# **Adaptive Deep Kernel Learning**

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#### **Abstract**

Deep kernel learning provides an elegant and principled framework for combining the structural properties of deep learning algorithms with the flexibility of kernel methods. By means of a deep neural network, it consists of learning a kernel operator which is combined with a differentiable kernel algorithm for inference. While previous work within this framework has mostly explored learning a single kernel for large datasets, we focus herein on learning a kernel family for a variety of tasks in few-shot regression settings. Compared to single deep kernel learning, our novel algorithm permits finding the appropriate kernel for each task during inference, rather than using the same for all tasks. As such, our algorithm performs more effectively with complex task distributions in few-shot learning, which we demonstrate by benchmarking against existing state-of-the-art algorithms using real-world, few-shot regression tasks related to drug discovery.

# 1 Introduction

Recent studies exploring few-shot learning have led to the creation of new algorithms that learn efficiently with very small samples and generalize beyond the training data [1, 2]. Most of these algorithms have adopted the meta-learning paradigm [3, 4], where some prior knowledge is learned across a large collection of diverse tasks and then transferred to new tasks for efficient learning with a limited amount of data. Despite many few-shot learning algorithms reporting improved performance over the state-of-the-art, considerable progress must still be made before such approaches can be adopted on a larger scale and in more practical settings. Notably, much of the work in this field has focused on classification and reinforcement learning [1], leaving the problem of few-shot regression largely unaddressed [5–7]. To the best of our knowledge, no real-world few-shot regression (FSR) benchmarks have been established in the literature. FSR methods, however, show great promise in addressing important problems encountered in fields plagued by small sample sizes such as drug discovery, where data acquisition is expensive and time consuming, limiting the amount of examples available to each study.

Meta-learning-based few-shot algorithms differ on two crucial aspects: the nature of the meta-knowledge captured and the amount of adaptation performed at test-time for new tasks/datasets. First, metric learning methods [8–12] accumulate meta-knowledge in high capacity covariance/distance functions and then combine them with simple base learners to produce the outputs. However, they do not adapt these covariance functions at test-time. Hence, few of the currently-used base learners have enough capacity to truly adapt [12, 13]. Second, initialization and optimization based methods [14, 6, 15] that learn the initialization point for gradient descent algorithms allow for more adaptation

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on new tasks, but remain time consuming and memory inefficient. To ensure optimal performance on FSR problems, it is crucial to combine the strengths of both types of methods.

In this study, we frame FSR as a deep kernel learning (DKL) problem, opposed to one in metric learning, allowing us to derive new algorithms. DKL methods combine the non-parametric flexibility of kernel methods with the structural properties of deep neural networks, which yields a more powerful way of learning input covariance functions and more adaptation capacity at test-time. We further improve over the general DKL algorithm for FSR by learning a family of covariance functions instead of a single one, resulting in greater adaptability. Our method selects an appropriate covariance function from this family at test-time, which makes it as adaptive as optimization and initialization-based methods.

Our Contributions: We first frame few-shot regression as a deep kernel learning problem and show why and how it is more expressive than classical metric learning methods. Next, we derive two DKL algorithms by combining set embedding techniques [16] and kernel methods [17, 18] to learn a family of kernels, allowing more adaptation at test-time while being sample efficient. We then propose two new real-world datasets for FSR, drawn from the drug discovery domain. Performance on these datasets as well as synthetic data shows that our model allows greater test adjustment than classical methods.

### 2 Preliminaries

In this section, we describe the DKL framework (introduced by Wilson et al. [19]) in more depth and show that it can be adapted to learn a covariance/kernel function for few-shot learning tasks.

**DKL Framework:** Let  $D^t_{trn} = \{(\mathbf{x}_i, y_i)\}_{i=1}^m \subset \mathcal{X} \times \mathbb{R}$ , a training dataset available for learning the regression task t ( $\mathcal{X}$  is the input space and  $\mathbb{R}$  is the output space). A DKL algorithm aims to obtain a non-linear embedding of inputs in the embedding space  $\mathcal{H}$ , using a deep neural network  $\phi_{\boldsymbol{\theta}}: \mathcal{X} \to \mathcal{H}$  of parameters  $\boldsymbol{\theta}$ . Then, it finds the minimal norm regressor  $h^t_*$  in the reproducing kernel Hilbert space (RKHS)  $\mathcal{R}$  on  $\mathcal{H}$ , that fits the training data, i.e.

$$h_*^t := \underset{h \in \mathcal{R}}{\operatorname{argmin}} \lambda \|h\|_{\mathcal{R}} + \ell(h, \mathcal{D}_{trn}^t)$$
 (1)

where  $\ell$  is a non-negative loss function that measures the loss of a regressor h; and  $\lambda$  weighs the importance of the norm minimization against the training loss. Following the representer theorem [17, 20],  $h_*^t$  can be written as a finite linear combination of kernel evaluations on training inputs, i.e.

