

1 **A Multisite and Multimodal Comparative Study of Cerebellar Connectome**
2 **Between Multiple Sclerosis and Neuromyelitis Optica Spectrum Disorders**

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4 Yuping Yang¹, Junle Li¹, Zhen Li¹, Yaou Liu^{2*}, Jinhui Wang^{1,3*}

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6 ¹Institute for Brain Research and Rehabilitation, Guangdong Key Laboratory of
7 Mental Health and Cognitive Science, Center for Studies of Psychological
8 Application, South China Normal University, Guangzhou, China

9 ²Department of Radiology, Beijing Tiantan Hospital, Capital Medical University,
10 Beijing, China

11 ³Key Laboratory of Brain, Cognition and Education Sciences (South China Normal
12 University), Ministry of Education

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14 **Running title:** Cerebellar connectome in MS and NMOSD

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16 ***Correspondence authors**

17 Jinhui Wang, Institute for Brain Research and Rehabilitation, Guangdong Key
18 Laboratory of Mental Health and Cognitive Science, Center for Studies of
19 Psychological Application, South China Normal University, Guangzhou 510631,
20 China. Email: jinhui.wang.1982@m.scnu.edu.cn; Yaou Liu, Department of
21 Radiology, Beijing Tiantan Hospital, Fengtai District, Capital Medical University,
22 No. 119 the West Southern 4th Ring Road, Beijing, 100070, China. Email:
23 liuyaou@bjtth.org.

24 **Abstract**

25 The cerebellum has been increasingly recognized to play key roles in the pathology of
26 multiple sclerosis (MS) and spectrum disorders (NMOSD), two main demyelinating
27 diseases with similar clinical presentations. Despite accumulating evidence from
28 neuroimaging research for cerebellar volumetric alterations in the diseases, however,
29 there have been no network-based studies examining convergent and divergent
30 alterations in cerebellar connectome between MS and NMOSD. This multisite and
31 multimodal study examined common and specific alterations in within-cerebellar
32 coordination and cerebello-cerebral communication between MS and NMOSD by
33 retrospectively collecting structural and resting-state functional MRI data from 208
34 MS patients, 200 NMOSD patients and 228 healthy controls (HCs) in seven sites in
35 China. Morphological brain networks were constructed by estimating interregional
36 similarity in cortical thickness and functional brain networks were formed by
37 calculating interregional temporal synchronization in functional signals. After
38 identifying cerebellar modular architecture and based on prior cerebral
39 cytoarchitectonic classification and functional partition, within-cerebellar and
40 cerebello-cerebral morphological and functional connectivity were compared among
41 the MS, NMOSD and HC groups. Five modules were identified within the cerebellum
42 including Primary Motor A (PMA), Primary Motor B (PMB), Primary Non-Motor
43 (PNM), Secondary Motor (SM) and Secondary Non-Motor (SNM) modules.
44 Compared with the HCs, the MS and NMOSD patients exhibited both increases and
45 decreases in within-cerebellar morphological connectivity that were mainly involved
46 in the PMA, PMB and SNM. Particularly, the two patient groups showed a common
47 altered pattern characterized by decreases between the PMA and SNM, both of which
48 were more densely connected with the PMB. For cerebello-cerebral morphological

49 connectivity, widespread reductions were found in both patient groups for the SM and
50 SNM with almost all cerebral cytoarchitectonic classes and functional systems while
51 increases were observed only in the NMOSD patients for the PMB with cerebral areas
52 involving motor and sensory domains. With regard to cerebellar functional
53 connectivity, fewer alterations were observed in the patients that were all
54 characterized by reductions and were mainly involved in cerebello-cerebral
55 interactions between cerebellar motor modules and cerebral association cortex and
56 high-order networks, particularly in the NMOSD patients. Cerebellar connectivity-
57 based classification achieved around 60% accuracies to distinguish the three groups to
58 each other with morphological connectivity as predominant features for
59 differentiating the patients from controls while functional connectivity for
60 discriminating the two diseases. Altogether, this study characterizes common and
61 specific circuit dysfunctions of the cerebellum between MS and NMOSD, which
62 provide novel insights into shared and unique pathophysiologic mechanisms
63 underlying the two diseases.

