A guide for fitting generalized linear models in Tensorflow

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Generalized linear model (GLM)

- Flexible generalization of ordinary linear regression
 - Response variable y follows a specific distribution (noise model)
 - A function of the mean response variable (*link function*) varies linearly with the predictors *X*

$$f(\mu) = Xw, y \sim Dist(\mu, ...)$$

Model type	Distribution	Link function	Mean function (inverse function, activation function)
Linear regression	Normal (Gaussian)	Identity $f(\mu) = \mu$	Linear μ = Xw
Poisson regression	Poisson	$\log f(\mu) = \log(\mu)$	Exponential $\mu = \exp(Xw)$
Logistic regression	Bernoulli	Logit $f(\mu) = \log(\mu/(1-\mu))$	Sigmoid $\mu = \exp(Xw) / (1 + \exp(Xw))$

Fitting GLM in Tensorflow

- Why Tensorflow?
 - GPU acceleration for massive parallelization
- A typical session with simultaneous recording
 - 200-300 neurons per imaging plane
 - ~30,000 timepoints
 - ~1000 predictors (after basis expansion)

$$f(Y) = XW \leftarrow 1000 \times 250$$

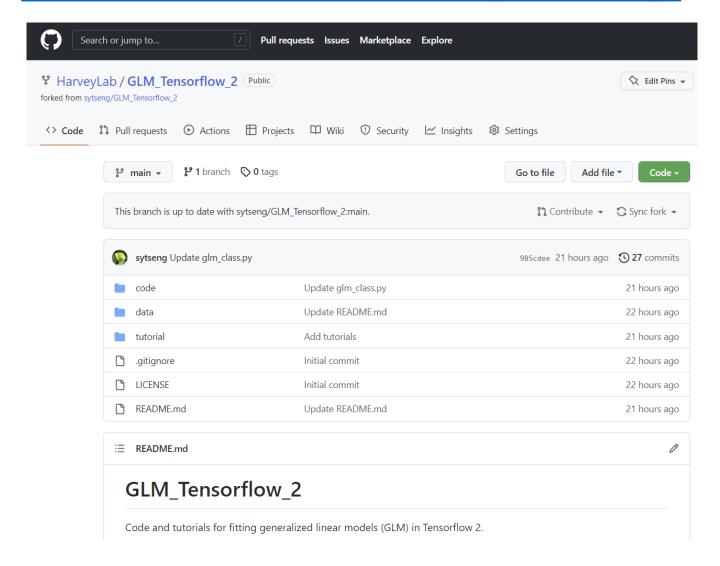
 \uparrow
 $30,000 \times 250$ $30,000 \times 1000$

- Formalized as a multi-output,
 one-layer feedforward neural network
- Combine all outputs for a single loss
- Gradient descent optimization

Fitting takes 5-10 min for Poisson GLMs (1-2 sec per neuron!) with 5-fold cross validation over >10 regularization values

Github repo for GLM in Tensorflow

https://github.com/HarveyLab/GLM Tensorflow 2



Tutorials for fitting GLM in Tensorflow

Two tutorials

- Tutorial_for_using_GLM_class.ipynb
- Tutorial_for_fitting_neural_calcium_imaging_data_with_
 GLM.ipynb

GLM class

- Written in Python, object-oriented
- Can be loaded as a module

```
import glm_class as glm
```

Initialization

```
model = glm.GLM() or model = glm.GLM CV()
```

Four class methods

```
model.fit()
model.select_model()
model.predict()
model.evaluate()
```

Model initialization

Four types of input arguments

- Model type:
 - activation: 'linear', 'exp', 'sigmoid', 'relu', 'softplus'
 - loss_type: 'gaussian', 'poisson', 'binominal'
- Regularization:
 - regularization: 'elastic_net', 'group_lasso'
 - lambda_series: list of regularization strengths in descending order
 - /1_ratio: 0 for L2/ridge, 1 for L1/Lasso, or btw 0-1 for elastic net
 - *smooth_strength:* add smoothness penalty
- Optimization: optimizer ('adam', 'sgdm'), learning_rate, momentum
- Convergence: min_iter_per_lambda, max_iter_per_lambda, num_iter_check, convergence_tol

