



Using Random Effects to Account for High-Cardinality Categorical Features and Repeated Measures in Deep Neural Networks

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Outline

- Motivation
- Linear Mixed Models
- Our approach: LMMNN
- Results
- Convergence
- Extensions

High cardinality categorical features

- Using EMR of hospital patients to predict hospital readmission, the feature disease could take 1 out of thousands of values (Lin et. al., 2019)
- Using the CelebA facial images dataset to develop a computer vision model to localize facial features, several images from the same person (a.k.a *repeated measures*), over 10K identities (Liu et. al., 2015)
- Predicting the price of a Airbnb rental, ~40K hosts in NYC alone (Kalehbasti et al., 2019)

Current solutions

- One-hot encoding (OHE) $\rightarrow q$ levels to q additional binary features
- Entity embeddings \rightarrow reduce dimensionality from q to d with a learned $D_{q \times d}$ dictionary
- Ignore!
- Other (Hancock and Khoshgoftaar, 2020):
 - Clustering (expert knowledge, "other" strategy, clustering algos)
 - Feature Hashing
 - Supervised numerical encoding

TL;DR

		Ta	ble 4: Real da	ta feat	tures sui	nmary (table			
Dataset	n	q	p	Cat	Categorical		y		ype	DNN
UKB PA Drugs CelebA Airbnb	96K 215K 202K 50K	350 3.6K 10K 40K	15 10K 218x178x3 196		g ntity t	PA rating noseX-Y log(price)		Tabular Text Images Tabular		MLP LSTM CNN MLP
Tab	Table 5: Real data 5-CV mean test MSEs and standard errors in parentheses.									
Dataset	Dataset Ignore			OHE Em		Embeddings MeNe		ets LM		MNN
UKB PA Drugs CelebA no CelebA no Airbnb	UKB PA0.812 (.008)Drugs2.74 (.032)CelebA noseX1.68 (.05)CelebA noseY1.64 (.09)Airbnb0.156 (.002)		08) 0.816 (.0 2) 2.77 (.00) –) – 02) –	009) 0.817 (.0 05) 2.72 (.0 3.6 (.3) 2.5 (.2) 0.158 (.0		(.010) 051)) (.003)	0) 0.811 (.009)) 2.81 (.031) 7.6 (.3) 12.3 (1.1) 3) 0.153 (.003)) 0.809 (.008) 2.66 (.006) 1.54 (.07) 1.39 (.04)) 0.142 (.002)	

Linear Mixed Models (LMM) (I)



Random Intercepts model:

- single categorical (RE) feature of q levels
- $D = \sigma_b^2 I_q$
- Z is a binary matrix
- s.t. $y_{ij} = \beta_0 + \beta_1 x_{ij,1} + \dots + \beta_1 x_{ij,p-1} + b_j + \varepsilon_{ij}, i = 1, \dots, n_i, j = 1, \dots, q$

Linear Mixed Models (LMM) (II)

Marginal distribution of y:

 $y \sim N(X\beta, V(\psi))$

where ψ are variance components to estimate and $V(\psi) = ZDZ' + \sigma_e^2 I_n$

Log-likelihood of \$\beta\$, \$\psi\$:

$$l(\beta, \psi) = -\frac{1}{2}(y - X\beta)'V(\psi)^{-1}(y - X\beta) - \frac{1}{2}\log|V(\psi)| - \frac{n}{2}\log 2\pi$$

Get \$\beta\$, \$\psi\$ via MLE/REML

- Predict (BLUP) $\hat{b} = DZ'V(\hat{\psi})^{-1}(y X\hat{\beta})$ (avoid inversion for random intercepts...)
- Predict $\hat{y}_{te} = X_{te}\hat{\beta} + Z_{te}\hat{b}$ or $\hat{y}_{te} = X_{te}\hat{\beta}$ for unknown levels

Random effects: what for

By treating high-cardinality categorical features as RE in DNN, we hope to:

- Model the correlation between clustered observations better (faces of the same person!)
- Scale to higher q (only 2 additional params to estimate for random intercepts, many DNN ignore such features altogether)
- Ultimately leading to better prediction performance (e.g. MSE)
- Bonus: scale non-linear MM as well!



^{*}In this work we focus on the random intercepts model for high-cardinality categorical features in a regression setting first, see extensions later.

