



# Identification of Electroencephalogram Signals in Alzheimer's Disease by Multifractal and Multiscale Entropy Analysis

Momo Ando<sup>1</sup>, Sou Nobukawa<sup>1,2\*</sup>, Mitsuru Kikuchi<sup>3,4</sup> and Tetsuya Takahashi<sup>4,5,6</sup>

<sup>1</sup> Graduate School of Information and Computer Science, Chiba Institute of Technology, Narashino, Japan, <sup>2</sup> Department of Computer Science, Chiba Institute of Technology, Narashino, Japan, <sup>3</sup> Department of Psychiatry and Behavioral Science, Kanazawa University, Ishikawa, Japan, <sup>4</sup> Research Center for Child Mental Development, Kanazawa University, Ishikawa, Japan, <sup>5</sup> Department of Neuropsychiatry, University of Fukui, Fukui, Japan, <sup>6</sup> Uozu Shinkei Sanatorium, Uozu, Japan

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> \***Correspondence:** Sou Nobukawa nobukawa@cs.it-chiba.ac.jp

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Ando M, Nobukawa S, Kikuchi M and Takahashi T (2021) Identification of Electroencephalogram Signals in Alzheimer's Disease by Multifractal and Multiscale Entropy Analysis. Front. Neurosci. 15:667614. doi: 10.3389/fnins.2021.667614 Alzheimer's disease (AD) is the most common form of dementia and is a progressive neurodegenerative disease that primarily develops in old age. In recent years, it has been reported that early diagnosis of AD and early intervention significantly delays disease progression. Hence, early diagnosis and intervention are emphasized. As a diagnostic index for AD patients, evaluating the complexity of the dependence of the electroencephalography (EEG) signal on the temporal scale of Alzheimer's disease (AD) patients is effective. Multiscale entropy analysis and multifractal analysis have been performed individually, and their usefulness as diagnostic indicators has been confirmed, but the complemental relationship between these analyses, which may enhance diagnostic accuracy, has not been investigated. We hypothesize that combining multiscale entropy and fractal analyses may add another dimension to understanding the alteration of EEG dynamics in AD. In this study, we performed both multiscale entropy and multifractal analyses on EEGs from AD patients and healthy subjects. We found that the classification accuracy was improved using both techniques. These findings suggest that the use of multiscale entropy analysis and multifractal analysis may lead to the development of AD diagnostic tools.

Keywords: EEG signal, Alzheimer's disease, multifractal, multiscale entropy, early diagnosis

# **1. INTRODUCTION**

Alzheimer's disease (AD) is the most common form of dementia and is a progressive neurodegenerative disease that primarily develops in old age (Liu et al., 2014). The World Health Organization estimates that the global prevalence of AD will increase to 0.6% in 2030 and 1.2% by 2046 (Brookmeyer et al., 2007). Although there is no effective treatment for AD, in recent years, it has been reported that early diagnosis of AD and early intervention significantly delay the progression of the disease. Hence, it would be ideal to diagnose AD early in its clinical course (Liu et al., 2014).

In AD, there are three significant anatomical changes: progressive neuronal death, neurofibrillary tangles, and senile plaques in extensive brain areas (Sims et al., 2017; Yamaguchi-Kabata et al., 2018). Positron emission tomography (PET) and magnetic resonance imaging (MRI)

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are often used to diagnose AD and detect neurotransmitter activity disorders, amyloid beta plaque deposition, and brain atrophy (Ewers et al., 2011; McKhann et al., 2011; Sperling et al., 2011). As methods focused on functional neural activity, studies based on the temporal behavior of neural activity were conducted using electroencephalography (EEG), magnetoencephalography (MEG), and functional magnetic resonance imaging (fMRI) (Greicius et al., 2004; Jeong, 2004; Stam, 2005; Dickerson and Sperling, 2008; Takahashi, 2013; Yang and Tsai, 2013; Wang et al., 2017; Nobukawa et al., 2020).

