Gates Are Not What You Need in RNNs^{*}

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Abstract. Recurrent neural networks have flourished in many areas. Consequently, we can see new RNN cells being developed continuously, usually by creating or using gates in a new, original way. But what if we told you that gates in RNNs are redundant? In this paper, we propose a new recurrent cell called *Residual Recurrent Unit* (RRU) which beats traditional cells and does not employ a single gate. It is based on the residual shortcut connection together with linear transformations, ReLU, and normalization. To evaluate our cell's effectiveness, we compare its performance against the widely-used GRU and LSTM cells and the recently proposed Mogrifier LSTM on several tasks including, polyphonic music modeling, language modeling, and sentiment analysis. Our experiments show that RRU outperforms the traditional gated units on most of these tasks. Also, it has better robustness to parameter selection, allowing immediate application in new tasks without much tuning. We have implemented the RRU in TensorFlow, and the code is made available on GitHub.

Keywords: Recurrent neural networks \cdot Residual neural networks \cdot Gates \cdot Robustness \cdot Deep learning.

1 Introduction

Recurrent neural networks (RNN) have achieved widespread use in sequence processing tasks such as language modeling, speech and music recognition. RNNs are composed of a single computation cell, called Recurrent Unit which is unrolled along the sequence dimension.

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In order to achieve stable training and the ability to store and make use of long-term dependencies, the recurrent units are designed in a special way. The most well-known units are LSTM and GRU, which use gating mechanism to store and extract information from the recurrent state. Two kinds of gates are used in these units – reset gates and update gates, where each of them is based on modulating the state information with the input information and is implemented by multiplying two values, one of which is often range-limited by the sigmoid function. Gates are considered crucial for propagation of long-term information [5] and virtually all proposals of Recurrent Units, including numerous recent developments, contain such multiplicative input-state interactions. Recent developments of improved RNN variants contain even more multiplicative interactions [12] that culminates in Mogrifier LSTM, where the input information is modulated through a series of reset gates [16]. They find that about five reset gates achieve the best performance yielding Mogrifier LSTM to be the top-performing RNN cell on several datasets so far.

In contrast, we show that gates are not essential at all to construct a wellperforming recurrent unit. To this end, we develop a recurrent cell not containing a single gate (see visualization in Figure 1). The proposed cell surpasses not only GRU and LSTM but also the so far best Mogrifier LSTM on many commonly used benchmark tasks. Our cell is based on a residual ReLU network employing normalization and ReZero [1] recurrent state update. If well-performing recurrent units without gates can be created, perhaps we shouldn't think of gates every time we think of recurrent networks and maybe this insight could help us create strong non-gated networks in the future.

2 Relation to Prior Work

A great number of RNN cells have been developed. They usually use gates to ensure stable training over many time-steps. The most well-known is LSTM [8] which has an extra cell state that can pass information directly forward and three gates (forget, input and output) which control the data flow. This specific structure has allowed it to become the state-of-the-art cell, still being used extensively to this day. Its success is considered to stem from the 3 gates at its core, and since then, it has been considered that RNN cells need gates for optimal performance.

Following the success of the LSTM, a new cell called GRU [4] was proposed, which is loosely based on LSTM and often referred to as the light version of the LSTM. This cell combines the forget and input gates of an LSTM, resulting in 2 gates total – reset gate and update gate. As a lighter cell, it can be computed faster while the results are often similar.

Based on the marvellous results of LSTM and GRU, many extensions and modifications have been proposed. Phased LSTM [17] introduce a new time gate. Gating also in the depthwise direction is explored in [10,24]. The power of multiplicative interactions, which is the central operation in gates, is explored in [12,23]. The number of gates is brought to the culmination in Mogrifier LSTM,



Fig. 1. The structure of the *Residual Recurrent Unit* (RRU) with two state-input linear transformations. The dotted brackets cover the layer which can be used zero or more times, in this case – once.

which does input and hidden state modulation by multiple rounds of reset gates [16]. Their results suggest that using about five rounds of reset gates gives the optimal performance and positions Mogrifier LSTM as the currently best RNN cell in WikiText-2, Penn Treebank word-level and Penn Treebank character-level tasks.

