Incomplete learning of multi-modal connectome for brain disorder diagnosis via modal-mixup and deep supervision

Abstract

Recently, the study of multi-modal brain networks has dramatically facilitated the efficiency in brain disorder diagnosis by characterizing multiple types of connectivity of brain networks and their intrinsic complementary information. Despite the promising performance achieved by multi-modal technologies, most existing multi-modal approaches can only learn from samples with complete modalities, which wastes a considerable amount of mono-modal data. Otherwise, most existing data imputation approaches still rely on a large number of samples with complete modalities. In this study, we propose a modal-mixup data imputation method by randomly sampling incomplete samples and synthesizing them into complete data for auxiliary training. Moreover, to mitigate the noise in the complementary information between unpaired modalities in the synthesized data, we introduce a bilateral network with deep supervision for improving and regularizing mono-modal representations with disease-specific information. Experiments on the ADNI dataset demonstrate the superiority of our proposed method for disease classification in terms of different rates of samples with complete modalities.

Keywords: Brain connectome, incomplete learning, deep supervision, brain disorder, missing modalities

1. Introduction

Neuroimaging developments have been widely adopted in medical scenarios such as disease diagnosis (Zhu et al., 2021; Liu et al., 2022; Yang et al., 2022) and lesion segmentation (Falk et al., 2019; Menze et al., 2014; Zhou et al., 2018). Advanced neuroimaging candidates, i.e., functional Magnetic Resonance Imaging (fMRI) and Diffusion Tensor Imaging (DTI), are powerful tools for brain disorder diagnosis by characterizing neural connections and information flow between brain regions (Yin et al., 2022; Li et al., 2021; Kawahara et al., 2017). Derived functional and structural connectomes are modeled as graphs by representing the activity of neurons as nodes interconnected by a set of edges, providing a more holistic view for relating abnormal discharge of neurons and brain dysfunction (Dadi et al., 2019). Analysis of brain connectome can contribute to the scientific understanding of cognitive process and potentially aid in diagnosing and treating neurological disorders (Gabrieli et al., 2015; Pu et al., 2015).

   In general, the brain networks can be categorized into two classes: functional networks derived from fMRI or EEG, and structural networks obtained from DTI or DSI (Bullmore and Bassett, 2011; Achard et al., 2006; Zalesky and Fornito, 2009). Combining functional and structural connectome enables the exploration of brain state by neuron activation and

connection in vivo by leveraging complementary information between functional and structural networks. And multi-modal brain networks provide a more constructive scene with distinctive biomarkers. Despite the promising performance achieved by multi-modal technologies, one of the core issues is that most methods can only use data with complete modalities. In practice, it is difficult to gather a large amount of complete data. Especially when diagnosing neurodegenerative diseases based on fMRI and DTI, only partial subjects have both images. In addition, missing modality is a common issue in real-world multi-modal scenarios, and the missingness can be caused by various reasons such as sensor damage, data corruption, and human mistakes in recording (Ding et al., 2018; Chen and Zhang, 2020). Unfortunately, most existing multi-modal methods may have to discard a large portion of the data collected.

One way to address the abovementioned issue is by introducing incomplete learning (Ma et al., 2021; Liu et al., 2021; Ding et al., 2021). The core of incomplete learning is to impute missing modalities (Wang et al., 2020). After data imputation, multi-modal learning methods can be used to embed multi-modal representations. Existing data imputation approaches such as generative adversarial networks (GAN) (Goodfellow et al., 2020) still depend on the amount of samples with complete modalities. Learning with insufficient samples may lead to model over-fitting and poor generalization, especially when there exist severe missing modalities. In this regard, we propose a modal-mixup data imputation approach for synthesizing samples with incompleteness into complete data for training. The idea behind this is by mix-up data augmentation approach (Zhang et al., 2017; Verma et al., 2019; Thulasidasan et al., 2019), where samples of different classes are randomly mixed into updated samples. The modal-mixup method randomly samples data with missing modalities to constitute complete data. The constructive samples are leveraged as auxiliary data for training.