Our approach (LMMNN) (II)



Our approach (LMMNN) (III)

Marginal distribution of y:

 $y \sim N(f(X), V(g, \psi))$

where ψ are variance components to estimate and $V(g, \psi) = g(Z)Dg(Z)' + \sigma_e^2 I_n$

• Negative log-likelihood (NLL) of f, g, ψ is a natural loss function at each gradient descent iteration on mini-batch ξ of size m:

$$\mathrm{NLL}_{\xi}(f,g,\psi) = \frac{1}{2} \left(y_{\xi} - f(X_{\xi}) \right)' V(g,\psi)_{\xi}^{-1} \left(y_{\xi} - f(X_{\xi}) \right) + \frac{1}{2} \log \left| V(g,\psi)_{\xi} \right| + \frac{m}{2} \log 2\pi$$

• Get \hat{f} , \hat{g} , $\hat{\psi}$ via DNN optimization (mini-batch SGD, auto-differentiation)

- Predict (BLUP) $\hat{b} = DZ'V(\hat{g},\hat{\psi})^{-1}(y \hat{f}(X))$ (avoid inversion for random intercepts...)
- Predict $\hat{y}_{te} = \hat{f}(X_{te}) + \hat{g}(Z_{te})\hat{b}$ or $\hat{y}_{te} = \hat{f}(X_{te})$ for unknown levels

Other approaches

MeNets (Xiong et. al., 2019)

$$y = f(X)\beta + f(Z)b + \varepsilon$$

DeepGLMM (Tran et. al., 2020)

$$g(\mu_{it}) = f\left(x_{it}^{(1)}, \omega, \beta^{(1)}\right) + (\beta^{(2)} + \alpha_i)' x_{it}^{(2)}$$

Results: Simulated data (I)

- $y = (X_1 + \dots + X_{10}) \cdot \cos(X_1 + \dots + X_{10}) + 2 \cdot X_1 \cdot X_2 + g(Z)b + \varepsilon$
- $X_l \sim U(-1, 1); l = 1, ..., 10$ • $\sigma_e^2 = 1$
- $q \in \{100, 1,000, 10,000\}; \sigma_b^2 \in \{0.1, 1.0, 10.0\}$
- n = 100,000, 80/20% train/test split

5 repetitions

Base DNN: 4-layers MLP with [100, 50, 25, 12, 1] neurons, 25% Dropout, ReLU activation

Results: Simulated data (II)

Table 1 results	Table 1: Simulated model with $g(Z) = Z$, mean test MSEs and standard errors in parentheses. Bold results are non-inferior to the best result in a paired t-test.										
σ_b^2	q	Ignore	OHE	Embeddings	lme4	MeNets	LMMNN				
0.1	$10^2 \\ 10^3 \\ 10^4$	1.25 (.012) 1.23 (.009) 1.22 (.004)	1.20 (.010) 1.31 (.008) 1.54 (.008)	1.18 (.006) 1.24 (.004) 1.56 (.007)	2.92 (.017) 2.96 (.022) 2.97 (.014)	1.15 (.013) 1.40 (.065) 1.51 (.133)	1.14 (.010) 1.14 (.009) 1.17 (.010)				
1	$10^2 \\ 10^3 \\ 10^4$	2.17 (.041) 2.16 (.015) 2.14 (.013)	1.23 (.008) 1.39 (.015) 1.68 (.013)	1.21 (.010) 1.32 (.014) 1.68 (.013)	2.93 (.013) 2.94 (.013) 3.17 (.021)	1.22 (.022) 1.42 (.091) 1.66 (.056)	1.09 (.010) 1.14 (.006) 1.27 (.014)				
10	$10^2 \\ 10^3 \\ 10^4$	10.45 (.38) 11.37 (.11) 11.31 (.04)	1.56 (.044) 1.75 (.022) 2.34 (.027)	1.57 (.039) 1.72 (.041) 2.20 (.033)	2.92 (.012) 2.95 (.024) 3.37 (.020)	1.86 (.156) 2.19 (.143) 3.29 (.423)	1.10 (.013) 1.12 (.015) 1.32 (.007)				

Results: Simulated data (III)