$$h_*^t(\mathbf{x}) = \sum_{(\mathbf{x}_i, y_i) \in D_{trn}^t} \alpha_i^t k_{\rho}(\phi_{\theta}(\mathbf{x}), \phi_{\theta}(\mathbf{x}_i)), \tag{2}$$

where  $\alpha^t = (\alpha_1^t, \dots, \alpha_m^t)$  are the combination weights; and  $k_{\rho} \colon \mathcal{H} \times \mathcal{H} \to \mathbb{R}_+$  is a reproducing kernel of  $\mathcal{R}$  with hyperparameters  $\rho$ . Candidates include the radial basis, polynomial, and linear kernels. Depending on the loss function  $\ell$ , the weights  $\alpha^t$  can be obtained by using a differentiable kernel method enabling to compute the gradients of the loss w.r.t. the parameters  $\theta$ . Such methods include Gaussian Process (GP), Kernel Ridge Regression (KRR), and Logistic Regression (LR).

DKL inherits benefits and drawbacks from deep learning and kernel methods. It follows that gradient descent algorithms are required to optimize  $\theta$  which can be high dimensional such that seeing a significant amount of training samples is thus essential to avoid overfitting. However, once the latter condition is met, scalability of the kernel method becomes limiting as the running time of kernel methods scales approximately in  $O(m^3)$  for a training set of m samples. Some approximations of the kernel [21, 22] are thus needed for the scalability of the DKL method.

Few-Shot DKL: In the setup of episodic meta learning, also known as few-shot learning, one has access to a meta-training collection  $\mathscr{D}_{meta-trn} := \left\{ (D_{trn}^{t_j}, D_{val}^{t_j}) \right\}_{j=1}^T$ , of T tasks to learn how to learn from few datapoints. Each task  $t_j$  has its own training (or support) set  $D_{trn}^{t_j}$  and validation (or query) set  $D_{val}^{t_j}$ . A meta-testing collection  $\mathscr{D}_{meta-tst}$  is also available to assess the generalization performances of the few-shot algorithm across unseen tasks. To learn a few-shot DKL method for FSR in such settings, one can share the parameters of  $\phi_{\theta}$  across all tasks, similar to metric learning

algorithms. Hence, for a given task  $t_j$ , the inputs are first transformed by the function  $\phi_{\theta}$  and then a kernel method is used to obtain the regressor  $h_*^{t_j}$ , which will be evaluated on  $D_{val}^{t_j}$ .

**KRR:** Using the squared loss and the L2-norm to compute  $||h||_{\mathcal{R}}$ , KRR gives the optimal regressor for a task t and its validation loss  $\mathcal{L}_{\theta,\rho,\lambda}^t$  as follows:

$$h_*^t(\mathbf{x}) = \alpha K_{\mathbf{x},trn}, \quad \text{with} \quad \alpha = (K_{trn,trn} + \lambda I)^{-1} \mathbf{y}_{trn}$$
 (3)

$$\mathcal{L}_{\theta,\rho,\lambda}^{t} = \underset{\mathbf{x},y \sim D_{val}^{t}}{\mathbf{E}} (\alpha K_{\mathbf{x},trn} - y)^{2}, \tag{4}$$

where  $K_{trn,trn}$  is the matrix of kernel evaluations and each entry  $K_{trn,trn:il} = k_{\rho}(\phi_{\theta}(\mathbf{x}_i),\phi_{\theta}(\mathbf{x}_l))$  for  $(\mathbf{x}_i,\cdot),(\mathbf{x}_l,\cdot)\in D_{trn}^{t_j}$ . An equivalent definition applies to  $K_{\mathbf{x},trn}$ .

**GP:** Using the negative log likelihood loss function instead, the GP algorithm gives a probabilistic regressor for which predictive mean is given by Equation 3 and the loss for a task t is:

$$\mathcal{L}_{\boldsymbol{\theta}, \boldsymbol{\rho}, \lambda}^{t} = -\ln \mathcal{N}(\mathbf{y}_{val}; \mathbb{E}[h_{*}^{t}], \operatorname{cov}(h_{*}^{t})), \tag{5}$$

$$\mathbb{E}[h_*^t] = K_{val,trn}(K_{trn,trn} + \lambda I)^{-1} \mathbf{y}_{trn}, \tag{6}$$

$$cov(h_*^t) = K_{val,val} - K_{val,trn}(K_{trn,trn} + \lambda I)^{-1}K_{trn,val}$$
(7)

Finally, the parameters  $\theta$  of the neural network, along with  $\lambda$  and the hyperparameters  $\rho$  of the kernel, are optimized using the expected loss on all tasks:

$$\underset{\boldsymbol{\theta}, \boldsymbol{\rho}, \lambda}{\operatorname{argmin}} \underset{t \sim \mathcal{D}_{meta-trn}}{\mathbf{E}} \mathcal{L}_{\boldsymbol{\theta}, \boldsymbol{\rho}, \lambda}^{t}. \tag{8}$$

For tractability of this expectation, we use a Monte-Carlo approximation with b tasks. Unless otherwise specified, we use b=32 and  $m:=|D_{trn}^{t_j}|=10$ , yielding a mini-batch of 320 samples. In our experiments, we have fixed  $\lambda$  to  $1/|D_{trn}^{t_j}|$ .