64

65 **Keywords:** cerebellum, functional connectivity, morphological network, cortical
66 thickness, multiple sclerosis, neuromyelitis optica

67 **Introduction**

68 While representing only about 10% of the size of the whole brain, the cerebellum
69 comprises much more neurons than the cerebrum (Andersen, Korbo, and Pakkenberg
70 1992; Herculano-Houzel n.d.; Pelvig et al. 2008) and contains almost 80% of the
71 surface area of the cerebral cortex (Sereno et al. 2020). The cerebellum has long been
72 considered to devote exclusively to motor control, the last decade has however
73 indicated that it is also engaged in various higher-level cognitive processes because of
74 its complex internal structure and tight interactions with motor and non-motor regions
75 of the cerebral cortex (Schmahmann et al. 2019; C. J. Stoodley and Schmahmann
76 2010). Moreover, cerebellar abnormalities are continuously demonstrated to be
77 associated with a variety of motor or non-motor dysfunctions, such as gait ataxia,
78 coordination deficiency, attention deficits, working memory decline, visual-spatial
79 function damage and information processing speed droop (Hoppenbrouwers et al.
80 2008; Schmahmann et al. 2019). Thus, the cerebellum has attracted considerable
81 research interests in recent years, particularly under different pathological conditions.

82 Multiple Sclerosis (MS) is an inflammatory and neurodegenerative disease
83 affecting the central nervous system such as the optic nerve, spinal cord, and the
84 cerebellum. Previous neuroimaging studies have shown that patients with MS exhibit
85 widespread alterations in cerebellar structure and function including reduced grey
86 matter volume (Anderson et al. 2009; Calabrese et al. 2010; Grothe et al. 2017;
87 Ramasamy et al. 2009; Weier et al. 2012), decreased cerebellar regional homogeneity
88 (Dogonowski et al., 2014) and altered cerebello-cerebral connectivity (Cerasa et al.
89 2012; Cocozza et al. 2018; Savini et al. 2019; Schoonheim et al. 2021; Tona et al.
90 2018). Neuromyelitis Optica spectrum disorders (NMOSD) is another inflammatory
91 central nervous system disease that primarily affects the optic nerves and spinal cord

92 with the IgG-antibody against the aquaporin-4 (AQP4) receptor as a specific
93 autoantibody marker (Chang and Chang 2020; Lennon et al. 2004). Since the
94 cerebellum has high AQP4 expression (Nico et al. 2002) and dense connections with
95 the optic nerves and spinal cord, it has been increasingly investigated in NMOSD and
96 is reported to show declined grey matter volume (Sun et al., 2019), disrupted
97 microstructural integrity (Liu et al., 2018), impaired functional connectivity (Han et
98 al., 2020; Liu et al., 2018) and abnormal spontaneous neural activity power (Liu et al.,
99 2020) in patients. These findings collectively indicate that the cerebellum is a
100 common, vulnerable structure to MS and NMOSD, which has important implications
101 for understanding the physiopathology and improving the evaluation of clinical
102 disability of the two diseases.

103 Despite the abovementioned advances, however, it is largely unknown whether
104 differential patterns of structural and functional alterations occur in the cerebellum
105 between MS and NMOSD. Currently, there are only a few studies that directly
106 compare MS and NMOSD with respect to cerebellar alterations. These studies
107 exclusively focused on cerebellar volume and consistently reported no significant
108 differences between MS and NMOSD (Duan et al. 2021; Lee et al. 2018; Schneider et
109 al. 2017; Weier et al. 2015). To date, there have been no network-based studies
110 examining whether and how cerebellar structural and functional connectivity are
111 differentially involved in MS and NMOSD and whether cerebellar connectivity have
112 the potential to distinguish the two diseases.

113 In this study, we examined convergent and divergent alterations in cerebellum
114 connectome by collecting structural and resting-state functional MRI data from a
115 large cohort of participants in 7 sites in China. First, a multiplex network model was
116 used to integrate morphological and functional connectivity for module detection

117 within the cerebellum via a multilayer community detection algorithm. Then,
118 cerebellar module-based morphological and functional connectivity were
119 characterized within the cerebellum and between the cerebellum and cerebral
120 cytoarchitectonic classes and functional systems. Finally, convergent and divergent
121 alterations in cerebellar connectivity and their clinical and behavioral relevance as
122 well as diagnostic power were examined.

