

SIMBA: SIMPLICITY BIAS FOR SCALING UP PARAMETERS IN DEEP REINFORCEMENT LEARNING

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ABSTRACT

Recent advances in CV and NLP have been largely driven by scaling up the number of network parameters, despite traditional theories suggesting that larger networks are prone to overfitting. These large networks avoid overfitting by integrating components that induce a *simplicity bias*, guiding models toward simple and generalizable solutions. However, in deep RL, designing and scaling up networks have been less explored. Motivated by this opportunity, we present *SimBa*, an architecture designed to scale up parameters in deep RL by injecting a simplicity bias. *SimBa* consists of three components: (i) an observation normalization layer that standardizes inputs with running statistics, (ii) a residual feedforward block to provide a linear pathway from the input to output, and (iii) a layer normalization to control feature magnitudes. By scaling up parameters with *SimBa*, the sample efficiency of various deep RL algorithms—including off-policy, on-policy, and unsupervised methods—is consistently improved. Moreover, solely by integrating *SimBa* architecture into SAC, it matches or surpasses state-of-the-art deep RL methods with high computational efficiency across DMC, MyoSuite, and HumanoidBench. These results demonstrate *SimBa*'s broad applicability and effectiveness across diverse RL algorithms and environments.

Explore videos at <https://anonymous-simba.github.io>

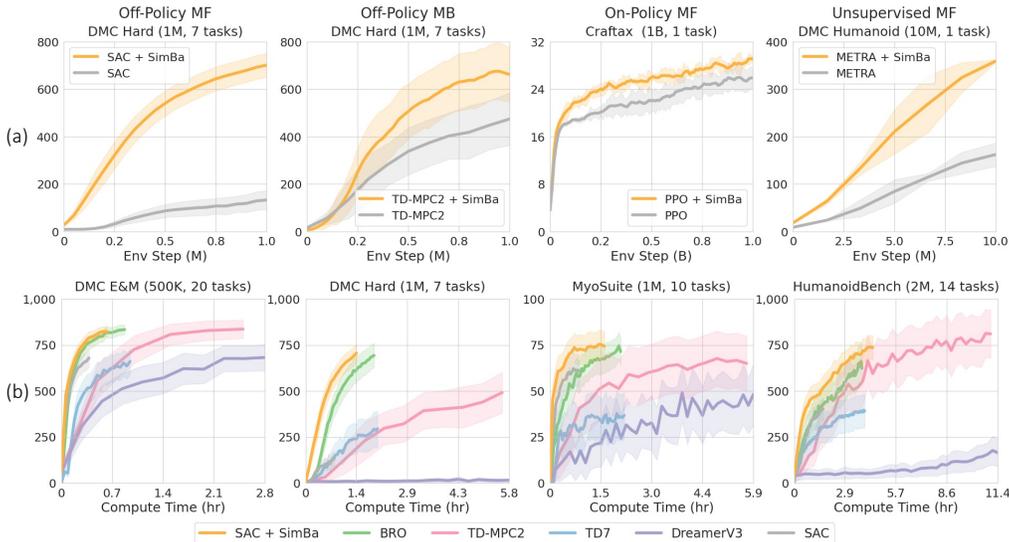


Figure 1: **Benchmark Summary.** *SimBa* infuses *simplicity bias* through architectural changes, without modifying the underlying deep RL algorithm. (a) **Sample Efficiency:** *SimBa* improves sample efficiency across various RL algorithms, including off-policy model-free (SAC), off-policy model-based (TD-MPC2), on-policy model-free (PPO), and unsupervised (METRA) RL. (b) **Compute Efficiency:** When applying *SimBa* with SAC, it matches or surpasses state-of-the-art off-policy RL methods with minimal computational overhead across 51 continuous control tasks, by only modifying the network architecture and scaling up the number of network parameters.

1 INTRODUCTION

Scaling up neural network sizes has been a key driver of recent advancements in computer vision (CV) (Dehghani et al., 2023) and natural language processing (NLP) (Google et al., 2023; Achiam et al., 2023). By increasing the number of parameters, neural networks gain enhanced expressivity, enabling them to cover diverse functions and discover effective solutions that smaller networks might miss. However, this increased capacity also heightens the risk of overfitting, as larger networks can fit intricate patterns in the training data that do not generalize well to unseen data.

Despite this risk of overfitting, empirical evidence shows that neural networks tend to converge toward simpler functions that generalize effectively (Kaplan et al., 2020; Nakkiran et al., 2021). This phenomenon is attributed to a *simplicity bias* inherent in neural networks, where standard optimization algorithms and architectural components guide highly expressive models toward solutions representing simple, generalizable functions (Shah et al., 2020; Berchenko, 2024). For instance, gradient noise in stochastic gradient descent prevents models from converging on sharp local minima, helping them avoid overfitting (Chizat & Bach, 2020; Gunasekar et al., 2018; Pesme et al., 2021). Architectural components such as ReLU activations (Hermann et al., 2024), layer normalization (Ba et al., 2016), and residual connections (He et al., 2020) are also known to amplify simplicity bias. These components influence the types of functions that neural networks represent at initialization, where networks that represent simpler function at initialization are more likely to converge to simple functions (Valle-Perez et al., 2018; Mingard et al., 2019; Teney et al., 2024).

While scaling up network parameters and leveraging simplicity bias have been successfully applied in CV and NLP, these principles have been underexplored in deep reinforcement learning (RL), where the focus has primarily been on algorithmic advancements (Hessel et al., 2018; Hafner et al., 2023; Fujimoto et al., 2023; Hansen et al., 2023). Motivated by this opportunity, we introduce the *SimBa* network, a novel architecture that explicitly embeds simplicity bias to effectively scale up parameters in deep RL. SimBa comprises of three key components: (i) an observation normalization layer that standardizes inputs by tracking the mean and variance of each dimension, reducing overfitting to high-variance features (Andrychowicz et al., 2020); (ii) a pre-layer normalization residual feedforward block (Xiong et al., 2020), which maintains a direct linear information pathway from input to output and applies non-linearity only when necessary; and (iii) a post-layer normalization before the output layer to stabilize activations, ensuring more reliable policy and value predictions.

To verify whether SimBa amplifies simplicity bias, we compared it against the standard MLP architecture often employed in deep RL. Following Teney et al. (2024), we measured simplicity bias by (i) sampling random inputs from a uniform distribution; (ii) generating network outputs; and (iii) performing Fourier decomposition on these outputs. A smaller sum of the Fourier coefficients indicates that the neural network represents a low-frequency function, signifying greater simplicity. We define the simplicity score as the inverse of this sum, meaning a higher score corresponds to a stronger simplicity bias. As illustrated in Figure 2.(a), our analysis revealed that SimBa has a higher simplicity score than the MLP (Further details are provided in Section 2).

To evaluate the benefit of leveraging simplicity bias on network scaling, we compared the performance of Soft Actor-Critic (Haarnoja et al., 2018, SAC) using both MLP and our SimBa architecture across 3 humanoid tasks from DMC benchmark (Tassa et al., 2018). We increased the width of both actor and critic networks, scaling from 0.1 to 17 million parameters. As shown in Figure 2.(b), SAC with MLP experiences performance degradation as the number of parameters increases. In contrast, SAC with the SimBa network consistently improves its performance as the number of parameter increases, highlighting the value of embedding simplicity bias when scaling deep RL networks.

To further evaluate SimBa’s versatility, we applied it to various RL algorithms by only changing the network architecture and scaling up parameters. The algorithms included off-policy (SAC (Haarnoja et al., 2018), TD-MPC2 (Hansen et al., 2023)), on-policy model-free (PPO (Schulman et al., 2017)),

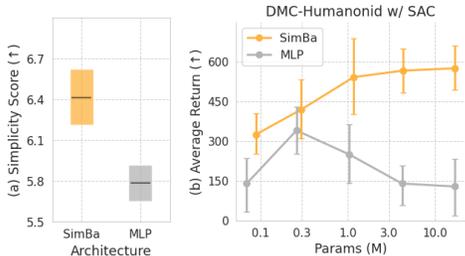


Figure 2: (a) SimBa exhibits higher simplicity bias than MLP. (b) SAC with SimBa improves its performance with increased parameters, whereas SAC with MLP degrades it. Each standard deviation is 95% CI.

and unsupervised RL (METRA (Park et al., 2023)). As illustrated in Figure 1.(a), SimBa consistently enhances the sample efficiency of these algorithms. Furthermore, as shown in Figure 1.(b), when SimBa is integrated into SAC, it matches or surpasses state-of-the-art off-policy methods across 51 tasks in DMC, MyoSuite (Caggiano et al., 2022), and HumanoidBench (Sferrazza et al., 2024). Despite the increased number of parameters, SAC with SimBa remains computationally efficient because it does not employ any computationally intensive components such as self-supervised objectives (Fujimoto et al., 2023), planning (Hansen et al., 2023), or replay ratio scaling (Nauman et al., 2024), which state-of-the-art methods rely on to achieve high performance.

2 PRELIMINARY

Simplicity bias refers to the tendency of neural networks to prioritize learning simpler patterns over capturing intricate details (Shah et al., 2020; Berchenko, 2024). In this section, we introduce metrics to quantify simplicity bias; in-depth definitions are provided in Appendix A.

2.1 MEASURING FUNCTION COMPLEXITY

We begin by defining a **complexity measure** $c : \mathcal{F} \rightarrow [0, \infty)$ to quantify the complexity of functions in $\mathcal{F} = \{f|f : \mathcal{X} \rightarrow \mathcal{Y}\}$ where $\mathcal{X} \subseteq \mathbb{R}^n$ denote the input space and $\mathcal{Y} \subseteq \mathbb{R}^m$ the output space.

Traditional complexity measures, such as the Vapnik–Chervonenkis dimension (Blumer et al., 1989) and Rademacher complexity (Bartlett & Mendelson, 2002), are well-established but often intractable for deep neural networks. Therefore, we follow Teney et al. (2024) and adopt Fourier analysis as our primary complexity measure. Given $f(x) := (2\pi)^{d/2} \int \tilde{f}(k) e^{ik \cdot x} dk$ where $\tilde{f}(k) := \int f(x) e^{-ik \cdot x} dx$ is the Fourier transform, we perform a discrete Fourier transform by uniformly discretizing the frequency domain, $k \in \{0, 1, \dots, K\}$. The value of K is chosen by the Nyquist-Shannon limit (Shannon, 1949) to ensure accurate function representation. Our complexity measure $c(f)$ is then computed as the frequency-weighted average of the Fourier coefficients:

$$c(f) = \frac{\sum_{k=0}^K \tilde{f}(k) \cdot k}{\sum_{k=0}^K \tilde{f}(k)}. \quad (1)$$

Intuitively, larger $c(f)$ indicates higher complexity due to a dominance of high-frequency components such as rapid amplitude changes or intricate details. Conversely, lower $c(f)$ implies a larger contribution from low-frequency components, indicating a lower complexity function.

2.2 MEASURING SIMPLICITY BIAS

In theory, simplicity bias can be measured by evaluating the complexity of the function to which the network converges after training. However, directly comparing simplicity bias across different architectures after convergence is challenging due to the randomness of the non-stationary optimization process, especially in RL, where the data distribution changes continuously.

Empirical studies suggest that the initial complexity of a network strongly correlates with the complexity of the functions it converges to during training (Valle-Perez et al., 2018; De Palma et al., 2019; Mingard et al., 2019; Teney et al., 2024). Therefore, for a given network architecture f with an initial parameter distribution Θ_0 , we define the **simplicity bias score** $s(f) : \mathcal{F} \rightarrow (0, \infty)$ as:

$$s(f) \approx \mathbb{E}_{\theta \sim \Theta_0} \left[\frac{1}{c(f_\theta)} \right] \quad (2)$$

where f_θ denotes the network architecture f parameterized by θ .

