
Evaluating Impact of Distribution Shifts on Preprocessing and Modeling Choices for Speech Decoding from MEG Signals

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Abstract

We take machine learning approaches to the second phase of the LibriBrain Competition 2025, which requires phoneme classification from MEG recordings. We compare preprocessing strategies, phonological label representations, and model architectures. We highlight the impact of distribution shifts between validation and hidden evaluation data caused by inconsistent standardization, and find that simple residual convolutional networks outperform more complex architectures under this shift.

1 Introduction

Decoding speech units from non-invasive neuroimaging is a central goal for brain–computer interfaces [Lotte et al., 2018]. The LibriBrain 2025 phoneme classification track offers a standardized benchmark with aligned MEG segments and pre-defined data splits [Özdogan et al., 2025, Landau et al., 2025]. Our experiments revealed large gaps between validation and leaderboard metrics, highlighting unresolved distribution shifts.

Our contributions are threefold: (i) we quantify how group averaging and careful feature scaling stabilize training, (ii) we benchmark three neural backbones designed for short MEG windows, and (iii) we analyze failure modes caused by inconsistent standardization between validation and hidden evaluation data. These insights aim to guide future works on decoding of speech from MEG signals.

2 Related Works

2.1 Neural decoding from non-invasive brain signals

Non-invasive modalities like electroencephalography (EEG) and magnetoencephalography (MEG) are key to brain-computer interfaces (BCIs) and cognitive neuroscience, offering millisecond-level temporal precision despite noise and individual variability [Lotte et al., 2018, Schirrmeister et al., 2017].

Early research used system identification for stimulus reconstruction, e.g., predicting attended speakers from single-trial EEG [O’sullivan et al., 2015]. Recent advances combine self-supervised and contrastive learning to align neural signals with pretrained speech representations (e.g., wav2vec 2.0), achieving up to 41% accuracy across 1,000+ segments [Défossez et al., 2023]. MEG generally outperforms EEG, while functional near-infrared spectroscopy (fNIRS) and functional magnetic resonance imaging (fMRI) balance spatial resolution, latency, and portability [Cao et al., 2021].

31 **2.2 Deep learning for time-series analysis**

32 Deep networks extract nonlinear, hierarchical features from raw time series, replacing handcrafted
33 engineering [Gamboa, 2017]. RNNs, LSTMs, and GRUs capture long-range dependencies for
34 EEG classification and seizure prediction [Karim et al., 2017]. CNNs and temporal convolutional
35 networks (TCNs) model local spatiotemporal patterns efficiently [Walther et al., 2023], while
36 Transformers exploit self-attention for global dependencies. Hybrid CNN-LSTM and CNN-
37 Transformer models combine both advantages, improving robustness [Sun et al., 2021]. These
38 architectures have been adapted for multi-channel neural data with spatial filtering and frequency
39 decomposition, supporting end-to-end decoding from sensors to cognition.

40 **2.3 Speech and phoneme decoding from brain signals**

41 EEG and MEG encode phonetic and speech representations. Early studies showed that MEG
42 distinguishes phonetic contrasts and speech onset [Lukka et al., 2000], and later models decoded
43 overt and imagined speech with over 90% accuracy under limited vocabularies [Dash et al., 2020,
44 LaRocco et al., 2023]. Despite progress, most datasets remain small. New corpora such as
45 *LibriBrain* and *SpeechImagery* enable cross-subject and cross-session generalization [Özdogan et al.,
46 2025, Moreira et al., 2025]. Several works leverage phonological or articulatory features (e.g.,
47 place and manner of articulation) as intermediate targets, providing interpretable and biologically
48 plausible representations Wang et al. [2012].

49 **2.4 Domain adaptation and distribution shift**

50 MEG decoding faces strong distribution shifts across sessions and subjects, degrading transfer
51 performance. Solutions include (1) **knowledge distillation**, using teacher-student frameworks
52 to align representations [Meng et al., 2019]; (2) **MMD alignment**, minimizing maximum mean
53 discrepancy between source and target domains [Gretton et al., 2012]; (3) **data augmentation**,
54 such as time-frequency perturbation and channel dropout; and (4) **adversarial or self-supervised**
55 **learning**, enforcing domain-invariant priors [Lin and Zhang, 2023, Wang, 2025]. Together, these
56 methods support robust and transferable neural decoding under realistic distribution shifts.