123

124 **Materials and Methods**

125 **Participants**

126 This study included a total of 256 patients with MS, 270 patients with NMOSD and
127 281 health controls (HCs) from 7 different sites in China including Beijing Tiantan
128 Hospital, Xuanwu Hospital, Tianjin Medical University General Hospital, Huashan
129 Hospital, the First Affiliated Hospital of Nanchang University, China-Japan Union
130 Hospital of Jilin University, and the First Affiliated Hospital of Chongqing Medical
131 University. All participants were selected according to the following inclusion
132 criteria: (1) the MS patients were diagnosed according to 2010 McDonald criteria
133 (Polman et al. 2011) or 2017 revision of McDonald criteria (Thompson et al. 2018)
134 and the NMOSD patients satisfied the recently 2015 NMOSD criteria (Wingerchuk et
135 al. 2015); (2) without other neurological or psychiatric disorder; (3) the patients were
136 either in relapsing phase (less than 4 weeks from the last relapse) or remitting phase
137 (more than 4 weeks from the last relapse); (4) right-handed. We excluded 171
138 participants due to (1) image artifacts; (2) incomplete image or clinical data; and (3)
139 excess head motion during the R-fMRI scan (see below for details). Finally, a total of
140 636 participants (208 MS patients, 200 NMOSD patients and 228 HCs) were included
141 in the final analysis. This study was approved by the institutional review board of

142 corresponding hospitals, and written informed consent was obtained from each
143 participant.

144

145 **Clinical and neuropsychological assessment**

146 Clinical variables included disease duration and Expanded Disability Status Scale
147 (EDSS). For neuropsychological assessment, the following tests were used: the
148 California Verbal Learning Test-Second Edition (CVLT; assessing the auditory or
149 verbal episodic memory), the Brief Visuospatial Memory Test-Revised (BVMT,
150 assessing the visual or spatial episodic memory) and the Paced Auditory Serial
151 Addition Task (PASAT; assessing auditory processing speed, attention and working
152 memory). Of note, participants in only 3 sites (Beijing Tiantan Hospital, Xuanwu
153 Hospital, and Tianjin Medical University General Hospital) completed the
154 neuropsychological tests.

155

156 **Multimodal MR image acquisition**

157 Multimodal images were acquired in this study, including T2 FLAIR (fluid-attenuated
158 inversion recovery) image, high-resolution 3D T1-weighted imaging and R-fMRI.
159 The imaging parameters are summarized in Table S1.

160

161 **WM lesion loads and structural image filling**

162 For each patient, a WM lesion mask was manually delineated using the 3D-slicer
163 software (<https://www.slicer.org>) if WM hyperintensity was evident on his or her T2
164 FLAIR image. Individual WM lesion loads were calculated as the total volumes
165 within the masks. Based on the masks, a SLF algorithm was used to refill WM lesions
166 on individual T1 images (<http://atc.udg.edu/salem/slftoolbox/software.html>).

167 **Structural MRI processing**

168 ***Cerebellar cortical thickness.*** Cortical thickness estimation of the cerebellum
169 was accomplished by a patch-based multi-atlas segmentation tool called CERES
170 (Romero et al. 2017) - the winner of the Medical Image Computing and Computer-
171 Assisted Intervention cerebellum segmentation challenge (Carass et al. 2018) - on the
172 web-based platform volBrain (<http://volbrain.upv.es>; (Manjón and Coupé 2016)).
173 Briefly, a spatially adaptive non-local means filter was first applied to reduce noise in
174 the structural images. Then, the N4 bias field correction (Tustison et al. 2010) was
175 performed to correct intensity inhomogeneity. The corrected images were further
176 linearly registered (affine transform) to the standard Montreal Neurological Institute
177 (MNI) space using the Advanced Normalization Tools, followed by the N4 bias field
178 correction again. Subsequent analyses were limited to the cerebellar areas to reduce
179 the computational burden. To achieve a better cerebellum anatomic matching between
180 the corrected images and the library of manually labeled templates, the non-linear
181 transformations were estimated for the corrected images and templates to the MNI152
182 atlas. By concatenating the forward non-linear transformation to the MNI152 atlas of
183 each library case and the inverse non-linear transformation of each corrected image, a
184 subject-specific library was obtained. Finally, a local intensity normalization based on
185 regions of interest (ROIs) derived from majority voting segmentation of the subject-
186 specific library was applied to ensure the same intensity of cerebellar tissues across
187 participants.