Among all these evaluations, EEG is cost-effective, widely available, and non-invasive, making it ideal for clinical applications (Vecchio et al., 2013; Kulkarni and Bairagi, 2018). AD's pathological progression alters EEG behavior, such as slow waves, low synchronization of neural activity among brain regions, and low temporal complexity. Complexity analysis is a good approach to detect cortical disconnection in AD because this state impairs mutual neural interaction among widespread brain regions. Studies assessing EEG signals' complexity in patients with AD focused on deterministic chaos and fractal dimensions, such as the correlation dimension and Lyapunov exponent (Kantz and Schreiber, 2004). These studies reported a reduction in the complexity of neural activity in AD patients (Woyshville and Calabrese, 1994; Besthorn et al., 1995; Jelles et al., 1999; Jeong, 2004; Smits et al., 2016; Al-Nuaimi et al., 2017). Moreover, EEG dynamics at each temporal scale and frequency band, such as theta, beta, and gamma bands, are associated with different memory function components, cognitive and perceptual function (Klimesch et al., 2007). Hence, as a diagnostic index for AD patients with various brain function defects, the evaluation of the complexity with temporal scale dependence in EEG signal is effective (Mizuno et al., 2010; Nobukawa et al., 2019, 2020).

Multiscale entropy (MSE) analysis and multifractal (MF) analysis are known as typical temporal scale complexity dependency analyses (Takahashi, 2013; Yang and Tsai, 2013). In addition to EEG's temporal dependency in AD, MSE analysis also showed lower complexity on a small temporal scale in the frontal region in AD; in comparison, higher complexity was observed across this brain region in AD on a larger temporal scale (Mizuno et al., 2010; Ni et al., 2016). Zorick *et al.* reported that a statistical model based on MF analysis could detect clinical stages of severity and degree of progress from cognitive impairment to AD (Zorick et al., 2020). As described above, MSE and MF analyses have a high ability to detect the complexity in EEG signals of AD. As such, these indices might become biomarkers for AD to evaluate the alteration of EEG complexity (Mizuno et al., 2010; Ni et al., 2016; Nobukawa et al., 2020).

Recent studies have focused on the enhancement of classification accuracy combining several feature values, including complexity indexes in EEG of AD patients (Wang et al., 2015; Gómez et al., 2017; Ieracitano et al., 2020; Nobukawa et al., 2020). Particularly, Wang et al. (2015) and Gómez et al. (2017) showed that combinations of spectrum and bispectrum entropy measures enhance the accuracy of EEG signals classification in AD. Therefore, these combinations of complexity measures are a new avenue for the diagnosis of AD EEG signals. Furthermore, Cukic *et al.* showed that multiscale

 TABLE 1 | Physical characteristics of healthy control (HC) and subjects with

 Alzheimer's disease (AD).

	HC participants	AD participants	p-values	
Male/female	7/11	5/11	0.72	
Age(year)	59.3 (5.3, 55–66)	57.5 (4.7, 43–64)	0.31	
MMSE score	NA	15.5 (4.7, 10–26)	NA	

analysis (MSE) and fractal dimension provide complementary information on brain activity in healthy subjects (Cukic et al., 2018). This complementary relationship may enhance the accuracy of AD identification. In this context, we hypothesize that the combination of MSE and fractal analysis may contribute to a better understanding of EEG dynamics' alteration in AD. In this study, we performed MSE analysis and multifractal analysis on the EEGs of patients with AD and healthy controls (HC).

# 2. MATERIALS AND METHODS

## 2.1. Subject

The subjects of this study were 16 patients with AD and 18 sexmatched and aged-matched healthy old individuals (see **Table 1**) (Mizuno et al., 2010; Nobukawa et al., 2019, 2020). The sample size of AD and HC groups was determined based on previous works on complexity analysis (Abásolo et al., 2008; Mizuno et al., 2010; Nobukawa et al., 2019, 2020). For this study, we defined healthy old individuals as nonsmokers and not on medication. Subjects with medical or neurological conditions, including epilepsy or head trauma in the past, and subjects with a history of alcohol or drug dependence were excluded. We recruited patients with AD or probable AD who met the NINCDS-ADRDA criteria and in a state before the onset of primary dementia based on DSM-IV criteria. Recruited patients with AD were not receiving medications that act on the central nervous system.

