

Bilinear Convolution Decomposition for Causal RL Interpretability

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Abstract

Efforts to interpret reinforcement learning (RL) models tend to target the activation space, and fewer recent studies target the weight space. Here we use a dual framework of both the weight and activation spaces in order to interpret and intervene in a RL network. To enhance RL interpretability, we enable linear decomposition via linearization of an IMPALA network : we replace nonlinear activation functions in both convolution and fully connected layers with bilinear variants (we term BIMPALA). Previous work on MLPs have shown that bilinearity enables quantifying functional importance through weight-based eigendecomposition to identify interpretable low rank structure [Pearce et al., 2024b]. By extending existing MLP decomposition techniques to convolution layers, we are able to analyze channel and spatial dimensions separately through singular value decomposition. We find BIMPALA networks to be feasible and competitive, as they perform comparably to their ReLU counterparts when we train them on various ProcGen games. Importantly, we find the bilinear approach in combination with activation-based probing provide advantages for interpretability and agent control. In a maze-solving agent, we find a set of orthonormal eigenvectors (we term *eigenfilters*), the top-2 of which act as cheese (solution target) detectors, and another pair of eigenfilters we can manipulate to control the policy.

Introduction

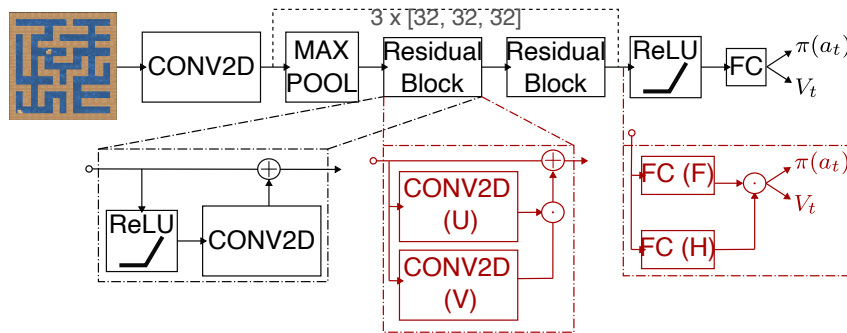


Figure 1: BIMPALA: a simplified IMPALA architecture (black) modified by replacing ReLU operations with bilinear gating (red) for both the convolution (CONV2D; Equation 3) and fully connected (FC; Equation 1) layers.

While recent advances in reinforcement learning have produced increasingly capable reasoning agents [Mnih et al., 2013, Gu et al., 2017, Baker et al., 2019], analyzing their internal mechanisms has

proven difficult. This challenge is particularly pronounced in multi-step reasoning tasks, where the relationship between model architecture and computational strategy is often opaque. Additionally, there is a general notion of the performance-interpretability trade-off [Assis et al., 2025], which argues that more transparent models tend to have lower performance.

In this work, we demonstrate that increased interpretability need not come at the cost of performance. We explore an approach embedded within mechanistic interpretability. Mechanistic interpretability has emerged as a promising framework for understanding neural networks by identifying and analyzing features - specific directions in activation space that encode meaningful computational patterns [Cunningham et al., 2023, Trenton Bricken et al., 2023, Adly Templeton et al., 2024, Rinsky et al., 2024]. Traditional approaches have focused primarily on activation patterns during inference, but recent work suggests that analyzing model weights directly may provide complementary insights.

Our work explores a subset of models where nonlinearities are replaced with linear counterparts. Bilinear MLPs [Dauphin et al., 2017] offer an architectural innovation that enables direct interpretation of model weights. While initially proposed for language modeling tasks [Pearce et al., 2024b], we show their benefits extend to understanding an agent’s spatial decision-making. As proof of concept that the bilinear approach can indeed benefit interpretability of RL models, we simplified a common RL agent, IMPALA [Espeholt et al., 2018], and compared it with its bilinear counterpart (Figure 1).

We argue the importance of studying weights and activations jointly. By analyzing both the weight space through eigenfilters and the activation space through targeted probes, we find interpretable features that track specific computational steps, from interpretable convolution features to the action features. Additionally, we find that while standard basis analyses can appear informative, they often mask the true computational structure of the network. Instead, we show that bases informed by action spaces and targeted probes provide more reliable insights into model behavior during multi-step tasks.

Our contributions (1) We introduce a bilinear architectures (BIMPALA) for RL and show that it trains well in "easy" ProcGen environments. (2) We show how bilinear convolution layers can be decomposed into bases of self interacting eigenfilters. (3) We show that the standard basis is often non-interpretable and less informative compared to basis derived from probes or the action/logits space. (4) We propose new techniques using weights alongside activations to analyze mechanisms in bilinear convolution networks. We validate our approach by finding a cheese detector on a maze solving agent and re-targeting the agent towards counterfactual cheese positions.

Background

The core benefit of the bilinear approach hinges on removing nonlinearities from the neural network, allowing spectral decomposition. Spectral decomposition of MLPs has revealed interpretable low-rank structure across multiple tasks Pearce et al. [2024a], and we extend this approach to convolution layers for an RL agent.

In this section, we briefly review multi-layer perceptrons (MLPs), convolutions, and bilinear gating. Throughout, we denote scalars like s , vectors like \mathbf{v} , matrices like M , tensors like \mathbf{T} , dot product with \cdot , pointwise product with \odot , and convolution with $*$.

Bilinear MLPs A conventional MLP is composed of 3 (or more) fully-connected (FC) layers, where inputs are up-projected into a hidden layer and then down-projected into the output layer. The hidden activations of a conventional MLP can be characterized as a $\mathbb{R}^n \rightarrow \mathbb{R}^m$ encoder which takes input \mathbf{x} and applies a learned linear transformation, with weights W and bias \mathbf{b} , followed by an activation function σ .