188 ***Cerebral cortical thickness.*** Cortical thickness estimation of cerebral cortex was
189 performed by the CAT12 toolbox (<http://dbm.neuro.uni-jena.de/cat12/>) based on the
190 SPM12 package (<https://www.fil.ion.ucl.ac.uk/spm/software/spm12/>). As an efficient
191 and reliable alternative to FreeSurfer, the CAT12 toolbox provides a volume-based

192 approach for cortical thickness estimation without extensive reconstruction of cortical
193 surface. The CAT12 started with an initial segmentation of individual structural
194 images into gray matter (GM), white matter (WM) and cerebrospinal fluid (CSF)
195 based on an adaptive Maximum A Posterior technique (Rajapakse, Giedd, and
196 Rapoport 1997). Then, a projection-based method (Dahnke, Yotter, and Gaser 2013)
197 was performed to estimate cortical thickness which could handle the partial volume
198 information, sulcal blurring and sulcal asymmetries. Subsequently, a central cortical
199 surface was created and reparametrized into a common coordinate system via
200 spherical mapping (Yotter, Thompson, and Gaser 2011). Finally, individual cortical
201 thickness maps were resampled into the common fsaverage template and smoothed
202 using a Gaussian kernel with 15-mm full width at half maximum.

203

204 **Functional MRI processing**

205 The R-fMRI data were processed with the GRETNA toolbox (J. Wang et al. 2015)
206 based on the SPM12 package (<http://www.fil.ion.ucl.ac.uk/spm/software/spm12/>).
207 First, following removal of the first 5 volumes for magnetic saturation, inter-volume
208 head motion was corrected via rigid transformation. After converting rotational
209 displacements from degrees to millimeters on the surface of a sphere of radius 50 mm
210 (Power et al. 2012), participants with excessive motion were excluded in terms of the
211 criteria of > 3 mm translation or > 0.5 mm mean frame-wise displacement. For the
212 remaining participants, there were no significant differences in the maximum [HCs =
213 0.675 (IQR = 0.620), MS = 0.688 (IQR = 0.600), and NMOSD = 0.641 (IQR =
214 0.552); $p = 0.287$, permutation test] and mean frame-wise [HCs = 0.150 (IQR =
215 0.102), MS = 0.137 (IQR = 0.099), and NMOSD = 0.139 (IQR = 0.116); $p = 0.529$,
216 permutation test] displacement of head motion among three groups. Then, the

217 corrected functional images underwent band-pass filtering (0.01 - 0.08 Hz) and
218 nuisance regression [24-parameter head motion profiles (Friston et al. 1996), white
219 matter signals, cerebrospinal fluid signals and global signals] in a single linear model
220 to avoid reintroducing artifacts (Lindquist et al. 2019). The white matter and
221 cerebrospinal fluid signals were calculated within subject-specific masks derived from
222 tissue segmentation of individual structural images (threshold = 0.9), which were co-
223 registered to the corresponding mean volume of the corrected functional images.
224 Finally, the functional images were normalized into standard MNI space by applying
225 deformation fields derived from the tissue segmentation of individual structural
226 images, and spatially smoothed by a Gaussian kernel (full width at half maximum = 6
227 mm).