Each patient was evaluated using the Function Assessment Stage (FAST) and Mini-Mental State Examination (MMSE). Three patients had mild dementia (FAST 3); seven moderate dementia (FAST 4); and six severe dementia (FAST 5). The MMSE score ranged from 10 to 26, with an average of 15.56. **Table 1** shows subjects' characteristics. All subjects provided informed consent prior to the start of the study. The research protocol was approved by the Ethics Committee of Kanazawa University. All procedures in this study were conducted in accordance with the Declaration of Helsinki.

## 2.2. EEG Recordings

As reported in previous studies, methods have been established to record and preprocess EEG data (Mizuno et al., 2010). When recording the EEG, the participants were seated in an electrically shielded and soundproof recording room, and the room lighting was controlled. For the EEG measurement, 16 electrodes (Fp1, Fp2, F3, F4, C3, C4, P3, P4, O1, O2, F7, F8, Fz, Pz, T5, and T6) were used in the electrode arrangement called the International 10–20 System. EEG activity was measured using the binaural connection as a reference.

EEG-4518 manufactured by Nihon Kohden Co., Ltd. Tokyo, Japan, was used for measurement. Eye movements were tracked using bipolar electrocardiography (EOG). The EEG signal was recorded using a sampling frequency of 200 Hz and bandpass filtered at 2.0-60 Hz. As pre-processing steps were not conducted (i.e., filtering except for bandpass, artifacts removal, or data reconstruction), because such processing may destroy the data's intrinsic dynamics, we visually selected epochs without artifacts. The electrode/skin conductance impedance was carefully controlled at each electrode to  $< 5k\Omega$ . Each subject's EEG signal was measured for 10-15 min in a resting state with eyes closed. A video surveillance system was used to visually inspect the subjects' alertness and to confirm that only epochs with closed eyes and a wakefulness state (not light sleep) were measured. Visual inspection of EEG and EOG records identified EEG time series segments recorded in a wakefulness state with closed eyes. Subjects were considered fully awake when predominant alpha activity appeared in the posterior region in response to the fast eye movements of the EOG channel (Wada et al., 1996). MSE analysis and MF analysis were conducted against a continuous 50-s(10000 data points) epoch.

#### 2.3. Multifractal Analysis

In MF analysis, wavelet readers derived from the coefficients of the discrete wavelet transform are widely used (Jaffard et al., 2006; Wendt and Abry, 2007). The discrete wavelet coefficient of the discrete signal X(t) is given by

$$d_X(j,k) = \int_R X(t)2^j, \psi_0(2^{-j}t - k)dt \quad (j = 1, 2, ..., k = 1, 2, ...),$$
(1)

where  $\psi_0$  is a compact-supported mother wavelet function. Onedimensional wavelet leaders were expressed by

$$L_{x}(j,k) = \sup_{\lambda' \subset \Im\lambda_{j,k}} |d_{X}(j,k)|.$$
(2)

Here,  $\lambda = \lambda_{j,k} = [k2^j, (k + 1)2^j]$  represents the time interval of scale  $2^j$ , and  $3\lambda_{j,k-1} = \bigcup \lambda_{j,k} \bigcup \lambda_{j,k+1}$  represents the adjacent time (Wendt and Abry, 2007). The singular value spectrum D(h), which is the distribution of the fractal dimension represented by the Hölder exponent *h*, is represented by wavelet leaders (Jaffard et al., 2006; Wendt and Abry, 2007):

$$D(h) = \inf_{q \neq 0} (1 + qh - \zeta_L(q)).$$
(3)