Several works have questioned the necessity of the high number of gates in RNNs. [5], through numerous experiments, find that the forget gate and the output activation function are the most critical components of the LSTM block and removing any of them impairs performance significantly. A similar conclusion was obtained in [22] and a new cell called JANET was proposed, which is based on the LSTM but uses just the forget gate. The minimalistic designs of recurrent cells with only one forget gate were proposed in [7, 26]. The need for a reset gate in GRU was questioned in [18], where they obtain improved accuracy in Automatic Speech Recognition with the unit having only an update gate, Batch Normalization and ReLU activation.

There has been some tinkering done with using a residual shortcut connection in RNNs. Consequently, a Residual RNN [25] has been proposed. Res-RNN and gRes-RNN were proposed, where the former is a purely residual recurrent network and the latter combines residual shortcut connection with a gate. Their units show improved speed but are not able to unequivocally beat LSTM in terms of accuracy. [11] proposes a Residual LSTM, which is a mix between a regular LSTM and a residual shortcut connection to improve performance in case of many layers. Even with the addition of the residual shortcut connection, it still contains a high number of gates.

3 Residual Recurrent Unit

We propose a new cell that does not contain a single gate but provides a competitive performance. We call this cell the *Residual Recurrent Unit*, RRU for short. The cell consists of a residual shortcut connection and several linear transformations employing Normalization and ReZero [1] recurrent state update. ReZero is a simple zero-initialized parameter that controls how much of the residual branch contributes to the updated state. ReZero helps us make the unit structure gateless, which makes it more simple. As it is initialized as 0, only the previous state is taken in effect at the first training steps, and the optimal weight for the residual branch emerges during training. ReZero ResNet has been shown to have non-vanishing gradient yielding stable training for very deep convolutional networks and Transformers [1] and achieving great results. Here we adapt it to replace gates in recurrent networks to see what benefits it could give to recurrent units.

The visualization of our cell can be seen in Figure 1. RRU takes input x_t at the current time-step t which is a vector of dimension m and the previous state h_{t-1} of dimension n and produces the updated state h_t and the output o_t of dimension p. The calculations done in the RRU, are described as:

$$j_t = \text{ReLU}(\text{Normalize}(W^x x_t + W^h h_{t-1} + b^j)) \tag{1}$$

$$[j_t = \operatorname{ReLU}(W^k j_t + b^k)]^*$$
(2)

$$d_t = \text{Dropout}(j_t) \tag{3}$$

$$c_t = W^c d_t + b^c \tag{4}$$

$$h_t = \sigma(S) \odot h_{t-1} + Z \odot c_t \tag{5}$$

$$o_t = W^o d_t + b^o \tag{6}$$

In the above equations, upper case letters depict the learnable parameters. Weight matrices with their respective dimensions are: $W^x \in \mathbb{R}^{m \times g}$; $W^h \in \mathbb{R}^{n \times g}$; $W^k \in \mathbb{R}^{g \times g}$; $W^o \in \mathbb{R}^{g \times p}$; $W^c \in \mathbb{R}^{g \times n}$, where g is the hidden size which we choose to be equal to q * (m + n), and q is the middle layer size multiplier (we typically use values from 0.1 to 8.0). Bias vectors and their dimensions are: $b^j \in \mathbb{R}^g$; $b^k \in \mathbb{R}^g$, $b^o \in \mathbb{R}^p$; $b^c \in \mathbb{R}^n$. There are two learnable scaling factors S and Z of dimension n. The sigmoid function is denoted as σ , $[\cdot]^*$ denotes the use of the equation inside the brackets zero or more times, \odot denotes scalar multiplication.

The structure of RRU is similar to the ReZero residual network [1]. At first, the previous hidden state h_{t-1} and the input of the current timestep x_t are passed through a linear transformation followed by L2 normalization and ReLU. Then, zero or more linear transformations follow employing ReLU activation. To make the formulas clearer, Equation 2 has two vectors named j_t , in reality, each layer has their own W^k and B^k . This value is a hyperparameter, for which each dataset has a specific value which works best, usually 1 or 2. Dropout [20] follows, and the result is linearly transformed to get two values – the output of the current timestep o_t and the hidden state candidate c_t . The next hidden state h_t is produced by a weighted sum of the previous state and the candidate. The candidate is scaled by a zero-initialized parameter Z according to the ReZero principle. There is a slight difference from the ReZero paper in that we use a separate scale for each feature map, but ReZero uses a common scaling factor for all maps, see Section 6 for evidence that our version works better.