Class methods

Fitting (with training data)

A list containing the size of each group For group_lasso or smoothness penalty

Model selection (with validation data)

```
model.select_model(X_val, Y_val, [min_lambda, make_fig])
```

Evaluation (with training, validation, and test data)

Prediction (with training, validation, and test data)

```
Y_pred = model.predict(X)
```

Model attributes

Loss & lambda trace

```
model.loss_trace and model.lambda_trace
```

Selected weights & intercepts

```
model.selected w and model.selected w0
```

Selected lambda & lambda indices

```
model.selected_lambda and model.selected_lambda_ind
```

GLM with cross validation

Model initialization

- Additional input arguments
 - *n_folds:* number of CV folds
 - auto_split: perform CV split automatically
 - split_by_group: split CV folds according to a third-party group provided during fitting time
 - split_random_state: Numpy random state for splitting

GLM with cross validation

Model fitting

```
Train and validation indices for each fold Third party group info if auto_split = False if split_by_group = True model_cv.fit(X, Y, [train_idx, val_idx, group_idx, initial_w0, initial_w, feature_group_size, verbose])
```

Model selection (no validation data required)

```
model_cv.select_model([se_fraction, min_lambda, make_fig])

Controls the tolerance of choosing models with smallest deviance vs. larger regularization;

O for model with minimal deviance, 1 for 1SE rule
```

Same methods for model evaluation and prediction

Useful utility functions

Null deviance (avg for each response, or for every datapoint)

```
null_dev = null_deviance(y, loss_type = 'poisson')
null_dev_pt = pointwise_null_deviance(y, loss_type = 'poisson')
```

Deviance (avg for each response, or for every datapoint)

```
frac_dev_expl, d_model, d_null = deviance(y_pred, y_true, loss_type = 'poisson')
dev_pt = pointwise_deviance(y_true, y_pred, loss_type = 'poisson')
```

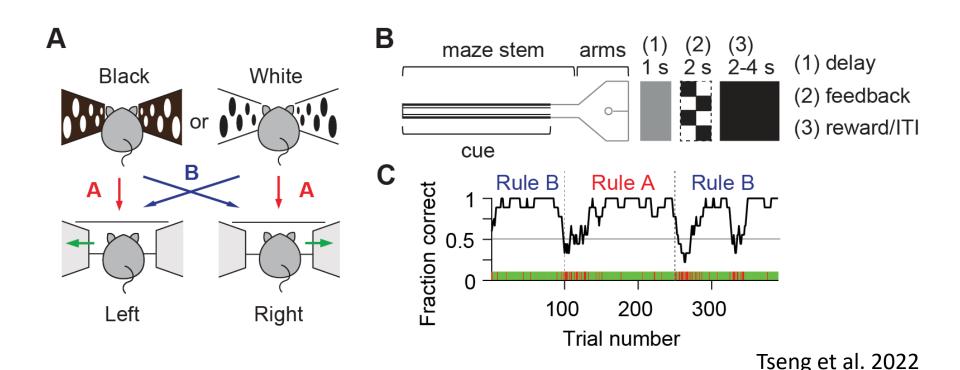
 Make prediction given design matrix, weights and intercepts, and activation function

```
prediction = make prediction(X, w, w0, activation = 'exp')
```