To summarize, this algorithm finds a representation common to all tasks such that the kernel method (in our case, GP and KRR) will generalize well from a small amount of samples. Interestingly, this alleviates two of the main limitations of single task DKL: i) the scalability of the kernel method is no longer an issue since we are in the few-shot learning regime<sup>2</sup>, and ii) the parameters  $\theta$  (and  $\rho$ ,  $\lambda$ ) are learned across a potentially large amount of tasks and samples, providing the opportunity to learn a complex representation without overfitting.

It is worth mentioning that using the linear kernel and the KRR algorithm, we recover the few-shot classification algorithm R2-D2 proposed by Bertinetto et al. [12]. Their intent was to show that KRR can be used for fast adaptation at test-time in classification settings as it is differentiable. In contrast, our intent is to formalize and adapt the DKL framework to FSR and justify how this powerful combination of kernel methods and deep networks can learn covariance functions.

#### 3 Proposed Method

#### 3.1 Adaptive Deep Kernel Method

As described earlier, a DKL algorithm for FSR learns a fixed kernel function  $k_{DKL}(\mathbf{x},\mathbf{x}') := k_{\rho}(\phi_{\theta}(\mathbf{x}),\phi_{\theta}(\mathbf{x}'))$  shared across all tasks of interest. While a regressor on a given task can obtain an arbitrarily small loss as the training set size increases, a fixed kernel might not learn well in the few-shot regime. To verify this hypothesis, we propose an adaptive deep kernel learning (ADKL) algorithm illustrated by Figure 1. It learns an adaptive, task-dependent, kernel for few-shot learning instead of a single fixed kernel. We define the adaptive kernel as follows:

$$k_{ADKL}(\mathbf{x}, \mathbf{x}'; \mathbf{z}_t) := k_{\rho}(\phi_{\theta}(\mathbf{x}; \mathbf{z}_t), \phi_{\theta}(\mathbf{x}'; \mathbf{z}_t))$$
(9)

<sup>&</sup>lt;sup>2</sup>Even with several hundred samples, the computational cost of embedding each example is usually higher than inverting the Gram matrix.

where  $\mathbf{z}_t = \psi_{\eta}(D_{trn}^t)$  represents a task embedding obtained by transforming the training set  $D_{trn}^t$  with the task encoding network  $\psi_{\eta}$ . We now describe in more detail the task encoding network  $\psi_{\eta}$  and the architecture of the adapted kernel for a given task t.

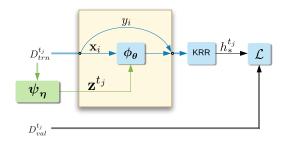


Figure 1: The ADKL algorithm. The green component corresponds to the adaptive part of the algorithm, which contrasts it with standard deep kernel learning approaches.

**Task Encoding** The challenge of the  $\psi_{\eta}$  network is to capture complex dependencies in the training set  $D^t_{trn}$  to provide a useful task encoding  $\mathbf{z}$ . Furthermore, the task encoder should be invariant to permutations of the training set and be able to encode a variable amount of samples. After exploring a variety of architectures, we found that more complex ones such as Transformers [23] tend to underperform. This is possibly due to overfitting or the sensitivity of training such architectures.

Consequently, we introduce slight modifications to DeepSets, an order invariant network proposed by Zaheer et al. [16]. It begins with the computation of the representation of each input-target pair  $\mathbf{x}\mathbf{y}_i = \mathbf{r}(Concat(\mathbf{u}(\mathbf{x}_i), \mathbf{v}(y_i)))$  for all  $(\mathbf{x}_i, y_i) \in D^t_{trn}$ , using neural networks  $\mathbf{u}, \mathbf{v}$ , and  $\mathbf{r}$ . The  $\mathbf{x}\mathbf{y}_i$  captures nonlinear interactions between the inputs and the targets if  $\mathbf{r}$  is a nonlinear transformation. Then, by computing  $\mu^t_{\mathbf{x}\mathbf{y}}$  and  $\sigma^t_{\mathbf{x}\mathbf{y}}$ , the empirical mean and standard deviation of the set  $\{\mathbf{x}\mathbf{y}_1, \mathbf{x}\mathbf{y}_2, \dots, \mathbf{x}\mathbf{y}_m\}$ , respectively, we obtain the task representation as follows:

$$\mathbf{z}^{t} = \boldsymbol{\psi}_{\boldsymbol{\eta}}(D_{trn}^{t}) := \mathbf{w}(Concat(\boldsymbol{\mu}_{\mathbf{x}\mathbf{y}}^{t}, \boldsymbol{\sigma}_{\mathbf{x}\mathbf{y}}^{t})), \tag{10}$$

where  ${\bf w}$  is a also a neural network. As  $\mu^t_{{\bf x}{\bf y}}$  and  $\sigma^t_{{\bf x}{\bf y}}$  are invariant to permutations in  $D^t_{trn}$ , it follows that  $\psi_{\eta}$  is also permutation invariant. Overall,  $\psi_{\eta}$  is just a nonlinear mapping of the first and second moment of the sample representations which were also nonlinear transformations of the original inputs and targets. The learnable parameters  $\eta$  of the task encoder include all the parameters of the networks  ${\bf u}, {\bf v}, {\bf r}$ , and  ${\bf w}$ , and are shared across all tasks.

**Adapted Kernel Computation** Once the task representation is obtained, we compute the conditional input embedding using the function  $\phi_{\theta}$ . Let  $\mathbf{u}'(\mathbf{x})$  be the non-conditional embedding of the input  $\mathbf{x}$  using a neural network  $\mathbf{u}'$ , whose parameters are shared with the network  $\mathbf{u}$  within the task encoder. We simply compute the conditional embedding of inputs as:

$$\phi_{\theta}(\mathbf{x}; \mathbf{z}_t) = \mathbf{o}(Concat(\mathbf{u}'(\mathbf{x}), \mathbf{z}_t)), \tag{11}$$

where  $\mathbf{o}$  is a nonlinear neural network that allows for capturing complex interactions between the task and the input representations. The adapted kernel for a given task is then obtained by combining Equations 9, 10, 11. The learnable parameters of  $\mathbf{o}$  and  $\mathbf{u}'$  together constitute  $\boldsymbol{\theta}$  and are shared across all tasks. Alternatively, different architectures such as Feature-wise Linear Modulation (FiLM) [24], Hypernetwork [25], or Bilinear transformation [26] could be used to compute  $\phi_{\boldsymbol{\theta}}(\mathbf{x}; \mathbf{z})$ , though we found that a simple concatenation was sufficient for our applications.

**Kernel Methods** To find the regressor that minimizes Equation 1 for each task t, we use GP and KRR described in Section 2. In the following, we use ADKL-GP and ADKL-KRR to refer to our algorithm when the task-level regressor is given by GP and KRR, respectively.

# 3.2 Meta-Regularization

To help the training of the ADKL algorithm, we maximize the mutual information between  $D_{trn}^{t_j}$  and  $D_{val}^{t_j}$ . This serves as a regularizer that helps the encoder  $\psi_{\eta}$  learn a useful representation for

describing the task. This is done using the MINE algorithm [27], which optimizes a lower bound on the mutual information. For two random variables  $r, s \sim p(r, s)$  and a similarity measure  $f_{\phi}$  between r and s, parameterized by  $\phi$ , the following inequality holds:

$$I[r,s] \ge \max_{\phi} \underset{r,s \sim p(r,s)}{\mathbf{E}} f_{\phi}(r,s) - \ln \underset{r \sim p(r)}{\mathbf{E}} \underset{s \sim p(s)}{\mathbf{E}} e^{f_{\phi}(r,s)}. \tag{12}$$

Using a mini-batch approximation of the expectations<sup>3</sup>, and the cosine distance c as the similarity measure between the two sets, this yields  $I[D_{trn}, D_{val}] \ge \tilde{I}_n$ , where

$$\tilde{I}_{\eta} \stackrel{\text{def}}{=} \frac{1}{b} \sum_{j=1}^{b} c(\psi_{\eta}(D_{trn}^{t_{j}}), \psi_{\eta}(D_{val}^{t_{j}})) - \ln \frac{1}{b(b-1)} \sum_{j=1}^{b} \sum_{i \neq j} e^{c(\psi_{\eta}(D_{trn}^{t_{j}}), \psi_{\eta}(D_{val}^{t_{i}}))}.$$
(13)

When adding this to our training objective with  $\gamma \geq 0$  as a tradeoff hyperparameter, we have:

$$\underset{\boldsymbol{\theta}, \boldsymbol{\eta}, \boldsymbol{\rho}, \lambda}{\operatorname{argmin}} \underset{t_j \sim B}{\mathbf{E}} \mathcal{L}_{\boldsymbol{\theta}, \boldsymbol{\eta}, \boldsymbol{\rho}, \lambda}^{t_j} - \gamma \tilde{I}_{\boldsymbol{\eta}}. \tag{14}$$

### 4 Related Work

Our study spans the research areas of deep kernel learning and few-shot learning. For a comprehensive overview of few-shot learning methods, we refer the reader to Wang and Yao [1], Chen et al. [2], as we focus on work related to DKL herein.