Table	3: Sin	nulated model	with $g(Z) = d$	ZW, mean test	MSEs and sta	ndard errors i	n parentheses.
σ_b^2	q	Ignore	OHE	Embeddings	lme4	MeNets	LMMNN
0.1	$10^2 \\ 10^3 \\ 10^4$	1.42 (.039) 4.92 (.345) 35.1 (.456)	1.22 (.018) 1.49 (.033) 3.25 (.086)	1.19 (.024) 1.43 (.035) 3.39 (.116)	2.91 (.021) 2.95 (.020) 3.42 (.036)	1.25 (.084) 1.44 (.061) 7.35 (1.9)	1.13 (.013) 1.16 (.013) 1.59 (.023)
1	$10^2 \\ 10^3 \\ 10^4$	4.13 (.626) 35.6 (2.78) 334 (18)	1.32 (.025) 2.49 (.151) 9.13 (2.29)	1.31 (.033) 2.56 (.355) 14.4 (2.89)	2.88 (.023) 2.96 (.045) 4.29 (.1)	1.40 (.106) 7.00 (1.9) 143.3 (32)	1.16 (.011) 1.19 (.028) 3.43 (.757)
10	$10^2 \\ 10^3 \\ 10^4$	32.5 (6.86) 324 (25) 3337 (134)	1.74 (.121) 8.94 (.79) 60.1 (4.6)	2.43 (.317) 9.27 (.92) 91.1 (4.5)	2.90 (.023) 2.96 (.031) 13.8 (1.2)	12.0 (3.03) 164 (18) 2880 (463)	1.12 (.012) 1.20 (.020) 13.3 (2.0)

Results: Simulated data (IV)

Tab	ole 2: 3	Simulat	ed mode	l, estin	nated vari	ance co	mponents	on aver	rage	
			g(Z)	= Z		g(Z) = ZW				
	_	ln	ne4	LM	MNN	h	me4	LMN	ANN	
σ_b^2	q	$\hat{\sigma}_e^2$	$\hat{\sigma}_b^2$	$\hat{\sigma}_e^2$	$\hat{\sigma}_b^2$	$\hat{\sigma}_e^2$	$\hat{\sigma}_b^2$	$\hat{\sigma}_e^2$	$\hat{\sigma}_b^2$	
0.1	10^{2}	2.92	0.09	1.12	0.10	2.92	0.49	1.09	0.15	
	10^{3}	2.91	0.10	1.16	0.10	2.91	3.52	0.92	0.16	
	10^{4}	2.90	0.10	1.17	0.17	2.91	33.8	0.19	0.16	
1	10^{2}	2.90	1.01	1.12	1.00	2.91	2.44	1.07	0.41	
	10^{3}	2.90	0.98	1.15	1.00	2.90	32.0	0.84	0.43	
	10^{4}	2.90	0.99	1.26	1.02	2.92	336.6	0.19	0.35	
10	10^{2}	2.90	10.13	1.04	9.24	2.89	32.9	1.06	2.21	
	10^{3}	2.91	10.02	1.12	10.01	2.89	337.8	0.75	1.30	
	10^4	2.91	10.01	1.34	9.72	2.90	3305.6	0.22	1.98	

Results: Simulated data (V)



Results: Real data

Table 4: Real data features summary table											
Dataset	n	q	p	Ca		Categorical y		Input		ype	DNN
UKB PA Drugs CelebA Airbnb Tab	96K 215K 202K 50K	350 3.6K 10K 40K al data 5	15 10K 218x178x3 196 5-CV mean test		job drug ider host	PA g rating ntity noseX t log(pri Es and standard en		g X-Y price) errors	Tabular Text Images Tabular in parenthese		MLP LSTM CNN MLP
Dataset Ignore				OHE		Embeddings		MeNe	ets	LMMNN	
UKB PA 0.812 (.008) Drugs2.74 (.032)CelebA noseX 1.68 (.05) CelebA noseY1.64 (.09)Airbnb0.156 (.002))8) 2) () () () () 2)	0.816 (.0 2.77 (.00 - - -)09))5)	9) 0.817 (.010) 5) 2.72 (.051) 3.6 (.3) 2.5 (.2) 0.158 (.003)		0.811 (.009) 2.81 (.031) 7.6 (.3) 12.3 (1.1) 0.153 (.003)		0.80 2.60 1.54 1.39 0.14	09 (.008) 5 (.006) 4 (.07) 9 (.04) 42 (.002)	