Another difference from the traditional ResNet architecture is that we have introduced a scaling factor S for the residual connection (ResNet has a fixed S = 1). Such change is motivated by the need for the recurrent network to forget some of the information from the previous time-steps. The scale S is limited to the range 0-1 by the sigmoid function to eliminate unstable behaviour in some cases, especially if S has become negative. Since S < 1, the cell's memories gradually fade out. Note that is a learnable parameter (for each feature map separately) so the network can choose the rate of forgetting. Such fade-out is not so flexible as a forget gate employed in other cells but the network can compensate for it using the residual branch c. We have observed that the value of S actually changes during training, mainly decreasing for most of the feature maps. This can be explained that the network initially uses residual connection to provide a stable gradient for training but later learns to rely on the c value more which provides greater control. The value of S is initialized in the way that after the sigmoid, it is uniformly distributed in the range 0-1. This idea is suggested by [6] and relieves us from the need for another hyperparameter that needs to be tuned. We also experimented with a constant initialization of S; that worked similarly but required an adequate constant, usually in the range 0.4 - 0.95 depending on the dataset (see Section 6). The ablation study also shows that S is important in general; for Penn Treebank removing the scaling produces a significantly worse result.

The initial hidden state h_0 is prepared to have all zeros except the first feature map, which is set to $\frac{1}{4}\sqrt{n}$. Initializing with all zeroes causes a blowup due to the employed normalization in the case when zeros are given also as the input values x_t in the first timesteps. Such inputs may occur if the input is padded by zeros from some shorter sequence.

Note that although Equation 5 resembles the update gate of LSTM, it is not because in the update gate, both of the multiplied values depend on the input or hidden state but in our case, S and Z are learnable parameters – effectively constants after training.

4 Experiments

To test our unit's performance against the chosen competitors (GRU, LSTM, Mogrifier LSTM), we run experiments on language modeling, music modeling, sentiment analysis, and MNIST image classification tasks. These tasks are com-

5

monly used for evaluating recurrent networks. All chosen units are equivalent in terms of trainable parameters in our experiments, to ensure a fair comparison.

For the polyphonic music modeling, the task is to predict the next note when knowing the previous ones. The model's performance is measured by negative log-likelihood (NLL) loss which sums the negative logarithm of all the correct prediction probabilities. For this task, we chose four datasets that are usually evaluated together – JSB Chorales, Nottingham, MuseData, Piano-midi.de [3]. They are already split into training, validation and testing portions. For wordlevel language modeling, we use the Penn Treebank dataset [15]. The aim of this dataset is to predict the next word by knowing the previous ones. The network's performance is measured in perplexity, which is the probability that a word will show up as the next word from the previous context. For character-level language modeling datasets we chose enwik8 [9], $text8^4$ and Penn Treebank [15]. The aim is to predict the next character from the previous ones. The performance is measured in BPC (bits per character), which is the number of bits used to represent a single character from the text. Although word-level and characterlevel tasks may seem similar, they evaluate different aspects of the cell. The word-level task has a large vocabulary, so the emphasis is put on evaluating RRU's ability to deal with diverse inputs, but the character-level task has a large window size requiring the cell to remember and process long history.

The goal for IMDB sentiment analysis dataset [14] is to predict the sentiment of the text as either positive or negative. This dataset is of interest because it is quite different from the previous ones. The performance of the network is measured as the accuracy of the predictions. The challenging part of the dataset is to learn the hidden state in a way that can understand even complicated structures, for example, double negatives, etc.

A frequently used dataset for evaluating recurrent networks is MNIST image classification. RNNs are not well suited for image classification; therefore, it puts every aspect of the network at stress. The aim here is to predict the number that each picture represents by processing a sequence that consists of all the pixels in the picture.

For music modeling and word-level language modeling, we run hyperparameter optimization on each cell, using Bayesian hyperparameter tuning from the HyperOpt library [2]. The optimized hyperparameters include: number of learnable parameters, dropout rate, learning rate and some of the most influential hyperparameters specific to each cell; for more detailed configurations see Appendix D. The RRU cell has a built-in dropout, so to get an equitable environment, we added recurrent dropout to GRU, LSTM and Mogrifier LSTM as described in [19]. For character-level Penn Treebank, we also run optimization, but we only grid search through the different dropout rates from 0.0 to 0.9 and the best testing result from these runs is shown in the results table. This experiment is further described in Appendix A. For the rest of the datasets, hyperparameter tuning would require computational power exceeding our budget, so we only run a singular experiment on each cell on as fair a configuration as

⁴ http://mattmahoney.net/dc/textdata

Table 1. Evaluation results on each task. The "+/-" notation means that we did grid search through ten different dropout values to find the one which performs the best. Every metric over the horizontal line is better if it is lower and every metric under it is better if it is higher.

