |\bar{x}_i - \bar{y}_i|^p \right)^{\frac{1}{p}}$$

$$= \left( \sum_{i=1}^{b} w_i |\bar{x}_i - \bar{y}_i| \cdot |\bar{x}_i - \bar{y}_i| ... |\bar{x}_i - \bar{y}_i| \right)^{\frac{1}{p}}$$

$$= \left( \sum_{i=1}^{b} w_i^\frac{1}{p} |\bar{x}_i - \bar{y}_i| \cdot w_i^\frac{1}{p} |\bar{x}_i - \bar{y}_i| ... w_i^\frac{1}{p} |\bar{x}_i - \bar{y}_i| \right)^{\frac{1}{p}}$$

$$w_i > 0 \quad \left( \sum_{i=1}^{b} w_i^\frac{1}{p} |\bar{x}_i - \bar{y}_i|^p \right)^{\frac{1}{p}}$$

With these weighted terms for $m_2$, it is possible to describe
all used aggregations and latent space difference methods.
The proposed method deals with multiple higher order ten-
sors instead of a single vector, so the weights $w_i$ additionally
depend on constants like the direction of the aggregations
and their position in the latent space tensors. But it is easy to
see that mapping a higher order tensor to a vector and keep-
ing track of additional constants still retains all properties
in the same way. As a result, the described architecture by
design yields a pseudometric that is suitable for comparing
simulation data in a way that corresponds to our intuitive
understanding of distances.

**B. Architectures**

The following sections provide details regarding the archi-
tecture of the base network and some experimental design.

**B.1. Base Network Design**

Fig. 7 shows the architecture of the base network for the
$LSiM$ metric. Its purpose is extracting features from both
inputs of the Siamese architecture that are useful for the
further processing steps. To maximise the usefulness and
to avoid feature maps that show overly similar features,
the chosen kernel size and stride of the convolutions is
important. Starting with larger kernels and strides means
the network has a big receptive field and can consider simple,
low-level features in large regions of the input. For the two
following layers, the large strides are replaced by additional
MaxPool operations that serve a similar purpose and reduce
the spatial size of the feature maps.

For the three final layers only small convolution kernels
and strides are used, but the number of channels is signifi-
cantly larger than before. These deep features maps typically
contain high-level structures, which are most important to
distinguish complex changes in the inputs. Keeping the
number of trainable weights as low as possible was an im-
portant consideration for this design, to prevent overfitting
to certain simulations types and increase generality. We
explored a weight range by using the same architecture and
only scaling the number of feature maps in each layer. The
final design shown in Fig. 7 with about 0.62 million weights
worked best for our experiments.

In the following, we analyze the contributions of the per-
layer features of two different metric networks to highlight
differences in terms of how the features are utilized for
the distance estimation task. In Fig. 8 the learned feature
map aggregation weights of our $LSiM$ network show a very
similar mean and a small standard deviation throughout the
five layers. This means, all feature maps similarly contribute
to establishing the distances, and the aggregation just fine-
tunes the relative importance of each feature. In addition, all
features receive a weight greater than zero, and as a result
no feature is excluded from contributing to the final distance
value.

Employing a fixed pre-trained feature extractor on the other
hand shows a very different picture: Although the mean
across the different network layers is similar, the contribu-

We can form a series of different features independently of the spatial dimensions. The unit length normalization only happens in the channel dimension, so all following operations accumulate values along the main paper). Let \( G \) denote a 3\(^{rd} \) order feature tensor with dimensions \( (g_x, g_z, g_y) \) from one layer of the base network. We can form a series \( G_0, G_1, \ldots \) for every possible content of this tensor across our training samples (computed as a preprocessing step). Below, we evaluate three different normalization methods and also consider not normalizing at all, denoted by \( \text{norm}_{\text{none}}(G) = G \).

The normalization only happens in the channel dimension, so all following operations accumulate values along \( (g_x, g_y) \) while keeping \( g_z \) constant, i.e., are applied independently of the spatial dimensions. The unit length normalization proposed by Zhang et al.

\[
\text{norm}_{\text{unit}}(G) = \frac{G}{\|G\|_2}
\]

only considers the current sample. In this case \( \|G\|_2 \) is a 2\(^{nd} \) order tensor with the Euclidean norms of \( G \) along the channel dimension. Combined with summation as the aggregation operation in channel direction, this results in a cosine distance which only measures angles of the latent space vectors. Using a learned average instead means the angles are no longer uniform, but warped according to the importance of each feature (i.e. the resulting angle changes differently for the same amount of change in two separate features). Extending this idea to consider other training samples as well, leads to a global unit length normalization

\[
\text{norm}_{\text{global}}(G) = \frac{G}{\max(\|G_0\|_2, \|G_1\|_2, \ldots)}
\]

where the maximum Euclidean norm of all available samples is employed. As a result, not only the angle of the latent space vectors, but also their magnitude compared to the largest feature vector is available in the aggregation. This formulation is not really robust yet, because the largest feature vector could be an outlier w.r.t. the typical content.
Instead, we can consider the full feature vector as a normal distribution and transform it to a standard normal distribution with the proposed

\[ \text{norm}_{\text{dist.}}(G) = \frac{G - \text{mean}(\|G_0\|_2, \|G_1\|_2, \ldots)}{\text{std}(\|G_0\|_2, \|G_1\|_2, \ldots)}. \]

In addition to the angle, this formulation allows for a robust comparison of the magnitude of each feature vector in the global magnitude distribution.