57 **3 Methods**

58 **3.1 Dataset and preprocessing**

59 Each example consists of a 500 ms window centered on a phoneme, recorded across 306 MEG
60 channels in 250 Hz sample rate [Özdogan et al., 2025]. We adopt the competition’s recommendation
61 to average windows within each class. Unless noted, we form groups of size 100 and repeat the
62 sampling procedure three times to balance denoising with data diversity.

63 **The train validation set is selected differently** from the PNPL library, as detailed in Appendix A.
64 Both the train and validation sets are standardized with training statistics; the test set, containing the
65 same examples as the validation set, is standardized with its own statistics to simulate the holdout
66 set conditions, where input features are standardized per split. This mismatch creates a distribution
67 shift analyzed in Section 5 and Appendix E.

68 **3.2 Label representations**

69 MEG labels are supplied as phoneme identifiers. We additionally map each phoneme to a 17-
70 dimensional ternary vector of articulatory features derived from PanPhon [Mortensen et al., 2016].
71 From the original 21 features, we remove non-contrastive dimensions (e.g., “sg” for “spread
72 glottis”) to reduce redundancy. Optionally, we binarize the remaining values from $\{-1, 0, 1\}$ to
73 $\{0, 1\}$ by treating -1 and 0 as absence of the feature. Table 1 illustrates the mapping for some
74 phonemes. During training, the auxiliary regression target is optimized with mean-squared error;
75 during inference, output is selected based on cosine similarity to the nearest phoneme.

Table 1: Example phonological vectors with their corresponding phonemes after removing non-contrastive features.

Phoneme	syl	son	cons	cont	delrel	lat	nas	strid	voi	ant	cor	lab	hi	lo	back	round	tense
/p/	—	—	+	—	—	—	—	0	—	+	—	+	—	—	—	0	
/b/	—	—	+	—	—	—	—	0	+	+	—	+	—	—	—	0	
/t/	—	—	+	—	—	—	—	0	—	+	+	—	—	—	—	0	
/d/	—	—	+	—	—	—	—	0	+	+	+	—	—	—	—	0	

76 3.3 Data augmentation

77 To mitigate class imbalance and sensor variability, we rely on lightweight augmentations applied
 78 randomly: additive Gaussian noise, temporal shifts, temporal masking, channel dropout, amplitude
 79 scaling, and frequency band perturbation. Detailed hyperparameters are provided in Appendix B.

80 3.4 Distribution alignment

81 We experiment with a small fully connected mapper trained with maximum mean discrepancy
 82 (MMD) to align standardized-then-averaged test features to the training distribution [Gretton et al.,
 83 2012]. The MMD is computed between source features X and target features Y as:

$$\text{MMD}(X, Y) = \frac{1}{n^2} \sum_{i,j} k(x_i, x_j) + \frac{1}{m^2} \sum_{i,j} k(y_i, y_j) - \frac{2}{nm} \sum_{i,j} k(x_i, y_j),$$

84 The mapper f_θ is composed of three linear layers (input and output dimensions 306×125 , hidden
 85 dimensions 512) with ReLU activations. Training minimizes MMD with a Gaussian kernel of
 86 bandwidth 10 using Adam (learning rate 10^{-4} , batch size 128, 50 epochs), where source features
 87 are drawn from the training set and target features from the test set after standardization and group
 88 averaging.

89 Appendix C visualizes a principal component projection of source, mapped, and target features.
 90 While the mapped features match the target mean, they collapse variance, explaining the reduced
 91 discrimination observed on the leaderboard.

92 4 Model architectures

93 **ResNet-style CNN.** Our primary model is based on the baseline model provided by Özdogan et al.
 94 [2025], which stacks temporal convolutions with residual connections and group normalization . A
 95 lightweight classifier head predicts phoneme logits and optional phonological features.

96 **STFT CNN.** We apply a short-time Fourier transform (STFT) to each channel (window size 25,
 97 hop 5) and share a 2D CNN across channels. The branch is trained jointly with the time-domain
 98 CNN but performs worse when evaluated on the hidden set.

99 **CNN-Transformer hybrid.** We append a 4-layer Transformer encoder with 8 attention heads to
 100 capture long-range dependencies across sensors and time. The module improves validation accuracy
 101 but amplifies overfitting when the evaluation statistics differ from training data.