Modern models, such as LLMs, feature an encoder variant called a Gated Linear Unit (GLU), comprised of the pointwise product of *two* linear up-projections, with learned weights F and G , and only one of the projections is passed through an activation function. Omitting biases for simplicity,

$$\text{Enc}_{\text{GLU}}(\mathbf{x}, F, H) = \sigma(\mathbf{x}F) \odot (\mathbf{x}H)$$

Bilinear encoders, and our bilinear FC in Figure 1, use an identity activation, keeping the overall transformation linear:

$$\text{FC}_{\text{Bilinear}}(\mathbf{x}, F, H) = (\mathbf{x}F) \odot (\mathbf{x}H) \tag{1}$$

This linearization allows spectral decomposition of the weights and activations, which can have interpretable value [Pearce et al., 2024b]. Importantly, Pearce et al. [2024b] show bilinear MLPs can be expressed as a third order tensor \mathbf{B} , comprised of interaction matrices for each output dimension, parameterizing the interactions between pairs of inputs. In Decomposing Convolutions, we provide an analog \mathbf{B} for convolution layers.

Convolution layers A 2D convolution layer (Conv2D) takes an input \mathbf{X} of shape $[\text{width}, \text{height}, c_{in}]$ where c_{in} is the number of input channels, and applies a learned kernel \mathbf{U} of width k with stride s followed by a pointwise activation function σ :

$$\text{Conv2D}(\mathbf{X}, \mathbf{U}) = \sigma(\mathbf{X} * \mathbf{U})$$

With $s = 1$, Conv2D outputs a tensor of shape $[\text{width}, \text{height}, c_{out}]$, where c_{out} is the number of output channels. Kernel \mathbf{U} of shape $[k, k, c_{in}, c_{out}]$ acts locally on $k \times k$ patches, and we denote the kernel for a given output channel i as $\mathbf{U}^{(i)}$. Assuming an identity activation and letting $\ell = \lfloor \frac{k}{2} \rfloor$, kernel weights $\mathbf{U}^{(i)}$, as illustrated in Appendix A Figure 12 (top left in blue), act on a local patch around spatial coordinates (α, β) via:

$$u(\alpha, \beta, i) = \sum_{j=1}^{c_{in}} \sum_{|k_1| \leq \ell} \sum_{|k_2| \leq \ell} U^{(i)}[j, k_1, k_2] \cdot X[j, \alpha + k_1, \beta + k_2] \quad (2)$$

Here, $u(\alpha, \beta, i)$ is a scalar, denoting output channel i 's entry at spatial location (α, β) , while $U^{(i)}[j, k_1, k_2]$ and $X[j, \alpha + k_1, \beta + k_2]$ are row and column vectors from $k \times k$ matrices representing the kernel and current input patch respectively for a single input-output channel combination.

Analogous to a bilinear FC (Equation 1), a bilinear convolution layer (BConv2D) would then require two convolutions. Assuming kernels \mathbf{U} and \mathbf{V} ,

$$\text{BConv2D}(\mathbf{x}, \mathbf{U}, \mathbf{V}) = (\mathbf{x} * \mathbf{U}) \odot (\mathbf{x} * \mathbf{V}) \quad (3)$$

Decomposing Convolutions

The main advantage of adopting the bilinear form for a convolution layer is decomposition into sets of orthonormal eigenvectors for each output channel, which we call *eigenfilters*. Analogous to Pearce et al. [2024b], we can express a BConv2D layer as a tensor \mathbf{B} , comprised of interaction matrices (B) for each (scalar) output. Specifically, B parameterizing the input channel interactions between pairs of inputs at a single spatial location for a single output channel (α, β, i) (Figure 12 in Appendix A).

Importantly, spectral decomposition is easily achievable because B has a symmetric form B^{sym} . In Appendix A, we derive B^{sym} for convolution layers and show it is equivalent to B . In short, for each of c_{out} output channels, we get a matrix B^{sym} of dimension $k^2 c_{in} \times k^2 c_{in}$. Hence, each spatial location of the input image contributes to \mathbf{B} with shape $[c_{out}, k^2 c_{in}, k^2 c_{in}]$.

Bilinear component decomposition protocol Similar to the decomposition approach in [Pearce et al., 2024b], we can fix an output vector $\mathbf{z} \in \mathbb{R}^{c_{out}}$ and multiply it by B^{sym} along the output channel dimension to produce a matrix $Q^z = \mathbf{z} B^{sym}$ of shape $[k^2 c_{in}, k^2 c_{in}]$, that functions as a quadratic form on the input space. Since convolution layers are *locally* connected rather than fully connected, the output vector \mathbf{z} is in output channel space, and the decomposition produces an eigenbasis for the filters that we call *eigenfilters*. That is, you get a basis consisting of $k^2 c_{in}$ *eigenfilters* of shape $[k, k, c_{in}]$. In spectral theorem terminology, we have $Q^z = F^T \Lambda F$, where F is an orthonormal matrix (satisfying $F^{-1} = F^T$) of eigenvectors, and Λ is a real, diagonal matrix of eigenvalues.

More generally, we can transform the entirety of the \mathbf{B} tensor to change the channel basis to another. For example, we may find that another basis of the output channels is more informative than the standard basis. In order to capture this, we can rotate the basis with a rotation matrix R . Briefly, just as we can compute individual interaction matrices Q^z for a vector $\mathbf{z} \in \mathbb{R}^{c_{out}}$, we can compute a full basis and multiply B^{sym} by R to get B_R^{sym} that operates in the basis rotated with R .