**Figure 9.** Performance on our test data for the proposed approach (LSiM) and a smaller model (AlexNet\textsubscript{frozen}) using different normalizations.

Fig. 9 shows a comparison of these normalization methods on the combined test data. Using no normalization is detrimental in both cases as succeeding operations cannot reliably compare the features. Interestingly, the unit length normalization works best for the AlexNet\textsubscript{frozen} metric (similar to LPIPS from Zhang et al.) that only uses learned aggregation weights with a fixed AlexNet feature extractor. This observation allows for a conclusion about the features extracted by AlexNet. For the original task of image classification, the magnitude of a feature vector does not seem to carry information about the feature. Interpreting the length as part of the feature for our task in norm\textsubscript{global} and norm\textsubscript{dist.}, obviously harms the performance of the metric. Therefore, training the feature extractor such that the magnitude of the feature vectors bears some meaning should improve the results for the complex normalizations. The performance of our approach with a fully trained feature extractor in Fig. 9 shows exactly this behaviour: A more complex normalization directly yields better results since the features can be adapted to utilize it.

**B.3. Recursive "Meta-Metric"**

Since comparing the feature maps is a central operation of the proposed metric calculations, we experimented with replacing it with an existing CNN-based metric. In theory, this would allow for a recursive, arbitrarily deep network that repeatedly invokes itself: first, the deep representations of inputs are used, then the deep representations of the deep representations, etc. In practice, however, using more than one recursion step is currently not feasible due to increasing computational requirements in addition to vanishing gradients.

Fig. 10 shows how our computation method can be modified for a CNN-based latent space difference, instead of an element-wise operation. Here we employ LPIPS (Zhang et al., 2018). There are two main differences compared to proposed method: First, the LPIPS latent space difference creates single distance values for a pair of feature maps instead of a spatial feature difference. As a result, the following aggregation is a single learned average operation and spatial or layer aggregations are no longer necessary. We also performed experiments with a spatial LPIPS version here, but due to memory limitations these were not successful. Second, the convolution operations in LPIPS have a lower limit for spatial resolution, and some feature maps of our base network are quite small (see Fig. 7). Hence, we

**Figure 10.** Adjusted distance computation for a LPIPS-based latent space difference. To provide sufficiently large inputs for LPIPS, small feature maps are spatially enlarged with nearest neighbor interpolation. In addition, LPIPS creates scalar instead of spatial differences leading to a simplified aggregation.
up-scale the feature maps below the required spatial size of 32 × 32 using nearest neighbor interpolation.

On our combined test data, such a metric with a fully trained base network achieves a performance comparable to \texttt{AlexNet\textsubscript{random}} or \texttt{AlexNet\textsubscript{frozen}}.

**B.4. Optical Flow Metric**

In the following, we describe our approach to compute a metric via optical flow (OF). For an efficient OF evaluation we employed a pre-trained network (Ilg et al., 2016). From an OF network \( f : \mathbb{I} \times \mathbb{I} \rightarrow \mathbb{R}^{\text{max} \times \text{max} \times 2} \) with two inputs data fields \( x, y \in \mathbb{I} \), we get the flow vector field \( f^{xy}(i, j) = (f_1^{xy}(i, j), f_2^{xy}(i, j))^T \), where \( i \) and \( j \) denote the location and \( f_1 \) and \( f_2 \) denote the components of the flow vectors. In addition, we have a second flow field \( f^{yx}(i, j) \) computed from the reversed input ordering. We can now define a function \( m : \mathbb{I} \times \mathbb{I} \rightarrow [0, \infty) \):

\[
m(x, y) = \sum_{i=0}^{\text{max}} \sum_{j=0}^{\text{max}} \sqrt{(f_1^{xy}(i, j))^2 + (f_2^{xy}(i, j))^2} + \sqrt{(f_1^{yx}(i, j))^2 + (f_2^{yx}(i, j))^2}
\]

Intuitively, this function computes the sum over the magnitudes of all flow vectors in both vector fields. With this definition, it is obvious that \( m(x, y) \) fulfills the metric properties of non-negativity and symmetry (see Eq. (6) and (7)). Under the assumption that identical inputs create a zero flow field, a relaxed identity of indiscernibles holds as well (see Eq. (9)). Compared to the proposed approach there is no guarantee for the triangle inequality though, so \( m(x, y) \) only qualifies as a pseudo-semimetric.

Fig. 11 shows flow visualizations on data examples produced by FlowNet2. The metric works relatively well for inputs that are similar to the training data from FlowNet2, like the shape data example in the top row. For data that provides some outline, for instance the smoke simulation example in the middle row or also liquid data, the metric does not work as well but still provides a reasonable flow field. But for full spatial examples like from the Burger’s or Advection-Diffusion equation (see bottom row) the network is no longer able to produce meaningful flow fields. The results are often a very uniform flow with similar magnitude and direction.

**B.5. Non-Siamese Architecture**

To compute a metric without the Siamese architecture outlined above, we use a network structure with a single output, as shown in Fig. 12. Thus, instead of having two identically feature extractors and combining the feature maps, here the distance is directly predicted from the stacked inputs with a single network with about 1.24 million weights. After using the same feature extractor as described in Section B.1, the final set of feature maps is spatially reduced with an adaptive MaxPool operation. Next, the result is flattened and three consecutive fully connected layers process the data to form the final prediction. Here, the last activation function is a sigmoid instead of ReLU. The reason is that a ReLU
would clamp every negative intermediate value to a zero distance, while a sigmoid compresses the intermediate value to a small distance that is more meaningful than directly clamping it.