Moreover, most multi-modal networks leverage inter-modal dependencies for classification, however, fail to tackle synthesized data, since there exists unwanted noise within the complementary information. Accordingly, the unpaired functional and structural connectome might decrease the performance. In terms of this, we propose a bilateral network, which discards inter-modal associations and leverages the concatenated features for classification directly. Apart from most existing multi-modal networks that apply complementary information, we implement the deep supervision modules to regularize the latent monomodal representations with disease-specific information for multi-modal learning. Experiments on the ADNI database with a cohort of 124 subjects demonstrate the superiority of our proposed approach in improving the classification performance of incomplete learning.

2. Related works

Multi-modal Brain Connectome study. Recently, deep learning methods have been state-of-the-art tools to embed high-order representations and achieve promising performances. For example, (Wang et al., 2018) performed a multi-layer convolution on fMRI and DTI data simultaneously. (Dsouza et al., 2021) regularized convolution on functional connectivity with structural graph Laplacian. A triplet attention network with a self-attention mechanism was introduced to map high-order multi-modal representations (Zhu et al., 2022). (Feng et al., 2019) proposed to perform hyperedge to perform heterogeneous

graph convolution on multi-modal data. However, these approaches have rarely explored learning with complementary information from incomplete multi-modal data.

Incomplete Learning. The exploration of incomplete learning with missing modalities has recently attracted much attention. In most cases, incomplete learning can be divided into two conditions: the test set is complete and incomplete, while the training samples are incomplete with missing modalities. In this study, we focus on studying with the complete test set. In this regard, more mono-modal neuroimages, which are easier to be collected, could be gathered for training, which might improve the model performance and generalizability. Besides, data imputation methods such as KNN are commonly used in most conditions (Campos et al., 2015). Advanced imputation methods, such as adversarial training with a similar structure as GAN, have also been proposed to deal with imputing the missing modalities (Cai et al., 2018). (Wang et al., 2020) proposed a knowledge distillation-based approach to integrating the supplementary information of multiple modalities. However, most of these studies still rely on a large amount of complete data to obtain a well-trained data imputation model.

3. Method

3.1. Problem formulation

The training sets included complete data and incomplete data, while the test sets were complete. For a multi-modal dataset with M modalities, there are $2^M - 1$ different combinations of missing modalities. In this study, functional and structural brain networks are studied with $M = 2$. Especially for the training sets, the samples with missing modalities are denoted as $X^{1u} \in R^{n_{1u} \times d_{1u}}, X^{2u} \in R^{n_{2u} \times d_{2u}}$. And the complete data are represented as $X^c \in R^{n_c \times d_c}$ with a total of $N_{train} = n_c + n_{1u} + n_{2u}$ training samples, as shown in Figure 1 A). In this study, given a collection of incomplete multi-modal data samples $\{X_i\}_{i=1}^N$ as input, where each sample consists of a set of available modalities $X_i = \{x_{i,m}\}$, our goal is to design a model to capture dependencies between modalities and fuse multi-modal data with missing modalities in an architecture.

3.2. Modal-Mixup

(Zhang et al., 2017) first proposed the Mixup method for image classification, where synthetic samples are generated by linearly interpolating a pair of training samples as well as their targets. Consider a pair of samples $(x^i; y^i)$ and $(x^j; y^j)$, a synthetic sample is geneated as $\hat{x}^{ij} = \lambda x^i + (1 - \lambda)x^j, y^{ij} = \lambda y^i + (1 - \lambda)y^j$, where $\lambda \in (0, 1)$ is the mixing ratio for the pair. In this study, we follow the mixup approach and propose a modal-mixup method by replacing the combined operations with concatenation layers. In detail, the synthetic data samples are obtained by:

$$\hat{x}^{ij} = ||\{x^i, x^j\}, x^i \in X^{1u}, x^j \in X^{2u} \tag{1}$$

where $||$ denotes the concatenatation operation, as shown in Figure 1 B). The updated label is obtained as $y^{ij} = \frac{y^i + y^j}{2}$. In this regard, the samples with modal-1 missing are mixed with those with modal-2 missing and constitute complete data $\hat{x}^{ij}$.