Convergence

- Main challenge in decomposing the full NLL gradient: $V(g, \psi)^{-1}$ and $\log |V(g, \psi)|$ where V is $n \times n$ and only a $m \times m$ "part" of it is used in each mini-batch of size m
- For the simple random intercepts model with g(Z) = Z we show in the paper how the full gradient can be written exactly as the sum of q sub-gradients where each batch consists of n_j level j observations:

$$\frac{\partial NLL}{\partial \psi} = \sum_{j=1}^{q} \left[-\frac{1}{2} \left(y_j - f(X_j) \right)' V_j^{-1} \frac{\partial V_j}{\partial \psi} V_j^{-1} \left(y_j - f(X_j) \right) + \frac{1}{2} tr \left(V_j^{-1} \frac{\partial V_j}{\partial \psi} \right) \right]$$

This:

- may not be realistic for some datasets as n_i could be large for some j
- is not the case in general when $g(Z) \neq Z$
- for other LMM scenarios, needs a block-diagonal $V(g, \psi)$ (see Extensions)
- For a more general structure of $V(g, \psi)$ see e.g. Chen et. al. 2020, but this is still WIP

Extensions

 A single high-cardinality categorical feature is just the start

- Many useful correlation structures D have already shown promising results (future work):
 - Multiple categorical features (see simulations)
 - Longitudinal data
 - Kriging over random fields, similar to GP
 - GLMM (e.g. classification setting)

Table 6: Simulated model with g(Z) = Z and two categorical features, mean test MSEs and standard errors in parentheses.

σ_{b1}^2	σ_{b2}^2	q_1	q_2	Ignore	OHE	Embeddings	lme4	LMMNN
0.5	0.5	10^{3}	10^{3}	2.18 (.03)	1.45 (.02)	1.34 (.01)	2.97 (.03)	1.13 (.01) 1.23 (.01)
		$10^{-10^{-10^{-10^{-10^{-10^{-10^{-10^{-$	10^{-1} 10^{4}	2.13 (.02)	1.83 (.02)	1.80 (.02)	3.12 (.03)	1.23 (.01) 1.30 (.00)
0.5	5.0	10^{3}	10^{3}	6.73 (.04)	1.66 (.03)	1.57 (.02)	3.00 (.04)	1.12 (.00)
		10^{3}	10^{4}	6.75 (.04)	2.20 (.03)	2.01 (.03)	3.31 (.02)	1.29 (.01)
		10^{4}	10^{3}	6.50 (.05)	1.88 (.03)	1.92 (.04)	3.15 (.01)	1.23 (.01)
		10^{4}	10^{4}	6.68 (.12)	2.48 (.03)	2.16 (.02)	3.43 (.01)	1.37 (.01)
5.0	5.0	10^{3}	10^{3}	11.26 (.19)	1.83 (.02)	1.80 (.07)	2.97 (.02)	1.14 (.02)
		10^{3}	10^{4}	11.33 (.19)	2.36 (.03)	2.11 (.02)	3.32 (.02)	1.30 (.01)
		10^4	10^{4}	11.24 (.09)	3.02 (.03)	2.55 (.000)	3.69 (.02)	1.49 (.02)

Code

- Python 3.8, Tensorflow-Keras (Chollet, 2015)
- lmmnn package, key feature:
 NLL custom Keras loss layer

All code in: <u>https://github.com/gsimchoni/lmmnn</u> # after importing all that is necessary for Keras
from lmmnn.layers import NLL

```
def cnn_lmmnn():
```

```
input layer = Input((IMG HEIGHT, IMG WIDTH, 3))
    y true input = Input(shape=(1, ),)
    Z input = Input(shape=(1, ), dtype=tf.int64)
    x = Conv2D(64, (2, 2), activation='relu')(input layer)
    x = MaxPool2D((2, 2))(x)
    x = Conv2D(32, (2, 2), activation='relu')(x)
    x = MaxPool2D((2, 2))(x)
    x = Conv2D(16, (2, 2), activation='relu')(x)
    x = MaxPool2D((2, 2))(x)
    x = Flatten()(x)
    x = Dense(100, activation='relu')(x)
    v \text{ pred output} = \text{Dense}(1)(x)
    nll = NLL('intercepts', 1.0, [1.0])(
        y true input, y pred output, [Z input]
        inputs = [input layer, y true input, Z input],
        outputs=nll
model = cnn lmmnn()
model.compile(optimizer= 'adam')
history = model.fit(
    [X train['images'], y train, X train[RE feature]],
    None,
    validation split = 0.1,
    batch size = batch size, epochs=epochs
```

References

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Thank you.