Here, *q* indicates the moment for scaling index  $\zeta_L(q)$ . The scaling index  $\zeta_L(q)$  and the structural function  $S_L(q, j)$  are represented by Equations (4, 5), respectively:

$$\zeta_L(q) = \liminf_{j \to 0} \left( \frac{\log_2 S_L(q, j)}{j}, \right)$$
(4)

$$S_L(q,j) = \frac{1}{n_j} \sum_{k=1}^{n_j} |L_X(j,k)|^q.$$
 (5)



**FIGURE 1** Singular value spectra D(h) in multi-tractal analysis for one healthy control (HC) subject. Here, *h* exhibits Hölder exponent.  $c_1$  shows *h*-value where D(h) = 1.0 (q = 0); absolute value of  $c_2$  corresponds to the range of D(h) distribution between q = -5 and 5.

Here,  $n_i$  indicates the number of samples of X when the scale is  $2^{j}$ . As Hölder exponent h approaches 1.0, the shape of the time series becomes more differentiable. In contrast, as Hölder exponent h approaches zero, the shape of the time series becomes nearly discontinuous. If the scaling index  $\zeta_L(q)$  is a linear function and D(h) converges to a particular h, then the signal is monofractal. On the other hand, in the scaling index, where  $\zeta_L(q)$  deviates from linearity and D(h) is distributed over a wide range of h, the signal is multifractal. In this study, to capture the profile of D(h), we used the primary cumulant  $c_1$  of D(h), which corresponds to a dominant component of D(h) as smoothness index estimated in the entire time-series. Moreover, we used the secondary cumulant  $c_2$ , which corresponds to the magnitude of fluctuation intermittently appearing as the index for multifractality. Figure 1 shows the results of multifractal analysis of one HC subject.  $c_1$  shows *h*-value where D(h) = 1.0 (q = 0), which corresponds to the degree of differentiability in the dominant component of the entire timeseries, i.e., smoothness. The absolute value of  $c_2$  corresponds to the range of D(h) distribution between q = -5 and 5. In the monofractal time-series, D(h) converges at a particular h (in the time-series with no multifractality, D(h) converges at h-value with q = 2), while in the multifractal time-series, the range of D(h) becomes wider (Ihlen, 2012; Mukli et al., 2015). Therefore, the degree of variation of D(h) corresponding to  $c_2$  reflects the multifractality. In this study, multifractal analysis was performed using the wavelet toolbox of MATLAB (https://jp.mathworks. com/products/wavelet.html).

#### 2.4. Multiscale Entropy Analysis

To perform the multiscale entropy (MSE) analysis, we used the dependence of the EEG time series complexity on the temporal scale (Costa et al., 2002). The sample entropy for the time-series

of random Z-scored variable  $\{x_1, x_2, ..., x_N\}$  is defined as

$$h(r,m) = -\log \frac{C_{m+1}(r)}{C_m(r)}.$$
 (6)

 $C_m(r)$  is the probability of  $|\mathbf{x}_i^m - \mathbf{x}_j^m| < r(i \neq j, i, j = 1, 2, ...)$ .  $\mathbf{x}_i^m$  indicates an *m*-dimensional vector  $\mathbf{x}_i^m = \{x_i, x_{i+1}, ..., x_{i+m-1}\}$ .  $\{x_i, x_{i+1}, ..., x_N\}$  is obtained course-grained process:

$$x_j = \frac{1}{\tau} \sum_{i=(j-1)\tau+1}^{j_\tau} y_i (1 \le j \le \frac{N}{\tau}).$$
(7)

where,  $\{y_1, y_2, ..., y_N\}$  is observed signals.  $\tau(\tau = 1, 2, ...)$  is the temporal scale. In this study, we set m = 2 and r = 0.2 (Costa et al., 2002). In this study, MSE analysis was performed using the Physio Toolkit, a toolbox of MATLAB (http://physionet.incor. usp.br/physiotools/sampen/).