In terms of metric properties, this architecture only provides non-negativity (see Eq. (6)) due to the final sigmoid function. All other properties can not be guaranteed without further constraints. This is the main disadvantage of a non-Siamese network. These issues could be alleviated with specialized training data or by manually adding constraints to the model, e.g., to have some amount of symmetry (see Eq. (7)) and at least a weakened identity of indiscernibles (see Eq. (9)). However, compared to a Siamese network that guarantees them by design, these extensions are clearly sub-optimal. As a result of the missing properties, this network has significant problems with generalization. While it performs well on the training data, the performance noticeably deteriorates for several of the test data sets.

**B.6. Skip Connections in Base Network**

As explained above, our base network primarily serves as a feature extractor to produce activations that are employed to evaluate a learned metric. In many state-of-the-art methods, networks with skip connections are employed (Ronneberger et al., 2015; He et al., 2016; Huang et al., 2017), as experiments have shown that these connections help to preserve information from the inputs. In our case, the classification "output" of a network such as the AlexNet plays no actual role. Rather, the features extracted along the way are crucial. Hence, skip connections should not improve the inference task for our metrics. To verify that this is the case, we have included tests with a base network similar to the popular UNet architecture (Ronneberger et al., 2015). For our experiments, we kept the early layers closely in line with the feature extractors that worked well for the base network (see Section B.1). Only the layers in the decoder part have an increased spatial feature map size to accom-
moderate the skip connections. As expected, this network can be used to compute reliable metrics for the input data without negatively affecting the performance. However, as expected, the improvements of skip connections for regular inference tasks do not translate into improvements for the metric calculations.

C. Impact of Data Difficulty

![Figure 14](image)

Figure 14. Impact of increasing data difficulty for a reduced training data set. Evaluations on training data for $L^2$ and LPIPS, and the test performance of models trained with the different reduced data sets ($LSiM_{reduced}$) are shown.

We shed more light on the aspect of noise levels and data difficulty via six reduced data sets, that consist of a smaller amount of Smoke and Advection-Diffusion data with differently scaled noise strength values. Results are shown in Fig. 14. Increasing the noise level creates more difficult data as shown by the dotted and dashed plots representing the performance of the $L^2$ and LPIPS metric on each data set. Both roughly follow an exponentially decreasing function. Each point on the solid line plot is the test result of a reduced $LSiM$ model trained on the data set with the corresponding noise level. Apart from the data, the entire training setup was identical. This shows that the training process is very robust to the noise, as the result on the test data only slowly decreases for very high noise levels. Furthermore, small amounts of noise improve the generalization compared to the model that was trained without any noise. This is somewhat expected, as a model that never saw noisy data during training can not learn to extract features which are robust w.r.t. noise.

D. Data Set Details

In the following sections the generation of each used data set is described. For each figure showing data samples (consisting of a reference simulation and several variants with a single changing initial parameter), the leftmost image is the reference and the images to the right show the variants in order of increasing parameter change. For the figures 15, 16, 17, and 18 the first subfigure (a) demonstrates that medium and large scale characteristics behave very non-chaotic for simulations without any added noise. They are only included for illustrative purposes and are not used for training. The second and third subfigure (b) and (c) in each case show the training data of $LSiM$, where the large majority of data falls into the category (b) of normal samples that follow the generation ordering, even with more varying behaviour. Category (c) is a small fraction of the training data and the shown examples are specifically picked, worst case examples to show how the chaotic behaviour can sometimes override the ordering intended by the data generation. In some cases, category (d) is included to show how normal data samples from the test set differ from the training data.

D.1. Navier-Stokes Equations

These equations describe the general behaviour of fluids with respect to advection, viscosity, pressure, and mass conservation. Eq. (10) defines the conservation of momentum and Eq. (11) the conservation of mass inside the fluid.

\[
\frac{\partial u}{\partial t} + (u \cdot \nabla)u = -\frac{\nabla P}{\rho} + \nu \nabla^2 u + g \quad (10)
\]

\[
\nabla \cdot u = 0 \quad (11)
\]

In this context, $u$ is the velocity, $P$ is the pressure the fluid exerts, $\rho$ is the density of the fluid (usually assumed to be constant), $\nu$ is the kinematic viscosity coefficient that indicates the thickness of the fluid, and $g$ denotes the acceleration due to gravity. With this PDE three data sets were created using a smoke and a liquid solver. For all data, 2D simulations were run until a certain step and useful data fields were exported afterwards.

SMOKE

For the smoke data, a standard Eulerian fluid solver using a preconditioned pressure solver based on conjugate gradient and a Semi-Lagrangian advection scheme was employed.

The general setup for every smoke simulation consists of a rectangular smoke source at the bottom with a fixed additive noise pattern to provide smoke plumes with more details. Additionally, there is a downwards directed, spherical force field area above the source which divides the smoke in two major streams along it. We chose this solution over an actual obstacle in the simulation, in order to avoid overfitting to a clearly defined black obstacle area inside the smoke data. Once the simulation reaches a predefined time step, the density, pressure, and velocity field (separated by dimension) is exported and stored. Some example sequences can be found in Fig. 15.
Figure 15. Various smoke simulation examples using one component of the velocity (top rows), the density (middle rows), and the pressure field (bottom rows).