Contributions of Eigenfilters Since Q^z is used in practice as a quadratic form, its contributions towards \mathbf{z} for a flattened patch $\mathbf{x}_{\text{patch}}$ centered around a given position are given by $Q^z(\mathbf{x}_{\text{patch}}) = \mathbf{x}_{\text{patch}}^T Q^{\mathbf{x}_{\text{patch}}} \mathbf{x}_{\text{patch}} = \mathbf{x}_{\text{patch}}^T F_{\mathbf{z}}^T \Lambda_{\mathbf{z}} F_{\mathbf{z}} \mathbf{x}_{\text{patch}} = (F_{\mathbf{z}} \mathbf{x}_{\text{patch}})^T \Lambda_{\mathbf{z}} (F_{\mathbf{z}} \mathbf{x}_{\text{patch}}) = \sum_i \lambda_{\mathbf{z}}^i (f_{\mathbf{z}}^i \mathbf{x}_{\text{patch}})^2$ and $Q^z = \sum_i \lambda_{\mathbf{z}}^i f_{\mathbf{z}}^i f_{\mathbf{z}}^{iT}$. Each f^i is an individual eigenfilter, and has shape $[k, k, c_{in}]$. As the eigenfilter activations are applied to every valid position uniformly, we can equivalently write $Q^z(X) = \sum_i \lambda_{\mathbf{z}}^i (f_{\mathbf{z}}^i * X)^2$. Note that when applying an eigenfilter to activations we treat the eigenfilter as a regular convolution filter.

Separating channels from spatial coordinates with SVD Given a weight or activation vector A with shape $[\text{width}, \text{height}, c_{out}]$, having both spatial and channel dimensions, A can be reshaped into $[c_{out}, \text{width} \cdot \text{height}]$ and decomposed via SVD:

$$A = S \Sigma V^T = \sum_i \sigma_i s_i v_i^T$$

where S has shape $[c_{out}, c_{out}]$, and V has shape $[\text{width} \cdot \text{height}, \text{width} \cdot \text{height}]$. The top left singular vectors s_i live in the channel space, and can be used as output vectors for a BConv2D layer.

Since the top singular vectors in channel space also have a singular value, we can aggregate the contributions of the eigenvalues and the eigenvectors together. We can derive an eigendecomposition of a BConv2D layer for each singular channel, to get the following:

$$\begin{aligned} Q^{\text{probe}}(A) &= \sum_{j=1}^{c_{out}} s_j Q^{\mathbf{z}_j}(A) \\ &= \sum_{j=1}^{c_{out}} s_j \sum_i \lambda_{\mathbf{z}_j}^i (f_{\mathbf{z}_j}^i * A)^2 \\ &= \sum_{j=1}^{c_{out}} \sum_i (s_j \lambda_{\mathbf{z}_j}^i) (f_{\mathbf{z}_j}^i * A)^2 \end{aligned}$$

The importance of an eigenfilter for its singular channel is parameterized by the joint term $s_j \lambda_{\mathbf{z}_j}^i$. Note that $s_j \lambda_{\mathbf{z}_j}^i$ is signed, as the eigenvalues can be negative.

Experiments

In order to evaluate the usefulness of the bilinear approach in the context of RL, we ran a series of experiments. We detail training procedures, experimental protocols, and key findings from both quantitative and qualitative perspectives. We ask (1) do bilinear architectures achieve competitive performance compared to standard models like ReLU-based IMPALA and (2) do bilinear layers provide interpretable representations through spectral decomposition and probe-based analyses?

In order establish feasibility, we first evaluate and compare performance between BIMPALA and IMPALA on a handful or randomly selected "easy" ProcGen environments.

We next train probes and propose a protocol to decompose probes in conjunction with convolution layers. This allows us to identify a cheese filter using the top-2 eigenfilters of a convolution layer.

We next explore methods without the need for training probes. First, we turn to the standard channel bases and perform ablation experiments. Unfortunately, we do not find the standard bases alone to be informative enough for interpretability.

We then decide to adopt two different approaches using both weights and activations without training probes. First, we decompose the full connected layer along each policy action and perform ablation experiments. We find action features to be faithful to actions needed to solve the maze. Finally, we perform steering experiments where we re-target the agent towards a counterfactual cheese position.

BIMPALA matches IMPALA performance

Architecture baseline We adapted the existing IMPALA framework Espeholt et al. [2018] by (1) simplifying the network by removing some convolution layers so that the residual block is a simple

gated convolution with a skip connection as well as removing the LSTM layer after the FC layer and (2) modifying the original structure to incorporate bilinear gating mechanisms in both Conv2D and FC layers (Figure 1). We refer to the bilinear variant as BIMPALA (Bilinear IMPALA).

Evaluation As proof of concept for the bilinear approach to RL, we picked a simplified architecture and trained on tasks within a simple established benchmark, the ProcGen environment [Cobbe et al., 2020], with proximal policy optimization (PPO) as PPO tends to be effective and easy to tune [Schulman et al., 2017]. We trained our simplified IMPALA and BIMPALA alongside each other on the "easy" distributions, which are computationally inexpensive and converge in less time steps than harder distributions, for a handful of environments, including **Maze**, **Heist**, **Plunder**, and **DodgeBall**. For full training parameters, see Table 1 in Appendix B.

We find BIMPALA matches and occasionally outperforms IMPALA across environments we tested (Figure 2), validating the feasibility of using bilinear layers for RL tasks. Specifically, BIMPALA generally demonstrates faster learning, higher final performance in terms of expected return, and maintaining lower entropy.

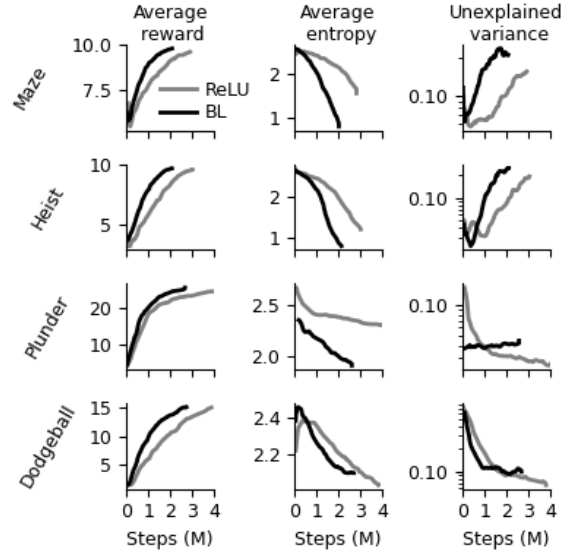


Figure 2: Performance comparison between ReLU and Bilinear IMPALA across different ProcGen environments.