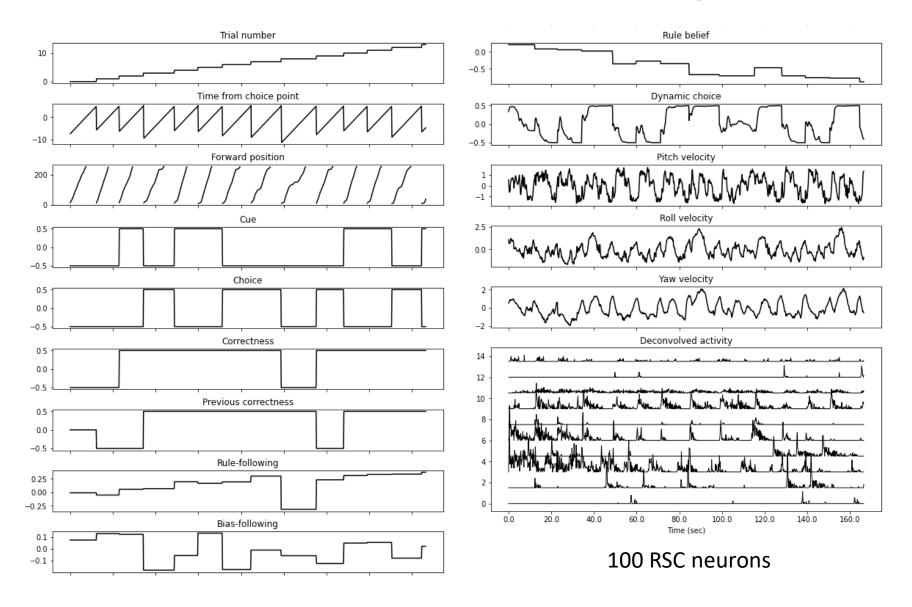
Fitting GLM to calcium imaging data

Tutorial_for_fitting_neural_calcium_imaging_data_with_GLM.ipynb

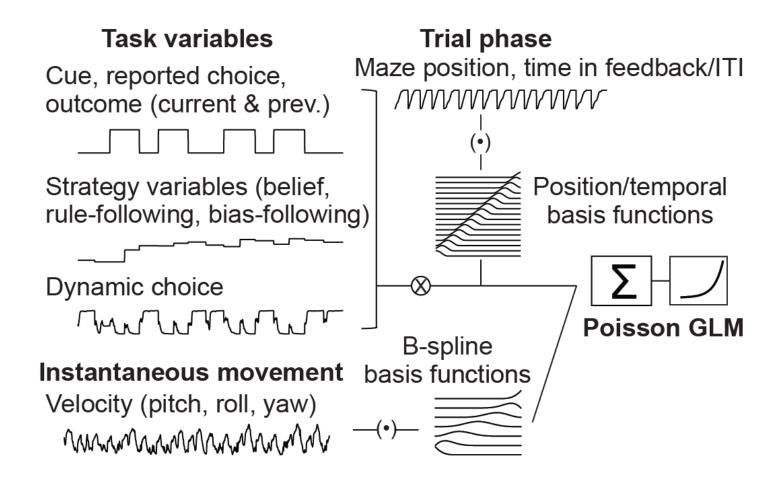
- Building design matrix (basis expansion)
- Model fitting, selection and evaluation
- Quantification of feature contribution