Across the spectrum of learning approaches, DKL methods lie between neural networks and kernel methods. While neural networks can learn from a very large amount of data without much prior knowledge, kernel methods learn from fewer data when given an appropriate covariance function that accounts for prior knowledge of the relevant task. In the first DKL attempt, Wilson et al. [19] combined GP with CNN to learn a covariance function adapted to a task from large amounts of data, though the large time and space complexity of kernel methods forced the approximation of the exact kernel using KISS-GP [22]. Dasgupta et al. [28] have demonstrated that such approximation is not necessary using finite rank kernels. Here, we also show that learning from a collection of tasks (FSR mode) does not require any approximation when the covariance function is shared across tasks. This is an important distinction between our study and other existing studies in DKL, which learn their kernel from single tasks instead of task collections.

On the spectrum between NNs and kernel methods, metric learning also bears mention. Metric learning algorithms learn an input covariance function shared across tasks, but rely only on the expressive power of DNNs. First, stochastic kernels are built out of shared feature extractors, simple pairwise metrics (e.g. cosine similarity [9], Euclidean distance [10]), or parametric functions (e.g. relation modules [29], graph neural networks [11]). Then, within tasks, the predictions consist of a distance-weighted combination of the training sample labels with the stochastic kernel evaluations no adaptation is done. The recently introduced Proto-MAML [13] method, which captures the best of Prototypical Networks [10] and MAML [14], allows within-task adaptation using MAML on a network built on top of the kernel function. Similarly, Kim et al. [6] have proposed a Bayesian version of MAML where a feature extractor is shared across tasks, while multiple MAML particles are used for the task-level adaptation. Bertinetto et al. [12] have also tackled this lack of adaptation for new tasks by using KRR and Logistic Regression to find the appropriate weighting of the training samples. This study can be considered the first application of DKL to few-shot learning. However, its contribution was limited to showing that simple differentiable learning algorithms can increase adaptation in the metric learning framework. Our work extends beyond by formalizing few-shot learning in the deep kernel learning framework where test-time adaptation is achieved through kernel methods. We also create another layer of adaptation by allowing task-specific kernels that are created at test-time.

Since ADKL-GP uses GPs, it has relations to neural processes [30], which proposes a scalable alternative to learning regression functions by performing inference on stochastic processes. Furthermore, in this family of methods, Conditional Neural Processes (CNP) [31] and Attentive Neural Processes

<sup>&</sup>lt;sup>3</sup>This yields a small bias on the gradient since the right hand side takes the log of the expectations. Since we are not interested in the precise value of the mutual information, this does not constitute a problem.

(ANP) [32] are even more relevant to our study as both methods learn conditional stochastic processes parameterized by conditions derived from training data points. While ANP imposes consistency with respect to some prior process, CNP does not and thus does not have the mathematical guarantees associated with stochastic processes. By comparison, our proposed ADKL-GP algorithm also learns conditional stochastic processes, but within the GP framework, thus benefiting from the associated mathematical guarantees.

## 5 Experiments

#### 5.1 Datasets

Previous work on few-shot regression has relied on toy datasets to evaluate performance. We instead introduce two real-world benchmarks drawn from the field of drug discovery. These benchmarks will allow us to measure the ability of few-shot learning algorithms to adapt in settings where tasks require a considerably different measure of similarity between the inputs. For instance, when predicting binding affinities between small molecules, the covariance function must learn characteristics of a binding site that changes from task to task. We describe each dataset used in our experiments below and, unless stated otherwise, their meta-training, meta-validation, and meta-testing contain, respectively, 56.25%, 18.75% and 25% of all of their tasks. A pre-processed version of these datasets is available with this work (URL to appear in camera-ready).