This measure indicates that networks with lower complexity at initialization are more likely to exhibit a higher simplicity bias, thereby converging to simpler functions during training.

3 RELATED WORK

3.1 SIMPLICITY BIAS

Initially, simplicity bias was mainly attributed to the implicit regularization effects of the stochastic gradient descent (SGD) optimizer (Soudry et al., 2018; Gunasekar et al., 2018; Chizat & Bach, 2020;

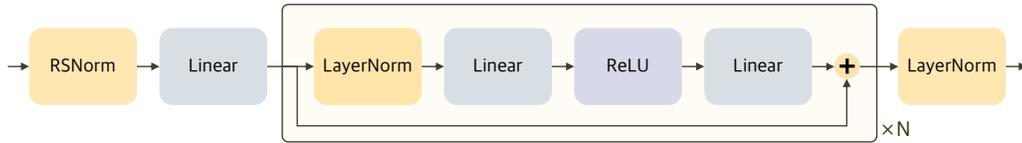


Figure 3: **SimBa architecture.** The network integrates Running Statistics Normalization (RSNorm), residual feedforward blocks, and post-layer normalization to embed simplicity bias into deep RL.

Pesme et al., 2021). During training, SGD introduces noise which prevents the model from converging to sharp minima, guiding it toward flatter regions of the loss landscape (Wu et al., 2022). Such flatter minima are associated with functions of lower complexity, thereby improving generalization.

However, recent studies suggest that simplicity bias is also inherent in the network architecture itself (Valle-Perez et al., 2018; Mingard et al., 2019). Architectural components such as normalization layers, ReLU activations (Hermann et al., 2024), and residual connections (He et al., 2020) promote simplicity bias by encouraging smoother, less complex functions. Fourier analysis has shown that these components help models prioritize learning low-frequency patterns, guiding optimization toward flatter regions that generalize better (Teney et al., 2024). Consequently, architectural design plays a crucial role in favoring simpler solutions, enabling the use of overparameterized networks.

3.2 DEEP REINFORCEMENT LEARNING

For years, deep RL has largely focused on algorithmic improvements to enhance sample efficiency and generalization. Techniques like Double Q-learning (Van Hasselt et al., 2016; Fujimoto et al., 2018), and Distributional RL (Dabney et al., 2018) have improved the stability of value estimation by reducing overestimation bias. Regularization strategies—including periodic reinitialization (Nikishin et al., 2022; Lee et al., 2024), Layer Normalization (Lee et al., 2023; Gallici et al., 2024), Batch Normalization (Bhatt et al., 2024), and Spectral Normalization (Gogianu et al., 2021)—have been employed to prevent overfitting and enhance generalization. Incorporating self-supervised objectives with model-based learning (Fujimoto et al., 2023; Hafner et al., 2023; Hansen et al., 2023) has further improved representation learning and sample efficiency.

Despite these advances, scaling up network architectures in deep RL remains underexplored, especially by leveraging the simplicity bias principle. Several recent studies have attempted to scale up network sizes—through ensembling (Chen et al., 2021; Obando-Ceron et al., 2024), widening (Schwarzer et al., 2023), residual connections (Espenholt et al., 2018) and deepening networks (Bjorck et al., 2021; Nauman et al., 2024). However, they often rely on computationally intensive layers like spectral normalization (Bjorck et al., 2021) or require sophisticated training protocols (Nauman et al., 2024), limiting their applicability.

In this work, we aim to design an architecture that amplifies simplicity bias, enabling us to effectively scale up parameters in RL, independent of using any other sophisticated training protocol.

4 SIMBA

This section introduces the SimBa network, an architecture designed to embed simplicity bias into deep RL. The architecture comprises Running Statistics Normalization, Residual Feedforward Blocks, and Post-Layer Normalization. By amplifying the simplicity bias, SimBa allows the model to avoid overfitting for highly overparameterized configurations.

Running Statistics Normalization (RSNorm). First, RSNorm standardizes input observations by tracking the running mean and variance of each input dimension during training, preventing features with disproportionately large values from dominating the learning process.

Given an input observation $\mathbf{o}_t \in \mathbb{R}^{d_o}$ at timestep t , we update the running observation mean $\mu_t \in \mathbb{R}^{d_o}$ and variance $\sigma_t^2 \in \mathbb{R}^{d_o}$ as follows:

$$\mu_t = \mu_{t-1} + \frac{1}{t} \delta_t, \quad \sigma_t^2 = \frac{t-1}{t} (\sigma_{t-1}^2 + \frac{1}{t} \delta_t^2) \quad (3)$$

where $\delta_t = \mathbf{o}_t - \mu_{t-1}$ and d_o denotes the dimension of the observation.

Once μ_t and σ_t^2 are computed, each input observation \mathbf{o}_t is normalized as:

$$\bar{\mathbf{o}}_t = \text{RSNorm}(\mathbf{o}_t) = \frac{\mathbf{o}_t - \mu_t}{\sqrt{\sigma_t^2 + \epsilon}} \quad (4)$$

where $\bar{\mathbf{o}}_t \in \mathbb{R}^{d_o}$ is the normalized output, and ϵ is a small constant for numerical stability.

While alternative observation normalization methods exist, RSNorm consistently demonstrates superior performance. A comprehensive comparison is provided in Section 7.1.

Residual Feedforward Block. The normalized observation $\bar{\mathbf{o}}_i$ is first embedded into a d_h -dimensional vector using a linear layer:

$$\mathbf{x}_t^l = \text{Linear}(\bar{\mathbf{o}}_t). \quad (5)$$

At each block, $l \in \{1, \dots, L\}$, the input \mathbf{x}_t^l passes through a pre-layer normalization residual feedforward block, introducing simplicity bias by allowing a direct linear pathway from input to output. This direct linear pathway enables the network to pass the input unchanged unless non-linear transformations are necessary. Each block is defined as:

$$\mathbf{x}_t^{l+1} = \mathbf{x}_t^l + \text{MLP}(\text{LayerNorm}(\mathbf{x}_t^l)). \quad (6)$$

Following Vaswani (2017), the MLP is structured with an inverted bottleneck, where the hidden dimension is expanded to $4 \cdot d_h$ and a ReLU activation is applied between the two linear layers.

Post-Layer Normalization. To ensure that activations remain on a consistent scale before predicting the policy or value function, we apply layer normalization after the final residual block:

$$\mathbf{z}_t = \text{LayerNorm}(\mathbf{x}_t^L). \quad (7)$$

The normalized output \mathbf{z}_t is then processed through a linear layer, generating predictions for the actor’s policy or the critic’s value function.

5 ANALYZING SIMBA

In this section, we analyze whether each component of SimBa amplifies simplicity bias and allows scaling up parameters in deep RL. We conducted experiments on challenging environments in DMC involving Humanoid and Dog, collectively referred to as DMC-Hard. Throughout this section, we used Soft Actor Critic (Haarnoja et al., 2018, SAC) as the base algorithm.

5.1 ARCHITECTURAL COMPONENTS

To quantify simplicity bias in SimBa, we use Fourier analysis as described in Section 2. For each network f , we estimate the simplicity bias score $s(f)$ by averaging over 100 random initializations $\theta \sim \Theta_0$. Following Teney et al. (2024), the input space $\mathcal{X} = [-100, 100]^2 \subset \mathbb{R}^2$ is divided into a grid of 90,000 points, and the network outputs a scalar value for each input which are represented as a grayscale image. By applying the discrete Fourier transform to these outputs, we compute the simplicity score $s(f)$ defined in Equation 2 (see Appendix B for further details).

Figure 4.(a) shows that SimBa’s key components—such as residual connections and layer normalization—increase the simplicity bias score $s(f)$, biasing the architecture toward simpler functional representations at initialization. When combined, these components induce a stronger preference for low-complexity functions (i.e., high simplicity score) than when used individually.

Figure 4.(b) reveals a clear relationship between simplicity bias and performance: architectures with higher simplicity bias scores lead to enhanced performance. Specifically, compared to the MLP, adding residual connections increases the average return by 50 points, adding layer normalization adds 150 points, and combining all components results in a substantial improvement of 550 points.



Figure 4: **Component Analysis.** (a) Simplicity bias scores estimated via Fourier analysis. Mean and 95% CI are computed over 100 random initializations. (b) Average return in DMC-Hard at 1M steps. Mean and 95% CI over 10 seeds, using SAC. Stronger simplicity bias correlates with higher returns for overparameterized networks.

5.2 COMPARISON WITH OTHER ARCHITECTURES

To assess SimBa’s scalability, we compared it to BroNet (Nauman et al., 2024), SpectralNet (Bjorck et al., 2021), and MLP. Detailed descriptions of each architecture are provided in Appendix F.

The key differences between SimBa and other architectures lie in the placement of components that promote simplicity bias. SimBa maintains a direct linear residual pathway from input to output, applying non-linearity exclusively through residual connections. In contrast, other architectures introduce non-linearities within the input-to-output pathway, increasing functional complexity. Additionally, SimBa employs post-layer normalization to ensure consistent activation scales across all hidden dimensions, reducing variance in policy and value predictions. Conversely, other architectures omit normalization before the output layer, which can lead to high variance in their predictions.

To ensure that performance differences were attributable to architectural design choices rather than varying input normalization, we uniformly applied RSNorm as the input normalization across all models. Our investigation focused on scaling up the critic network’s parameters by increasing the hidden dimension, as scaling up the actor-network showed limited benefits (see Section 7.2).

As illustrated in Figure 5.(a), SimBa achieves the highest simplicity bias score compared to the experimented architectures, demonstrating its strong preference for simpler solutions. In Figure 5.(b), while the MLP failed to scale with increasing network size, BroNet, SpectralNet, and SimBa all showed performance improvements as their networks scaled up.

Importantly, the scalability of each architecture was strongly correlated with its simplicity bias score, where a higher simplicity bias led to better scalability. SimBa demonstrated the best scalability, supporting the hypothesis that simplicity bias is a key factor in scaling deep RL models.

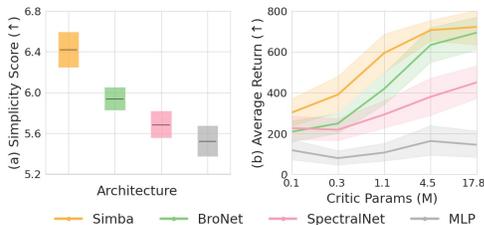


Figure 5: **Architecture Comparison.** (a) SimBa consistently exhibits a higher simplicity bias score. (b) SimBa demonstrates strong scaling performance in terms of average return for DMC-Hard compared to the other architectures. The results are from 5 random seeds.

6 EXPERIMENTS

This section evaluates SimBa’s applicability across various deep RL algorithms and environments. For each baseline, we either use the authors’ reported results or run experiments using their recommended hyperparameters. Detailed descriptions of the environments used in our evaluations are provided in Appendix H. In addition, we also include learning curves and final performance for each task in Appendix J and K, along with hyperparameters used in our experiments in Appendix I.