Predictors vs. neural activity



Model schematic



Position basis expansion

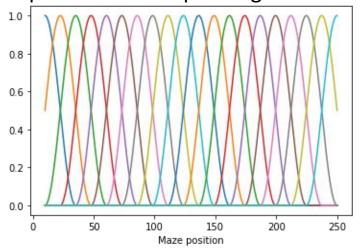
"Place field-like" tuning

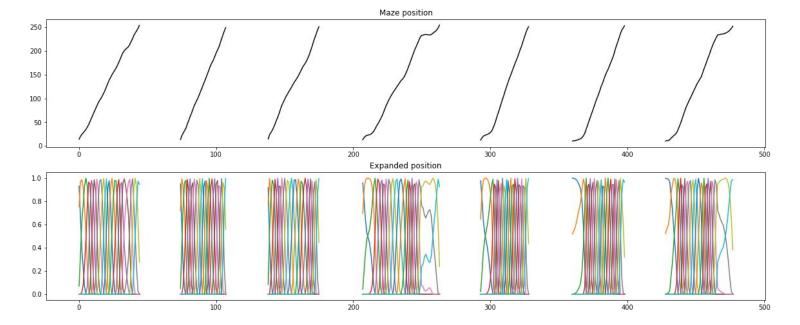
Raised cosine bumps

$$f_i(x) = \begin{cases} \frac{1}{2}\cos(ax - \varphi_i) + \frac{1}{2} & \text{if } \varphi_i - \pi < ax \le \varphi_i + \pi \\ 0 & \text{otherwise} \end{cases}$$

if
$$\varphi_i - \pi < ax \le \varphi_i + \pi$$
 otherwise

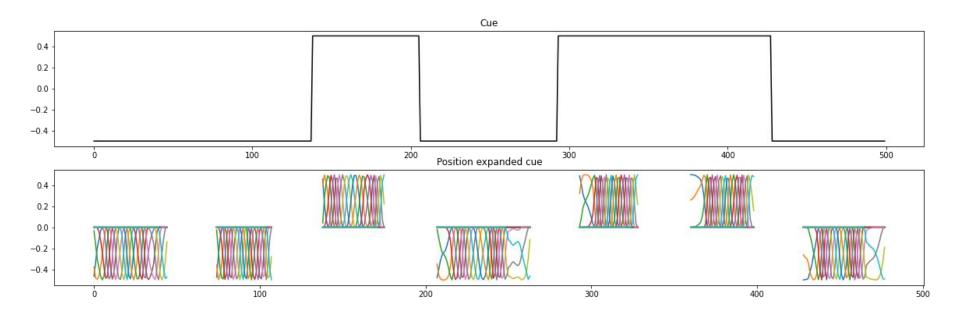






Load task variables onto position basis functions

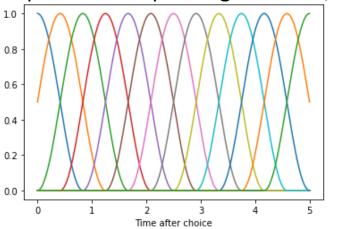
Simply take the **product** of the time series of task variables and each of the position basis functions, i.e. create **interaction terms**

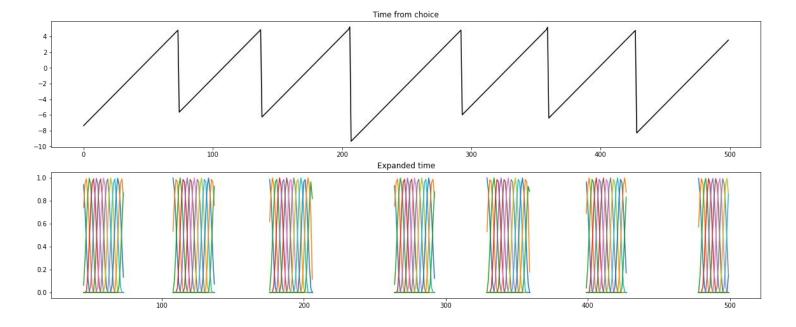


Temporal basis expansion

For time elapsed in feedback/ITI, follow the same principle to create temporal basis functions and load task variables onto them