A) Formulation of samples

B) Modal-mixup

C) Bilateral network with deep-supervision

Figure 1: Illustration of A) the data samples, B) proposed modal-mixup, and C) bilateral network with deep supervision.

3.3. Bilateral network with deep supervision

As is shown in Figure 1 C), our network is embedded with bilateral encoders for parsing heterogeneous multi-modal brain network representations. In detail, a two-layer multi-layer perception (MLP) is used for each branch. And the MLP layers take the vectorized brain connectome features into 32 features, followed by ReLU activation and dropout. The parsed representations are further concatenated and fed into a classifier. In addition, one core challenge of data imputation by modal-mixup mentioned above is the unpaired inter-modal complementary information that would introduce unwanted noise and decrease performance. In order to reduce the effect of interactions of inter-modal representations, in this study, we propose to discard inter-modal dependency learning modules proposed by other studies. Moreover, we improve the mono-modal representations and regularize them with disease-specific information for classification. In detail, the intermediate representations of the bilateral branches are fed into a 3-layer MLP for prediction. The deep-supervision loss L_{DS} is represented by a summation of two cross-entropy losses as:

$$L_{DS} = \sum_{s=1}^{m} L_{ce}(y_s, y) \tag{2}$$

3.4. Optimization

In the training process, the synthetic data and complete samples are fed into the framework. The objective function is constructed by a combination of cross-entropy loss from the target output and the deep supervision as:

$$L_{DS} = L_{target} + L_{DS} = L_{CE}(y_m, y) + \sum_{s=1}^{m} L_{CE}(y_s, y) \tag{3}$$

where $L_{CE}(y_m, y)$ denotes the cross-entropy loss between multi-modal prediction outputs and the ground truth and $L_{CE}(y_s, y)$ represents that between the s-th deep supervision output and the ground truth.

4. Experiments

4.1. Datasets and preprocessing

In this study, the ADNI database (http://www.adni-info.org/) was used to form the cohort, where fMRI and DTI images are collected. In this study, we collected 124 subjects that were diagnosed at the baseline for evaluation, including 61 healthy controls (NC) and 63 with mild cognitive impairment (MCI). Notably, MCI is considered to be a significant stage for the preclinical diagnosis of AD. The patients were diagnosed at baseline, and the HCs were healthy at their first examination.

All the fMRI images were pre-processed by reference to the Configurable Pipeline for the Analysis of Connectomes (CPAC) pipeline (Craddock et al., 2013), including skull striping, slice timing correction, motion correction, global mean intensity normalization, nuisance signal regression with 24 motion parameters, and band-pass filtering (0.01-0.08Hz). The functional images were finally registered into standard anatomical space (MNI152). The mean time series for a set of regions were computed and normalized into zero mean and unit variance. Pearson Coefficient Correlation was applied to measure functional connectivity. The DTI images were pre-processed by image denoising, head motion, eddy-current, susceptibility distortion, and field inhomogeneity correction by MRtrix 3 (Tournier et al., 2012). The streamline count was reconstructed to 5 million. In this study, the fMRI and DTI images were segmented by the Schaefer atlas (Schaefer et al., 2018) that identified 100 cortical parcels.

4.2. Implementation details

In our implementation, the number of layers for deep supervision is 3, with an output size of 128, 30, and 2, respectively. Each layer is followed by a leaky ReLU activation function and a dropout layer. The learning rate is set as 3e-4, and the weight decay is 5e-5. All the models in this study are trained for 600 epochs and would be stopped early when the loss has not been decreased for 50 epochs. We trained the models with PyTorch on one NVIDIA 2080-Ti GPU. 10-fold cross-validation was applied for evaluation, where 10% samples were randomly selected for testing for each fold. For all experiments, we evaluated the performance in terms of diagnosis accuracy (Acc), sensitivity (Sen), and F1-score (F1).

Figure 2: Classification accuracy of five models (MLP, BrainNetCNN, MGCN, TAN, and HGCN) by training with A) complete data, B) adversarial-based imputation, C) KNN-based imputation, D) KD-based imputation. Our proposed method is shown in a solid black line in all four figures for comparison.