Tool		DDII	CPU	TOTM	Mogrifier
TASK	runea	nnu	GNU	LSIM	LSTM
Music JSB Chorales (NLL)	+	7.72	8.20	8.33	8.18
Music Nottingham (NLL)	+	2.92	3.22	3.25	3.28
Music MuseData (NLL)	+	7.03	7.31	7.24	7.22
Music Piano-midi.de (NLL)	+	7.38	7.58	7.54	7.52
Word-level Penn Treebank (Perplexity)	+	102.56	122.21	140.35	126.88
Character-level Penn Treebank (BPC)	+/-	1.27	1.28	1.34	1.29
Character-level enwik8 (BPC)	-	1.37	1.53	1.49	1.36
Character-level text8 (BPC)	-	1.35	1.55	1.44	1.37
Sentiment analysis IMDB (Accuracy)	-	87.20	87.04	85.89	86.23
Sequential MNIST (Accuracy)	-	98.74	98.36	92.88	98.14
Permuted MNIST (Accuracy)	-	97.67	98.68	97.39	97.81

possible, that is, we try to use the same configuration but sometimes changes to the configuration have to be made, for more details see Appendix D.

Each experiment was done on a single machine with 16 GB RAM and a single NVIDIA T4 GPU, time spent during a single, full training varies from averagely 4 minutes (JSB Chorales) to averagely 115 hours (enwik8). We use RAdam optimizer [13] with gradient clipping. We train each dataset until the validation loss stops decreasing for 3-11 epochs, depending on the dataset.

Results of all of these experiments can be seen in Table 1. RRU is a definite leader giving the best performance on almost all of the datasets. Notably, RRU scores on top for all fully hyperparameter-optimized cases. In the two cases where RRU is outperformed, it still comes close, and it is probable that RRU would achieve the best performance if full hyperparameter optimization were performed for these datasets. We suspect that the RRU could reach a state-ofthe-art performance score on specific datasets, but we didn't have the time and resources to spend on tuning to achieve this. Convergence graphs of some of these experiments can be seen in Appendix C, in which RRU usually converges first and with better results.

5 Robustness against Hyperparameter Choice

One of the main reasons for LSTM's and GRU's wide usage is their robustness against hyperparameter choice; that is, usually, you can use them with some default parameters, and they will give decent results. Such property has been described in several works, one being [21]. We will test the robustness for each cell to the selection of learning rate and learnable parameter count – the two most important hyperparameters. We run a grid search on Nottingham music

modeling dataset with a reasonably wide hyperparameter range for all the cells. We plot the validation NLL and the number of epochs needed to reach the best accuracy as heatmaps in Figure 2. The figure shows that our cell has a much broader range of parameters in which it works well, which also means that our cell can be used successfully without much parameter tuning. Further robustness experiments can be seen in Appendix B.



Fig. 2. Robustness against learning rate and parameter count on the Nottingham dataset. The top row gives the testing NLL, and the bottom row shows the number of epochs needed to reach the best NLL. Good results are depicted with blue colour, poor results – with yellow. The red squares correspond to runs where the training failed, and the pink frames show the run with the best NLL. We observe that RRU consistently produces the best NLL across the entire range and achieves that within a small number of epochs.

6 Ablation Study

In this section, we investigate the influence of the key elements in our cell by replacing them with simpler ones to determine whether our cell's structure is optimal. We compare the RRU performance against versions with:

- Removed normalization.
- Using a single scalar multiplier Z for all feature maps as proposed in the ReZero paper [1]. Our cell uses a different multiplier for each map.
- Added ReLU over Equations 4 and 6. Such setup is employed in some ResNet architectures [29, 30] and could have more expressive power.
- Residual weight S set as a constant with a scalar value 1 i.e. using an unscaled residual shortcut connection, expressing Equation 5 as $h_t = h_{t-1} + Z \odot c_t$.

 Residual weight S initialization with a scalar value 0.95 instead of a random one.