6.1 OFF-POLICY RL

Experimental Setup. We evaluate the algorithms on 51 tasks across three benchmarks: DMC (Tassa et al., 2018), MyoSuite (Caggiano et al., 2022), and HumanoidBench (Sferrazza et al., 2024). The

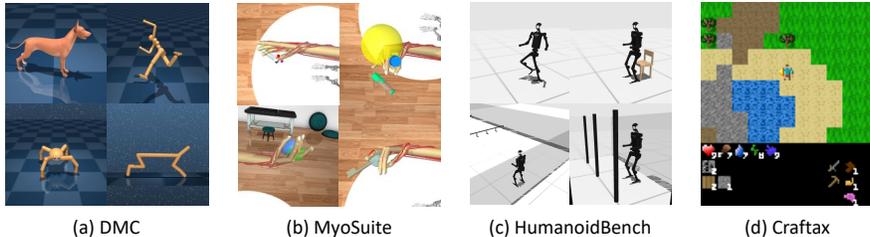


Figure 6: **Environment Visualizations.** SimBa is evaluated across four diverse benchmark environments: DMC, MyoSuite, and HumanoidBench, which feature complex locomotion and manipulation tasks, and Craftax, which introduces open-ended tasks with varying complexity.

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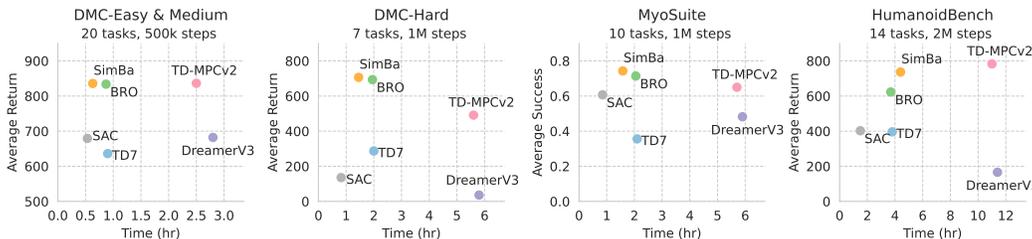


Figure 8: **Off-policy RL Benchmark.** Average episode return for DMC and HumanoidBench and average success rate for MyoSuite across 51 continuous control tasks. SimBa (with SAC) achieves high computational efficiency by only changing the network architecture.

DMC tasks are further categorized into DMC-Easy&Medium and DMC-Hard based on complexity. We vary the number of training steps per benchmark according to the complexity of the environment: 500K for DMC-Easy&Medium, 1M for DMC-Hard and MyoSuite, and 2M for HumanoidBench.

Baselines. We compare SimBa against state-of-the-art off-policy RL algorithms. (i) SAC (Haarnoja et al., 2018), an actor-critic algorithm based on maximum entropy RL; (ii) DDPG (Lillicrap, 2015), a deterministic policy gradient algorithm with deep neural networks; (iii) TD7 (Fujimoto et al., 2023), an enhanced version of TD3 incorporating state-action representation learning; (iv) BRO (Nauman et al., 2024), which scales the critic network of SAC while integrating distributional Q-learning, optimistic exploration, and periodic resets. For a fair comparison, we use BRO-Fast, which is the most computationally efficient version; (v) TD-MPC2 (Hansen et al., 2023), which combines trajectory planning with long-return estimates using a learned world model; and (vi) DreamerV3 (Hafner et al., 2023), which learns a generative world model and optimizes a policy via simulated rollouts.

Integrating SimBa into Off-Policy RL. The SimBa architecture is algorithm-agnostic and can be applied to any deep RL method. To demonstrate its applicability, we integrate SimBa into two model-free (SAC, DDPG) and one model-based algorithm (TD-MPC2), evaluating their performance on the DMC-Hard benchmark. For SAC and DDPG, we replace the standard MLP-based actor-critic networks with SimBa. For TD-MPC2, we substitute the shared encoder from MLP to SimBa while matching the number of parameters as the original implementation.

Figure 7 shows consistent benefits across all algorithms: integrating SimBa increased the average return by 570, 480, and, 170 points for SAC, DDPG, and TD-MPC2 respectively. These results demonstrate that scaling up parameters with a strong simplicity bias significantly enhances performance in deep RL.

Comparisons with State-of-the-Art Methods. Here, we compare SAC + SimBa against state-of-the-art off-policy deep RL algorithms, which demonstrated the most promising results in previous experiments. Throughout this section, we refer to SAC + SimBa as SimBa.

Figure 8 presents the results, where the x-axis represents computation time using an RTX 3070 GPU, and the y-axis denotes performance. Points in the upper-left corner (👉) indicate higher compute efficiency. Here, SimBa consistently outperforms most baselines across various environments while requiring less computational time. The only exception is TD-MPC2 on HumanoidBench; however, TD-MPC2 requires 2.5 times the computational resources of SimBa to achieve better performance, making SimBa the most efficient choice overall.

The key takeaway is that SimBa achieves these remarkable results without the bells and whistles often found in state-of-the-art deep RL algorithms. It is easy to implement, which only requires modifying the network architecture into SimBa without additional changes to the loss functions (Schwarzer et al., 2020; Hafner et al., 2023) or using domain-specific regularizers (Fujimoto et al., 2023; Nauman et al., 2024). Its effectiveness stems from an architectural design that leverages simplicity bias and scaling up the number of parameters, thereby maximizing performance.

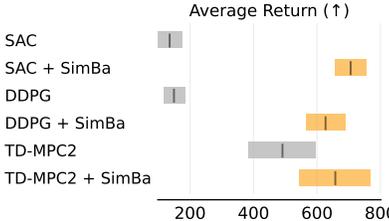


Figure 7: **Off-Policy RL with SimBa.** Replacing MLP with SimBa leads to substantial performance improvements across various off-policy RL methods. Mean and 95% CI are averaged over 10 seeds for SAC and DDPG, and 3 seeds for TD-MPC2 in DMC-Hard.

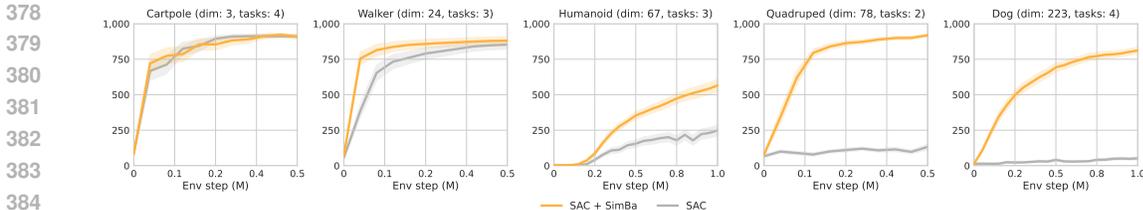


Figure 9: **Impact of Input Dimension.** Average episode return for DMC tasks plotted against increasing state dimensions. Results show that the benefits of using SimBa increase with higher input dimensions, effectively alleviating the curse of dimensionality.



Figure 10: **On-policy RL with SimBa.** Average return of achievements and task success rate for three different tasks comparing PPO + SimBa and PPO on Craftax. Integrating SimBa enables effective learning of complex behaviors.

Impact of Input Dimension. To further identify in which cases SimBa offers significant benefits, we analyze its performance across DMC environments with varying input dimensions. As shown in Figure 9, the advantage of SimBa becomes more pronounced as input dimensionality increases. We hypothesize that higher-dimensional inputs exacerbate the curse of dimensionality, and the simplicity bias introduced by SimBa effectively mitigates overfitting in these high-dimensional settings.

6.2 ON-POLICY RL

Experimental Setup. We conduct our on-policy RL experiments in Craftax (Matthews et al., 2024), an open-ended environment inspired by Crafter (Hafner, 2022) and NetHack (Küttler et al., 2020). Craftax poses a unique challenge with its open-ended structure and compositional tasks. Following Matthews et al. (2024), we use Proximal Policy Optimization (Schulman et al., 2017, PPO) as the baseline algorithm. When integrating SimBa with PPO, we replace the standard MLP-based actor-critic networks with SimBa and train both PPO and PPO + SimBa on 1024 parallel environments for a total of 1 billion environment steps.

Results. As illustrated in Figure 10, integrating SimBa into PPO significantly enhances performance across multiple complex tasks. With SimBa, the agent learns to craft iron swords and pickaxes more rapidly, enabling the early acquisition of advanced tools. Notably, by using these advanced tools, the SimBa-based agent successfully defeats challenging adversaries like the Orc Mage using significantly fewer time steps than an MLP-based agent. These improvements arise solely from the architectural change, demonstrating the effectiveness of the SimBa approach for on-policy RL.

6.3 UNSUPERVISED RL

Experimental Setup. In our unsupervised RL study, we incorporate SimBa for online skill discovery, aiming to identify diverse behaviors without relying on task-specific rewards. We focus our experiments primarily on METRA (Park et al., 2023), which serves as the state-of-the-art algorithm in this domain. We evaluate METRA and METRA with SimBa on the Humanoid task from DMC, running 10M environment steps. For a quantitative comparison, we adopt state coverage as our main metric. Coverage is measured by discretizing the x and y axes into a grid and counting the number of grid cells covered by the learned behaviors at each evaluation epoch, following prior literature (Park et al., 2023; Kim et al., 2024a;b).

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Results. As illustrated in Figure 1.(a) and 11, integrating SimBa into METRA significantly enhances state coverage on the Humanoid task. The high-dimensional input space makes it challenging for METRA to learn diverse skills. By injecting simplicity bias to manage high input dimensions and scaling up parameters to facilitate effective diverse skill acquisition, SimBa effectively leads to improved exploration and broader state coverage.

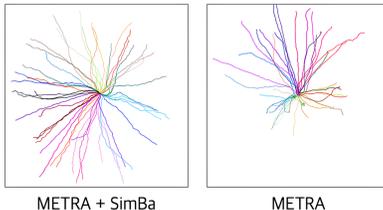


Figure 11: **URL with Simba.** Integrating SimBa to METRA enhances state coverage.

7 ABLATIONS

We conducted ablations on DMC-Hard with SAC + SimBa, running 5 seeds for each experiment.

7.1 OBSERVATION NORMALIZATION

A key factor in SimBa’s success is using RSNorm for observation normalization. To validate its effectiveness, we compare 6 alternative methods: (i) Layer Normalization (Ba et al., 2016); (ii) RMS Normalization (Zhang & Sennrich, 2019); (iii) Batch Normalization (Ioffe & Szegedy, 2015); (iv) Env Wrapper RSNorm, which tracks running statistics for each dimension, normalizes observations upon receiving from the environment, and stores them in the replay buffer; (v) Initial N Steps: fixed statistics derived from initially collected transition samples, where we used $N = 5,000$; and (iv) Oracle Statistics, which rely on pre-computed statistics from previously collected expert data.

As illustrated in Figure 12, layer normalization and batch normalization offer little to no performance gains. While the env-wrapper RSNorm is somewhat effective, it falls short of RSNorm’s performance. Although widely used in deep RL frameworks (Dhariwal et al., 2017; Hoffman et al., 2020; Raffin et al., 2021), the env-wrapper introduces inconsistencies in off-policy settings by normalizing samples with different statistics based on their collection time. This causes identical observations to be stored with varying values in the replay buffer, reducing learning consistency. Fixed initial statistics also show slightly worse performance than RSNorm, potentially due to their inability to adapt to the evolving dynamics during training. Overall, only RSNorm matched the performance of Oracle statistics, making it the most practical observation normalization choice for deep RL.

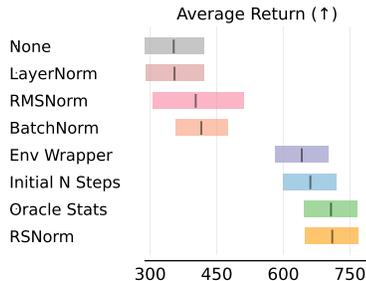


Figure 12: **Obs Normalization.** RSNorm consistently outperforms alternative normalization methods. Mean and 95% CI over 5 seeds.

7.2 SCALING THE NUMBER OF PARAMETERS

Here, we investigate scaling parameters in the actor-critic architecture with SimBa by focusing on two aspects: (i) scaling the actor versus the critic network, and (ii) scaling network width versus depth. For width scaling, the actor and critic depths are set to 1 and 2 blocks, respectively. For depth scaling, the actor and critic widths are fixed at 128 and 512, respectively, following our default setup.