4.3. Competitive baseline

In this study, we compare our proposed method with different combinations of data imputation approaches with various multi-modal learning frameworks. The data imputation methods include training with only complete data (C), missing modality imputation by K-nearest neighbors (KNN) (Campos et al., 2015), adversarial-based imputation (ADV) (Cai et al., 2018), and knowledge distillation-based information integration (KD) (Wang et al., 2020). And the competitive multi-moodal networks involve BrainNetCNN (Kawahara et al., 2017), Triplet Attention Network (TAN) (Zhu et al., 2022), M-GCN (Dsouza et al., 2021), HGCN (Feng et al., 2019).

5. Results

5.1. Comparison results on incomplete learning

Figure 2 demonstrates the results of competitive networks with different data imputation approaches in various ratios of complete data. The results of TAN with KD-based imputation are missing since the network is proposed for multi-modal data, which is not available for the KD-based method. In the figure, the results of MLP, BrainNetCNN, MGCN, HGCN,

and TAN are shown in dark blue, red, green, light blue, and orange, respectively, while those of our bilateral framework are shown in black. It can be seen that as the rate of complete data decreases, the performance becomes smaller. To note that, as fewer complete samples are used for training, the performance of these methods does not always degrade. For example, training with 70% complete data by BrainNetCNN performs worse than that with 50% complete data. We hyperthesis that the number of samples enrolled in our study is relatively small, and there might be fluctuations.

Table 1: Comparison results on incomplete learning with 10% complete data in terms of accuracy (Acc %), sensitivity (Sen %), and F1-score (F1 %). The average and standard deviation (Mean±Std) across ten folds are displayed.

Type	Model	ADNI		
		ACC	Sen	F1
C	MLP	75.08±6.64	75.66±7.94	75.08±6.64
	BrainNetCNN	77.95±4.84	81.67±12.62	77.95±4.84
	M-GCN	74.77±10.73	77.644±9.89	75.91±12.57
	Triplenet	71.14±8.51	76.17±9.92	71.14±8.51
	HGCN	76.44±10.09	74.38±9.47	80.35±9.18
ADV	MLP	82.58±9.44	81.13±8.82	85.15±8.11
	BrainNetCNN	80.56±4.11	79.80±7.03	80.56±4.11
	M-GCN	76.09±10.91	76.56±13.33	76.09±10.91
	Triplenet	77.20± 11.08	77.35±14.84	81.90±7.21
	HGCN	79.92±8.61	79.18±12.67	79.92±8.61
KNN	MLP	81.67±9.02	81.93±10.25	83.99±8.48
	BrainNetCNN	79.77±7.87	88.10±10.43	79.77±7.87
	M-GCN	76.59±12.51	81.82±14.47	78.71±10.62
	Triplenet	80.76±8.33	82.23±10.63	80.76±8.34
	HGCN	80.83±8.43	84.52±11.39	80.83±8.43
KD	MLP	81.80±8.76	80.47±9.97	81.80±8.76
	BrainNetCNN	78.03±4.49	86.85±11.15	78.03±4.49
	M-GCN	73.64±11.20	81.43±15.97	73.64±11.20
	HGCN	76.71±7.88	84.83±9.50	76.71±7.88
Modal-mixup	Deep-supervision (ours)	84.21±7.52	89.79±10.84	84.21±7.52

In addition, from Figure 2 A), we can see that TAN performs the best with more than 50% complete data among five well-estimated networks while achieving the worst results when there is severe incomplete data, i.e., the ratio of complete data < 50%. In this regard, TAN depends more on the number of training samples. Figure 2 B), C), and D) display the results of the networks using adversarial-based, KNN-based, and KD-based approaches. With these methods, the five networks generally outperform those trained with only complete data. Notably, the data imputation methods might bring in noise and decrease performance. In particular, when there is only 10% data for training, MLP outperforms other architectures in most cases. This indicates that such highly complex multi-modal architectures depend on learning with complementary information between modalities and are limited to decoupling unpaired inter-modal associations. Moreover,

compared with these approaches, our proposed method is feasible to achieve a slight drop in performance, even if there are severe incomplete samples.