Protocol to enhance interpretability for RL with probes

Having established that the bilinear approach can perform competitively in RL environments, we next want to use this architecture to enhance interpretability. We suggest a protocol to connect bottom-up mechanistic approaches to top-down concept based approaches.

1. Train a linear probe for a concept of interest on a Conv2D activation space with shape $[\text{width}, \text{height}, c_{out}]$, reshaped as $[c_{out}, \text{width} \cdot \text{height}]$
2. Rewrite the probe's weights using SVD, and use the top left channel-space singular vectors as output directions for a last BConv2D layer. Determine the number of singular components needed, based on the distribution of singular values.
3. Perform an eigendecomposition towards the top left singular vectors in channel space, to identify directions in the filter weights that write to the probe (similar to Pearce et al. [2024b]).

This protocol will yield a full basis of eigenvectors for each output direction. Note that it's possible for the important eigenvectors between output directions to not be fully orthogonal, especially if interpreting multiple probes in parallel. Although we do not investigate overlapping filters here, analyzing the cosine similarity between important eigenvectors relating to different singular channels, where importance is measured by $|s_j \lambda_{u_j}^i|$, may further inform the function of the eigenvectors.

Training concept probes With the protocol defined, the next step is to implement it by training concept probes for specific features and analyzing their decomposition. For the remainder of the paper, we focus on ProcGen's Maze environment, where the player, a mouse, must navigate a maze to find the sole piece of cheese and earn a reward. We trained linear probes to detect the presence of the cheese at position (8, 14) in the maze by creating a dataset comprising 2000 mazes with the cheese at position (8, 14) and 2000 mazes without a cheese.

We see that probes trained on the outputs of the residual blocks get about 99% accuracies and F_1 scores (Table 2 in Appendix C).

Dominant singular probe channels Next, we apply singular value decomposition to the probes. The top singular component alone explains 30% of the variance, and 16 components are needed to explain $\geq 90\%$ of the variance (Figure 3).

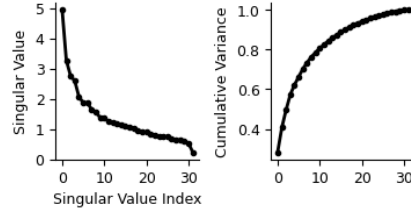


Figure 3: Singular values and explained variance for cheese probe

Eigenfilter decomposition for singular probe channels We then decomposed the last BConv2D layer towards the top singular channel. Whereas the channel basis spectra have just two eigenvalues, indicating that it may not be an informative basis, the singular spectrum is nondegenerate, and hence is more likely to be informative about the task. (Figure 4).

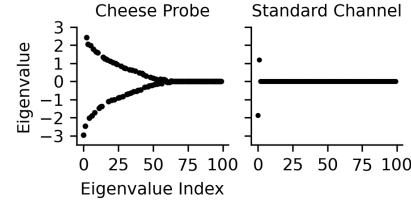


Figure 4: Last BConv2D’s eigenfilter spectrums towards cheese probe’s top singular channel (left) and in the standard channel basis (right)

In order to verify the importance of the singular spectrum for solving the maze, we repeated the decomposition for the first and second BConv2D layers and performed ablation experiments. In these and all following ablation experiments, we ensure that we run each set of ablations on the same mazes by using 2020 seeded environments, capping steps per rollout at 200200 to save runtime. We ablated all but the topk eigenfilters for each output channel of each of the BConv2D layers. We reconstructed our networks and ran each of ablated models in the Maze environment (Figure 5).

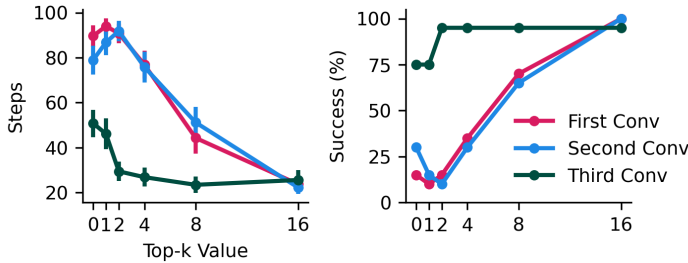


Figure 5: Maze metrics when ablating parts (bottom - $k+1$) of different convolution layers’ eigenfilter spectrum associated with the cheese probe. Here and subsequent plots, error bars are SEM

We find that maze performance recovers close to 100% with just the top-2 eigenfilters in the last BConv2D layer. In this last convolution layer, additional eigenfilters help solve the maze in less steps. For the first and second BConv2D layers, we see a different trend, where it takes 16 eigenfilters for maze performance to be recovered, and the contributions of each added eigenfilter is less step-like and more continuous. This adds credence to the notion that decomposing the layers towards the cheese probe’s top singular channel is informative.

Furthermore, we find information about the cheese location from even just the top positive eigenfilter. In Figure 6, we visualize the top positive and negative eigenfilter activations for a set of pairs of mazes, one with the cheese at the selected position and the other without the cheese. While the positive filter activates on non-cheese patterns, the negative filter down-weights non-cheese patterns without erasing the cheese activation. The positive and negative activations of the respective filters result in a cheese detector filter.