228

229 **Construction of within-cerebellar networks**

230 ***Cerebellar parcellation.*** In the CERES, the cerebellum was segmented using a
231 multi-atlas segmentation technique called non-local patch-based label fusion (Coupé
232 et al. 2011). Specifically, an Optimized PatchMatch algorithm (Giraud et al. 2016; Ta
233 et al. 2014) was performed to speed up the patch matching process and an adaptive
234 multi-scale approach was used for better accuracy. To avoid slightly irregular edges in
235 the resultant label maps, a convolution with a smoothing kernel of $5 \times 5 \times 5$ voxels
236 was further applied to the label maps. Finally, the cerebellum of each participant was
237 divided into 24 regions including the lobule I_II, lobule III, lobule IV, lobule V, lobule
238 VI, Crus I, Crus II, lobule VIIb, lobule VIIIa, lobule VIIIb, lobule IX and lobule X in
239 each hemisphere (Figure 1).

240 ***Morphological connectivity estimation.*** For a given pair of regions within the
241 cerebellum, morphological connectivity was estimated by calculating Jensen-Shannon

242 divergence based similarity in the distribution of intraregional cortical thickness.
243 Details on the calculation of morphological connectivity are described elsewhere (Li
244 et al. 2021; H. Wang et al. 2016).

245 ***Functional connectivity estimation.*** For a given pair of regions within the
246 cerebellum, functional connectivity was estimated by calculating Pearson correlation
247 in intraregional mean time series.

248

249 **Construction of cerebello-cerebral networks**

250 ***Cerebral parcellation.*** The cerebrum was divided into 400 ROIs using a
251 functionally defined atlas, which was derived from R-fMRI data from 1489 subjects
252 (Schaefer et al. 2018).

253 ***Morphological connectivity estimation.*** For each pair of regions between the
254 cerebellum and cerebrum, morphological connectivity was estimated using the same
255 methods as for the cerebellar morphological networks.

256 ***Functional connectivity estimation.*** For each pair of regions between the
257 cerebellum and cerebrum, functional connectivity was estimated using the same
258 methods as for the cerebellar functional networks.

259

260 **Removal of multicenter effects**

261 For multicenter studies, a crucial step is to remove site effects to avoid spurious
262 findings (Han et al. 2006). In this study, we utilized a harmonization approach called
263 Combat (J. P. Fortin et al. 2018) to moderate site effects. The Combat harmonization
264 is demonstrated to successfully remove inter-site technical variability while
265 preserving inter-site biological variability in image-based measurements. Specifically,
266 the Combat model can be written as:

267
$$y_{ijv} = \alpha_v + X_{ij}\beta_v + \gamma_{iv} + \delta_{iv}\varepsilon_{ijv}$$

268 where y_{ijv} represents connectivity strength of edge v (or cortical thickness of
269 region v) for subject j in site i , α_v is the average connectivity strength (or cortical
270 thickness) for edge (or region) v , X is a design matrix for the covariates of interest
271 (e.g., age, sex and group), β_v is a vector of regression coefficients corresponding to
272 covariates in X , and ε_{ijv} is the residual term that is assumed to follow a normal
273 distribution with zero mean. The terms γ_{iv} and δ_{iv} represent the additive and
274 multiplicative site effects of site i on edge (or region) v , respectively, and are
275 estimated by conditional posterior means as described in (J.-P. Fortin et al. 2017;
276 Johnson, Li, and Rabinovic 2007). The final ComBat-harmonized connectivity
277 strength (or cortical thickness) for edge (or region) v is calculated as:

278
$$y_{ijv}^{ComBat} = \frac{y_{ijv} - \hat{\alpha}_v - X_{ij}\hat{\beta}_v - \gamma_{iv}^*}{\delta_{iv}^*} + \hat{\alpha}_v + X_{ij}\hat{\beta}_v$$

279 where γ_{iv}^* and δ_{iv}^* are the empirical Bayes estimates of γ_{iv} and δ_{iv} , respectively.