Table 1 further demonstrates the detailed results with 10% complete data for training, where the accuracy (Acc), sensitivity (Sen), and F1-score (F1) are displayed. In most cases, networks with the KNN-based data imputation method achieve better performances than other approaches. This is because the other data imputation methods are still limited by the number of complete data. For example, the adversarial-based method depends on the complete data to generate accurate and robust data for missing modalities. Overall, even with only 10% complete data, our proposed method could achieve a promising performance with an accuracy of 84.21%, a sensitivity of 89.79%, and an F1-score of 84.21%.

5.2. Ablation studies

Table 2 demonstrates the results of ablation studies by training complete data with different ratios of complete data. From the results, we can see that although our method with modal-mixup outperforms other models, the deep supervision could further improve the accuracy by 1.56%, 3.56%, 3.09%, 3.56% and 1.81% with 10%, 30%, 50%, 70%, and 90% complete data. The results demonstrate that the deep supervision facilitates to achieve robust representations that are less dependent of the complementary information between modalities.

Table 2: Ablation studies on the modal-mixup and deep-supervision of different ratios of complete data. The average and standard deviation (Mean±Std) of accuracy across 10 folds are listed. The components evolved are shown in the table with ✓.

Modal-mixup	deep-supervision	10%	30%	50%	70%	90%
✓		82.65±9.98	83.41±5.76	84.39±9.85	85.15±5.34	86.14±7.64
✓	✓	84.21±7.52	86.97±6.79	87.48±8.87	88.71±6.55	87.95±10.01

6. Conclusion

In this study, we propose two strategies for incomplete learning with missing modalities: Modal-mixup is introduced for data imputation, which is less dependent of the amount of data with complete modalities; A bilateral network with deep-supervision is investigated for regularizing mono-modal representations and rebalancing inter-modal dependencies. Experimental results on the ADNI dataset demonstrate that our proposed method is feasible to model incomplete data and outperforms other combinations of architectures and data imputation methods. Especially compared with 90% complete data, our proposed method could only achieve a 3.74% drop in terms of accuracy with 10% complete data. In this regard, our proposed method provides novel insights for multi-modal learning, i.e., in some medical scenarios more mono-modal samples can be gathered for multi-modal study to improve performance and generalizability.

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References

Sophie Achard, Raymond Salvador, Brandon Whitcher, John Suckling, and ED Bullmore. A resilient, low-frequency, small-world human brain functional network with highly connected association cortical hubs. *Journal of Neuroscience*, 26(1):63–72, 2006.

Edward T Bullmore and Danielle S Bassett. Brain graphs: graphical models of the human brain connectome. *Annual review of clinical psychology*, 7:113–140, 2011.

Lei Cai, Zhengyang Wang, Hongyang Gao, Dinggang Shen, and Shuiwang Ji. Deep adversarial learning for multi-modality missing data completion. In *Proceedings of the 24th ACM SIGKDD international conference on knowledge discovery & data mining*, pages 1158–1166, 2018.

Sergio Campos, Luis Pizarro, Carlos Valle, Katherine R Gray, Daniel Rueckert, and Héctor Allende. Evaluating imputation techniques for missing data in adni: a patient classification study. In *Iberoamerican Congress on Pattern Recognition*, pages 3–10. Springer, 2015.

Jiayi Chen and Aidong Zhang. Hgmf: heterogeneous graph-based fusion for multimodal data with incompleteness. In *Proceedings of the 26th ACM SIGKDD International Conference on Knowledge Discovery & Data Mining*, pages 1295–1305, 2020.

Cameron Craddock, Sharad Sikka, Brian Cheung, Ranjeet Khanuja, Satrajit S Ghosh, Chaogan Yan, Qingyang Li, Daniel Lurie, Joshua Vogelstein, Randal Burns, et al. Towards automated analysis of connectomes: The configurable pipeline for the analysis of connectomes (c-pac). *Front Neuroinform*, 42:10–3389, 2013.