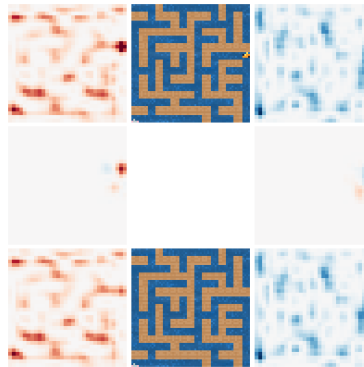


Figure 6: Activations for the top positive (left) and negative (right) eigenfilters in the second BConv2D layer, for the cheese probe’s top singular channel. Activations for a maze with cheese (top) vs without cheese (bottom). Middle plots show the difference between the activations with and without cheese.

244 Ablation within the standard bases

245 While the probe approach is rather promising and we are
 246 successful in finding a cheese detector in the second con-
 247 volution layer, it may not be feasible nor scalable to train
 248 probes for each feature we may want to interpret. This is
 249 especially true once we move beyond toy-like tasks such
 250 as the Maze environment. Hence, despite the degenerate
 251 spectrum (Figure 4), we turn back to the standard basis.

252 We ask how many eigenfilters are necessary for perfor-
 253 mance? We ablated all but the topk eigenfilters for the FC
 254 layer, all the BConv2D layers, or just the last BConv2D
 255 layer.

256 As we may have predicted from Figure 4, the spectrum
 257 of the last BConv2D layer is not informative and is
 258 marginally necessary for full performance. The agent,
 259 when compared to the BIMPALA, has a similar success
 260 rate, in a relatively low number of extra steps, and receives
 261 similar rewards when we ablate the last BConv2D entirely
 262 (topk=0), (Figure 7).

263 When we ablate all the BConv2D layers together, we see
 264 that the top-2 eigenfilters (per output channel) are sufficient to recover full performance (Figure 7).
 265 Full performance in the FC layer is achieved with the top eigenfilter. This eigenfilter is more important
 266 for performance than all the BConv2D layers combined, resulting in lower success in more steps with
 267 less reward when ablated.

268 While the contribution of the FC layer relative to the BConv2D layer may be an important insight, it
 269 is also expected. We could expect the FC , as the last layer of the network that outputs the policy and
 270 value, to contain most the information about the next step and therefore be the largest contributor
 271 to performance. Beyond that, the standard channel basis *by itself* may not be very fruitful for
 272 decomposing the network for interpretability. For example, while we can deduce that in the last
 273 BConv2D layer, the top positive and negative eigenfilters work together (Figure 4, 7), we do not know
 274 anything more granular.

275 Still, decomposition may be useful for interpretability beyond the standard bases and without training
 276 probes.

277 Interpretability based on action features

278 The FC layer outputs the directions of movement for the policy (UP,
 279 DOWN, RIGHT, LEFT). Instead of training probes, we could alter-
 280 natively decompose the directions relevant for actions directly by
 281 decomposing in the direction of each action output. For example, Figure
 282 8 shows the eigenvalues in the action spectrum for the UP action.
 283 We can see many UP action eigenvectors in the FC layer, with one
 284 very large positive eigenvalue. We next used these action spectra of
 285 the FC layer in ablation experiments.

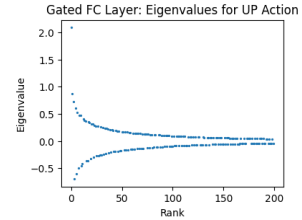


Figure 8: UP spectrum

286 **Ablation** In ablation experiments, we found that, despite the dense
 287 spectrum (Figure 8), preserving the top eigenvector for each action in
 288 the FC layer was sufficient for a 100% success rate (Figure 13 in Appendix E).

289 In Figure 9, we see how a single UP eigenvector proves sufficient to encode the optimal path through
 290 the maze. Specifically, we see that the upward logit values are selectively increased along the solution
 291 path and suppressed near dead ends.

292 Similarly, in Figure 10 we see the effect of the LEFT action spectrum. Note that the mouse, who
 293 is typically located at the bottom left corner at the start of an episode, can solve the maze without
 294 any LEFT eigenvectors (Figure 10, Top 0 LEFT). Yet, we see that increasing the number of LEFT
 295 eigenvectors allows the agent to reach the cheese from other locations, such as at the top. As we

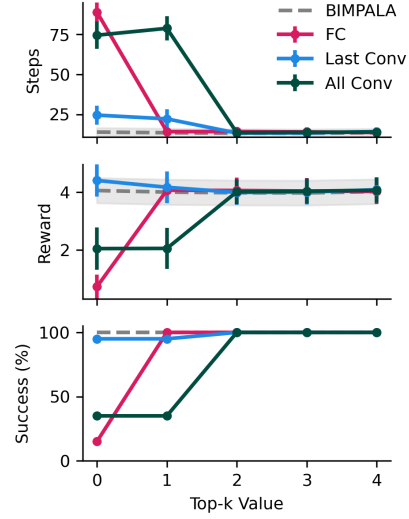


Figure 7: Maze metrics during standard basis ablation.

add more left eigenvectors, we see that the misleading right arrows diminish and the agent gradually reconstructs its left arrows (Figure 10, Diff: Top 4 from ALL LEFT), making the maze solvable for more configurations.