13 temporal bases spanning feedback/ITI





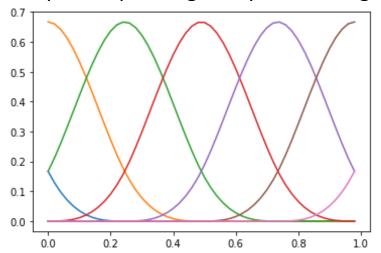
B-spline expansion for velocities

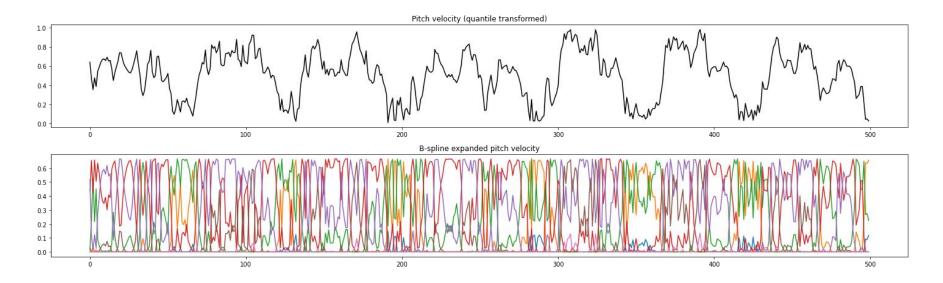
For movement variables, we assume consistent relationship btw neural activity and velocities (in maze vs. ITI)

Non-linear velocity tuning:

B-spline expansion (degree 3 polynomial) Quantile transformation: easy comparison

7 b-splines spanning the quantile range

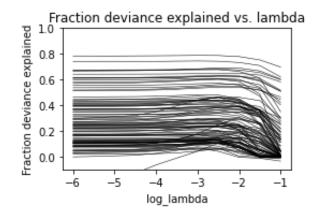


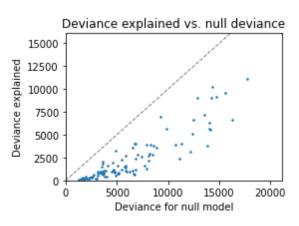


Fitting

- Poisson GLM, 5-fold cross validation
 - $Y = exp(XW + W_0)$
 - Multiply deconvolved activity by 10 -> mimic spike count
- Regularization: group lasso
 - L2 for features belong to the same basis expanded groups
 - L1 between basis expanded groups
- Evaluation

fraction deviance explained =
$$1 - \frac{\text{model deviance}}{\text{null deviance}}$$





Quantification of feature contribution

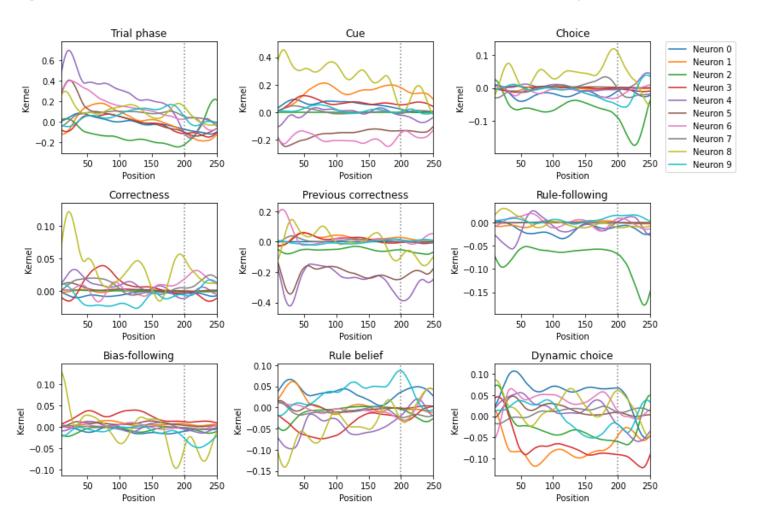
Model weights (kernels)