As illustrated in Figure 13, scaling up the width or depth of the critic (➡) generally improves performance, while scaling up the actor’s width or depth (⬆) tends to reduce performance. This contrast suggests that the target complexity of the actor may be lower than that of the critic, where scaling up the actor’s parameters might be ineffective. These findings align with previous study (Nauman et al., 2024), which highlights the benefits of scaling up the critic while showing limited advantages in scaling up the actor.

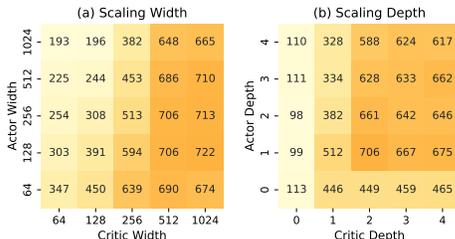


Figure 13: **Scaling Actor and Critic Network.** Performance of SAC with SimBa by varying width and depth for the actor and critic network.

486 Furthermore, for the critic network, scaling up the width is generally more effective than scaling up
 487 the depth. While both approaches can enhance the network’s expressivity, scaling up the depth can
 488 decrease the simplicity bias as it adds more non-linear components within the network. Based on
 489 our findings, we recommend width scaling as the default strategy.
 490

491 **7.3 SCALING REPLAY RATIO**

492
 493 According to scaling laws (Kaplan et al., 2020), performance can be enhanced not only by scaling
 494 up the number of parameters but also by scaling up the computation time, which can be done by
 495 scaling up the number of gradient updates per collected sample (i.e., replay ratio) in deep RL.

496 However, in deep RL, scaling up the replay ratio has been shown to decrease performance due to the
 497 risk of overfitting the network to the initially collected samples (Nikishin et al., 2022; D’Oro et al.,
 498 2022; Lee et al., 2023). To address this issue, recent research has proposed periodically reinitializing
 499 the network to prevent overfitting, demonstrating that performance can be scaled with respect to the
 500 replay ratio. In this section, we aim to assess whether the simplicity bias induced by SimBa can
 501 mitigate overfitting under increased computation time (i.e., higher replay ratios).

502 To evaluate this, we trained SAC with SimBA using 2, 4, 8, and 16 replay ratios, both with and without peri-
 503 odic resets. For SAC with resets, we reinitialized the entire network and optimizer every 500,000 gradient steps.
 504 We also compared our results with BRO (Nauman et al., 2024), which incorporates resets.

508 Surprisingly, as illustrated in Figure 14, SimBa’s performance consistently improves as the replay ratio in-
 509 creases, even without periodic resets. We have excluded results for BRO without resets, as it fails to learn
 510 meaningful behavior for all replay ratios (achieves lower than 300 points). Notably, when resets are included for
 511 SimBa, the performance gains become even more pronounced, with a replay ratio of 8 outperforming the most
 512 computationally intensive BRO algorithm (RR10).
 513
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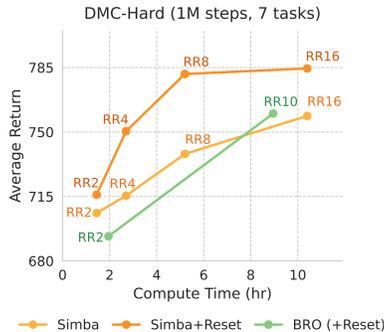


Figure 14: **Scaling Replay Ratio.** Performance of SimBa with and without periodic resets for various replay ratios.

517
 518 **8 LESSONS AND OPPORTUNITIES**

519
 520 **Lessons.** Deep reinforcement learning has historically struggled with overfitting, requiring complex
 521 training protocols and tricks to mitigate these issues (Hessel et al., 2018; Fujimoto et al., 2023).
 522 These complexities can hinder practitioners with limited resources. In this paper, we improved
 523 performance solely by modifying the network architecture while keeping the underlying algorithms
 524 unchanged, simplifying SimBa’s implementation and making it easy to adopt. By incorporating
 525 simplicity bias through architectural design and scaling up parameters, our network converges to
 526 simpler, generalizable functions, matching or surpassing state-of-the-art methods. This aligns with
 527 Richard Sutton’s Bitter Lesson (Sutton, 2019): while task-specific designs may offer immediate
 528 gains, scalable approaches provide more sustainable long-term benefits.

529
 530 **Opportunities.** While our exploration has focused on network architecture, optimization techniques
 531 like dropout (Hiraoka et al., 2021), data augmentation (Kostrikov et al., 2020), and advanced opti-
 532 mization algorithms (Foret et al., 2020) are also crucial for promoting convergence to simpler func-
 533 tions. Extending our insights to vision-based reinforcement learning presents an exciting future
 534 direction; designing convolutional architectures based on simplicity bias principles could enhance
 535 learning efficiency in visual tasks. Additionally, applying SimBa to multi-task reinforcement learn-
 536 ing could improve performance across diverse tasks by sharing simple, generalizable representations
 537 across tasks. Furthermore, although we concentrated on the model-free algorithm SAC, model-
 538 based algorithms have shown considerable success (Schwarzer et al., 2020; Hansen et al., 2023;
 539 Hafner et al., 2023), and preliminary investigations applying SimBa to model-based RL, specifically
 TD-MPC2 (Hansen et al., 2023), are promising. We encourage the research community to col-
 laboratively explore these avenues to advance deep RL architectures and facilitate their successful
 application in real-world scenarios.

540 REPRODUCIBILITY STATEMENT

541
542 To ensure the reproducibility of our experiments, we provide the complete source code to run SAC
543 with SimBa. Moreover, we have included a Dockerfile to simplify the testing and replication of our
544 results. The raw scores of each experiment including all baselines are also included. The code is
545 available at this [anonymous github link](#).

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864 A DEFINITION OF SIMPLICITY BIAS

865 This section provides more in-depth definition of simplicity bias for clarity.

866 A.1 COMPLEXITY MEASURE

867 Let $\mathcal{X} \subseteq \mathbb{R}^n$ denote the input space and $\mathcal{Y} \subseteq \mathbb{R}^m$ the output space. Consider the function space
868 $\mathcal{F} = \{f | f : \mathcal{X} \rightarrow \mathcal{Y}\}$. Given a training dataset $D = \{(x_i, y_i)\}_{i=1}^N$ and a tolerance $\epsilon > 0$, define the
869 set of functions $\mathcal{I} \subseteq \mathcal{F}$ achieving a training loss below ϵ :

$$870 \mathcal{I} = \left\{ f \in \mathcal{F} \mid \mathcal{L}_{\text{train}}(f) = \frac{1}{N} \sum_{i=1}^N \ell(f(x_i), y_i) < \epsilon \right\}, \quad (8)$$

871 where $\ell : \mathcal{Y} \times \mathcal{Y} \rightarrow [0, \infty)$ is a loss function such as mean squared error or cross-entropy.

872 To quantify the complexity of functions within \mathcal{F} , we adopt a Fourier-based complexity measure as
873 described in Section 2. Specifically, for a function f with a Fourier series representation:

$$874 f(x) = (2\pi)^{d/2} \int \tilde{f}(k) e^{ik \cdot x} dk, \quad (9)$$

875 where $\tilde{f}(k) = \int f(x) e^{-ik \cdot x} dx$ is the Fourier transform of f .

876 Given a Fourier series representation \tilde{f} , we perform a discrete Fourier transform by uniformly dis-
877 cretizing the frequency domain. Specifically, we consider frequencies $k \in \{0, 1, \dots, K\}$, where K
878 is selected based on the Nyquist-Shannon sampling theorem (Shannon, 1949) to ensure an accurate
879 representation of f . This discretization transforms the continuous frequency domain into a finite set
880 of frequencies suitable for computational purposes.

881 Then, the **complexity measure** $c : \mathcal{F} \rightarrow [0, \infty)$ is defined as the weighted average of the Fourier
882 coefficients from discretization:

$$883 c(f) = \frac{\sum_{k=0}^K \tilde{f}(k) \cdot k}{\sum_{k=0}^K \tilde{f}(k)}. \quad (10)$$

884 Equation 10 captures the intuition that higher complexity arises from the dominance of high-
885 frequency components, which correspond to rapid amplitude changes or intricate details in the func-
886 tion f . Conversely, a lower complexity measure indicates a greater contribution from low-frequency
887 components, reflecting simpler, smoother functions.

888 A.2 SIMPLICITY BIAS SCORE

889 Let f be a neural network, and let $\text{alg} \in \mathcal{A}$ denote the optimization algorithm used for training, such
890 as stochastic gradient descent. We quantify the **simplicity bias score** $s : \mathcal{F} \times \mathcal{A} \rightarrow (0, \infty)$ as:

$$891 s(f, \text{alg}) = \mathbb{E}_{f^* \sim P_{f, \text{alg}}} \left[\frac{1}{c(f^*)} \right] \quad (11)$$

892 where $P_{f, \text{alg}}$ denotes the distribution over \mathcal{I} induced by training f with alg . A higher simplicity bias
893 score indicates a stronger tendency to converge toward functions with lower complexity.

894 Then, from equation 11, we can compute the simplicity bias score with respect to the architecture
895 by marginalizing over the distribution of the algorithm as:

$$896 s(f) = \mathbb{E}_{\text{alg} \sim \mathcal{A}} \left[\mathbb{E}_{f^* \sim P_{f, \text{alg}}} \left[\frac{1}{c(f^*)} \right] \right]. \quad (12)$$

897 Directly measuring this bias by analyzing the complexity of the converged function is challenging
898 due to the non-stationary nature of training dynamics, especially in reinforcement learning where
899 data distributions evolve over time. Empirical studies (Valle-Perez et al., 2018; De Palma et al.,
900 2019; Mingard et al., 2019; Teney et al., 2024) suggest that the initial complexity of a network is
901 strongly correlated with the complexity of the functions it learns after training.

Therefore, as a practical proxy for simplicity bias, we define the approximate of the simplicity bias score $s(f)$ for a network architecture f with an initial parameter distribution Θ_0 as:

$$s(f) \approx \mathbb{E}_{\theta \sim \Theta_0} \left[\frac{1}{c(f_\theta)} \right]. \quad (13)$$

where f_θ denotes the network f parameterized by θ .

A higher simplicity bias score indicates that, on average, the network initialized from Θ_0 represents simpler functions prior to training, suggesting a predisposition to converge to simpler solutions during optimization. This measure aligns with the notion that networks with lower initial complexity are more likely to exhibit a higher simplicity bias.

B MEASURING SIMPLICITY BIAS

To quantify the simplicity bias defined in Equation 13, we adopt a methodology inspired by the Neural Redshift framework (Teney et al., 2024). We utilize the original codebase provided by the authors¹ to assess the complexity of random neural networks. This approach evaluates the inherent complexity of the architecture before any optimization, thereby isolating the architectural effects from those introduced by training algorithms such as stochastic gradient descent.

Following the methodology outlined in Teney et al. (2024), we perform the following steps to measure the simplicity bias score of a neural network $f_\theta(\cdot)$:

Sampling Initialization. For each network architecture f , we generate N random initializations $\theta \sim \Theta_0$. This ensemble of initial parameters captures the variability in complexity introduced by different random seeds. In this work, we used $N = 100$.

Sampling Input. We define the input space $\mathcal{X} = [-100, 100]^2 \subset \mathbb{R}^2$ and uniformly divide it into a grid of 90,000 points, achieving 300 divisions along each dimension. This dense sampling ensures comprehensive coverage of the input domain.

