



We consider a standard **multi-task Bayesian Optimization** (BO) framework, where one aims to optimize an expensive blackbox function $f_{T+1} : \mathcal{X} \to \mathbb{R}$ with a minimal number of function evaluations:

 $\mathbf{x}_{T+1}^{\star} = \operatorname{argmin}_{\mathbf{x} \in \mathcal{X}} f_{T+1}(\mathbf{x}),$

where $\mathcal{X} \subset \mathbb{R}^D$ denotes the configuration space. We assume that we have already **completed** *T* **related Hyperpa**rameter Optimization (HPO) tasks $\{f_1, \ldots, f_T\}$ that share the same configuration space \mathcal{X} and we have access to $\mathcal{D} = \{ D_t : D_t = \{ (\mathbf{x}_{tn}, y_{tn}) \}_{n=1}^{N_t} \}_{t=1}^{T},$ the data collected while optimizing the set of black-box functions

 ${f_t}_{t=1}^T, y_{tn} = f_t(\mathbf{x}_{tn}).$

Contributions

We resolve a deficiency of popular Adaptive Bayesian Linear Regression (ABLR) [1] applied to multi-task BO, which is that **the number** of non-linear basis functions is not adapted to the target task, where the number of observations is typically smaller than in the previous tasks, making it prone to overfitting. This issue is due to the direct use of conventional multi-task models in the context of sequential decision making problems such as BO. To solve this issue:

- We use **nested dropout** [2] to learn an **ordered set of features for transfer learning** in the context of BO. To the best of our knowledge, no other multi-task learning approach proposed in the literature is able to learn features that take the adaptive complexity of the target task into account.
- We use Automatic Relevance Determination (ARD) to automatically determine which basis functions to activate at transfer in a data-driven fashion. Hence, the resulting transfer learning model is able to adapt its capacity to the amount of data available in the target task.
- We show that we can improve the sample efficiency of multi-task BO and **avoid overfitting** in low data regimes without hurting the transfer learning performance in high data regimes.

Hyperparameter Transfer Learning with Adaptive Complexity

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ABLR with Adaptive Complexity (ABRAC)

Algorithm 1 ABRAC

- 1: Input: number of initial points n_0 , budget N, feature net $\phi_z(\cdot)$: $\mathbb{R}^P \to \mathbb{R}^d$ parametrized by z, filter F^k , previous evaluations $\{\{(x_{ti}, y_{ti})\}_{i=1}^{N_t}\}_{t=1}^T.$
- 2: Fit $\phi_{\mathbf{z}}(\cdot)$ using $\{\{(x_{ti}, y_{ti})\}_{i=1}^{N_t}\}_{t=1}^T$ using random truncation (nested dropout) in the final layer.
- 3: Observe f_{T+1} at n_0 randomly selected points $x_1, x_2, \ldots, x_{n_0} \in \mathcal{X}$.
- 4: $\mathcal{C} = \{(x_i, y_i)\}_{i=1}^{n_0}$, where $y_i = f_{T+1}(x_i)$.
- 5: Set $n = n_0$.
- 6: while n < N do
- 7: Fit probabilistic model g via ARD.
- 8: $x_n = \operatorname{argmax}_{x \in \mathcal{X}} A_q(x)$, where A is a given acquisition function.
- 9: **Observe** $y_n = f_{T+1}(x_n)$.
- 10: Update $\mathcal{C} \leftarrow \mathcal{C} \cup \{(x_n, y_n)\}, n \leftarrow n+1$
- 11: end while
- 12: Output: $\hat{x} = \operatorname{argmin}_{i=1,2,...,N} f_{T+1}(x_i)$

Fit Visualization on Forrester Functions



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Figure: Tabular benchmarks by Klein and Hutter (2019). Top: Protein Structure, bottom: Slice Localization.

- Scalable hyperparameter transfer learning. In NeurIPS, 2018.
- [2] Oren Rippel, Michael Gelbart, and Ryan Adams. Learning ordered representations with nested dropout. In *ICML*, 2014.





References

[1] Valerio Perrone, Rodolphe Jenatton, Matthias Seeger, and Cédric Archambeau.