 Fraction null deviance or fraction explained deviance through model breakdown

Fraction null deviance through model comparison with re-fitting

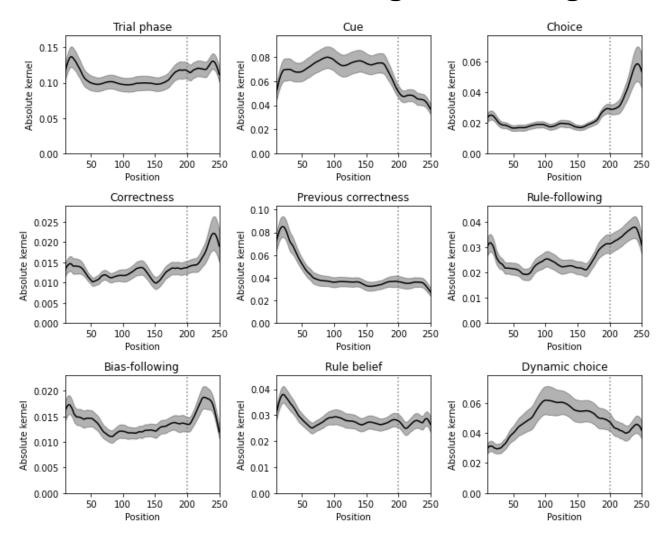
Model weights (kernels)

Weight vector x basis functions -> sum up into kernels



Model weights (kernels)

Mean absolute kernels: strength of tuning



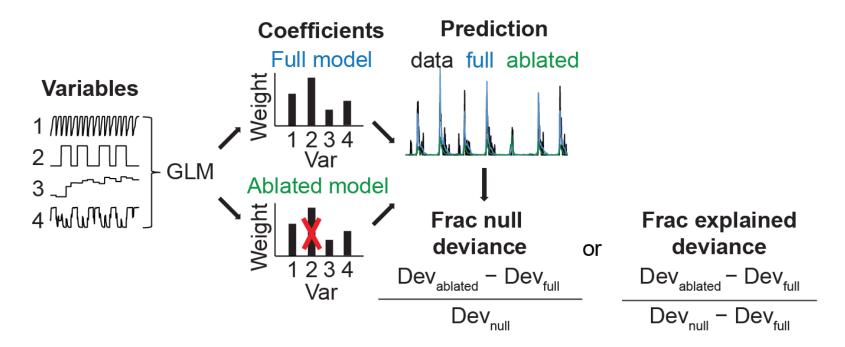
Model weights (kernels)

- Multiplicative factor on top of baseline activity
 - $y = \exp(w_0 + w_1x_1 + w_2x_2 + ...)$
 - Exponentiate to obtain "gain"
- Informative of directionality (signed)
- Robust to scaling of y
- Hard to interpret the magnitude and compare across variables
 - Better with "fraction null deviance" or "fraction explained deviance"

Model breakdown

"Model breakdown"

Quantification of variable contribution (without re-fitting)

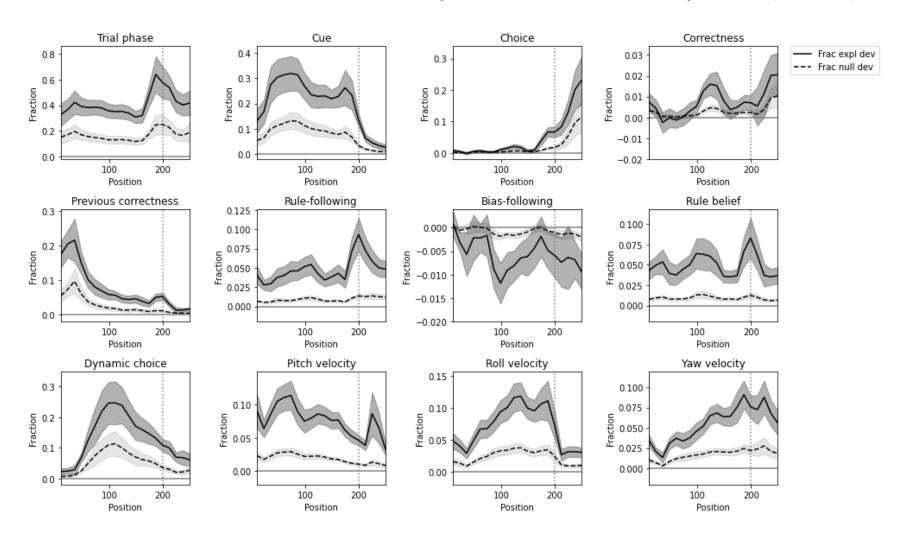


Zero weights = zero variables (must be zero-centered) = shuffle variables

Evaluate on CV held-out data (85%) instead of test data (15%)