#### 2.5. Statistical Analysis

For  $c_1$  and  $c_2$ , repeated measures analysis of variance (ANOVA) with the groups (HC vs. AD) as the between-subject factor and the electrodes (16 electrodes from Fp1 to T6) as the within-subject factors was performed to test for group differences. The result of ANOVA is represented by *F*-value based on a comparison of variances within/between groups. The Greenhouse-Geisser adjustment was applied in degrees of freedom. The  $\alpha$  bilateral level of 0.05 was used, considered a statistically significant criterion to avoid type I errors. *Post-hoc t*-tests were used to assess the significant main effects of group and per-electrode interactions. Benjamini–Hochberg false discovery rate (FDR) correction was applied to the *t*-score for multiple comparisons in  $c_1$  and  $c_2$  (q < 0.05) (16 *p*-values: 16 electrodes).

For sample entropy, repeated measures ANOVA with groups (HC vs. AD) as the between-subject factor and electrode (16 electrodes from Fp1 to T6) and temporal scale (30 temporal scales) as within-subject factors, was performed to test for group differences. The Greenhouse-Geisser adjustment and  $\alpha$  bilateral level of 0.05 were applied. The result of ANOVA is represented by *F*-value based on a comparison of variances within/between groups. *Post-hoc t*-tests were used to assess the significant main effects of the group and per-electrode and per-temporal-scale interactions. The FDR correction was applied to the *t*-score for multiple comparisons (q < 0.05) (480 *p*-values: 16 electrode × 30 scales).

Receiver operating characteristic (ROC) curves were used to evaluate the ability to identify AD. To identify AD, a logistic regression model based on the sample entropy,  $c_1$  and  $c_2$ , was used. Here, the logistic regression model outputs the identification probability of AD for each subject. Subsequently, the true positive rate/false positive rate at each threshold of identification probability from 0 to 1.0 in both groups are measured. Principal component analysis is used as a preprocess for dimensionality reduction. Logistic regression was applied to the 1st–3rd principal components of each evaluation index. The identification accuracy was evaluated by measuring the area under the ROC curve (AUC), which is an index of identification accuracy. Subsequently, according to AUC values,



**TABLE 2** | AD vs. HC repeated measure ANOVA analysis results [*F*-value (p value)] in multifractal (MF) analysis results, *F* and *p* value with p < 0.05 are

	Group	Group × nodes
c1	$F = 9.088 \ (p = 0.005)$	F = 1.460 (p = 0.204)
c2	$F = 0.654 \ (p = 0.425)$	$F = 1.981 \ (p = 0.072)$

the classification accuracy is graded in logistic regression models based on the sample entropy,  $c_1$  and  $c_2$ . Here, AUC = 1.0 corresponds to complete identification, and AUC = 0.5 corresponds to random identification.

## 3. RESULTS

represented by bold characters.

#### 3.1. Multifractal Analysis

We performed MF analysis on both HC and AD groups. **Figure 2** shows the mean and standard deviation for each group of D(h) and h. Since the distribution is wide, it reflects the multi-fractal property (Sikdar et al., 2018) of both groups' EEG signal. **Table 2** shows the repeated measures ANOVA results of 1st  $(c_1)$  and 2nd  $(c_2)$  cumulants of singular spectrum. The significant main effect in  $c_1$  was confirmed. The mean values of  $c_1$  and  $c_2$  in the AD and HC groups and the result of the *post-hoc t*-test between AD and HC are shown in **Figure 3**. The significantly higher  $c_1$  values in the AD group (q < 0.050 corresponding to p < 0.012) was confirmed at F3, Fz, F4, C3, C4, P3, Pz, and P4.