- 1. **Sinusoids**: This meta-dataset was recently proposed by Kim et al. [6] as a challenging few-shot synthetic regression benchmark. It consists of 5,000 tasks defined by a sinusoidal functions of the form:  $y = A \sin(wx + b) + \epsilon$ . The parameters A, w, b characterize each task and are drawn from the following intervals:  $A \in [0.1, 5.0], b \in [0.0, 2\pi], w \in [0.5, 2.0]$ . Samples for each task are generated by sampling inputs  $x \in [-5.0, 5.0]$  and observational noise from  $\epsilon \sim N(0, (0.01A)^2)$ . Every model on this task collection uses a fully connected network of 2 layers of 120 hidden units as its input feature extractor. Visuals of ground truths and predictions for some functions are shown in Appendix 3.
- 2. **Binding**: The goal here is to predict the binding affinity of small, drug-like molecules to proteins. This task collection was extracted from the public database BindingDB<sup>4</sup> and encompasses 7,620 tasks, each containing between 7 and 9,000 samples. Each task corresponds to a protein sequence, which thus defines a separate distribution on the input space of molecules and the output space of (real-valued) binding readouts.
- 3. **Antibacterial**: The goal here is to predict the antimicrobial activity of small molecules for various bacteria. The task collection was extracted from the public database PubChem<sup>5</sup> and contains 3,842 tasks, each consisting of 5 to 225 samples. A task corresponds to a combination of a bacterial strain and an experimental setting, which define different data distributions.

For both real-world datasets, the molecules are represented by their SMILES<sup>6</sup> encoding, which are descriptions of the molecular structure using short ASCII strings. All models evaluated on these collections share the same input feature extractor configuration: a 1-D CNN of 2 layers of 128 hidden units each and a kernel size of 5. We use CNN instead of LSTM or advanced graph convolutions methods for scalability reasons. Moreover, the targets were scaled linearly between 0 and 1.

# 5.2 Benchmarking analysis

We evaluate model performance against R2-D2 [12], CNP[31], and MAML[14]. R2-D2 is a natural comparison to ADKL-KRR (when the latter uses the linear kernel) to show whether the adapted deep kernel provides more test-time adaptation. CNP is also a natural comparison to ADKL-GP and will help measure performance differences between the task-level Bayesian models generated within the GP and CNP frameworks. MAML is considered herein for its fast-adaptation at test-time and as the representative of initialization and optimization based models. In the following experiments, all DKL methods use the linear kernel.

<sup>&</sup>lt;sup>4</sup>Original data available at www.bindingdb.org

<sup>&</sup>lt;sup>5</sup>Available at https://pubchem.ncbi.nlm.nih.gov/

 $<sup>^6\</sup>mathrm{See}\ \mathrm{https://en.wikipedia.org/wiki/Simplified\_molecular-input\_line-entry\_system.}$ 

Our first set of experiments evaluates performance on both the real-world and toy tasks. We train each method using support and query sets of size m=10. During meta-testing, the support set size is also m=10, but the query set consists of the remaining samples of each task. For the datasets lacking sufficient samples (the Binding and Antibacterial collections), we use half of the samples in the support set and the remaining in the query set. For each task, during meta-testing, we average the Mean Squared Error (MSE) over 20 random partitions of the query and support sets. We refer to this value as the task MSE. Figure 2 illustrates the task MSE distributions over tasks for each collection and algorithm (the best hyperparameters were chosen on the meta-validation set). In general, we observe that the real-world datasets are challenging for all methods but ADKL methods consistently outperform R2-D2 and CNP. The gap between ADKL-KRR and R2-D2 shows the importance of adapting the kernel to each task rather than sharing a single kernel. Furthermore, that ADKL-GP outperforms CNP shows the effectiveness of the ADKL approach in comparison with conditional neural processes. Finally, both adaptive deep kernel methods (ADKL-KRR and ADKL-GP) seem to perform comparably, despite different objective functions.

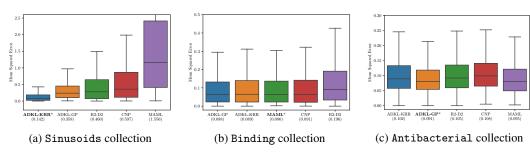


Figure 2: MSE distributions for tasks in the meta-testing sets. Only the best model for each dataset is marked with an asterisk and the mean MSE for the whole meta-test is shown below the algorithm name.

A second experimental set is used to measure the across-task generalization and within-task adaptation capabilities of our methods relative to others. We do so by controlling the number of training tasks ( $T \in \{100, 1000\}$ ) and the size of the support sets during meta-training and meta-testing ( $m \in \{5, 10, 15, 20, 25, 30\}$ ). Only the Sinusoids collection was used, as experiments with the real-world collections were deemed too time-consuming for the scope of this study. One would expect the algorithms to generalize poorly to new tasks for lower values of T, and their task-level models to adapt poorly to new samples for small values of m. However, as illustrated in Figure 3, all DKL methods generalize better across tasks than others, as their overall performance is robust against the number of training tasks. They also demonstrate improved within-task generalization using as few as 15 samples, while other methods require more samples to achieve the same. Moreover, for small support sets, ADKL-KRR shows better within-task generalization than ADKL-GP and R2-D2. Once again, the difference in performance between ADKL-KRR and R2-D2 can be attributed to the kernel adaptation at test-time as it is the only difference between both methods. This difference for small m between ADKL-GP and ADKL-KRR can be attributed to larger predictive uncertainty in GP as the number samples gets smaller.