Model breakdown

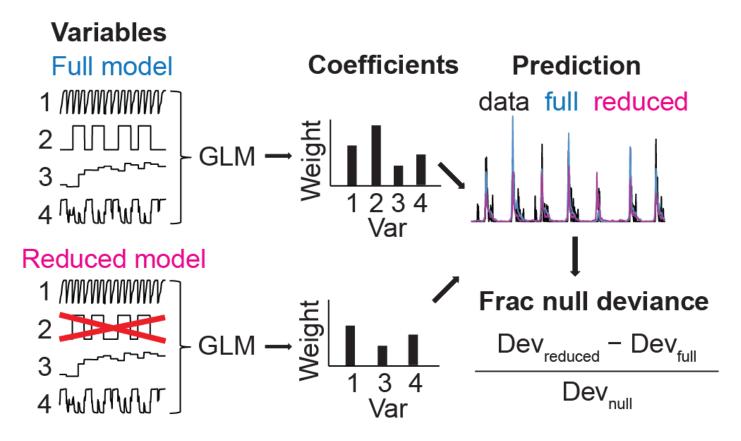
Evaluate fraction null deviance or fraction explained deviance at each position (or time) bin



Model comparison (with re-fitting)

"Model comparison"

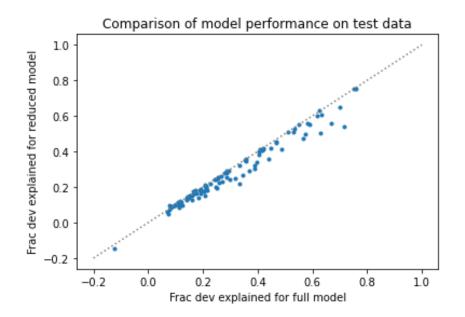
Quantification of variable contribution (with re-fitting)



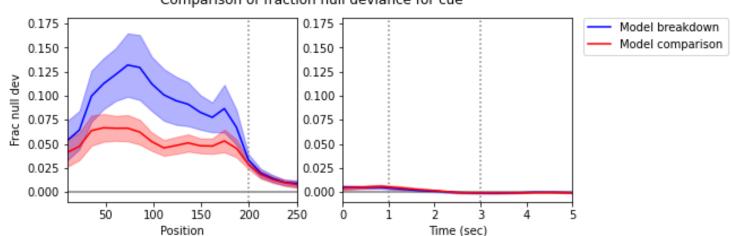
Remove variable = shuffle variable

Model comparison (with re-fitting)









Model comparison vs. breakdown

Model breakdown (fraction null deviance and fraction explained deviance)

- The metrics are easy to interpret and compare across variables
- Does not tell you the directionality of tuning
- Does not require re-fitting the model (fast to do)
- Not a linear breakdown for Poisson GLMs, but linear for Gaussian
- Can be less biased in assigning contribution of correlated variables, if the model is not terribly mis-specified

Model comparison (fraction null deviance)

- The metric is easy to interpret and compare across variables
- Does not tell you the directionality of tuning
- Require re-fitting of a model for each variable (computationally demanding)
- Extract the "unique" portion of deviance explained, can be biased (and underestimate the contribution of a variable) when there are correlated variables and you don't know which is actually contributing to response

References

- Tseng, S.-Y., Chettih, S.N., Arlt, C., Barroso-Luque, R., and Harvey, C.D. (2022). Shared and specialized coding across posterior cortical areas for dynamic navigation decisions. Neuron 110, 2484–2502.e16. [link]
- Minderer, M., Brown, K.D., and Harvey, C.D. (2019). The spatial structure of neural encoding in mouse posterior cortex during navigation. Neuron 102, 232–248.e11. [link]