Evaluating Function. For each sampled input point $x \in \mathcal{X}$, we compute the corresponding output $f_\theta(x)$. The collection of these output values forms a 2-D grid of scalars, effectively representing the network’s function as a grayscale image.

Discrete Fourier Transform. We apply a discrete Fourier transform (DFT) to the grayscale image obtained from the function evaluations. This transformation decomposes $f_\theta(\cdot)$ into a sum of basis functions of varying frequencies.

Measuring Function Complexity. Utilizing the Fourier coefficients obtained from the DFT, we compute the complexity measure $c(f_\theta)$ as defined in Equation 10. Specifically, we calculate the frequency-weighted average of the Fourier coefficients:

$$c(f_\theta) = \frac{\sum_{k=0}^K \tilde{f}_\theta(k) \cdot k}{\sum_{k=0}^K \tilde{f}_\theta(k)}, \quad (14)$$

where $\tilde{f}_\theta(k)$ denotes the Fourier coefficient at frequency k .

Estimating Simplicity Bias Score. For each network f , we estimate the simplicity bias score $s(f)$ by averaging the inverse of the complexity measures over all N random initializations:

$$s(f) \approx \frac{1}{N} \sum_{i=1}^N \frac{1}{c(f_{\theta_i})}, \quad (15)$$

where θ_i represents the i -th random initialization.

¹<https://github.com/ArmandNM/neural-redshift>

C ADDITIONAL ABLATIONS

C.1 ARCHITECTURAL ABLATION

Experimental Setup. Following an architectural component analysis in Section 5.1, we further conducted ablation studies by removing each key component from SimBa to observe how the simplicity score and average return differ with each ablation. Using the same training setup as Section 5.1, we conducted these experiments on DMC-Hard, training for 1 million environment steps.

Results. As shown in Figure 15.(a), we observe that, except for RSNorm, removing key components of SimBa—such as residual connections, pre-layer normalization, or post-layer normalization—leads to a reduction in the simplicity bias score. This indicates that the functional representation at initialization becomes biased toward more complex functions when these components are absent. This finding further emphasizes the critical role of each component in promoting simplicity bias within SimBa’s design. Additionally, Figure 15.(b) demonstrates that removing any key component significantly impacts SimBa’s performance, underscoring the necessity of each architectural element.

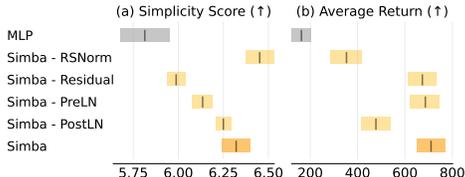


Figure 15: Additional Component Analysis. (a) Simplicity bias scores estimated via Fourier analysis. Mean and 95% CI are computed over 100 random initializations. (b) Average return in DMC-Hard at 1M steps. Mean and 95% CI over 10 seeds, using SAC.

Regarding RSNorm, although its absence significantly impacts the agent’s performance negatively, the overall simplicity bias score of the architecture increases rather than decreases. We speculate that this discrepancy arises because our Fourier Transform-based complexity measure relies on evaluating the network’s outputs over a uniformly distributed input space. RSNorm is designed to normalize activations based on the statistics of non-uniform or dynamically changing input distributions, excelling with inputs that exhibit variance and complexity not captured by a uniform distribution. Therefore, since our simplicity analysis measure is based on a uniform input distribution, it may not fully capture RSNorm’s practical contribution to promoting simplicity bias.

C.2 EXTENDED TRAINING

Experimental Setup. We assess SimBa’s robustness by evaluating its performance on the DeepMind Control Suite (DMC) Hard tasks over an extended training period of 5 million timesteps. We compared three methods: SAC (Haarnoja et al., 2018), TD-MPC2 (Hansen et al., 2023), and SAC with SimBa. All models are trained under identical hyperparameters and configurations as in our main experiments to ensure a fair comparison. The results are averaged over 5 random seeds.

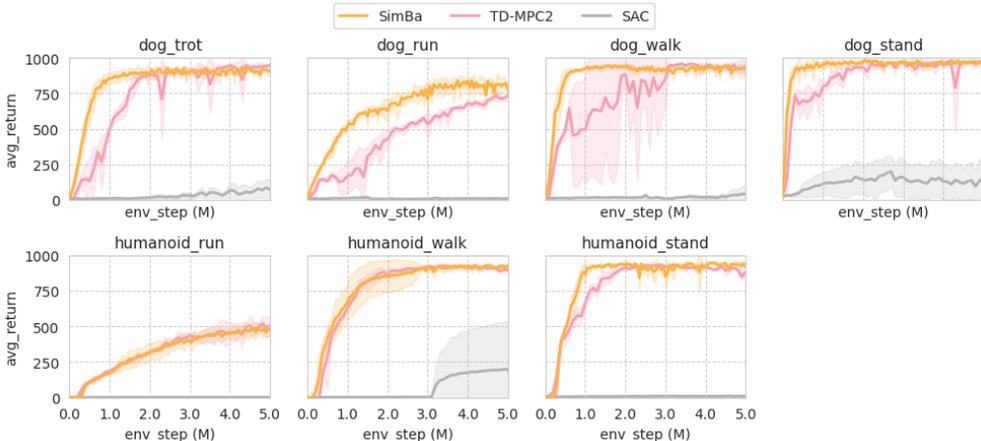


Figure 16: Per task learning curve for DMC Hard. Mean and 95% CI are computed over 5 seeds for SimBa, 3 seeds for TD-MPC2 and SAC.

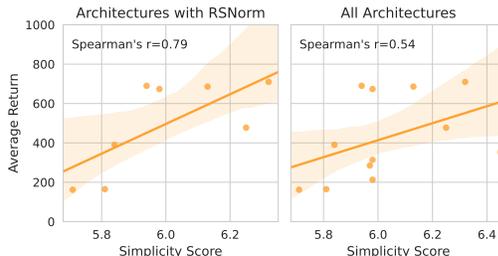
1026 **Results.** Figure 16 presents the learning curve of each method for 5 million timesteps. The learning
 1027 curves indicate that SimBa + SAC not only learns faster but also maintains more stable and consistent
 1028 performance improvements over time, demonstrating better sample efficiency and reduced variance
 1029 compared to the baselines.
 1030

1031 **D CORRELATION BETWEEN SIMPLICITY BIAS AND PERFORMANCE**
 1032

1033 To investigate the relationship between simplicity bias score and performance in deep reinforcement
 1034 learning, we computed the correlation between simplicity bias scores and the average returns
 1035 of various architectures evaluated in our study. Our analysis includes 12 distinct architectures, en-
 1036 compassing 4 primary types: MLP, SimBa, BroNet (Nauman et al., 2024), and SpectralNet (Bjorck
 1037 et al., 2021). Additionally, we examined 4 SimBa variants with component reductions and 4 SimBa
 1038 variants with component additions transitioning from MLP to SimBa.
 1039

1040 As discussed in Appendix F, the current simplicity bias measure may not accurately capture the
 1041 influence of RSNorm. To address this limitation, we separately evaluated the correlation for archi-
 1042 tectures incorporating RSNorm. All twelve architectures have nearly identical parameter counts,
 1043 approximately 4.5 million parameters each, with a maximum variation of 1%. Performance was
 1044 assessed on the DMC-Hard benchmark, trained for one million steps.

1045 **Results.** As shown in As illustrated in Figure 17(a), architectures employing RSNorm exhibit a
 1046 Pearson correlation coefficient of 0.79, indicating a strong positive correlation between simplicity bias
 1047 score and average return in scaled architectures. In contrast, Figure 17(b) shows a Pearson correlation
 1048 coefficient of 0.54, representing a moderate correlation across all architectures. We attribute the lower
 1049 correlation in the broader set to the current limitations of the simplicity bias measure. We believe these
 1050 findings support our hypothesis that higher simplicity bias scores are associated with better perfor-
 1051 mance in deep reinforcement learning with scaled-up architectures.
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Figure 17: Correlation Analysis. Pearson correlation between simplicity bias score and average return on DMC-Hard. (a) Architectures incorporating RSNorm show a strong correlation, $\rho = 0.79$. (b) All evaluated architectures exhibit a moderate correlation, $\rho = 0.54$.

E PLASTICITY ANALYSIS

Recent studies have identified the *loss of plasticity* as a significant challenge in non-stationary training scenarios, where neural networks gradually lose their ability to adapt to new data over time (Nikishin et al., 2022; Lyle et al., 2023; Lee et al., 2024). This phenomenon can severely impact the performance and adaptability of models in dynamic learning environments such as RL. To quantify and understand this issue, several metrics have been proposed, including stable-rank (Kumar et al., 2022), dormant ratio (Sokar et al., 2023), and L2-feature norm (Kumar et al., 2022).

E.1 METRICS

The **stable-rank** assesses the rank of the feature matrix, reflecting the diversity and richness of the representations learned by the network. This is achieved by performing an eigen decomposition on the covariance matrix of the feature matrix and counting the number of singular values σ_j that exceed a predefined threshold τ :

$$\text{s-Rank} = \sum_{j=1}^m \mathbb{I}(\sigma_j > \tau) \quad (16)$$

where $F \in \mathbb{R}^{d \times m}$ is the feature matrix with d samples and m features, σ_j are the singular values, and $\mathbb{I}(\cdot)$ is the indicator function.

The **dormant ratio** measures the proportion of neurons with negligible activation. It is calculated as the number of neurons with activation norms below a small threshold ϵ divided by the total number of neurons D :

$$\text{Dormant Ratio} = \frac{|\{i \mid \|a_i\| < \epsilon\}|}{D} \quad (17)$$

where a_i represents the activation of neuron i .

The **L2-feature norm** represents the average L2 norm of the feature vectors across all samples:

$$\text{Feature Norm} = \frac{1}{N} \sum_{i=1}^N \|F_i\|_2 \quad (18)$$

where F_i is the feature vector for the i -th sample. Large feature norms can signal overactive neurons, potentially leading to numerical instability.

E.2 RESULTS

To evaluate the impact of different architectures on network plasticity, we compared the Soft Actor-Critic (SAC) algorithm implemented with a standard Multi-Layer Perceptron (MLP) against SAC with the SimBa architecture on the DMC-Easy & Medium and DMC-Hard tasks. Additionally, we conducted an ablation study to analyze how each component of SimBa contributes to plasticity measures. Each configuration was evaluated across five random seeds.

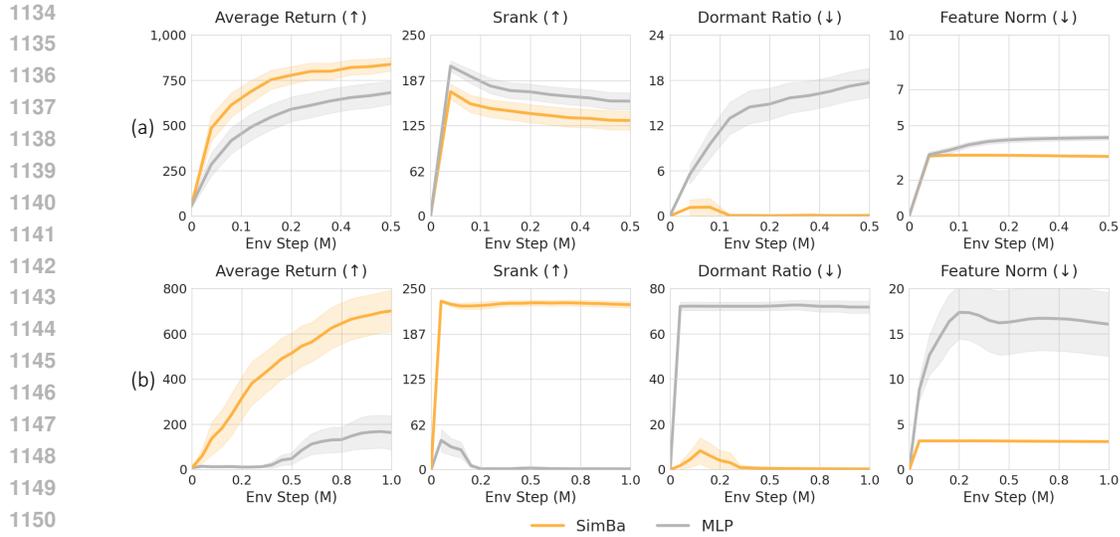


Figure 18: **Plasticity Analysis of SimBa versus MLP.** Metrics compared include dormant ratio, s-rank, and feature norm. Figures (a) and (b) represent DMC-Easy & Medium and DMC-Hard environments, respectively. A higher dormant ratio and feature norm, along with a lower s-rank, indicate a greater loss of plasticity.