For these experiments, we have chosen character-level Penn Treebank, IMDB, Sequential MNIST datasets, on which we do a single run, and Nottingham, wordlevel Penn Treebank, on which we optimize the hyperparameters as described in Section 4. For each dataset's configuration see the respective dataset's configuration for RRU in Section 4, except for character-level Penn Treebank for which the configuration differs by the dropout rate which isn't tuned, but set as 0.3. The results can be seen in Table 2. We conclude that the current structure of our cell gives the best performance. Interestingly, the considered simplifications produce only slightly worse results, suggesting that the RRU's structure is robust to changes. Although great results can be achieved without normalization, it seems to give the RRU more stability during the training and contributes to robustness (see Appendix 5). We notice that our approach of using a different ReZero multiplier for each hidden map gives better performance than using only the singular scalar ReZero parameter. Adding another ReLU over the vectors c_t and o_t does not help. Passing the full state without a scaling factor S decreases performance. Using constant initialization seems to give similar performance, but in these experiments, the random initialization gave us better results with the added bonus of fewer hyperparameters to tune.

Version	Character PTB (BPC)	IMDB (Accuracy)	Sequential MNIST (Accuracy)	Nottingham (NLL)	Word PTB (Perplexity)
RRU	1.32	87.45	98.74	2.92	102.70
no normalization	1.32	87.21	98.03	2.92	104.60
single scalar ReZero	1.32	87.08	98.45	2.97	104.44
ReLU over c and o_t	1.34	87.17	86.38	3.07	120.76
S = 1	2.25	86.78	98.43	3.56	110.74
${\cal S}$ initialized with 0.95	1.33	86.71	98.60	2.96	103.09

Table 2. Ablation study results for all of the datasets. "Character-level Penn Treebank" is denoted as "Character PTB" and "Word-level Penn Treebank" is denoted as "Word PTB" to save space.

7 Conclusion

By developing a new RNN cell without any gates and showing that it outperforms the gated cells on many tasks, we have demonstrated that gates are not necessary for RNNs. Our proposed RRU cell is robust to parameter selection and can be used in new tasks without much tuning. RRU has roughly the same speed as pure TensorFlow implementations of LSTM or GRU, and we look forward

to low-level optimized CUDA RRU implementation that matches the speed of optimized LSTM and GRU implementations. We expect that the insights gained in this work will contribute to further improvements in the designs of recurrent and residual networks.

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11

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A Appendix – Dropout Analysis

While running experiments, we noticed that our cell seems to handle dropout better than the other cells. To test this observation, we ran a grid search on Nottingham and character-level Penn Treebank datasets for each cell through different dropout rates — from 0.0 to 0.9, for more details of the configurations used see Appendix D. The results from these experiments can be seen in Figures 3 and 4. From these results, we can see that all cells benefit from dropout, but for RRU, its impact is much more pronounced, and RRU ultimately reaches better final results than the other cells. We can also see that the RRU works best with a dropout rate of around 0.7. Possibly, dropout is better suited for ReLU networks, as is RRU, rather than for gated networks.



Fig. 3. NLL (lower is better) depending on the dropout rate for each cell on the Nottingham dataset.



Fig. 4. BPC (lower is better) depending on the dropout rate for each cell on the character-level Penn Treebank dataset.

B Appendix – Further Robustness Experiments

In Section 5 we tested the robustness of our cell. Here we will do the same experiment on a different dataset, the word-level Penn Treebank, to furthermore test the robustness. We plot the validation perplexity and the number of epochs needed to reach the best accuracy as heatmaps in Figure 5. The figure shows that our cell has a much broader range of parameters in which it works well, which also means that our cell can be used successfully without much parameter tuning. We also notice that the LSTM and the Mogrifier LSTM had trouble learning anything at all on very small learning rates.



Fig. 5. Robustness against learning rate and parameter count for different cells on the word-level Penn Treebank dataset. The top row gives the testing perplexity, and the bottom row shows the number of epochs needed to reach the best perplexity. Good results are depicted with blue colour, poor results – with yellow. The red squares correspond to runs where the training failed, and the pink frames show the run with the best perplexity. We observe that RRU consistently produces the best perplexity across the entire range and achieves that within a small number of epochs.