(a) Data samples generated without noise: tiny output changes following generation ordering

(b) Normal training data samples with noise: larger output changes but ordering still applies

(c) Outlier data samples: noise can override the generation ordering by chance
Figure 16. Several liquid simulation examples using the binary indicator flags (top rows), the extrapolated level set values (middle rows), and one component of the velocity field (bottom rows) for the training data and only the velocity field for the test data.
With this setup the following initial conditions were varied in isolation:

- Smoke buoyancy in x- and y-direction
- Strength of noise added to the velocity field
- Amount of force in x- and y-direction provided by the force field
- Orientation and size of the force field
- Position of the force field in x- and y-direction
- Position of the smoke source in x- and y-direction

Overall, 768 individual smoke sequences were used for training, and the validation set contains 192 sequences with different initialization seeds.

L I Q U I D

For the liquid data, a solver based on the fluid implicit particle (FLIP) method proposed by Zhu & Bridson was employed. It is a Eulerian-Lagrangian hybrid approach that replaces the Semi-Lagrangian advection scheme with particle based advection to achieve higher accuracy and prevent the loss of mass. Still, this method is not optimal as we experienced problems with mass loss, especially for larger noise values.

The simulation setup consists of a large breaking dam and several smaller liquid areas for more detailed splashes. After the dam hits the simulation boundary a large, single drop of liquid is created in the middle of the domain that hits the already moving liquid surface. Then, the extrapolated level set values, binary indicator flags, and the velocity field (separated by dimension) are saved, with some examples shown in Fig. 16. The list of varied parameters include:

- Radius of the liquid drop
- Position of the drop in x- and y-direction
- Amount of additional gravity force in x- and y-direction
- Strength of noise added to the velocity field

The liquid training set consists of 792 sequences and the validation set of 198 sequences with different random seeds.

D.2. Advection-Diffusion and Burger’s Equation

For these PDEs our solvers only discretized and solve the corresponding equation in 1D. Afterwards, the different time steps of the solution process are concatenated along a new dimension to form 2D data with one spatial and one time dimension.

A D V E C T I O N - D I F F U S I O N E Q U A T I O N

This equation describes how a passive quantity is transported inside a velocity field due to the processes of advection and diffusion. Eq. (12) is the simplified Advection-Diffusion equation with constant diffusivity and no sources or sinks.

\[
\frac{\partial d}{\partial t} = \nu \nabla^2 d - u \cdot \nabla d \tag{12}
\]

Here, \(d\) denotes the density, \(u\) is the velocity, and \(\nu\) is the kinematic viscosity (also known as diffusion coefficient) that determines the strength of the diffusion. Our solver employed a simple implicit time integration and a diffusion solver based on conjugate gradient without preconditioning. The initialization for the 1D fields of the simulations was created by overlaying multiple parameterized sine curves with random frequencies and magnitudes.

In addition, continuous forcing controlled by further parameterized sine curves was included in the simulations over time. In this case, the only initial conditions to vary are the forcing and initialization parameters of the sine curves and the strength of the added noise. From this PDE only the passive density field was used as shown in Fig. 17. Overall, 798 sequences are included in the training set and 190 sequences with a different random initialization in the validation set.

For the Advection-Diffusion test set the noise was instead added directly to the passive density field of the simulations. This results in 190 sequences with more small scale details, as shown in Fig. 17(d).

B U R G E R ’ S E Q U A T I O N

This equation is very similar to the Advection-Diffusion equation and describes how the velocity field itself changes due to diffusion and advection.

\[
\frac{\partial u}{\partial t} = \nu \nabla^2 u - u \cdot \nabla u \tag{13}
\]

Eq. (13) is known as the viscous form of the Burger’s equation that can develop shock waves, and again \(u\) is the velocity and \(\nu\) denotes the kinematic viscosity. Our solver for this PDE used a slightly different implicit time integration scheme, but the same diffusion solver as used for the Advection-Diffusion equation.

The simulation setup and parameters were also the same; the only difference is that the velocity field instead of the density is exported. As a consequence, the data in Fig. 18 looks relatively similar to the results from the Advection-Diffusion equation. The training set features 782 sequences, and the validation set contains 204 sequences with different random seeds.
(a) Data samples generated without noise: tiny output changes following generation ordering

(b) Normal training data samples with noise: larger output changes but ordering still applies

(c) Outlier data samples: noise can override the generation ordering by chance

(d) Data samples from test set with additional background noise

*Figure 17.* Various examples from the Advection-Diffusion equation using the density field.
Figure 18. Different simulation examples from the Burger’s equation using the velocity field.

Figure 19. Examples from the shapes data set using a field with only binary shape values (first row), shape values with additional noise (second row), smoothed shape values (third row), and smoothed values with additional noise (fourth row).
D.3. Other

The remaining data sets are not based on PDEs and thus not generated with the proposed method. The data is only used to test the generalization of the discussed metrics and not for training or validation. The Shapes test set contains 160 sequences, the Video test set consists 131 sequences, and the TID test set features 216 sequences.

**Shapes**

This data set tests if the metrics are able to track simple, moving geometric shapes. To create it, a straight path between two random points inside the domain is generated and a random shape is moved along this path in steps of equal distance. The size of the used shape depends on the distance between start and end point, such that a significant fraction of the shape overlaps between two consecutive steps. It is also ensured that no part of the shape leaves the domain at any step, by using a sufficiently big boundary area when generating the path.

With this method, multiple random shapes for a single data sample are produced and their paths can overlap, such that they occlude each other to provide an additional challenge. All shapes are moved in their parametric representation and only when exporting the data, they are discretized onto a fixed binary grid. To add more variations to this simple approach, we also apply them in a non-binary way with smoothed edges and include additive gaussian noise over the entire domain. Examples for the different exports can be seen in Fig. 19.