102 Detailed model diagrams are provided in Appendix D.

103 5 Results

104 Table 2 summarizes representative submissions. All models are implemented in PyTorch, trained
 105 using a single RTX 5090 GPU by 10 epochs, and use group averaging with batches of 100 windows.
 106 Extended results with ablations are provided in Appendix E.

107 Some configurations are not submitted to the holdout set, denoted by dashes. The PanPhon feature
 108 experiments omit training F1-macro because optimization happens on continuous articulatory
 109 vectors with an MSE loss. Producing phoneme predictions requires a nearest-neighbor projection

110 back into the discrete inventory, which we only execute on validation and leaderboard splits to avoid
111 repeatedly decoding the entire training corpus each epoch.

Table 2: Representative F1-macro scores (%) for LibriBrain phoneme classification. Aug. denotes stochastic augmentations.

Configuration	Train	Validation	Test	Holdout	Aug.	Notes
CNN + label-balancing	90.81	71.95	47.47	35.40	No	Repeated grouping $\times 10$
CNN + LayerNorm	96.62	44.49	43.17	24.40	No	Layer normalization
CNN baseline	34.55	45.08	39.53	13.20	No	No group averaging
CNN + Augmentation	48.55	49.31	34.03	18.80	Yes	As in Section 3.3
CNN-Transformer	85.83	68.02	30.70	3.90	No	
STFT CNN	62.63	43.62	15.91	—	No	STFT ($N_{\text{fft}} = 25, H = 5$)
CNN-Transformer	—	51.68	3.33	—	No	Ternary PanPhon
CNN baseline + Dist. Mapper	—	—	0.06	—	No	

112 Simple CNNs with label balancing and group averaging achieve the best generalization; the
113 performance gap widens when models depend on fine-grained normalization such as the CNN-
114 Transformer hybrid. Adding phonological supervision yields minor validation gains but performs
115 poorly on the test set, suggesting failure to generalize under distribution shift. Similarly, STFT
116 preprocessing improves validation but degrades test performance. However, it remains unknown
117 whether CNN variants with phonological targets would outperform time-domain models if evaluated
118 on the holdout set.

119 Train-to-validation gaps highlight generalization ability. CNN with layer normalization achieves
120 96.62% train F1 but drops to 44.49% on validation, indicating overfitting. Label-balanced CNN
121 maintains 71.95% validation F1 from 90.81% training, showing better robustness. Ill-formed
122 normalization layers may overfit training statistics, exacerbating domain shifts. CNN-Transformer
123 hybrid retains 68.02% validation F1 from 85.83% training, suggesting attention mechanisms capture
124 more invariant features.

125 Validation-to-test comparisons reveal the impact of misaligned standardization. All models we've
126 evaluated show significant drops from validation to test. In comparison, test-to-holdout gaps
127 are fairly consistent (around 20%) across configurations, suggesting that the primary distribution
128 shift arises from differing standardization statistics rather than other latent factors. Attempts to
129 compensate with the distribution mapper (Section 3.4) improved qualitative alignment but reduced
130 class separability, offering limited practical benefit.

131 6 Discussion and conclusion

132 **Contributions.** Our study highlights that how we aggregate MEG signals matters more than
133 architectural novelty: label-balanced, group-averaged CNNs retain two-thirds of their training F1
134 on the leaderboard, while more expressive models collapse once the statistic mismatch is introduced.
135 The most reliable recipe couples conservative preprocessing with restrained capacity, indicating that
136 consistency across splits is a stronger signal of leaderboard success than raw training accuracy.

137 **Limitations.** Although it's tempting to attribute performance drops solely to distribution shifts,
138 the reality is more complex. Factors such as model capacity, training dynamics, and the inherent
139 variability of MEG signals all play a role.

140 Also, it remains unclear how well our findings generalize to other well-performed configurations
141 and preprocessing pipelines from other teams. Future work should systematically evaluate a broader
142 range of architectures and training strategies under controlled distribution shifts to validate the
143 robustness of our conclusions.

144 **Future directions.** For future progress, combining distribution-aware normalization, adaptation
145 methods that preserve intra-class spread, and supervision that retains the richness of phonological
146 features while remaining compatible with the leaderboard metric may yield more robust speech
147 decoding from MEG signals. Additionally, exploring self-supervised pretraining on large-scale
148 unlabeled MEG data could help models learn invariant representations that generalize better across
149 sessions and subjects.