In Appendix 6 we looked at different RRU versions from which we concluded that our version tops the other versions, but we want to show in more detail why normalization is beneficial in the RRU. We will compare the RRU with and without normalization with the robustness tests as described in Section 5. We plot the validation loss and the number of epochs needed to reach the best accuracy as heatmaps in Figure 6 and 7. We observe that RRU with normalization consistently produces the best loss across the entire range and achieves that within a small number of epochs which shows that normalization helps gain a



much broader range of parameters in which it works well and helps avoid failed training sessions.

Fig. 6. Robustness against leaning rate and parameter count for different cells on the Nottingham dataset. The top row gives the testing NLL, and the bottom row shows the number of epochs needed to reach the best loss. Color depictions are the same as in Figure 5.



Fig. 7. Robustness against leaning rate and parameter count for different cells on the word-level Penn Treebank dataset. The top row gives the testing perplexity, and the bottom row shows the number of epochs needed to reach the best loss. Color depictions are the same as in Figure 5.

C Appendix – Convergence Speed

Convergence speed is another important factor in neural networks, for that reason, we will display convergence graphs of three datasets in this section, one for each training mode – JSB Chorales (tuned), character-level Penn Treebank (half-tuned) and IMDB (not tuned). The graphs are taken from the experiments done in Section 4.

Convergence graph of the JSB Chorales dataset with the best parameters found in tuning for each cell can be seen in Figure 8. We can observe that RRU in this dataset trains much faster, beating every other cell's best NLL at 17th epoch, which implies that early stopping could be done.



Fig. 8. Validation NLL on JSB Chorales dataset for each epoch and tuned cell.

Convergence graph of the character-level Penn Treebank dataset after halftuning (going through 10 different dropout values and taking the best one) has been done can be seen in Figure 9. We see that the RRU beats all the other cells here, while doing it in a similar epoch count as others, even when it has a larger dropout rate, which usually increases epoch count tremendously.

Convergence graph of the IMDB dataset can be seen in Figure 10. In this graph we once again see that the RRU is capable of reaching it's best accuracy first and beating the other cells with it.



Fig. 9. Validation BPC on character-level Penn Treebank dataset for each epoch and half-tuned cell. In this Figure RRU has 0.7 dropout, GRU has 0.5 dropout, LSTM has 0.4 dropout and Mogrifier LSTM has 0.5 dropout.



Fig. 10. Validation Accuracy on the IMDB dataset for each epoch and cell.

D Appendix – Experiment Configurations

This appendix is intended for elaboration of the configurations of the experiments that are written in Section 4, but for specific configurations we suggest contacting the author(s).

D.1 Dataset Configurations

Polyphonic music modeling datasets contain sequences of played piano-roll elements (MIDI note numbers between 21 and 108 inclusive), which we represent as a binary mask for each time-step, where ones are placed in the positions corresponding to the MIDI notes played at that time step. We trim or pad the sequences to length 200, similarly to how it was done by [27].

Word-level Penn Treebank dataset is already split into training, validation and testing portions. We use a context length of 64 and a vocabulary of 10 thousand, which includes all of the words in the dataset.

Character-level Penn Treebank also is already split into training, validation and testing portions. However, enwik8 and text8 are not, so we use the standard split for each of them, which is 90% training, 5% validation, 5% testing. We use a window size of 512 for Penn Treebank and 256 for enwik8 and text8. The training is run in a stateful manner, meaning that we pass the old state forward to the next window (with a 10 % chance of passing a state filled with zeros).

The IMDB dataset consists of 25 thousand training sequences and 25 thousand testing sequences. We take 5 thousand sequences from the end of the training set as a validation set. We trim the sequences to length 500 and use a vocabulary of size 10 thousand created from the most frequently occurring words (there are approximately 25 thousand different words in total) to be able to fit the sequences into our GPU memory.

We use the MNIST image classification dataset in a specific way: we use Sequential MNIST in which we take the image by pixel rows and get a single sequence and Permuted MNIST in which the pixels of each row are randomly permuted according to some fixed permutation, which makes the task harder. Each 28×28 image is transformed into a sequence of 784 elements. The dataset consists of 60 thousand training images and of 10 thousand testing images. We reserve 10 thousand images from the training data for validation.

D.2 Main Experiments

All four polyphonic music datasets use the same configuration. We set context size as 200 (as done in [27]), batch size as 16, number of layers as 1, for RRU we use 1 ReLU layer and an output size of 64, all of these values we experimentally found to work well. The training stops when there hasn't been a lower NLL for 7 epochs.