**Video**

For this data set, different publicly available video recordings were acquired and processed in three steps. First, videos with abrupt cuts, scene transitions or camera movements were discarded, and afterwards the footage was broken down into single frames. Then, each frame was resized to match the spatial size of our other data by linear interpolation. Since directly using consecutive frames is no challenge for any analyzed metric and all of them recovered the ordering almost perfectly, we achieved a more meaningful data set by skipping several intermediate frames. For the final data set, we defined the first frame of every video as the reference and subsequent frames in an interval step of ten frames as the increasingly different variations. Some data examples can be found in Fig. 20.

**TID2013**

This data set was created by Ponomarenko et al. and used without any further modifications. It consists of 25 reference images with 24 distortion types in five levels. As a result it is not directly comparable to our data sets, so it is excluded from the test set aggregations. The distortions focus on various types of noise, image compression and color changes. Fig. 21 contains examples from the data set.

D.4. Data Augmentation

To add more variation to our data, we include different data augmentation techniques. These augmentations help the network to become more invariant to typical data transformations. Examples for the augmented data can be found in Fig. 22. In order to compare to existing natural image feature extractors trained for RGB images, we focus on three channel inputs. As a first step, the scalar, i.e. grey-scale, data is converted to a three channel RGB input with a random color map (out of five fixed variations) or no color map at all by copying the data directly to each channel.

Next, the data is randomly flipped on either axis and rotated in increments of $90^\circ$ to provide robustness to rotations. The
Learning Similarity Metrics for Numerical Simulations

Figure 21. Examples from the TID2013 data set proposed by Ponomarenko et al.. Displayed are a change of contrast, three types of noise, denoising, jpg2000 compression, and two color quantizations (from left to right and top to bottom).

Figure 22. Augmented samples from the training sets in groups with Smoke, Advection-Diffusion Equation, Burger’s Equation, and Liquid data. The upper row in each group shows the same reference simulation, and the lower row contains variations with different ground truth distances (increasing from left to right).

rotated data fields are then cropped from their simulation size \(256 \times 256\) to a size of \(224 \times 224\) which is the typical input size for existing feature extractors. Finally, each input is normalized to a standard normal distribution. The mean and standard deviation are computed from all available training data without augmentations in a pre-processing step with an online algorithm from Welford. Note that each data sample gets a new augmentation every time it is used, and that the corresponding reference receives the identical transformations the keep comparability (see Fig. 22). For all validation and test data, only a bilinear interpolation to the correct input size and the final normalization is performed.

D.5. Hardware

Data generation, training, and metric evaluations were performed on a machine with an Intel i7-6850 (3.60Ghz) CPU and an NVIDIA GeForce GTX 1080 Ti GPU.

E. Real-World Data

Below we give details of the three data-sets used for the evaluation in Section 6.3 of the main paper.

E.1. ScalarFlow

The ScalarFlow data set from Eckert et al. contains 3D velocities of real-world scalar transport flows reconstructed from multiple camera perspectives. For our evaluation, we cropped the volumetric \(100 \times 178 \times 100\) grids to \(100 \times 160 \times 100\) such that they only contain the area of interest and convert them to 2D with two variants: either by using the center slice or by computing the mean along the z-dimension. Afterwards, the magnitudes of the velocity vectors are computed, linearly interpolated to \(256 \times 256\), and then normalized. Variations for each reconstructed plume are acquired by using frames in equal temporal intervals. We
employed the velocity field reconstructions from 30 plumes (with simulation IDs 0 – 29) for both compression methods. Fig. 23 shows some example sequences.

E.2. Johns Hopkins Turbulence Database

The Johns Hopkins Turbulence Database (JHTDB) (Perlman et al., 2007) features various data sets of 3D turbulent flow fields created with direct numerical simulations (DNS). Here, we used three forced isotropic turbulence data sets with different resolutions (isotropic1024coarse, isotropic1024fine, and isotropic4096), two channel flows with different Reynolds numbers (channel and channel-5200), the forced magneto-hydrodynamic isotropic turbulence data set (mhd1024), and the rotating stratified turbulence data set (rotstrat4096).

For the evaluation, five 256 × 256 reference slices in the x/y-plane from each of the seven data sets are used. The spatial and temporal position of each slice is randomized within the bounds of the corresponding simulation domain. We compute velocity magnitudes, and normalize the value range. Variants for each reference are created by gradually varying the x- and z-position of the slice in equal intervals. The temporal position of each slice is varied as well, if a sufficient amount of temporally resolved data is available (for isotropic1024coarse, isotropic1024fine, channel, and mhd1024). This leads to 90 sequences in total. Fig. 24 shows examples from six of the JHTDB data sets.

E.3. WeatherBench

The WeatherBench repository from Rasp et al. represents a collection of various weather measurements of different atmospheric quantities such as precipitation, cloud coverage, wind velocities, geopotential, and temperature. The data ranges from 1979 to 2018 with a fine temporal resolution and is stored on a Cartesian latitude-longitude grid of the earth. In certain subsets of the data, an additional dimension such as altitude or pressure levels are available. As all measurements are available as scalar fields, only a linear interpolation to the correct input size and a normalization was necessary in order to prepare the data. We used the low-resolution geopotential data set at 500hPa (i.e., at around 5.5km height) with a size of 32 × 64 yielding smoothly changing features when upsampling the data. In addition, the high-res temperature data with a size of 128 × 256 for small scale details was used. For the temperature field, we used the middle atmospheric pressure level at 850hPa corresponding to an altitude of 1.5km in our experiments.