#### 3.2. Multi Scale Entropy Analysis

We performed an MSE analysis in the HC and AD groups. **Table 3** shows the repeated measures ANOVA results of MSE analysis. Significant group  $\times$  scale interactions without the main effect of sample entropy were confirmed. As *post hoc t*-test, the mean values of sample entropy in HC and AD groups and the *t*-value between HC and AD are shown in **Figure 4**. The results demonstrated a significantly lower sample entropy



of AD (q < 0.050 corresponding to p < 0.002) in the temporal scale region 1 to 5 (0.005 to 0.025 s). The result of MSE analysis was reported in our previous studies (Mizuno et al., 2010; Nobukawa et al., 2020). Particularly, in the study by

Mizuno et al. (2010), multiscale entropy analysis against AD EEG signals was reported, while our recent study (Nobukawa et al., 2020) showed the relationship between functional connectivity characterized by phase synchronization and multiscale entropy in AD EEG signals.

**TABLE 3** | AD vs. HC repeated measure ANOVA results [*F*-value (p-value)] in multi scale entropy (MSE) analysis results, *F* and *p* value with p < 0.05 are represented by bold characters.

Group	$\textbf{Group} \times \textbf{node}$	$\textbf{Group} \times \textbf{scale}$	Group × node × scale
F = 1.233	F = 1.860	F = 11.457	F = 0.979
(p = 0.275)	(p = 0.129)	( <b>p = 0.003</b> )	(p = 0.451)

## 3.3. ROC Curve

To evaluate the classification ability in  $c_1$  and  $c_2$ , we evaluated ROC curves. **Figure 5** shows the result of ROC in the case with 1st–3rd principal components in each evaluated index. In the sample entropy case, the values are averaged in 1 to 5 temporal scale. AUC in the  $c_1$  case exhibits the highest classification ability (AUC = 0.85 in the case  $c_1$ ; AUC = 0.78, in the case of  $c_2$ ; AUC = 0.82 in the case sample entropy). Furthermore,



**FIGURE 4** [Multi-scale entropy analysis in HC and AD group. The horizontal axis represents the temporal-scale factor,  $\tau$ . (A) Mean values of sample entropy from 1 (0.005 s) to 30 (0.15 s) scale factors in HC (left part) and AD (right part). (B) *t*-value between the AD and HC groups(left part). The warm (cold) color represents a higher (smaller) sample entropy value for AD than that for HC. The *t*-value satisfying the FDR correction criteria q < 0.050 corresponding to (p < 0.002). Significantly smaller sample entropy of AD low temporal scale regions 1 to 5 (0.005 to 0.025 s).



we evaluated ROC using all these values; the results showed an enhancement of classification ability (AUC = 1.00). To investigate why the combination of  $c_1$ ,  $c_2$ , and sample entropy enhances classification ability, we evaluated their relationship by correlation analysis. Figure 6 shows a scatterplot among the 1st component of  $c_1$ ,  $c_2$  and sample entropy used for ROC evaluation in Figure 5. The correlation coefficients *R* are shown in Table 4. The results show a high negative and positive correlation between  $c_1$  and sample entropy, a positive correlation between  $c_2$  and sample entropy, and a relatively low negative correlation between  $c_1$  and  $c_2$ . This relatively low correlation between  $c_1$  and  $c_2$ suggests that c<sub>2</sub> includes complementary information regarding multifractality in the classification. Moreover, to investigate the correlation between  $c_1$  and  $c_2$ , not the principal components, the correlation coefficient R between  $c_1$  and  $c_2$  in HC and AD groups is demonstrated in Figure 7. The results show the spatial dependency of correlation coefficient *R*, which might contribute to the enhancement of classification accuracy shown in Figure 7.

To demonstrate that the decision region for AD is determined by  $c_1$ ,  $c_2$ , and sample entropy, decision regions for AD with decision probability > 0.9 by logistic regression model were depicted on the plane between the 1st principal component of  $c_1$ and the 1st principal component of  $c_2$  and the plane between the 1st principal component of  $c_1$  and the 1st principal component of the sample entropy (see **Figure 8**). Here, all other components except the plane axis were set to average among subjects in both the HC and AD groups. As a result, we confirmed that the decision region exhibits dependent on all of them.