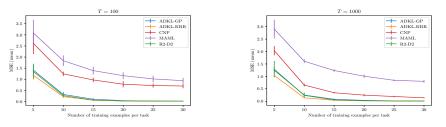


Figure 3: Average MSE over the testing tasks of the Sinusoids collection. The error bars represent the uncertainty on the average over 5 runs with different seeds.

#### 5.3 Active Learning

Here we report the results of active learning experiments. Our intent is to measure the effectiveness of the uncertainty captured by the predictive distribution of ADKL-GP for active learning. CNP, in comparison, serves to measure which of CNP and GP better captures the data uncertainty for improving FSR under active sample selection. For this purpose, we meta-train both algorithms using support and query sets of size m=5 (and  $T\in\{100,1000\}$  for the Sinusoids collection). During meta-test time, five samples are randomly selected to constitute the support set  $D_{trn}$  and build the initial hypothesis for each task. Then, from a pool U of unlabeled data, we choose the input  $\mathbf{x}^*$  of maximum predictive entropy, i.e.,  $\mathbf{x}^* = \operatorname{argmax}_{\mathbf{x} \in U} \mathbb{E}\left[\log p(y|\mathbf{x}, D_{trn})\right]$ . The latter is removed from U and added to  $D_{trn}$  with its predicted label. The within-task adaptation is performed on the new support set to obtain a new hypothesis which is evaluated on the query set  $D_{val}$  of the task. This process is repeated until we reach the allowed budget of 20 queries.

Figure 4 highlights that, in the active learning setting, ADKL-GP consistently outperforms CNP. Very few samples are queried by ADKL-GP to capture the data distribution, while CNP performance is far from optimal, even when allowed the maximum number of queries. Also, since using the maximum predictive entropy strategy is better than querying samples at random for ADKL-GP (solid vs. dashed line), these results suggest that the predictive uncertainty obtained with GP is informative and more accurate than that of CNP. Moreover, when the number of queries is greater than 10, we observe a performance degradation for CNP, while ADKL-GP remains consistent. This observation highlights the generalization capacity of DKL methods, even outside the few-shot regime where they have been trained — this same property does not hold true for CNP. We attribute this property of DKL methods to their use of kernel methods. In fact, their role in adaptation and generalization increases as we move away from the few-shot training regime.

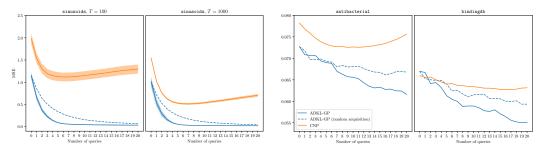


Figure 4: Average MSE performance on the meta-test during active learning. The width of the shaded regions denotes the uncertainty over five runs for the sinusoidal collection. No uncertainty is shown for the real-world tasks as they were too time consuming.

#### 5.4 Regularization and Kernel Impact

In our final set of experiments, we take a closer look at the impact of the base kernel and the metaregularization factor on the generalization during meta-testing. We do so by evaluating ADKL-KRR on the Sinusoids collection with different hyperparameter combinations. Figure 5 (left) shows the performances obtained by varying the meta-regularization parameter  $\gamma$  over different hyperparameter configurations (listed in Appendix 1). Looking at  $\gamma \in \{0, 0.1, 0.01\}$ , one can observe that non-zero values help in most cases, meaning that our added regularizer slightly improves the task encoder learning as intended. We also measure how different base kernels impact the learning process, as choosing the appropriate kernel function is crucial for kernel methods. We test the linear and RBF kernels and their normalized versions, where the normalized version of a kernel k is given by:  $k'(\mathbf{x}, \mathbf{x}') := k(\mathbf{x}, \mathbf{x}') / \sqrt{k(\mathbf{x}, \mathbf{x})k(\mathbf{x}', \mathbf{x}')}$ . Over different hyperparameter combinations, one observes that the linear kernel yields better generalization performances than the RBF kernel (see Figure 5 right). Such a result is within expectation as the scaling parameter of the RBF kernel is shared across tasks, making it more difficult to adapt the deep kernel. We could explore learning the base kernel hyperparameters using a network similar to  $\psi$  in future work. It is also worth nothing that although the kernel normalization impacts outcomes, there is no clear conclusion to be drawn. We therefore advise treating the kernel function and its normalization as hyperparameters of the DKL methods.