Kamalaker Dadi, Mehdi Rahim, Alexandre Abraham, Darya Chyzhyk, Michael Milham, Bertrand Thirion, Gaël Varoquaux, Alzheimer's Disease Neuroimaging Initiative, et al. Benchmarking functional connectome-based predictive models for resting-state fmri. *NeuroImage*, 192:115–134, 2019.

Yuhang Ding, Xin Yu, and Yi Yang. Rfnet: Region-aware fusion network for incomplete multi-modal brain tumor segmentation. In *Proceedings of the IEEE/CVF International Conference on Computer Vision*, pages 3975–3984, 2021.

Zhengming Ding, Handong Zhao, and Yun Fu. Learning representation for multi-view data analysis: models and applications. 2018.

Niharika Shimona Dsouza, Mary Beth Nebel, Deana Crocetti, Joshua Robinson, Stewart Mostofsky, and Archana Venkataraman. M-gcn: A multimodal graph convolutional network to integrate functional and structural connectomics data to predict multidimensional phenotypic characterizations. In *Medical Imaging with Deep Learning*, pages 119–130. PMLR, 2021.

Thorsten Falk, Dominic Mai, Robert Bensch, Özgün Çiçek, Ahmed Abdulkadir, Yassine Marrakchi, Anton Böhm, Jan Deubner, Zoe Jäckel, Katharina Seiwald, et al. U-net: deep learning for cell counting, detection, and morphometry. *Nature methods*, 16(1): 67–70, 2019.

Yifan Feng, Haoxuan You, Zizhao Zhang, Rongrong Ji, and Yue Gao. Hypergraph neural networks. In *Proceedings of the AAAI conference on artificial intelligence*, volume 33, pages 3558–3565, 2019.

John DE Gabrieli, Satrajit S Ghosh, and Susan Whitfield-Gabrieli. Prediction as a humanitarian and pragmatic contribution from human cognitive neuroscience. *Neuron*, 85(1): 11–26, 2015.

Ian Goodfellow, Jean Pouget-Abadie, Mehdi Mirza, Bing Xu, David Warde-Farley, Sherjil Ozair, Aaron Courville, and Yoshua Bengio. Generative adversarial networks. *Communications of the ACM*, 63(11):139–144, 2020.

Jeremy Kawahara, Colin J Brown, Steven P Miller, Brian G Booth, Vann Chau, Ruth E Grunau, Jill G Zwicker, and Ghassan Hamarneh. Brainnetcnn: Convolutional neural networks for brain networks; towards predicting neurodevelopment. *NeuroImage*, 146: 1038–1049, 2017.

Xiaoxiao Li, Yuan Zhou, Nicha Dvornek, Muhan Zhang, Siyuan Gao, Juntang Zhuang, Dustin Scheinost, Lawrence H Staib, Pamela Ventola, and James S Duncan. Braingnn: Interpretable brain graph neural network for fmri analysis. *Medical Image Analysis*, 74: 102233, 2021.

Yanbei Liu, Lianxi Fan, Changqing Zhang, Tao Zhou, Zhitao Xiao, Lei Geng, and Dinggang Shen. Incomplete multi-modal representation learning for alzheimer's disease diagnosis. *Medical Image Analysis*, 69:101953, 2021.

Yanbei Liu, Henan Li, Tao Luo, Changqing Zhang, Zhitao Xiao, Ying Wei, Yaozong Gao, Feng Shi, Fei Shan, and Dinggang Shen. Structural attention graph neural network for diagnosis and prediction of covid-19 severity. *IEEE Transactions on Medical Imaging*, 2022.

Mengmeng Ma, Jian Ren, Long Zhao, Sergey Tulyakov, Cathy Wu, and Xi Peng. Smil: Multimodal learning with severely missing modality. In *Proceedings of the AAAI Conference on Artificial Intelligence*, volume 35, pages 2302–2310, 2021.

Bjoern H Menze, Andras Jakab, Stefan Bauer, Jayashree Kalpathy-Cramer, Keyvan Farahani, Justin Kirby, Yuliya Burren, Nicole Porz, Johannes Slotboom, Roland Wiest, et al. The multimodal brain tumor image segmentation benchmark (brats). *IEEE transactions on medical imaging*, 34(10):1993–2024, 2014.