To create sequences with variations for a single time step of the weather data we used frames in equal time intervals, similar to the ScalarFlow data. Due to the very fine temporal discretization of the data, we consider a temporal interval of two hours as the smallest interval step of one in Fig. 26. We sampled three random starting points in time from each of the 40 years of measurements, resulting in 120 sequences overall. Fig. 25 shows a collection of example sequences.

E.4. Detailed Results

For each of the variants explained in the previous sections, we create test sets with six different spatial and temporal intervals. Fig. 26 shows the combined correlation of the sequences for different interval spacing when evaluating LSiM. For the results in the main paper, all correlation values shown here are aggregated by data source via mean and standard deviation.

While our metric reliably recovers the increasing distances within the data set, the individual measurements exhibit
Figure 24. Data samples extracted from the Johns Hopkins Turbulence Database with a spatial or temporal interval of ten. From top to bottom: mhd1024 and isotropic1024coarse (varied time step), isotropic4096 and rotstrat4096 (varied z-position), channel and channel5200 (varied x-position).

Figure 25. Examples of the processed WeatherBench data: high-res temperature data 1.40625deg/temperature (upper two rows) and low-res geopotential data 5.625deg/geopotential_500 (lower two rows). The temporal interval spacing between the images is ten.

Figure 26. Detailed breakdown of the results when evaluating LSiM on the individual data sets of ScalarFlow (30 sequences each), JHTDB (90 sequences each), and WeatherBench (120 sequences each) with different step intervals.
interesting differences in terms of their behavior for varying
distances. As JHTDB and WeatherBench contain relatively
uniform phenomena, a larger step interval creates more
difficult data as the simulated and measured states contain
changes that are more and more difficult to analyze along a
sequence. For ScalarFlow, on the other hand, the difficulty
decreases for larger intervals due to the large-scale motion of
the reconstructed plumes. As a result of buoyancy forces the
observed smoke rises upwards into areas where no smoke
has been before. For the network this makes predictions
relatively easy, as the large-scale translations are indicative
of the temporal progression, and small scale turbulence
effects can be largely ignored. For this data set smaller
intervals are more difficult as the overall shape of the plume
barely changes while the complex evolution of small scale
features becomes more important.

Overall, the LSiM metric recovers the ground truth ordering
of the sequences very well as indicated by the consistently
high correlation values in Fig. 26.

F. Additional Evaluations

In the following, we demonstrate other ways to compare the
performance of the analyzed metrics on our data sets. In
Tab. 2 the Pearson correlation coefficient is used instead of
Spearman’s rank correlation coefficient. While Spearman’s
correlation measures monotonic relationships by using rank-
ing variables, it directly measures linear relationships.

The results in Tab. 2 match very closely to the values com-
puted with Spearman’s rank correlation coefficient. The
best performing metrics in both tables are identical, only
the numbers slightly vary. Since a linear and a monotonic
relation describes the results of the metrics similarly well,
there are no apparent non-linear dependencies that can not
be captured using the Pearson correlation.

In the Tables 4 and 3 we employ a different, more intuitive
approach to determine combined correlation values for each
data set using the Pearson correlation. We no longer analyz-
ing the entire predicted distance distribution and the ground
truth distribution at once as done above. Instead, we individ-
ually compute the correlation between the ground truth and
the predicted distances for the single data samples of the
data set. From the single correlation values, we compute the
mean and standard deviations shown in the tables. Note that
this approach potentially produces less accurate compar-
ison results, as small errors in the individual computations
can accumulate to larger deviations in mean and standard
deviation. Still, both tables lead to very similar conclusions:
The best performing metrics are almost the same and low
combined correlation values match with results that have a
high standard deviation and a low mean.

Fig. 27 shows a visualization of predicted distances $c$ against
ground truth distances $d$ for different metrics on every sam-
ple from the test sets. Each plot contains over 6700 individ-
ual data points to illustrate the global distance distributions
created by the metrics, without focusing on single cases.
A theoretical optimal metric would recover a perfectly nar-
row distribution along the line $c = d$, while worse metrics
recover broader, more curved distributions. Overall, the
sample distribution of an $L^2$ metric is very wide. LPIPS
manages to follow the optimal diagonal a lot better, but our
approach approximates it with the smallest deviations, as
also shown in the tables above. The $L^2$ metric performs
very poorly on the shape data indicated by the too steeply

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Table 2. Performance comparison on validation and test data sets measured in terms of the Pearson correlation coefficient. **Bold** values show the best performing metric for each data set and **bold+italic** values are within a 0.01 error margin of the best performing. On the right a visualization of the combined test data results is shown for selected models.