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204 **A Dataset details**

205 Our dataset splits differ from those in the PNPL library. Table 3 summarizes the number of
206 trials, phoneme counts, and standardization parameters for each split. The validation set for this
207 competition phase is composed of 14 specific trials, listed in Table 4 [Landau et al., 2025]. The
208 test set contains the same trials but is standardized with its own statistics to simulate the holdout set
209 conditions.

Table 3: Summary of LibriBrain phoneme classification dataset splits.

Split	Trials	Phonemes	Standardization
Train	76	1,336,606	Train statistics
Validation	14	283,690	Train statistics
Test	14	283,690	Test statistics
Holdout	1	2,382	Holdout statistics
Total	91	1,622,678	–

Table 4: Validation trials for the dataset splits.

Subject	Session	Task	Trial
0	11	Sherlock1	2
0	12	Sherlock1	2
0	11	Sherlock2	1
0	12	Sherlock2	1
0	11	Sherlock3	1
0	12	Sherlock3	1
0	11	Sherlock4	1
0	12	Sherlock4	1
0	14	Sherlock5	1
0	15	Sherlock5	1
0	13	Sherlock6	1
0	14	Sherlock6	1
0	13	Sherlock7	1
0	14	Sherlock7	1

210 **B Augmentation recipes**

211 For reproducibility we detail the augmentation hyperparameters:

- 212 • Gaussian noise with standard deviation 0.01 relative to the sample standard deviation.
213 • Temporal shift uniformly sampled in ± 40 ms with circular padding.
214 • Temporal masking zeros out at most 80 ms of contiguous samples.
215 • Channel dropout randomly zeros 10% of sensors per step.
216 • Amplitude scaling multiplies the waveform by a factor drawn from $\mathcal{U}(0.9, 1.1)$.
217 • Frequency band perturbation performs a Fourier transform of the input signal in 100 Hz,
218 randomly selects one or more frequency bands from the spectrum, and scales their
219 amplitudes by a random 0.8 to 1.2 factor within a specified range to emulate varying
220 spectral conditions. The modified spectrum is then transformed back into the time domain.
221 • Each augmentation is applied independently with probability 0.3.

222 **C Distribution alignment**

223 The distribution shift between each trials is visualized as in Figure 1, where the pairwise channel
224 mean cosine similarities are computed with raw features without standardization or group averaging.

225 While some pairs of trials show high similarity (e.g. Sherlock5, Session 11 to 13), most trials differ
 226 significantly with cosine similarities around 0.5. This variance likely contributes to the distribution
 227 shift observed between validation and hidden evaluation data.

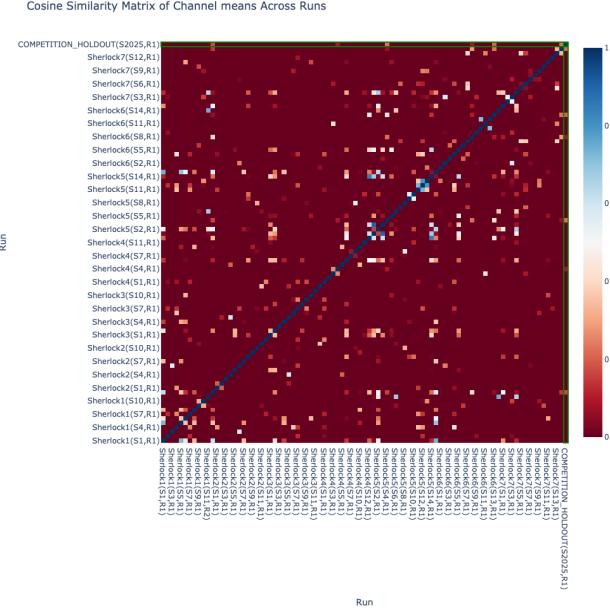


Figure 1: Pairwise cosine similarities between trials based on channel means. Trials used in holdout are highlighted with green boxes.

228 Figure 2 visualizes a PCA projection of source, mapped, and target features after training the
 229 MMD-based mapper. While the mapped features align with the target mean, they collapse variance,
 230 explaining the reduced discrimination observed on the leaderboard.