As illustrated in Figures 18.(a) and (b), SAC with the MLP architecture exhibits high dormant ratios and large feature norms across both DMC-Easy & Medium and DMC-Hard tasks, indicating significant loss of plasticity. While MLP achieves higher s-rank values than SimBa in the DMC-Easy & Medium tasks, SimBa outperforms MLP in s-rank for the DMC-Hard tasks. Importantly, SimBa consistently maintains lower dormant ratios, balanced feature norms, and overall higher s-rank values in more complex environments.

These findings indicate that SimBa effectively preserves plasticity, avoiding the degenerative trends observed with the MLP-based SAC.

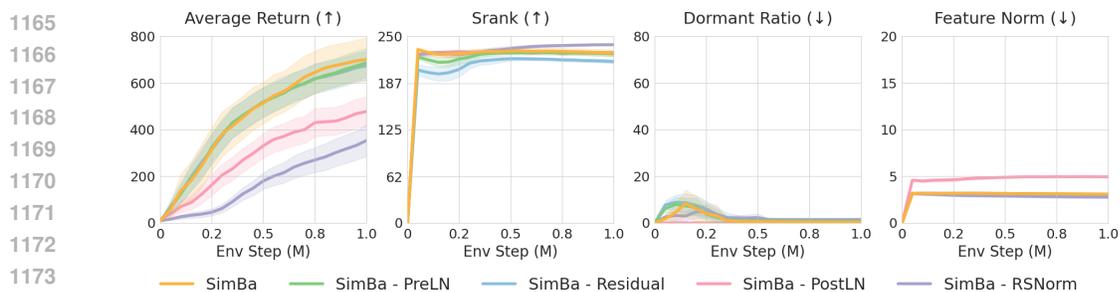


Figure 19: **Plasticity Analysis of SimBa Components.** Metrics compared include dormant ratio, s-rank, and feature norm. Figures (a) and (b) represent DMC-Easy & Medium and DMC-Hard environments, respectively. A higher dormant ratio and feature norm, along with a lower s-rank, indicate a greater loss of plasticity.

Furthermore, for the component ablation study (Figure 19), integrating each component of SimBa individually shows improved average returns. However, ablating each component does not significantly reduce plasticity measures, with only Post Layer Normalization showing an improved feature norm. This suggests that while individual components may not significantly impact plasticity measures with their sole use, their integration within the SimBa architecture is crucial for effectively mitigating plasticity loss.

F COMPARISON TO EXISTING ARCHITECTURES

Here we provide a more in-depth discussion related to the architectural difference between SimBa, BroNet, and SpectralNet. As shown in Figure 20, SimBa differs from BroNet in three key aspects: (i) the inclusion of RSNorm, (ii) the implementation of pre layer-normalization, (iii) the utilization of a linear residual pathway, (iv) and the inclusion of a post-layer normalization layer. Similarly, compared to SpectralNet, SimBa incorporates RSNorm, employs a linear residual pathway, and leverages spectral normalization differently.

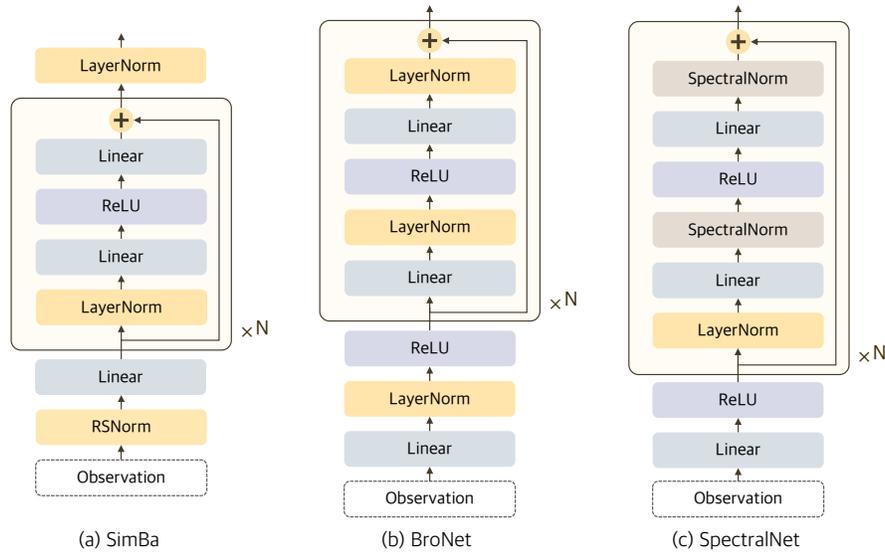


Figure 20: **Architecture Comparison.** Illustration of SimBa, BroNet, and SpectralNet.

G COMPUTATIONAL RESOURCES

The training was performed using NVIDIA RTX 3070, A100, or H100 GPUs for neural network computations and either a 16-core Intel i7-11800H or a 32-core AMD EPYC 7502 CPU for running simulators. Our software environment included Python 3.10, CUDA 12.2, and Jax 4.26.

When benchmarking computation time, experiments were conducted on a same hardware equipped with an NVIDIA RTX 3070 GPU and 16-core Intel i7-11800H CPU.

Table 1: **Environment details.** We list the episode length, action repeat for each domain, total environment steps, and performance metrics used for benchmarking SimBa.

	DMC	MyoSuite	HumanoidBench	Craftax
Episode length	1,000	100	500 - 1,000	—
Action repeat	2	2	2	1
Effective length	500	50	250 - 500	—
Total env. steps	500K-1M	1M	2M	1B
Performance metric	Average Return	Average Success	Average Return	Average Score

H ENVIRONMENT DETAILS

This section details the benchmark environments used in our evaluation. We list all tasks from each benchmark along with their observation and action dimensions. Visualizations of each environment are provided in Figure 6, and detailed environment information is available in Table 1.

H.1 DEEPMIND CONTROL SUITE

DeepMind Control Suite (Tassa et al., 2018, DMC) is a standard benchmark for continuous control, featuring a range of locomotion and manipulation tasks with varying complexities, from simple low-dimensional tasks ($s \in \mathbb{R}^3$) to highly complex ones ($s \in \mathbb{R}^{223}$). We evaluate 27 DMC tasks, categorized into two groups: DMC-Easy&Medium and DMC-Hard. All Humanoid and Dog tasks are classified as DMC-Hard, while the remaining tasks are grouped under DMC-Easy&Medium. The complete lists of DMC-Easy&Medium and DMC-Hard are provided in Tables 2 and 3, respectively.

For our URL experiments, we adhere to the protocol in Park et al. (2023), using an episode length of 400 steps and augmenting the agent’s observations with its x , y , and z coordinates. For the representation function ϕ (i.e., reward network) in METRA, we only used x , y , and z positions as input. This approach effectively replaces the colored floors used in the pixel-based humanoid setting of METRA with state-based inputs.

H.2 MYOSUITE

MyoSuite (Caggiano et al., 2022) simulates musculoskeletal movements with high-dimensional state and action spaces, focusing on physiologically accurate motor control. It includes benchmarks for complex object manipulation using a dexterous hand. We evaluate 10 MyoSuite tasks, categorized as “easy” when the goal is fixed and “hard” when the goal is randomized, following Hansen et al. (2023). The full list of MyoSuite tasks is presented in Table 4.

H.3 HUMANOIDBENCH

HumanoidBench (Sferrazza et al., 2024) is a high-dimensional simulation benchmark designed to advance humanoid robotics research. It features the Unitree H1 humanoid robot with dexterous hands, performing a variety of challenging locomotion and manipulation tasks. These tasks range from basic locomotion to complex activities requiring precise manipulation. The benchmark facilitates algorithm development and testing without the need for costly hardware by providing a simulated platform. In our experiments, we focus on 14 locomotion tasks, simplifying the setup by excluding the dexterous hands. This reduces the complexity associated with high degrees of freedom and intricate dynamics. All HumanoidBench scores are normalized based on each task’s target success score as provided by the authors. A complete list of tasks is available in Table 5.

H.4 CRAFTAX

Craftax (Matthews et al., 2024) is an open-ended RL environment that combines elements from Crafter (Hafner, 2022) and NetHack (Küttler et al., 2020). It presents a challenging scenario requiring the sequential completion of numerous tasks. A key feature of Craftax is its support for vectorized parallel environments and a full end-to-end GPU learning pipeline, enabling a large number

1296 of environment steps at low computational cost. The original baseline performance in Craftax is
 1297 reported as a percentage of the maximum score (226). In our experiments, we report agents' raw
 1298 average scores instead.

1300 Table 2: **DMC Easy & Medium.** We consider a total of 20 continuous control tasks for the DMC
 1301 Easy & Medium benchmark. We list all considered tasks below and baseline performance for each
 1302 task is reported at 500K environment steps.

1303	Task	Observation dim	Action dim
1304			
1305	Acrobot Swingup	6	1
1306	Cartpole Balance	5	1
1307	Cartpole Balance Sparse	5	1
1308	Cartpole Swingup	5	1
1309	Cartpole Swingup Sparse	5	1
1310	Cheetah Run	17	6
1311	Finger Spin	9	2
1312	Finger Turn Easy	12	2
1313	Finger Turn Hard	12	2
1314	Fish Swim	24	5
1315	Hopper Hop	15	4
1316	Hopper Stand	15	4
1317	Pendulum Swingup	3	1
1318	Quadruped Run	78	12
1319	Quadruped Walk	78	12
1320	Reacher Easy	6	2
1321	Reacher Hard	6	2
1322	Walker Run	24	6
1323	Walker Stand	24	6
1324	Walker Walk	24	6

1325 Table 3: **DMC Hard.** We consider a total of 7 continuous control tasks for the DMC Hard bench-
 1326 mark. We list all considered tasks below and baseline performance for each task is reported at 1M
 1327 environment steps.

1328	Task	Observation dim	Action dim
1329			
1330	Dog Run	223	38
1331	Dog Trot	223	38
1332	Dog Stand	223	38
1333	Dog Walk	223	38
1334	Humanoid Run	67	24
1335	Humanoid Stand	67	24
1336	Humanoid Walk	67	24

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1350 Table 4: **MyoSuite**. We consider a total of 10 continuous control tasks for the MyoSuite benchmark,
 1351 which encompasses both fixed-goal ('Easy') and randomized-goal ('Hard') settings. We list all
 1352 considered tasks below and baseline performance for each task is reported at 1M environment steps.

1353	Task	Observation dim	Action dim
1354	Key Turn Easy	93	39
1355	Key Turn Hard	93	39
1356	Object Hold Easy	91	39
1357	Object Hold Hard	91	39
1358	Pen Twirl Easy	83	39
1359	Pen Twirl Hard	83	39
1360	Pose Easy	108	39
1361	Pose Hard	108	39
1362	Reach Easy	115	39
1363	Reach Hard	115	39

1364
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 1367 Table 5: **HumanoidBench**. We consider a total of 14 continuous control locomotion tasks for the
 1368 UniTree H1 humanoid robot from the HumanoidBench domain. We list all considered tasks below
 1369 and baseline performance for each task is reported at 2M environment steps.