For word-level Penn treebank we set context size as 64, batch size as 128, number of layers as 2, embedding size of 64, for RRU we use 1 ReLU layer and

an output size of 128, all of these values we experimentally found to work well. The training stops when there hasn't been a lower perplexity for 7 epochs.

For polyphonic music modeling and word-level language modeling, we run hyperparameter optimization on each cell, where we tune the learning rate, the dropout rate and the number of parameters in 100 runs, with ranges as described in Table 3. We also tune the following parameters of the RNN cells: for RRU we tune the middle layer size multiplier q in range [0.1; 8.0]; for the LSTM we tune the forget bias in range [-3.0; 3.0]; for the Mogrifier LSTM we tune the feature rounds in range [5; 6] and the feature rank in range [40; 90], as these were mentioned as the optimal ranges in the Mogrifier LSTM paper. Optimal values for the tuned parameters for each dataset and cell can be seen in Table 4.

Table 3. Parameter ranges for main experiments that used tuning. ^{*u*} means the value is taken from the uniform scale and ^{*l*} means from the logarithmic scale. If '.' is present in the range, it means it includes float numbers, else the range is only from integers. 'M' means that the value is multiplied by 10^6 . These ranges were picked because they seemed sufficient to include all of the optimal values, which were approximately identified in the initial experiments.

Dataset	Learning rate	Dropout r	rate Number of	parameters
JSB Chorales				
Nottingham			[1]]	$1 \in M[u]$
MuseData	$[1e-4; 1e-2]^l$	[0.0; 0.8]	$]^u$ $[1M -$	15M
Piano-midi.de				
Word-level Penn Treebank			[10M -	$-30M]^{u}$

For character-level Penn Treebank we set number of parameters as 24 million, learning rate as 1e - 3, we set context size as 512, batch size as 64, number of layers as 2, embedding size of 16, for RRU we use 2 ReLU layers, middle layer size multiplier q as 4.0 and an output size of 128, for LSTM we set forget bias as 1.0, for Mogrifier LSTM we use 4 feature mask rounds of rank 24, all of these values we experimentally found to work well. The training stops when there hasn't been a lower BPC for 11 epochs. We run this configuration through ten different dropout rates from 0.0 to 0.9 and report the one with the lowest BPC.

The remaining datasets – enwik8, text8, IMDB, Sequential MNIST, P-MNIST – are run with no tuning. We tried to use the same configuration for each of the cells, but it wasn't always possible, because for some datasets some cells were unable to train with the common configuration. The configuration that was the same for each of the cells can be seen in Table 5 and the slight differences can be seen in Table 6. All of these parameter values were experimentally found to work well. enwik8 number of parameters (48 million) were taken from the Mogrifier LSTM paper and Sequential MNIST, P-MNIST number of parameters (70 thousand) were taken from [28].

Table 4. Optimal values for the main experiments that used tuning. 'M' means that the value is multiplied by 10⁶. "JSB Chorales" is denoted as "JSB", "MuseData" is denoted as "Muse", "Piano-midi.de" is denoted as "Piano" and "Word-level Penn treebank" is denoted as "Word PTB" to save space.

Parameter	JSB	Nottingham	Muse	Piano	Word PTB			
RRU								
Learning rate	0.00495	0.00029	0.00030	0.00077	0.00683			
Dropout rate	0.79362	0.77186	0.77518	0.71577	0.66035			
Number of parameters	$6.9 \mathrm{M}$	2.6M	8.6M	$5.9 \mathrm{M}$	$25.9 \mathrm{M}$			
Middle layer size multiplier	q 1.76958	7.31994	6.85699	3.68371	6.86655			
	GRU							
Learning rate	0.00460	0.00454	0.00260	0.00932	0.00012			
Dropout rate	0.67605	0.33787	0.64839	0.75069	0.27062			
Number of parameters	1.8M	3.4M	1.9M	2.4M	10.0M			
	LSTM							
Learning rate	0.01000	0.00882	0.00493	0.00727	0.00162			
Dropout rate	0.30081	0.20310	0.35383	0.22951	0.10384			
Number of parameters	1.3M	1.3M	1.8M	$11.9 \mathrm{M}$	14.2M			
Forget bias	0.76843	-0.04104	1.85746	0.59993	-2.33114			
Mogrifier LSTM								
Learning rate	0.00255	0.00169	0.00135	0.00012	0.00074			
Dropout rate	0.47521	0.73808	0.55499	0.77291	0.46872			
Number of parameters	1.3M	6.0M	10.3M	$2.9 \mathrm{M}$	24.2M			
Feature mask rounds	6	5	5	5	5			
Feature mask rank	58	48	76	63	74			