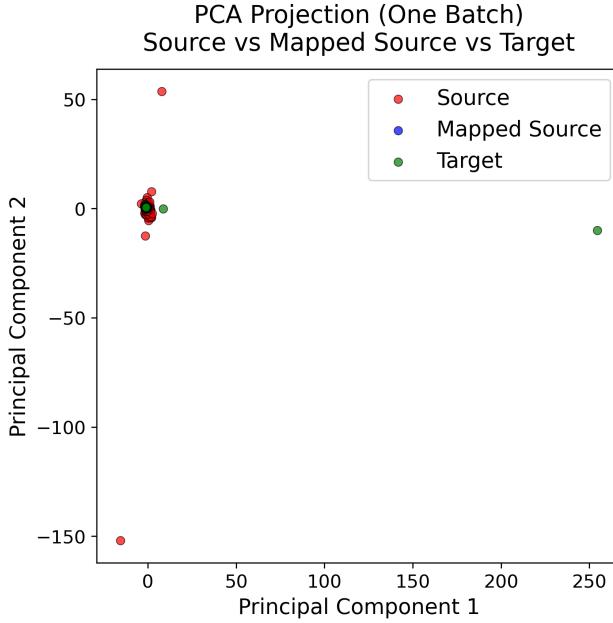


Figure 2: PCA projection of one batch where the mapper pulls source features (red) toward the target distribution (green) but compresses variance.

231 **D Detailed model configurations**

232 Figure 3 illustrates the three model architectures evaluated. The CNN backbone (a) mainly consists
 233 of several 1D convolutional blocks with channel size $D = 256$ and 2 residual connections. An
 234 optional normalization layer (layer norm or batch norm) is added as shown in (a). The STFT-
 235 CNN model (b) applies STFT to each channel independently before feeding into a ResNet-based
 236 architecture with 2D convolutions. The CNN-Transformer hybrid (c) appends a 4-layer Transformer
 237 encoder after a CNN “filtering” stage, attempting to capture long-range dependencies across sensors
 238 and time.

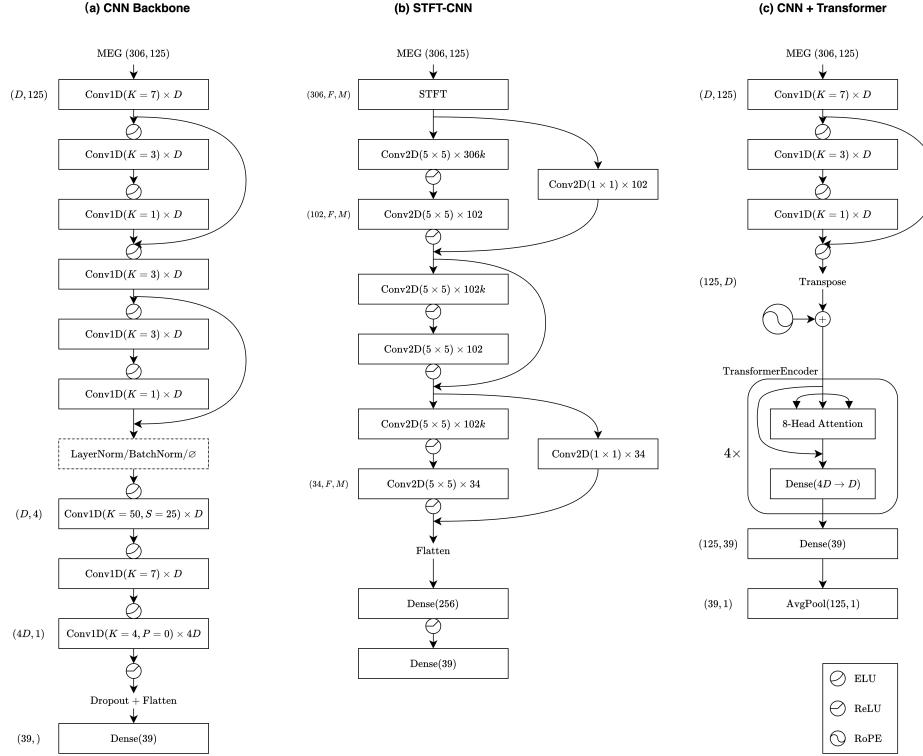


Figure 3: Overall model architecture. For the convolutional blocks, the stride S is 1 and padding P is set to maintain the same temporal dimension if not specified. (a) shows the basic CNN backbone, with optional layer norm or batch norm layer. (b) shows the STFT-CNN architecture, which applies STFT to each channel independently before feeding into the specialized ResNet-based model. (c) shows the CNN-Transformer architecture, which has a 4-layer Transformer encoder after several convolutional layers.