280

281 **Cerebellar module detection**

282 Cerebellar modular architecture was identified by applying a multilayer community
283 detection algorithm to a group-level multiplex network, which integrated both
284 morphological and functional connectivity within the cerebellum for the HCs. First, a
285 group-level mean morphological network and functional network were separately
286 obtained for the HCs. As two layers, these two networks were then interconnected via
287 adding edges to link each node in one layer with replica of the node in the other layer,
288 therefore forming a multiplex network. Thereafter, a nonparametric method of locally
289 adaptive network sparsification (Foti, Hughes, and Rockmore 2011) was utilized such
290 that only those locally significant connections that could not be explained by random

291 variation were retained in the multiplex network. Module detection is to find a
292 specific partition of network nodes that yields the largest modularity, Q , which is
293 defined for a multilayer network as (Mucha, Richardson, Macon, Porter, & Onnela,
294 2010):

$$295 \quad Q = \frac{1}{2\mu} \sum_{ijsr} \left[\left(A_{ijs} - \gamma_s \frac{k_{is} k_{js}}{2m_s} \right) \delta_{sr} + \delta_{ij} C_{jsr} \right] \delta(g_{is}, g_{jr})$$

296 where i and j represent nodes, s and r represent layers, A_{ijs} denotes intra-layer
297 connectivity strength between node i and node j in layer s , C_{jsr} denotes inter-
298 layer connectivity strength of node j between layer s and layer r , $k_{is} = \sum_j A_{ijs}$
299 denotes total intra-layer connectivity strength of node i in layer s , m_s is total
300 connectivity strength of all edges in layer s , γ_s is module resolution in layer s (the
301 larger γ , the more small-size communities), g_{is}/g_{jr} represents module assignment
302 of node i/j in layer s/r , 2μ is equal to $\sum_{jr} (k_{js} + c_{js})$ with $c_{js} = \sum_r C_{jsr}$
303 indicating total inter-layer connectivity strength node j from layer s to other layers,
304 and δ is a binary notation (1 if the same node for δ_{ij} , the same layer for δ_{sr} and the
305 same community assignment $\delta(g_{is}, g_{jr})$ and 0 otherwise). In this study, the module
306 detection was performed using the Louvain algorithm (Blondel et al. 2008).

307 During the processes mentioned above, there were two parameters that may
308 affect final module identification: the inter-layer connectivity strength, ω , and the
309 module resolution, γ . To identify stable cerebellar modular architecture, we
310 calculated the modularity, Q across a large range in the 2D parameter space of
311 (ω, γ) : $\omega = [0.01 - 1]$ and $\gamma = [0.01 - 3]$ both with an increment of 0.01.
312 Specifically, at each (ω, γ) in the 2D parameter space, we performed module
313 detection 1,000 times and obtained a consensus matrix in which the elements
314 indicated the proportions of each pair of nodes that were assigned to the same module

315 over the 1,000 repetitions (Bassett et al. 2013). If there existed any pair of nodes that
316 were not consistently assigned to the same module, the module detection was further
317 conducted on the consensus matrix (1,000 times) to derive a new consensus matrix.
318 Such procedures were iterated until a consensus partition was obtained. That is, the
319 final consensus matrix included only 0 and 1. After determining cerebellar partition at
320 each (ω, γ) , we employed the variation of information (Meilă 2003) to evaluate
321 stability of the cerebellar partition over different (ω, γ) . More specifically, we
322 computed the mean variation of information of cerebellar partition between a given
323 (ω_i, γ_j) and its eight contiguous neighbors $(\omega_{i-1}, \gamma_{j-1}, \omega_{i-1}, \gamma_j, \omega_{i-1}, \gamma_{j+1},$
324 $\omega_i, \gamma_{j-1}, \omega_i, \gamma_{j+1}, \omega_{i+1}, \gamma_{j-1}, \omega_{i+1}, \gamma_j$ and $\omega_{i+1}, \gamma_{j+1})$. The smaller the mean
325 variation of information was, the more stable the cerebellar partition was when subtle
326 (ω, γ) fluctuations occurred. Finally, we searched the widest continuous (ω, γ) range
327 in which the mean variation of information was equal to the minimum, and the
328 corresponding cerebellar partition was regarded as canonical cerebellar modular
329 architecture.