## 4. DISCUSSIONS

This study evaluated AD identification accuracy by focusing on the complementary relationship between two complexity analyses. The MF and MSE of EEG signals in HC and AD were evaluated, and classification accuracies quantified by the AUC of logistic regression models were compared. The results of  $c_1$  as the index for the smoothness of the EEG time series by MF analysis showed that  $c_1$  of AD significantly increased. However, the results of  $c_2$ , as the index for the EEG time series' multifractal nature by MF analysis, show that no significant AD alteration was observed. MSE analysis showed a significant region-specific reduction of small-temporal-scale sample entropy of AD (corresponding to the complexity of faster temporal EEG behaviors). In the comparison of classification accuracy between  $c_1$ ,  $c_2$ , and small-temporal-scale sample entropy,  $c_1$  exhibits the highest classification accuracy. Moreover, the classification accuracy with  $c_1$  was enhanced by considering the complementary relationship of  $c_2$  and sample entropy.

We must discuss the reason why  $c_1$  as the degree of smoothness increase in AD. In the alteration of EEG/MEG signals in AD, a reduction in temporal complexity has been widely observed (Woyshville and Calabrese, 1994; Besthorn et al., 1995; Jelles et al., 1999; Jeong, 2004; Wickramasinghe and Geisler, 2008; Smits et al., 2016; Kulkarni, 2018; Smailovic et al., 2019). Correspondingly, our results of sample entropy at a small temporal scale also exhibited a reduction in complexity. Considering the negative correlation between  $c_1$  and the smalltemporal-scale sample entropy (see Table 4), the increase in  $c_1$ , that is, the enhancement of EEG signal smoothness in AD, was caused by the decrease in small-temporal-scale complexity. Therefore, the enhancement of  $c_1$  reflects the loss of temporal complexity of neural activity in AD. This finding agrees with previous studies on MF analysis in AD (Jaffard et al., 2006; Wendt and Abry, 2007).

Furthermore, we must consider why small temporal scale complexity decreases in AD. Dysfunction of the gammaaminobutyric acid (GABA) signaling system caused by deposition of amyloid  $\beta$  and tau protein have been reported. These changes lead to the reduced oscillation of the gamma band activity produced by GABA signaling (Nava-Mesa et al., 2014; Govindpani et al., 2017; Calvo-Flores Guzmán et al., 2018). Consequently, dysfunction of the mutual interaction of gamma band activity can reduce the complexity more on the faster than on the slower temporal scales (Ahmadlou et al., 2011; Nobukawa et al., 2019).

Next, it is necessary to consider why the classification accuracy was highest when  $c_1$ ,  $c_2$ , and sample entropy were used. According to Cukic *et al.*, sample entropy and fractal dimension by mono-fractal analysis show a complementary relationship among temporal scales (Cukic et al., 2018), and this relationship can enhance the ability to detect an alteration of complexity. Our results (see **Figure 8**) also showed that a decision region for AD with decision probability > 0.9 by logistic regression model exhibits a dependency on  $c_1$ ,  $c_2$ , and sample entropy. Therefore, the combination of  $c_1$  corresponding to the fractal dimension and sample entropy might enhance the accuracy of AD detection. In addition to these findings, a recent MF analysis in AD showed that the multifractal degree reflects disease-specific alterations of complexity (Zorick et al., 2020). Although the classification ability in case of separate use of multifractals measured by  $c_2$ 



**TABLE 4** | Correlation coefficient values (R) for each combination of  $c_1$ ,  $c_2$ , and sample entropy in HC and AD.

C <sub>1</sub> vs. C <sub>2</sub>	$c_1$ vs. Sample entropySample entropy vs. $c_2$				
Correlation coefficient(HC	R = -0.56	R = -0.77	R = 0.82		
Correlation coefficient(AD)	R = -0.42	R = -0.85	<i>R</i> = 0.78		

is relatively low, the combination with  $c_2$  may contribute to the improvement of classification accuracy.