Jian Pu, Jun Wang, Wenwen Yu, Zhuangming Shen, Qian Lv, Kristina Zeljic, Chencheng Zhang, Bomin Sun, Guoxiang Liu, and Zheng Wang. Discriminative structured feature engineering for macroscale brain connectomes. *IEEE Transactions on Medical Imaging*, 34(11):2333–2342, 2015.

Alexander Schaefer, Ru Kong, Evan M Gordon, Timothy O Laumann, Xi-Nian Zuo, Avram J Holmes, Simon B Eickhoff, and BT Thomas Yeo. Local-global parcellation of the human cerebral cortex from intrinsic functional connectivity mri. *Cerebral cortex*, 28(9):3095–3114, 2018.

Sunil Thulasidasan, Gopinath Chennupati, Jeff A Bilmes, Tanmoy Bhattacharya, and Sarah Michalak. On mixup training: Improved calibration and predictive uncertainty for deep neural networks. *Advances in Neural Information Processing Systems*, 32, 2019.

J-Donald Tournier, Fernando Calamante, and Alan Connelly. Mrtrix: diffusion tractography in crossing fiber regions. *International journal of imaging systems and technology*, 22(1): 53–66, 2012.

Vikas Verma, Alex Lamb, Christopher Beckham, Amir Najafi, Ioannis Mitliagkas, David Lopez-Paz, and Yoshua Bengio. Manifold mixup: Better representations by interpolating hidden states. In *International Conference on Machine Learning*, pages 6438–6447. PMLR, 2019.

Qi Wang, Liang Zhan, Paul Thompson, and Jiayu Zhou. Multimodal learning with incomplete modalities by knowledge distillation. In *Proceedings of the 26th ACM SIGKDD International Conference on Knowledge Discovery & Data Mining*, pages 1828–1838, 2020.

Yan Wang, Yanwu Yang, Xin Guo, Chenfei Ye, Na Gao, Yuan Fang, and Heather T Ma. A novel multimodal mri analysis for alzheimer's disease based on convolutional neural network. In *2018 40th Annual International Conference of the IEEE Engineering in Medicine and Biology Society (EMBC)*, pages 754–757. IEEE, 2018.

Yanwu Yang, Xutao Guo, Zhikai Chang, Chenfei Ye, Yang Xiang, and Ting Ma. Multimodal dynamic graph network: Coupling structural and functional connectome for disease diagnosis and classification. *arXiv preprint arXiv:2210.13721*, 2022.

Wutao Yin, Longhai Li, and Fang-Xiang Wu. Deep learning for brain disorder diagnosis based on fmri images. *Neurocomputing*, 469:332–345, 2022.

Andrew Zalesky and Alex Fornito. A dti-derived measure of cortico-cortical connectivity. *IEEE transactions on medical imaging*, 28(7):1023–1036, 2009.

Hongyi Zhang, Moustapha Cisse, Yann N Dauphin, and David Lopez-Paz. mixup: Beyond empirical risk minimization. *arXiv preprint arXiv:1710.09412*, 2017.

Zongwei Zhou, Md Mahfuzur Rahman Siddiquee, Nima Tajbakhsh, and Jianming Liang. Unet++: A nested u-net architecture for medical image segmentation. In *Deep learning in medical image analysis and multimodal learning for clinical decision support*, pages 3–11. Springer, 2018.

Qi Zhu, Heyang Wang, Bingliang Xu, Zhiqiang Zhang, Wei Shao, and Daoqiang Zhang. Multi-modal triplet attention network for brain disease diagnosis. *IEEE Transactions on Medical Imaging*, 2022.

Wenyong Zhu, Liang Sun, Jiashuang Huang, Liangxiu Han, and Daoqiang Zhang. Dual attention multi-instance deep learning for alzheimer's disease diagnosis with structural mri. *IEEE Transactions on Medical Imaging*, 40(9):2354–2366, 2021.