1371	Task	Observation dim	Action dim
1372	Balance Hard	77	19
1373	Balance Simple	64	19
1374	Crawl	51	19
1375	Hurdle	51	19
1376	Maze	51	19
1377	Pole	51	19
1378	Reach	57	19
1379	Run	51	19
1380	Sit Simple	51	19
1381	Sit Hard	64	19
1382	Slide	51	19
1383	Stair	51	19
1384	Stand	51	19
1385	Walk	51	19

1386 Table 6: **Craftax**. We consider the symbolic version of Craftax. Baseline performance is reported
 1387 at 1B environment steps.

1389	Task	Observation dim	Action dim
1390	Craftax-Symbolic-v1	8268	43

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I HYPERPARAMETERS

I.1 OFF-POLICY

Table 7: **SAC+SimBa hyperparameters.** The hyperparameters listed below are used consistently across all tasks when integrating SimBa with SAC. For the discount factor, we set it automatically using heuristics used by TD-MPC2 (Hansen et al., 2023).

Hyperparameter	Value
Critic block type	SimBa Residual
Critic num blocks	2
Critic hidden dim	512
Critic learning rate	1e-4
Target critic momentum (τ)	5e-3
Actor block type	SimBa Residual
Actor num blocks	1
Actor hidden dim	128
Actor learning rate	1e-4
Initial temperature (α_0)	1e-2
Temperature learning rate	1e-4
Target entropy (\mathcal{H}^*)	$ \mathcal{A} /2$
Batch size	256
Optimizer	AdamW
Optimizer momentum (β_1, β_2)	(0.9, 0.999)
Weight decay (λ)	1e-2
Discount (γ)	Heuristic
Replay ratio	2
Clipped Double Q	HumanoidBench: True Other Envs: False

Table 8: **DDPG+SimBa hyperparameters.** The hyperparameters listed below are used consistently across all tasks when integrating SimBa with DDPG. For the discount factor, we set it automatically using heuristics used by TD-MPC2 (Hansen et al., 2023).

Hyperparameter	Value
Critic block type	SimBa Residual
Critic num blocks	2
Critic hidden dim	512
Critic learning rate	1e-4
Target critic momentum (τ)	5e-3
Actor block type	SimBa Residual
Actor num blocks	1
Actor hidden dim	128
Actor learning rate	1e-4
Exploration noise	$\mathcal{N}(0, 0.1^2)$
Batch size	256
Optimizer	AdamW
Optimizer momentum (β_1, β_2)	(0.9, 0.999)
Weight decay (λ)	1e-2
Discount (γ)	Heuristic
Replay ratio	2

1458 Table 9: **TDMPC2+SimBa hyperparameters.** We provide a detailed list of the hyperparameters
 1459 used for the shared encoder module in TD-MPC2 (Hansen et al., 2023). Aside from these, we
 1460 follow the hyperparameters specified in the original TD-MPC2 paper. The listed hyperparameters
 1461 are applied uniformly across all tasks when integrating SimBa with TD-MPC2.

Hyperparameter	Value
Encoder block type	SimBa Residual
Encoder num blocks	2
Encoder hidden dim	256
Encoder learning rate	1e-4
Encoder weight decay	1e-1

1470 I.2 ON-POLICY

1471 Table 10: **PPO+SimBa hyperparameters.** We provide a detailed list of the hyperparameters when
 1472 integrating SimBa with PPO (Schulman et al., 2017). Aside from these, we follow the hyperparam-
 1473 eters specified in the Craftax paper (Matthews et al., 2024).

Hyperparameter	Value
Critic block type	SimBa Residual
Critic num blocks	2
Critic hidden dim	512
Critic learning rate	1e-4
Actor block type	SimBa Residual
Actor num blocks	1
Actor hidden dim	256
Actor learning rate	1e-4
Optimizer	AdamW
Optimizer momentum (β_1, β_2)	(0.9, 0.999)
Optimizer eps	1e-8
Weight decay (λ)	1e-2

1490 I.3 UNSUPERVISED RL

1491 Table 11: **METRA+SimBa hyperparameters.** We provide a detailed list of the hyperparameters
 1492 when integrating SimBa with METRA (Park et al., 2023). Aside from these, we follow the hyper-
 1493 parameters specified in the original METRA paper.

Hyperparameter	Value
Critic block type	SimBa Residual
Critic num blocks	2
Critic hidden dim	512
Actor block type	SimBa Residual
Actor num blocks	1
Actor hidden dim	128

J LEARNING CURVE

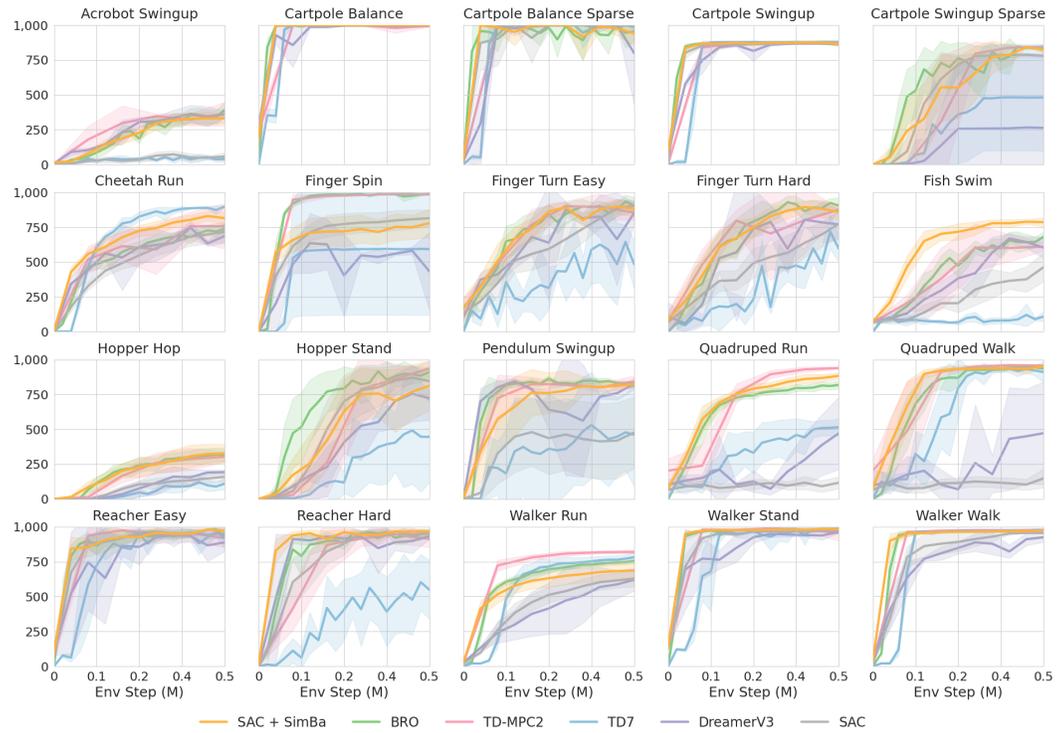


Figure 21: **Per task learning curve for DMC Easy&Medium.** Mean and 95% CI over 10 seeds for SimBa and BRO, 5 seeds for TD7 and SAC, 3 seeds for TD-MPC2 and DreamerV3.

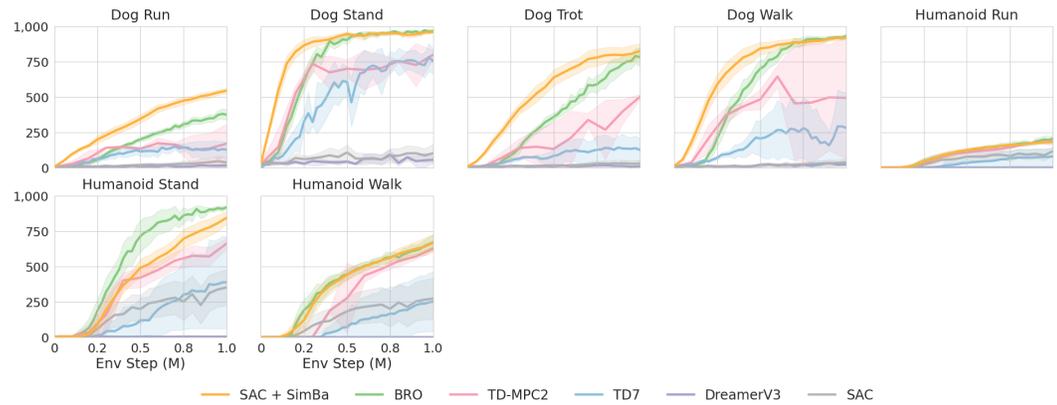


Figure 22: **Per task learning curve for DMC Hard.** Mean and 95% CI over 10 seeds for SimBa and BRO, 5 seeds for TD7 and SAC, 3 seeds for TD-MPC2 and DreamerV3.

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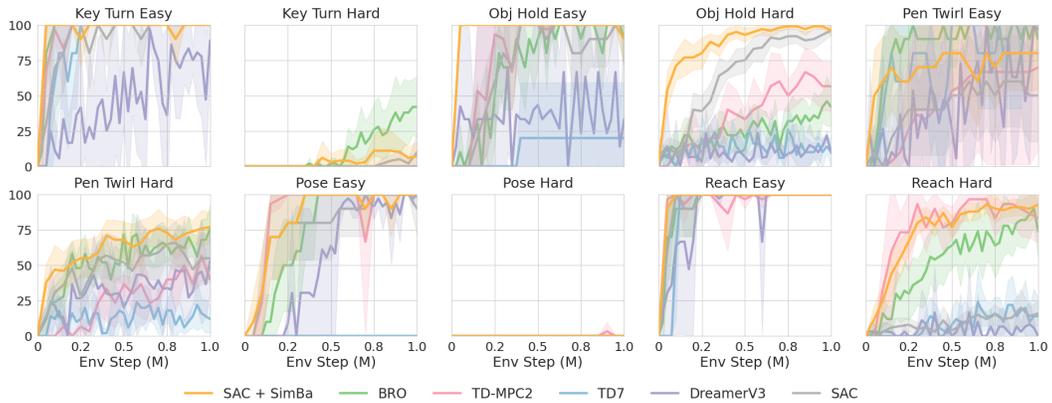


Figure 23: **Per task learning curve for MyoSuite.** Mean and 95% CI over 10 seeds for SimBa and BRO, 5 seeds for TD7 and SAC, 3 seeds for TD-MPC2 and DreamerV3.