To investigate whether the high heterogeneity of severity in patients with AD affects classification, we investigated distributions of  $c_1$ ,  $c_2$ , and mean sample entropy in a scale from 1 to 5 according to severity as classified by FAST (3 (mild dementia), 4 (moderate dementia), and 5 (severe dementia), through repeated measures ANOVA with severity as a betweensubject factor and electrode as a within-subjects factor. The results showed that severity did not have any significant main effect or interaction in  $c_1$  (F = 0.412, p = 0.671),  $c_2$  (F = 0.706, p = 0.512), and sample entropy (F = 0.532, p = 0.6); while a significant interaction between severity and electrodes in  $c_1$ (F = 2.103, p = 0.036) appeared. Therefore, although in larger AD groups a severity-dependent effect may appear, the bias of high heterogeneity of severity is limited. Additionally, in patients with mild dementia, no differences in the distribution of  $c_1$ ,  $c_2$ , and sample entropy compared to more severe patients appeared in the repeated measures ANOVA. Therefore, the classification accuracy may not change in case of a classification between HC and patients with mild dementia, which corresponds to the condition assumed for early diagnosis.

Finally, we must consider the limitations of this study. First, EEG signals do not always reflect the neural activity directly under the electrode. In this study, 16 electrodes were used to measure EEG, but the spatial resolution was too low to identify AD's complex functional connection structure. However, it is possible to use MEGs with a high spatial resolution and cortical positioning to solve this problem. Second, pre-processing for EEG signals was not adopted except for a band-pass filter. However, a recent study by Racz et al. (2018) indicated that appropriate pre-processing is needed for complexity analysis. Artifacts and noise are to be avoided, especially at the stage of clinical application. Therefore, this pre-processing for complexity analysis must be developed and adopted in future studies. Third, we consider that for our EEG data set, the multifractal analysis method proposed by Jaffard et al. (2006) and Wendt and Abry (2007) is sufficient, because a corrupted/inversed D(h) distribution did not arise (see Figure 2). Additionally,





this study was conducted on the assumption of multifractality in EEG signals (Takahashi, 2013; Yang and Tsai, 2013; Sikdar et al., 2018). However, several studies highlighted the issues of incorrect estimation of multifractal indexes in time-series without multifractality (Grech and Pamula, 2012; Mukli et al., 2015). Therefore, multifractal analysis methods with higher robustness (Mukli et al., 2015) are desired at the stage of clinical application, since proper validation of EEG multifractality (Mukli et al., 2015; Racz et al., 2018) is an important issue. Fourth, the AD group had high heterogeneity of severity, and the sample size of the AD group was small, which could have influenced the classification accuracy. Therefore, the classification ability of our proposed method must be evaluated in larger AD groups. Additionally, a Bayesian statistic approach is more suitable for small size and high sample heterogeneity than that based on frequentist inference.

# 5. CONCLUSION

In this study, both MSE and MF analysis showed a reduction in EEG complexity in AD patients. Classification accuracy is better by combining MSE analysis and MF analysis than when applying each one individually. Despite its limitations, this study shows that MSE and MF analysis play complementary roles in detecting the alteration of neural activity in AD. The use of both MSE and MF analysis may facilitate the development of AD diagnostic tools.

# DATA AVAILABILITY STATEMENT

The datasets presented in this article are not readily available because the informed consent did not include the declaration regarding publicity of clinical data. Requests to access the datasets should be directed to Sou Nobukawa, nobukawa@cs.it-chiba.ac.jp.

# ETHICS STATEMENT

The studies involving human participants were reviewed and approved by Ethics Committee of Kanazawa University. The patients/participants provided their written informed consent to participate in this study.

## **AUTHOR CONTRIBUTIONS**

MA, SN, MK, and TT conceived the methods. MA and SN analyzed the results, wrote the main manuscript text, and prepared all the figures. MK conducted the experiments. All authors reviewed the manuscript.

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**Conflict of Interest:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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