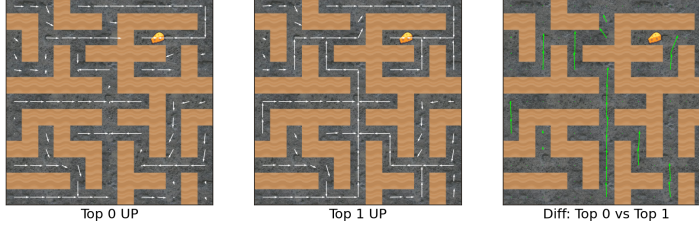


Figure 9: Vector field visualizing maze navigation without any (left) and with a single (middle) UP eigenvector, and the difference highlighted in green (right). The top UP eigenvector is sufficient to solve the maze and without UP eigenvectors, the agent does not move upwards

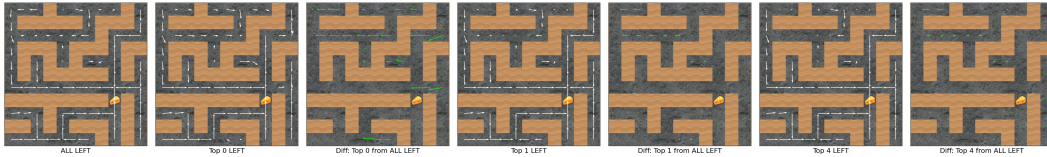
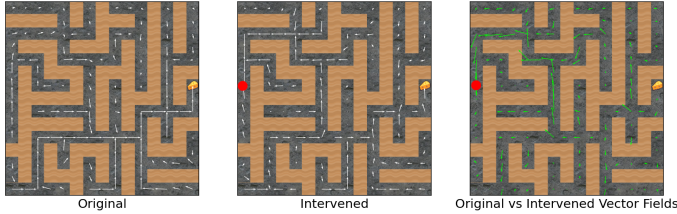


Figure 10: LEFT action spectrum visualization in a maze environment. While the maze is solvable without any LEFT eigenvector, adding LEFT eigenvectors allows solving the maze from other starting positions (e.g. top or right of the cheese)

Steering Experiments



Having seen the effect of action spectra on maze solving and the importance of the FC layer to solving the maze in general, we wondered if it was possible to redirect the agent while leaving the FC intact. In particular, we wondered if we could redirect the agent by intervening in the convolution layers.

Figure 11: Re-targeting the agent by intervening to redirect towards a counterfactual cheese position (red dot)

Rather than averaging activation spaces together, we directly alter the weight contributions from hidden layers (Res in Equation 4). We obtain the activations for the maze’s cheese position (x_{cheese} in Equation 4) by subtracting the activations for the maze without the cheese from the activations for the maze with the cheese. Similarly, we get the activations for the counterfactual cheese position (y_{cheese} in Equation 4) by subtracting the activations for the maze without the cheese from the activations for the maze with the cheese in the counterfactual position. We intervene using the top-2 eigenfilters (eig in Equation 4) and overwrite the contributions, using the equation:

$$Res' = Res - eig * (x_{cheese}) + eig * (y_{cheese}) \quad (4)$$

In essence, we are trying to re-target the agent towards a counterfactual cheese position (Figure 11) on the opposite side of the maze. While the mouse can still solve the maze, we can see that the vector fields indicating movement are altered. Specifically, we can see arrows pointing towards the counterfactual cheese position during intervention (Figure 11, middle). And if we look at the difference between the original and intervened mazes, we can see that the green arrows draw paths away from the real cheese towards the counterfactual cheese position.

Discussion

Summary We introduce an approach to interpreting convolution neural networks, by replacing nonlinearities with bilinear variants that achieve comparable and occasionally superior performance, although this was not our aim. Our approach allows us to find a closed form for self-interacting convolution features that can be combined with a top down concept based approach to derive causally relevant mechanisms used by RL agents in their decision making process. Therefore, we see great value in bilinear variants that offer more interpretability prospects while achieving competitive performance to its non-analytic variants.

Limitations We found significant challenges in interpreting the units of computation in an entirely data independent fashion. Instead, we found that top activating dataset examples for eigenfilters tend not to be informative. Still, the decomposition allows us to break concept probes into more granular units of computation.

We considered only one architecture, IMPALA, for our policy, although we expect the general approach of replacing nonlinearities with bilinear variants to be widely applicable.

Due to computational requirements, we trained on the "easy" mode of a handful of ProcGen environments and we only analyzed the BIMPALA network for interpretability in the context of the Maze environment. It is not clear if the methods we presented here will transfer well to more complex environments with multiple objectives.

Studying activations of probes, eigenvectors and eigenfilters across the temporal dimension may help in identifying interesting phenomenon such as reasoning and planning in RL environment. However, this might not be tractable with our current method as interactions between eigenfilters grow exponentially with each time step. Additionally, we do not concretely show how to derive insights specifically for multi-step reasoning, and aim to address this in future work.

We do not address a range of components often found in convolution neural networks, such as batch norm, dropout, or pooling. While we do not examine these components here, their implications, such as the performance tradeoffs between different pooling strategies, should be considered when evaluating architecture variants in the future.

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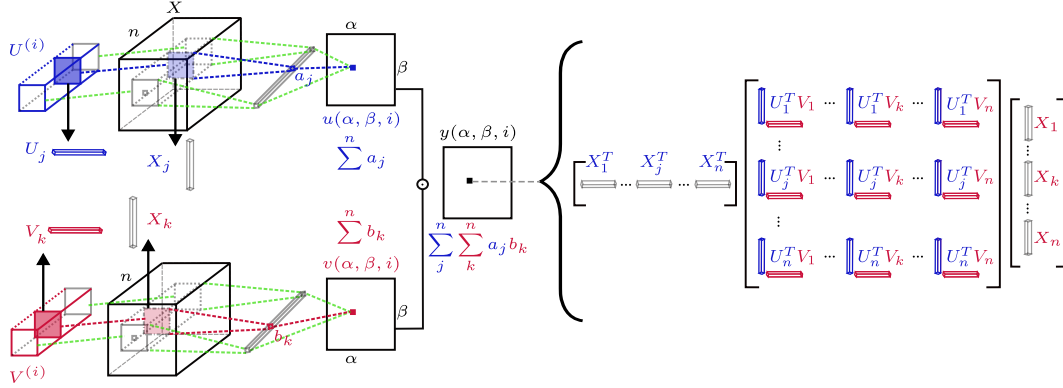


Figure 12: Transformation from spatial convolution operations (left) to a bilinear interaction matrix B (right) associated with scalar entry $y(\alpha, \beta, i)$ in output channel i . (Left, top, blue) Computation of spatial convolutions $U^{(i)}$ with input X_j , producing terms a_j . (Left, bottom, red) Computes convolutions $V^{(i)}$ with input X_k , producing terms b_k . (Right) The previous operations can be reformulated as a product of three block matrices, where the outer product of channel responses ($U^T V$) forms a symmetric bilinear matrix. The diagram emphasizes how local spatial convolutions (shown in the cubes) are transformed into a bilinear form B .