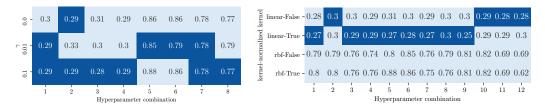


Figure 5: Impact of  $\gamma$  and the kernel on the average meta-test MSE for different hyperparameter combinations (dark blue is best). The combinations are detailed in the Appendix 1, 2

### 6 Conclusion

In this paper, we investigate the benefits of DKL methods for FSR. By comparing methods on both real-world and toy task collections, we have demonstrated the effectiveness of the DKL framework in FSR. Both ADKL-GP and ADKL-KRR outperform the single kernel DKL method, providing evidence that they add more adaptation capacity at test-time through adaptation of the kernel. Given its Bayesian nature, ADKL-GP also allows for improvement of the learned models at test-time, providing great value in settings such as drug discovery. By making our drug discovery task collections publicly available, we hope that the community will leverage these advances to propose FSR algorithms that are ready to be deployed in real-life settings, in turn having a positive impact on the drug discovery process.

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# 7 Appendix

# A Regularization impact

Table 1 presents the hyperparameter combinations used in the experiments to assess the impact of the joint training parameter  $\gamma$ .

Table 1: Effect of using joint training (parameter  $\gamma$ ) on the MSE performance

			<u> </u>	U 1/			
				$\gamma$	0.0	0.01	0.1
s	Run	kernel	hidden size (target FE)	hidden size (input FE)			
	1	linear	[32, 32]	[128, 128, 128]	0.297	0.290	0.290
	2	linear	[32, 32]	[128, 128]	0.288	0.328	0.288
	3	linear		[128, 128, 128]	0.306	0.300	0.285
	4	linear		[128, 128]	0.290	0.297	0.287
	5	rbf	[32, 32]	[128, 128, 128]	0.864	0.851	0.876
	6	rbf	[32, 32]	[128, 128]	0.859	0.787	0.864
	7	rbf		[128, 128, 128]	0.780	0.778	0.778
_	8	rbf		[128, 128]	0.769	0.793	0.768

Note that the performance is significantly worse when using RBF kernel.

# **B** Kernel impact

Table 2 shows the hyperparameter combinations we used to assess the effect of using different kernels, as well as the impact of normalizing them.

Table 2: MSE performance depending on the kernel choice and normalization

Run	encoder arch	target FE	kernel normalized input FE	Linear False	True	RBF False	True
1	CNP	[32, 32]	[128, 128, 128]	0.279	0.274	0.786	0.795
2	CNP	[32, 32]	[128, 128]	0.299	0.300	0.789	0.799
3	CNP	[]	[128, 128, 128]	0.304	0.289	0.761	0.755
4	CNP	[]	[128, 128]	0.295	0.293	0.742	0.757
5	DeepSet	[32, 32]	[128, 128, 128]	0.313	0.269	0.804	0.877
6	DeepSet	[32, 32]	[128, 128]	0.301	0.277	0.849	0.856
7	DeepSet	[]	[128, 128, 128]	0.292	0.273	0.764	0.754
8	DeepSet	[]	[128, 128]	0.303	0.298	0.788	0.763
9	KRR	[32, 32]	[128, 128, 128]	0.296	0.246	0.815	0.815
10	KRR	[32, 32]	[128, 128]	0.289	0.290	0.824	0.824
11	KRR	[]	[128, 128, 128]	0.279	0.291	0.690	0.694
12	KRR	[]	[128, 128]	0.275	0.299	0.685	0.622

### C Prediction curves on the Sinusoids collection

Figure C.6 presents a visualization of the results obtained by each model on three tasks taken from the meta-test set. We provide the model with ten examples from an unseen task consisting of a slightly noisy sine function (shown in blue), and present in orange the the approximation made by the network based on these ten examples.

Note that contrary to others, CNP and ADKL-GP give us access to the uncertainty.

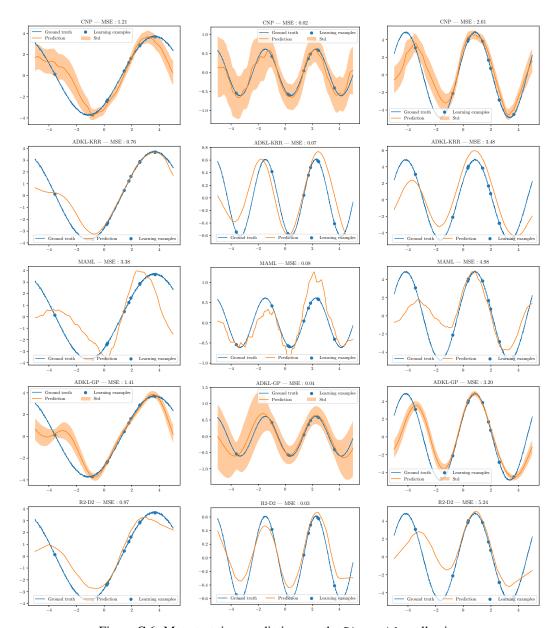


Figure C.6: Meta-test time predictions on the Sinusoids collection