330

331 **Cortical cytoarchitectonic classification and functional partition**

332 The cerebrum can be divided into several canonical modules based on different
333 approaches. In this study, two partitions - cytoarchitectonic classification and
334 functional partition - were used to divide the cerebrum since a recent study showed
335 that structural and functional connectivity followed cytoarchitectonic and functional
336 hierarchies in the cerebrum (Vázquez-Rodríguez et al. 2019). 1) Cytoarchitectonic
337 classification of the cerebrum (cerebrum^C hereafter). The cerebral cortex displays
338 substantial variation in cytoarchitecture, according to which the cerebrum is
339 categorized into seven classes (Seidlitz et al. 2018; Vértes et al. 2016): primary motor

340 cortex (PM), association cortex (AC1), association cortex (AC2), primary/secondary
341 sensory (PSS), primary sensory cortex (PS), limbic regions (LB) and insular cortex
342 (IC). We manually assigned each of the 400 cerebral regions to one of the seven
343 cytoarchitectonic classes. 2) Functional partition of the cerebrum (cerebrum^F
344 hereafter). In terms of intrinsic functional connectivity patterns, the second partition
345 divides the cerebrum into seven functional systems (Thomas Yeo et al. 2011): visual
346 network (VN), somatomotor network (SMN), dorsal attention network (DAN), ventral
347 attention network (VAN), limbic network (LN), fronto-parietal network (FPN) and
348 default mode network (DMN). Each of the 400 cerebral regions belonged to one of
349 the seven functional systems.

350

351 **Construction of cerebellar module-based networks within the cerebellum and**
352 **between the cerebellum and cerebrum**

353 Based on the methods mentioned above, we constructed 4 networks for each
354 participant at the region level: one 24×24 within-cerebellar morphological network,
355 one 24×24 within-cerebellar functional network, one 24×400 cerebello-cerebral
356 morphological network and one 24×400 cerebello-cerebral functional network.

357 Based on the cerebellar modular architecture (5 modules; see Results for details) and
358 cerebrum^C and cerebrum^F, the 4 region-level networks were transformed to 6
359 cerebellar module-based networks: one 5×5 within-cerebellar morphological
360 network, one 5×5 within-cerebellar functional network, one 5×7 cerebello-
361 cerebral^C morphological network, one 5×7 cerebello-cerebral^F morphological
362 network, one 5×7 cerebello-cerebral^C functional network and one 5×7 cerebello-
363 cerebral^F functional network. This was achieved by averaging connectivity weights of
364 all connections linking any pair of cerebellar modules or linking one cerebellar

365 module and one cerebral cytoarchitectonic class/functional system.

366

367 **Statistical analysis**

368 ***Group effects.*** Chi-squared tests were used to compare dichotomous variables
369 including sex (male vs. female), disease state (acute vs. chronic) and WM lesion
370 (presence vs. absence). For continuous variables (demographic: age; clinical: disease
371 duration, EDSS and WM lesion volume; neuropsychological: CVLT, BVMT and
372 PASAT; imaging-based: cortical thickness, morphological connectivity and functional
373 connectivity), non-parametric permutation tests were utilized after examining their
374 normality (Lilliefors test). Age, sex and mean framewise displacement of head motion
375 (if applicable) were treated as covariates in the analyses of neuropsychological
376 variables and imaging-based measures. Briefly, for a given continuous variable, we
377 initially calculated a statistic (T or F) through two-sample t test (for disease duration,
378 EDSS and WM lesion volume), ANOVA (for age) or ANCOVA (for CVLT, BVMT,
379 PASAT, cortical thickness, morphological connectivity and functional connectivity).
380 To obtain an empirical distribution of the statistic, we randomly reshuffled the data
381 and re-computed the statistic (10,000 times). Based on the empirical distribution, a *p*
382 value was calculated as the proportion of permutations that generated the statistic
383 equal to or greater or less than (for T) or equal to or greater than (for F) the real
384 observation. Notably, the covariates were not reshuffled during the permutation
385 process. To correct for multiple comparisons for imaging-based measures, the false
386 discovery rate (FDR) procedure was used at the level of $q < 0.05$ (cortical thickness: 5
387 cerebellar modules; morphological/functional connectivity: 15 within-cerebellar and
388 35 cerebello-cerebral connections). For significant differences among the three
389 groups, post hoc pairwise comparisons were further performed by permutation tests.