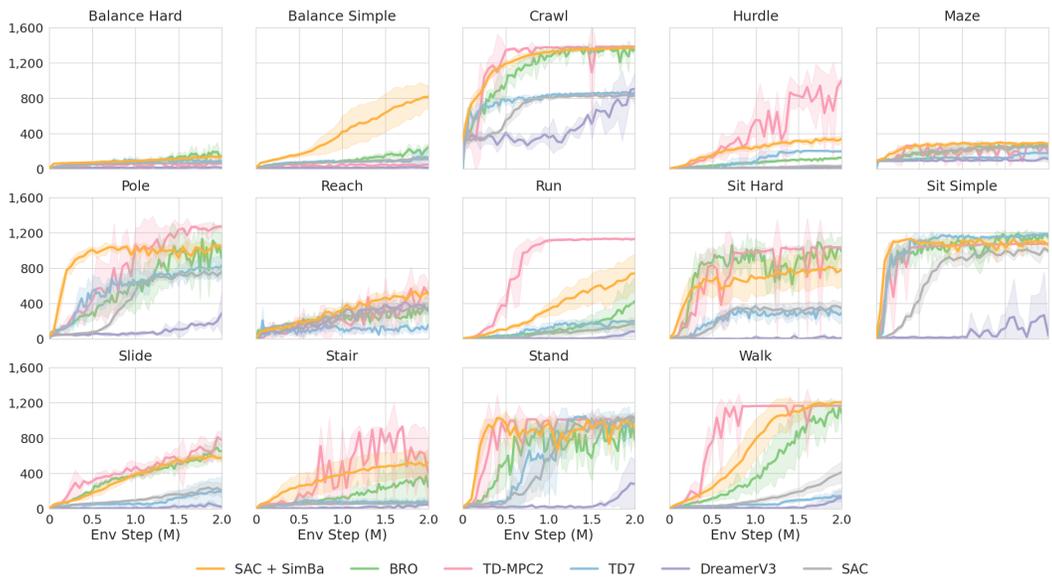


Figure 24: **Per task learning curve for HumanoidBench.** Mean and 95% CI over 10 seeds for SimBa, 5 seeds for BRO, TD7, and SAC, 3 seeds for TD-MPC2 and DreamerV3. We no

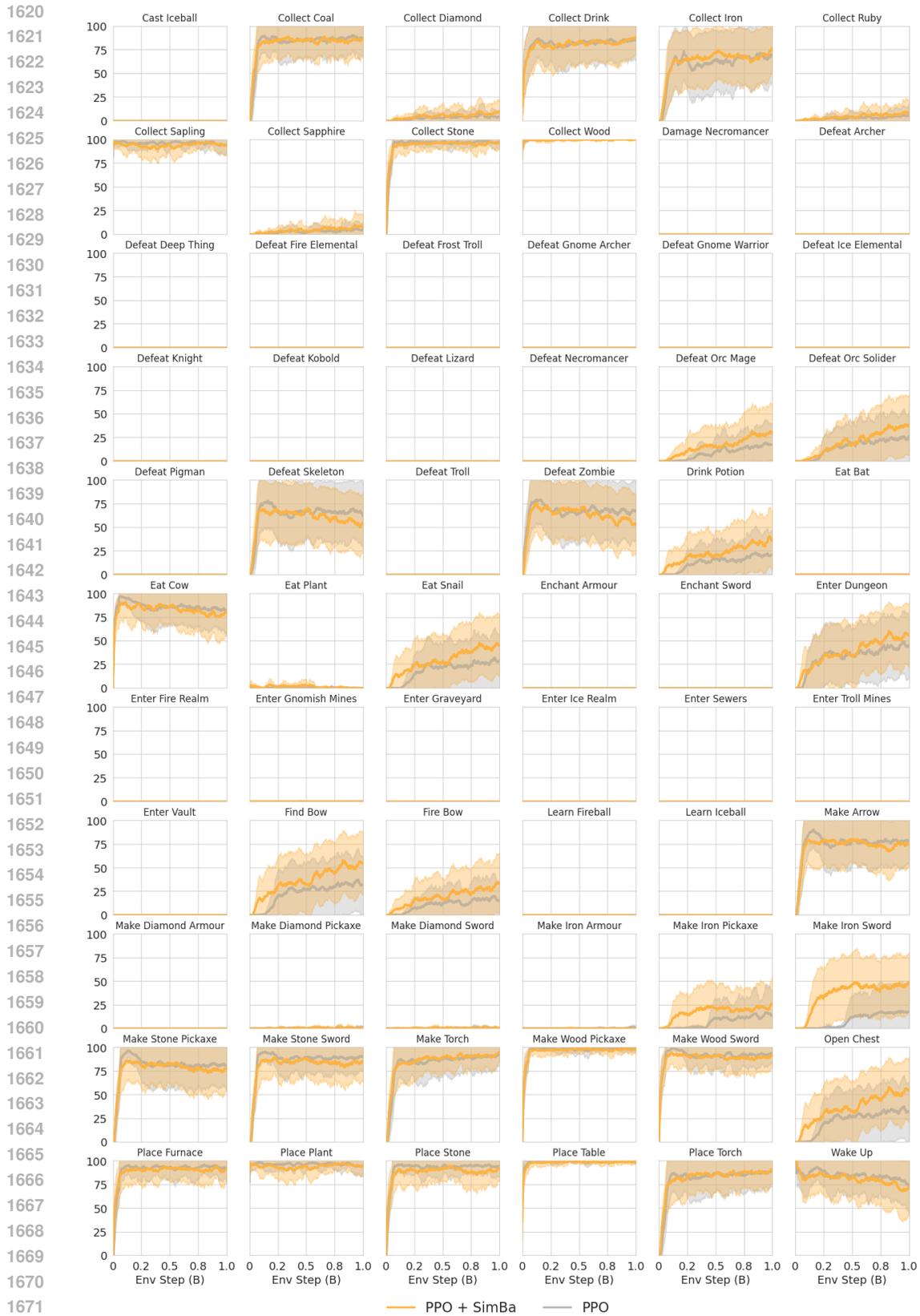


Figure 25: Per task learning curve for Crafttax. We visualize the success rate learning curve for 66 tasks in Crafttax. Mean and 95% CI over 5 seeds for PPO + SimBa and PPO.

K FULL RESULT

Table 12: **Per task results for DMC Easy&Medium.** Results for SimBa and BRO are averaged over 10 seeds, for TD7 and SAC over 5 seeds, and for TD-MPC2 and DreamerV3 over 3 seeds.

Method	SimBa	BRO	TD7	SAC	TD-MPC2	DreamerV3
Acrobot Swingup	331.57	390.78	39.83	57.56	361.07	360.46
Cartpole Balance	999.05	998.66	998.79	998.79	993.93	994.95
Cartpole Balance Sparse	940.53	954.21	988.58	1000.00	1000.00	800.25
Cartpole Swingup	866.53	878.69	878.09	863.24	876.07	863.90
Cartpole Swingup Sparse	823.97	833.18	480.74	779.99	844.77	262.69
Cheetah Run	814.97	739.67	897.76	716.43	757.60	686.25
Finger Spin	778.83	987.45	592.74	814.69	984.63	434.06
Finger Turn Easy	881.33	905.85	485.66	903.07	854.67	851.11
Finger Turn Hard	860.22	905.30	596.32	775.21	876.27	769.86
Fish Swim	786.73	680.34	108.84	462.67	610.23	603.34
Hopper Hop	326.69	315.04	110.27	159.43	303.27	192.70
Hopper Stand	811.75	910.88	445.50	845.89	936.47	722.42
Pendulum Swingup	824.53	816.20	461.40	476.58	841.70	825.17
Quadruped Run	883.68	818.62	515.13	116.91	939.63	471.25
Quadruped Walk	952.96	936.06	910.55	147.83	957.17	472.31
Reacher Easy	972.23	933.77	920.38	951.80	919.43	888.36
Reacher Hard	965.96	956.52	549.50	959.59	913.73	935.25
Walker Run	687.16	754.43	782.32	629.44	820.40	620.13
Walker Stand	983.02	986.58	984.63	972.59	957.17	963.28
Walker Walk	970.73	973.41	976.58	956.67	978.70	925.46
IQM	885.70	888.02	740.19	799.75	895.47	748.58
Median	816.78	836.62	623.18	676.59	836.93	684.18
Mean	823.12	833.78	636.18	679.42	836.34	682.16
OG	0.1769	0.1662	0.3638	0.3206	0.1637	0.3178

Table 13: **Per task results for DMC Hard.** Results for SimBa and BRO are averaged over 10 seeds, for TD7 and SAC over 5 seeds, and for TD-MPC2 and DreamerV3 over 3 seeds.

Method	SimBa	BRO	TD7	SAC	TD-MPC2	DreamerV3
Dog Run	544.86	374.63	127.48	36.86	169.87	15.72
Dog Stand	960.38	966.97	753.23	102.04	798.93	55.87
Dog Trot	824.69	783.12	126.00	29.36	500.03	10.19
Dog Walk	916.80	931.46	280.87	38.14	493.93	23.36
Humanoid Run	181.57	204.96	79.32	116.97	184.57	0.91
Humanoid Stand	846.11	920.11	389.80	352.72	663.73	5.12
Humanoid Walk	668.48	672.55	252.72	273.67	628.23	1.33
IQM	773.28	771.50	216.04	69.03	527.11	9.63
Median	706.39	694.20	272.62	159.36	528.26	17.13
Mean	706.13	693.40	287.06	135.68	491.33	16.07
OG	0.2939	0.3066	0.7129	0.8643	0.5087	0.9839

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Table 14: **Per task results for MyoSuite.** Results for SimBa and BRO are averaged over 10 seeds, for TD7 and SAC over 5 seeds, and for TD-MPC2 and DreamerV3 over 3 seeds.

Method	SimBa	BRO	TD7	SAC	TD-MPC2	DreamerV3
Key Turn Easy	100.00	100.00	100.00	100.00	100.00	88.89
Key Turn Hard	7.00	42.00	0.00	10.00	0.00	0.00
Object Hold Easy	90.00	90.00	20.00	90.00	100.00	33.33
Object Hold Hard	96.00	42.00	10.00	96.00	56.67	9.44
Pen Twirl Easy	80.00	90.00	100.00	50.00	70.00	96.67
Pen Twirl Hard	77.00	76.00	12.00	55.00	40.00	53.33
Pose Easy	100.00	100.00	0.00	90.00	100.00	100.00
Pose Hard	0.00	0.00	0.00	0.00	0.00	0.00
Reach Easy	100.00	100.00	100.00	100.00	100.00	100.00
Reach Hard	93.00	74.00	14.00	16.00	83.33	0.00
IQM	95.20	87.60	22.31	71.40	77.50	46.56
Median	77.00	72.00	34.00	62.50	65.00	47.00
Mean	74.30	71.40	35.60	60.70	65.00	48.17
OG	99.93	99.93	99.96	99.94	99.94	99.95

Table 15: **Per task results for HumanoidBench.** Results for SimBa are averaged over 10 seeds, for BRO, TD7 and SAC over 5 seeds, and for TD-MPC2 and DreamerV3 over 3 seeds.

Method	SimBa	BRO	TD7	SAC	TD-MPC2	DreamerV3
Balance Hard	137.20	145.95	79.90	69.02	64.56	16.07
Balance Simple	816.38	246.57	132.82	113.38	50.69	14.09
Crawl	1370.51	1373.83	868.63	830.56	1384.40	906.66
Hurdle	340.60	128.60	200.28	31.89	1000.12	18.78
Maze	283.58	259.03	179.23	254.38	198.65	114.90
Pole	1036.70	915.89	830.72	760.78	1269.57	289.18
Reach	523.10	317.99	159.37	347.92	505.61	341.62
Run	741.16	429.74	196.85	168.25	1130.94	85.93
Sit Hard	783.95	989.07	293.96	345.65	1027.47	9.95
Sit Simple	1059.73	1151.84	1183.26	994.58	1074.96	40.53
Slide	577.87	653.17	197.07	208.46	780.82	24.43
Stair	527.49	249.86	77.19	65.53	398.21	49.04
Stand	906.94	780.52	1005.54	1029.78	1020.59	280.99
Walk	1202.91	1080.63	143.86	412.21	1165.42	125.59
IQM	747.43	558.03	256.77	311.46	885.67	72.30
Median	733.09	632.93	385.41	399.93	796.18	160.49
Mean	736.30	623.05	396.33	402.31	787.64	165.55
OG	0.3197	0.4345	0.6196	0.6016	0.2904	0.8352