405 Consider the output of a convolution layer at location (α, β) for the i -th output channel:

$$u(\alpha, \beta, i) = \sum_j \sum_{k_1}^K \sum_{k_2}^K U^{(i)}[j, k_1, k_2] \cdot X[j, \alpha + k_1, \beta + k_2]$$

We can define the contribution of the filter applied to the j th input channel as:

$$a_j = \sum_{k_1}^K \sum_{k_2}^K U^{(i)}[j, k_1, k_2] \cdot X[j, \alpha + k_1, \beta + k_2]$$

406 This allows us to rewrite the output as:

$$u(\alpha, \beta, i) = \sum_j a_j$$

407 We can flatten the input tensor $X[j, \alpha : \alpha + k, \beta : \beta + k]$ into a K^2 -dimensional vector for each spatial
 408 location (α, β) . Let us denote this flattened version as $X[j, :, :]_f$. Similarly, we can write a flattened
 409 version of the filter $U^{(i)}[j, :, :]$, which we'll call $U^{(i)}[j, :, :]_f$. Note that the filter is independent of the
 410 position (α, β) .

411 Using these flattened representations, we can express a_j as:

$$a_j = U^{(i)}[j, :, :]_f \cdot X[j, :, :]_f$$

412 For readability, we can simplify the notation of the flattened vectors. We will also remove the notation
 413 for the output channel (i) , as all the operations we discuss here are for a single output channel.
 414 Simplifying the notation, we get:

$$a_j = U_j \cdot X_j$$

415 Note that U_j is a K^2 row vector, and X_j is a K^2 column vector. The gated operation is given by:

$$u(\alpha, \beta, i) \odot v(\alpha, \beta, i)$$

416 where v is the output of another Conv2D block. We perform a pointwise multiplication of the outputs
 417 u and v :

$$u(\alpha, \beta, i) = \sum_j a_j, \text{ where } a_j = U_j X_j$$

418 and using the same simplified notation for $V^{(i)}[k, :, :]_f$:

$$v[\alpha, \beta, i] = \sum_k^n b_k, \text{ where } b_k = V_k X_k$$

419 Therefore, for any α and β :

$$\begin{aligned} u[\alpha, \beta, i] \odot v[\alpha, \beta, i] &= \left(\sum_j^n a_j \right) \left(\sum_k^n b_k \right) \\ &= \sum_j^n \sum_k^n (a_j b_k) \end{aligned}$$

420 Interaction of channel k with channel j is given by:

$$\begin{aligned} a_j b_k &= (U_j X_j)(V_k X_k) \\ &= (X_j^T U_j^T)(V_k X_k) \quad (\text{since } a_j \text{ is a scalar}) \end{aligned}$$

421 Note that we can write the following sum:

$$\sum_j^n \sum_k^n X_j^T U_j^T V_k X_k$$

422 as a product of three block matrices Figure 12: $\begin{bmatrix} X_1^T & X_2^T & \dots & X_n^T \end{bmatrix} \begin{bmatrix} U_1^T V_1 & \dots & U_1^T V_n \\ \vdots & & \vdots \\ U_n^T V_1 & & U_n^T V_n \end{bmatrix} \begin{bmatrix} X_1 \\ X_2 \\ \vdots \\ X_n \end{bmatrix}$

423 The bilinear matrix B has a symmetric form given by

$$B^{sym} = \begin{bmatrix} \frac{U_1^T V_1 + (U_1^T V_1)^T}{2} & \dots & \frac{U_1^T V_n + (U_n^T V_1)^T}{2} \\ \vdots & & \vdots \\ \frac{U_n^T V_1 + (U_1^T V_n)^T}{2} & \dots & \frac{U_n^T V_n + (U_n^T V_n)^T}{2} \end{bmatrix}$$

424 We show that B^{sym} is indeed symmetric over all possible input pairs:

$$\begin{aligned} X_j^T B^{sym} X_k &= X_j^T \left(\frac{U_j^T V_k + (U_k^T V_j)^T}{2} \right) X_k \\ X_k^T B^{sym} X_j &= X_k^T \left(\frac{U_k^T V_j + (U_j^T V_k)^T}{2} \right) X_j \\ &= X_j^T \left(\frac{V_j^T U_k + U_j^T V_k}{2} \right) X_k \\ &= X_j^T \left(\frac{U_j^T V_k + (U_k^T V_j)^T}{2} \right) X_k \end{aligned}$$

425 Additionally, we can see that:

$$\begin{aligned} &X_j^T \left(\frac{\textcolor{red}{U}_j^T \textcolor{red}{V}_k + (\textcolor{blue}{U}_k^T \textcolor{blue}{V}_j)^T}{2} \right) X_k \\ &+ X_k^T \left(\frac{\textcolor{blue}{U}_k^T \textcolor{blue}{V}_j + (\textcolor{red}{U}_j^T \textcolor{red}{V}_k)^T}{2} \right) X_j \\ &= X_j^T \textcolor{red}{U}_j^T \textcolor{red}{V}_k X_k + X_k^T \textcolor{blue}{U}_k^T \textcolor{blue}{V}_j X_j \end{aligned}$$

426 for all j and k . The respective red and blue terms are compatible, because each term in the expansion
427 is a scalar and is thus equal to its transpose. Therefore, B^{sym} agrees with B on every input.