| Metric               | Validation data sets | Test data sets |
|----------------------|----------------------|----------------|
|                      | Smo   | Liq   | Adv   | Bur   | TID   | LiqN | AdvD | Sha   | Vid   | All   |
| $L^2$                | 0.66  | 0.81  | 0.71  | 0.58  | **0.85** | 0.72  | 0.57  | 0.55  | 0.76  | 0.56  |
| SSIM                 | 0.67  | 0.74  | 0.75  | 0.68  | 0.83  | 0.25  | **0.69** | 0.34  | 0.66  | 0.47  |
| LPIPS v0.1           | **0.71** | 0.75  | **0.76** | **0.72** | 0.79  | 0.63  | 0.61  | 0.82  | **0.80** | 0.65  |
| AlexNet*random       | 0.63  | 0.75  | 0.66  | 0.64  | 0.80  | 0.63  | 0.65  | 0.68  | 0.77  | 0.60  |
| AlexNet*frozen       | 0.67  | 0.70  | 0.68  | 0.70  | 0.79  | 0.40  | 0.63  | 0.84  | **0.80** | 0.61  |
| Optical flow         | 0.63  | 0.56  | 0.37  | 0.39  | 0.49  | 0.45  | 0.28  | 0.61  | 0.74  | 0.48  |
| Non-Siamese          | **0.71** | **0.82** | 0.75  | 0.69  | 0.26  | 0.72  | 0.62  | 0.65  | 0.68  | 0.63  |
| Siamese scratch      | 0.65  | **0.83** | 0.74  | 0.66  | 0.72  | 0.78  | 0.59  | 0.83  | 0.78  | **0.70** |
| LSiM*noiseless       | 0.64  | **0.82** | 0.74  | 0.60  | 0.69  | **0.80** | 0.58  | 0.83  | 0.75  | 0.68  |
| LSiM*strong noise    | 0.63  | 0.81  | 0.71  | 0.61  | 0.70  | 0.78  | 0.50  | 0.80  | 0.76  | 0.64  |
| LSiM (ours)          | 0.68  | **0.82** | **0.76** | 0.70  | 0.71  | **0.79** | 0.61  | **0.86** | 0.76  | **0.71** |
Table 3. Performance comparison on validation data sets measured by computing mean and standard deviation (in brackets) of Pearson correlation coefficients from individual data samples. **Bold** values show the best performing metric for each data set and **bold+italic** values are within a 0.01 error margin of the best performing. On the right a visualization of the combined test data results is shown for selected models.

| Metric              | Validation data sets |
|---------------------|----------------------|
|                     | Smo | Liq | Adv | Bur |
| $L^2$               |     |     |     |     |
| SSIM                |     |     |     |     |
| LPIPS v0.1          | **0.77 (0.28)** | **0.79 (0.24)** | **0.81 (0.26)** | **0.77 (0.32)** |
| AlexNet\_random     | **0.68 (0.36)** | **0.79 (0.28)** | **0.71 (0.36)** | **0.69 (0.36)** |
| AlexNet\_frozen     | **0.72 (0.31)** | **0.74 (0.29)** | **0.73 (0.35)** | **0.75 (0.33)** |
| Optical flow        | **0.66 (0.38)** | **0.59 (0.47)** | **0.38 (0.52)** | **0.41 (0.49)** |
| Non-Siamese         | **0.76 (0.27)** | **0.87 (0.19)** | **0.80 (0.24)** | **0.75 (0.33)** |
| Skip from scratch   | **0.69 (0.34)** | **0.87 (0.19)** | **0.79 (0.26)** | **0.72 (0.34)** |
| LSiM\_noiseless     | **0.68 (0.33)** | **0.85 (0.24)** | **0.78 (0.30)** | **0.66 (0.37)** |
| LSiM\_strong noise  | **0.67 (0.36)** | **0.85 (0.22)** | **0.76 (0.33)** | **0.67 (0.39)** |
| LSiM (ours)         | **0.72 (0.31)** | **0.85 (0.22)** | **0.81 (0.23)** | **0.75 (0.33)** |

Table 4. Performance comparison on test data sets measured by computing mean and std. dev. (in brackets) of Pearson correlation coefficients from individual data samples. **Bold** values show the best performing metric for each data set and **bold+italic** values are within a 0.01 error margin of the best performing.

| Metric              | Test data sets |
|---------------------|---------------|
|                     | TID | LiqN | AdvD | Sha | Vid | All |
| $L^2$               | **0.98 (0.19)** |   |     |     |     |     |
| SSIM                | **0.92 (0.40)** |   | **0.73 (0.36)** |     |     |     |
| LPIPS v0.1          | **0.94 (0.34)** | **0.66 (0.38)** | **0.65 (0.41)** | **0.83 (0.24)** | **0.85 (0.30)** | **0.74 (0.36)** |
| AlexNet\_random     | **0.96 (0.27)** | **0.68 (0.34)** | **0.69 (0.38)** | **0.68 (0.26)** | **0.82 (0.33)** | **0.71 (0.34)** |
| AlexNet\_frozen     | **0.94 (0.33)** | **0.41 (0.49)** | **0.67 (0.39)** | **0.85 (0.21)** | **0.85 (0.29)** | **0.70 (0.40)** |
| Optical flow        | **0.74 (0.67)** | **0.50 (0.34)** | **0.32 (0.53)** | **0.63 (0.45)** | **0.78 (0.45)** | **0.53 (0.49)** |
| Non-Siamese         | **0.47 (0.88)** | **0.76 (0.24)** | **0.66 (0.41)** | **0.67 (0.28)** | **0.76 (0.42)** | **0.71 (0.35)** |
| Skip from scratch   | **0.99 (0.14)** | **0.85 (0.15)** | **0.61 (0.42)** | **0.84 (0.23)** | **0.82 (0.33)** | **0.76 (0.33)** |
| LSiM\_noiseless     | **0.98 (0.19)** | **0.86 (0.15)** | **0.61 (0.41)** | **0.84 (0.26)** | **0.79 (0.38)** | **0.76 (0.34)** |
| LSiM\_strong noise  | **0.99 (0.14)** | **0.83 (0.19)** | **0.52 (0.45)** | **0.81 (0.23)** | **0.82 (0.35)** | **0.73 (0.36)** |
| LSiM (ours)         | **0.97 (0.23)** | **0.83 (0.22)** | **0.64 (0.42)** | **0.86 (0.23)** | **0.80 (0.37)** | **0.77 (0.34)** |

Figure 27. Distribution evaluation of ground truth distances against normalized predicted distances for $L^2$, LPIPS and LSiM on all test data (color coded).
increasing blue lines that flatten after a ground truth distance of 0.3. \textit{LPIPS} already significantly reduces this problem, but \textit{LSiM} still works slightly better.