Table 5. Parameter values for parameters that are the same for each cell in nontuned main experiments. "Sequential MNIST", "P-MNIST" are denoted together as "MNISTs" and "text8", "enwik8" are denoted together as "wiki8s". 'M' means that the value is multiplied by 10^6 . [*RRU*] means the parameter is used in the RRU cell (same denotation is used for the other cells).

Parameter	wiki8s	IMDB	MNISTs
Number of parameters	48M	20M	70K
Context size	256	512	784
Batch size	64	64	64
Number of layers	2	2	2
Embedding size	32	64	-
Breaks after epochs with no performance gain	3	5	5
[RRU] ReLU layers	2	1	1
[RRU] Middle layer size multiplier q	4.0	2.0	2.0
[RRU] Output size	128	64	64
[LSTM] Forget bias	1.0	1.0	1.0
[MogrifierLSTM] Feature mask rounds	6	5	6
[MogrifierLSTM] Feature mask rank	79	40	50

Dataset	Cell	Learning rate	Dropout rate
enwik8	*	1e-3	0.7
	RRU	1e-3	
text8	GRU	1e-4	0.7
	LSTM	1e-3	0.7
	Mogrifier LSTM	1e-3	
IMDB	*	1e-3	0.5
	RRU	1e-3	0.7
Sequential MNIST	GRU	1e-3	0.7
& P-MNIST	LSTM	1e-4	0.7
	Mogrifier LSTM	5e-5	0.5

Table 6. Parameters that differ between cells on some datasets in non-tuned main experiments. '*' in "Cell" field means that all cells have the same configuration.

D.3 Robustness Experiments

For Nottingham here we use the same configuration as the configuration for Nottingham in the main experiments (Section D.2), except no tuning is done, so we set middle layer size multiplier q as 2 and dropout rate as 0.5. There are 100 runs in total from a full grid search from learning rates 1e - 4, 2e - 4, 3e - 4, 5e - 4, 8e - 4, 13e - 4, 22e - 4, 36e - 4, 6e - 3, 1e - 2 (ten values from 1e - 4 to 1e - 2 in logarithmic scale) – and number of parameters – 1.0M, 2.6M, 4.1M, 5.7M, 7.2M, 8.8M, 10.3M, 11.9M, 13.4M, 15.0M (ten values from 1.0M to 15.0M in uniform scale).

For word-level Penn Treebank here we use the same configuration as the configuration for word-level Penn Treebank in the main experiments (Section D.2), except no tuning is done, so we set middle layer size multiplier q as 2 and dropout rate as 0.5. There are 100 runs in total from a full grid search from learning rates -1e-4, 2e-4, 3e-4, 5e-4, 8e-4, 13e-4, 22e-4, 36e-4, 6e-3, 1e-2 (ten values from 1e-4 to 1e-2 in logarithmic scale) – and number of parameters – 10.0M, 12.2M, 14.4M, 16.7M, 18.9M, 21.1M, 23.3M, 25.6M, 27.8M, 30.0M (ten values from 10.0M to 30.0M in uniform scale).

D.4 Dropout Experiments

For Nottingham here we use the same configuration as the configuration for Nottingham in the main experiments (Section D.2), except no tuning is done, so we set the number of parameters to 5 million, learning rate as 1e - 3, middle layer size multiplier q as 2.0. We run this configuration through different dropout rates — from 0.0 to 0.9.

For character-level Penn Treebank experiment configuration see characterlevel Penn Treebank experiment configuration in Section D.2.

E Appendix - Appendix References

[27] Y Cem Subakan and Paris Smaragdis, "Diagonal rnns in symbolic music modeling," in 2017 IEEE Workshop on Applications of Signal Processing to Audio and Acoustics (WASPAA). IEEE, 2017, pp. 354–358.

[28] Shaojie Bai et al., "An empirical evaluation of generic convolutional and recurrent networks for sequence modeling," *arXiv preprint arXiv:1803.01271*, 2018.