428 For each of m output channels, we get a matrix B^{sym} of dimension $nK^2 \times nK^2$, making its total
429 shape $[nK^2, nK^2, m]$.

Parameter	Type	Default Value	Description
Distribution Mode	String	easy	Difficulty or type of environment distribution. Choices: easy, hard, exploration, memory, extreme.
Environment Name	String	maze	Name of the environment to train on.
Number of Environments	Integer	64	Number of environments to use in parallel during training.
Number of Levels	Integer	100,000	Number of unique levels available for training.
Start Level	Integer	0	Starting level of the environment.
Method Label	String	hazelnut	Label or identifier for the method used.
GPU ID	Integer	7	GPU ID to use for training. Default is set to target GPU 7.
Learning Rate	Float	0.0001	Learning rate for the optimizer.
Entropy Coefficient	Float	0.01	Coefficient controlling entropy regularization.
Value Function Coefficient	Float	0.5	Coefficient balancing value function loss during training.
Discount Factor (γ)	Float	0.999	Discount factor for future rewards.
Lambda (λ)	Float	0.95	Generalized advantage estimation (GAE) discount factor.
Clip Range	Float	0.2	PPO clip range for policy loss updates.
Maximum Gradient Norm	Float	0.5	Maximum allowable gradient norm for clipping.
Steps per Update	Integer	256	Number of environment steps per policy update.
Batch Size	Integer	8	Batch size used for training.
Number of Epochs per Update	Integer	3	Number of training epochs per policy update.
Maximum Training Steps	Integer	12,800,000,000	Maximum number of total environment steps for training.
Pooling Method	String	avg	Pooling method used in the architecture. Options: avg, max, etc.

Table 1: Training parameters for ProcGen training

Layer	Sequence 0 (%)	Sequence 1 (%)	Sequence 2 (%)	Fully Connected (%)
Initial Conv	99.88			-
Conv	100.00	100.00	100.00	-
MaxPool	99.88	100.00	100.00	-
ResBlock0	99.88	100.00	100.00	-
ResBlock0 Gated Conv	59.28	42.75	69.07	-
ResBlock1	100.00	100.00	100.00	-
ResBlock1 Gated Conv	69.07	0.00	40.12	-
Gated FC	-	-	-	68.36
Logits FC	-	-	-	81.20
Value FC	-	-	-	2.73

Table 2: F_1 scores for position probes trained on the output of different layers of the network. The scores indicate how well each layer preserves cheese position information.

432 D Training resources

433 Each ProcGen environment and model combination trained in 15-20 GPU hours on a 48GB VRAM
434 4xA40 gpu node

435 E Ablation supplemental figure

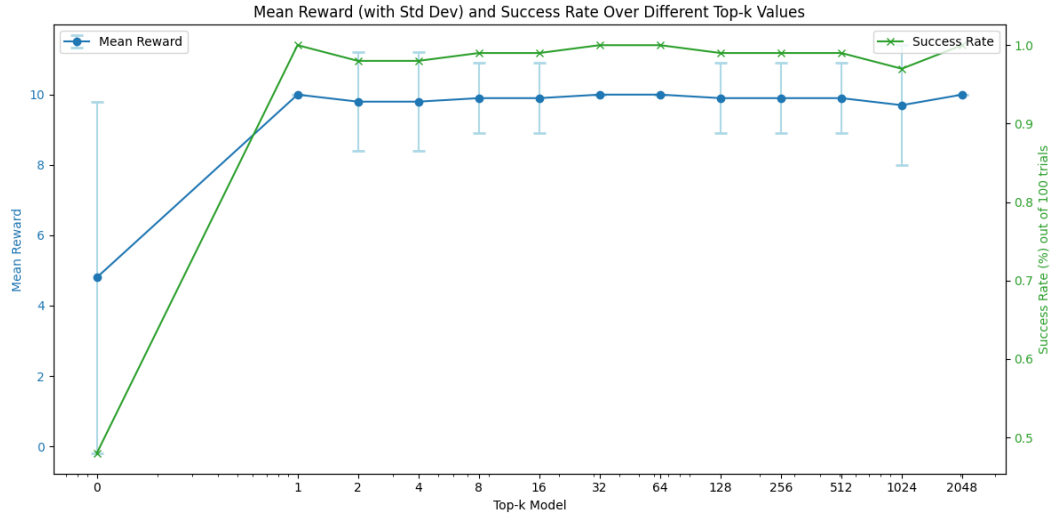


Figure 13: Keeping just first eigenvector for each output action in the final FC layer is enough to preserve near 100% success rate in solving mazes.

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For (1), see Figure 1, Figure 2, and BIMPALA matches IMPALA performance

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For (3), see Ablation within the standard bases and Figure 7

For (4), see Protocol to enhance interpretability for RL with probes, Interpretability based on action features, Steering Experiments and corresponding figures in each subsection

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748 such risks were disclosed to the subjects, and whether Institutional Review Board (IRB)
749 approvals (or an equivalent approval/review based on the requirements of your country or
750 institution) were obtained?

751 Answer: [NA]

752 Justification: No human or living subjects were used in the study.

753 Guidelines:

- 754 • The answer NA means that the paper does not involve crowdsourcing nor research with
755 human subjects.
- 756 • Depending on the country in which research is conducted, IRB approval (or equivalent)
757 may be required for any human subjects research. If you obtained IRB approval, you
758 should clearly state this in the paper.
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765 Question: Does the paper describe the usage of LLMs if it is an important, original, or
766 non-standard component of the core methods in this research? Note that if the LLM is used
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768 scientific rigor, or originality of the research, declaration is not required.

769 Answer: [NA]

770 Justification: We did not use LLMs beyond getting feedback on certain, limited lines of
771 code. The manuscript was written without LLM assistance.

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