A similar issue is visible for the Advection-Diffusion data, where for \( L^2 \) a larger number of red samples is below the optimal \( c = d \) line, than for the other metrics. \textit{LPIPS} has the worst overall performance for liquid test set, indicated by the large number of fairly chaotic green lines in the plot. On the video data, all three metrics perform similarly well.

A fine-grained distance evaluation in 200 steps of \( L^2 \) and our \textit{LSiM} metric via the mean and standard deviation of different data samples is shown in Fig. 28. Similar to Fig. 27, the mean of an optimal metric would follow the ground truth line with a standard deviation of zero, while the mean of worse metrics deviates around the line with a high standard deviation. The plot on the left combines eight samples with different seeds from the \textit{Sha} data set, where only a single shape is used. Similarly, the center plot aggregates eight samples from \textit{Sha} with more than one shape. The right plot shows six data samples from the \textit{LiQ} test set that vary by the amount of noise that was injected into the simulation.

The task of only tracking a single shape in the example on the left is the easiest of the three shown cases. Both metrics have no problem to recover the position change until a variation of 0.4, where \( L^2 \) can no longer distinguish between the different samples. Our metric recovers distances with a continuously rising mean and a very low standard deviation. The task in the middle is already harder, as multiple shapes can occlude each other during the position changes. Starting at a position variation of 0.4 both metrics have a quite high standard deviation, but the proposed method stays closer to the ground truth line. \( L^2 \) shows a similar issue as before, because it flattens relatively fast. The plot on the right features the hardest task. Here, both metrics perform similar as each has a different problem in addition to an unstable mean. Our metric stays close to the ground truth, but has a quite high standard deviation starting at about a variation of 0.4.

The standard deviation of \( L^2 \) is lower, but instead it starts off with a big jump from the first few data points. To some degree this is caused by the normalization of the plots, but it still overestimates the relative distances for small variations in the simulation parameter.

These findings also match with the distance distribution evaluations in Fig. 27 and the tables above: Our method has a significant advantage over shallow metrics on shape data, while the differences of both metrics become much smaller for the liquid test set.

\section{G. Notation}

In this work, we follow the notation suggested by Goodfellow et al. Vector quantities are displayed in bold and tensors use a sans-serif font. Double-barred letters indicate sets or vector spaces. The following symbols are used:

| \( \mathbb{R} \) | Real numbers |
| \( i, j \) | Indexing in different contexts |
| \( \mathbb{I} \) | Input space of the metric, i.e., color images/field data of size \( 224 \times 224 \times 3 \) |
| \( a \) | Dimension of the input space \( \mathbb{I} \) when flattened to a single vector |
| \( x, y, z \) | Elements in the input space \( \mathbb{I} \) |
| \( \mathbb{L} \) | Latent space of the metric, i.e., sets of \( 3^{rd} \) order feature map tensors |
| \( b \) | Dimension of the latent space \( \mathbb{L} \) when flattened to a single vector |
| \( x, y, z \) | Elements in the latent space \( \mathbb{L} \), corresponding to \( x, y, z \) |
| Symbol | Description |
|--------|-------------|
| $w$    | Weights for the learned average aggregation (1 per feature map) |
| $p_0, p_1, \ldots$ | Initial conditions / parameters of a numerical simulation |
| $n$    | Number of steps when varying a simulation parameter, thus length of the network input sequence |
| $o_0, o_1, \ldots, o_n$ | Series of outputs of a simulation with increasing ground truth distance to $o_0$ |
| $\Delta$ | Amount of change in a single simulation parameter |
| $t_1, t_2, \ldots, t_t$ | Time steps of a numerical simulation |
| $s$    | Strength of the noise added to a simulation |
| $c$    | Ground truth distance distribution, determined by the data generation via $\Delta$ |
| $d$    | Predicted distance distribution (supposed to match the corresponding $c$) |
| $\bar{c}, \bar{d}$ | Mean of the distributions $c$ and $d$ |
| $\| \cdot \|_2$ | Euclidean norm of a vector |
| $m(x, y)$ | Entire function computed by our metric |
| $m_1(x, y)$ | First part of $m(x, y)$, i.e., base network and feature map normalization |
| $m_2(\tilde{x}, \tilde{y})$ | Second part of $m(x, y)$, i.e., latent space difference and the aggregations |
| $G$    | 3rd order feature tensor from one layer of the base network |
| $g_c, g_x, g_y$ | Channel dimension ($g_c$) and spatial dimensions ($g_x, g_y$) of $G$ |
| $f$    | Optical flow network |
| $f^{xy}, f^{yx}$ | Flow fields computed by an optical flow network $f$ from two inputs in $\mathbb{I}$ |
| $f_1^{xy}, f_2^{xy}$ | Components of the flow field $f^{xy}$ |
| $\nabla, \nabla^2$ | Gradient ($\nabla$) and Laplace operator ($\nabla^2$) |
| $\partial$ | Partial derivative operator |
| $t$    | Time in our PDEs |
| $u$    | Velocity in our PDEs |
| $\nu$  | Kinematic viscosity / diffusion coefficient in our PDEs |
| $d, \rho$ | Density in our PDEs |
| $P$    | Pressure in the Navier-Stokes Equations |
| $g$    | Gravity in the Navier-Stokes Equations |