239 **E Extended results and visualizations**

240 Table 5 reports the full leaderboard history. All models use group averaging with window
 241 size 100. “Repeat” specifies the number of iterations through the dataset that is then group-averaged.
 242 Additionally, the 2nd configuration in Table 2 is used to generate the confusion matrix in Figure ??.

243 Figure 4 shows confusion matrices for the best-performing validation and leaderboard submissions.
 244 The same model sharply degrades on the leaderboard, confirming the distribution shift.

Table 5: Extended comparison of model and data configurations. Performance is reported in F1-macro (%). A dash denotes runs not submitted to the holdout set.

Train	Validation	Test	Holdout	Backbone	Group Avg.	Label Bal.	Repeat	Note
68.16	64.27	47.74	3.60	CNN	Yes	Yes	1	+Augmentation
90.81	71.95	47.47	35.40	CNN	Yes	Yes	10	
78.42	70.09	45.02	—	CNN	Yes	Yes	3	
80.06	71.77	44.28	21.60	CNN	Yes	Yes	1	
96.62	44.49	43.17	24.40	CNN + LayerNorm	Yes	Yes	1	
91.50	44.31	42.84	—	CNN + BatchNorm	Yes	Yes	1	
34.55	45.08	39.53	13.20	CNN	No	No	1	
48.55	49.31	34.03	18.80	CNN	Yes	No	1	+Augmentation
85.83	68.02	30.70	3.90	CNN + Transformer	Yes	Yes	1	
62.63	43.62	15.91	—	STFT CNN	Yes	Yes	5	STFT, $N_{\text{ft}} = 25$
62.05	40.00	15.78	—	STFT CNN	Yes	Yes	5	STFT, $N_{\text{ft}} = 50$
41.97	30.01	8.98	—	STFT CNN	Yes	Yes	5	STFT, $N_{\text{ft}} = 25, H = 20$
74.37	64.86	4.29	—	CNN + Transformer	Yes	Yes	1	
—	51.68	3.33	—	CNN + Transformer	Yes	Yes	1	Ternary PanPhon,
75.22	63.82	0.98	—	CNN + Transformer	Yes	Yes	1	
—	57.21	0.88	—	CNN + Transformer	Yes	Yes	1	Binary PanPhon,
—	—	0.06	—	CNN	Yes	Yes	1	w/ Distribution Mapper
—	—	0.12	—	CNN + Transformer	Yes	Yes	1	w/ Distribution Mapper

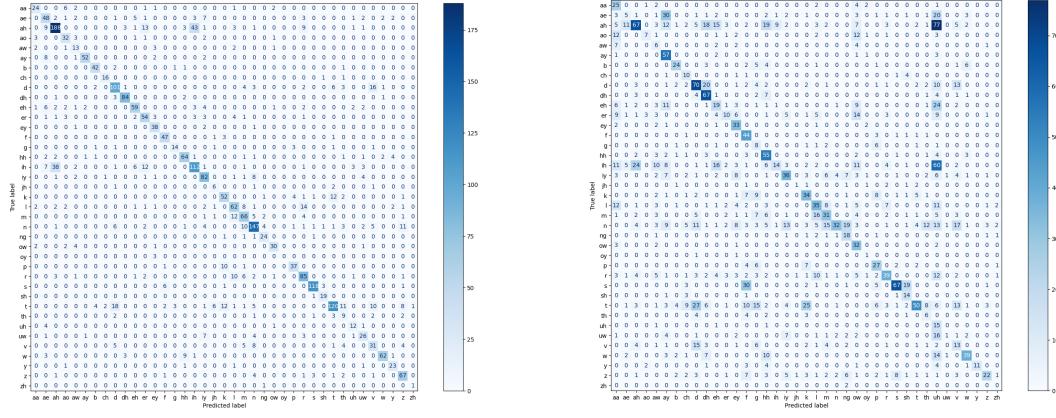


Figure 4: Confusion matrices for the 2nd configuration in Table 5 on validation (left) and leaderboard (right) data.