390 ***Relationship between cerebellar alterations and clinical/neuropsychological***
391 ***variables.*** For imaging-based metrics showing significant group effects, non-
392 parametric Spearman partial correlation was used to evaluate their relationships with
393 clinical and neuropsychological variables in the MS and/or NMOSD group in terms of
394 patterns of post hoc comparisons. Age, sex and mean framewise displacement of head
395 motion (if applicable) were regarded as covariates in the correlation analyses. Again,
396 the FDR procedure was employed to correct for multiple comparisons.

397

398 **Classification analysis**

399 To examine the potential of cerebellar connectivity in distinguishing the three groups
400 from each other, we trained linear SVM classifiers with all within-cerebellar and
401 cerebello-cerebral morphological and functional connectivity as initial features. Out-
402 of-sample classification performance of the classifiers was evaluated using a 10-fold
403 cross-validation procedure. Specifically, in each fold of classification between two
404 groups, feature selection was first performed to exclude irrelevant features via non-
405 parametric permutation tests on the training dataset ($p < 0.05$; 1,000 permutations).
406 Based on the selected features, a linear SVM classifier was then trained and applied to
407 the testing dataset. After the 10-fold cross-validation procedure, a classification
408 accuracy was calculated. To obtain robust estimation of the classification accuracy,
409 the above classification process was repeated 100 times and the resultant mean across
410 all repeats was reported. Meanwhile, the consensus features that were consistently
411 selected in at least 90% of all folds and repeats ($10 \times 100 = 1,000$) were recorded
412 together with their mean weights. Finally, to further evaluate whether the classifiers
413 performed significantly better than random operations, an empirical null distribution
414 of classification accuracy was obtained by reshuffling the group labels of participants

415 (1,000 times) followed by 10-fold cross-validated classification. A p value was
416 computed as the proportion of values in the empirical null distribution that were larger
417 than the real observation. Notably, effects of age, sex, and mean frame-wise head
418 motion (if applicable) were removed from all features via multiple linear regression
419 before the classification procedure.

420

421 **Results**

422 **Demographic, clinical, and neuropsychological evaluations**

423 Table 1 summarizes demographic, clinical, and neuropsychological information of the
424 participants. Significant group effects were found on age ($p = 0.003$) and sex ($p <$
425 0.001) separately due to an older age of the NMOSD patients than the MS patients (p
426 < 0.001) and HCs ($p = 0.020$) and a higher female-to-male ratio of the NMOSD
427 patients than the MS patients ($p < 0.001$) and HCs ($p < 0.001$) as well as of the MS
428 patients than the HCs ($p = 0.025$). For clinical data, no significant differences were
429 found in the acute-to-chronic phase ratio or disease duration between the two patient
430 groups ($p > 0.05$). However, the NMOSD patients exhibited lower incidence of WM
431 lesion ($p < 0.001$) and less lesion volume ($p < 0.001$) than the MS patients. In
432 addition, the NMOSD patients showed significantly higher EDSS than the MS
433 patients ($p < 0.001$). Regarding neuropsychological variables, both the patient groups
434 performed significantly worse than the HCs in the CVLT and PASAT (both $p <$
435 0.001). In addition, the NMOSD patients showed lower BVMT than the HCs ($p <$
436 0.001). No differences were found in any neuropsychological variable between the
437 two patient groups ($p > 0.05$).

438

439 **Cerebellar modular architecture**

440 In the searched 2D parameter space of inter-layer connectivity strength, ω , and
441 resolution coefficient, γ , we identified a widest range ($\omega = 0.06 \sim 0.50$; $\gamma =$
442 $1.39 \sim 2.00$) wherein cerebellar modular architecture maintained stable when the
443 parameters fluctuated (Figure 2). Specifically, for each combination of the parameter
444 pair in the identified range, the cerebellum was subdivided into five spatially
445 contiguous and bilaterally symmetrical modules with regions resembling largely
446 between morphological and functional networks ($Q = 0.566$; Figure 3). The modules
447 had a good correspondence with the well-established cerebellar double-motor/triple-
448 non-motor organization, and thus were termed Primary Motor A module (PMA,
449 including bilateral Lobule I-II and Lobule III, with additional bilateral Lobule X for
450 morphological networks), Primary Motor B module (PMB, including bilateral Lobule
451 IV, Lobule V and Lobule VI), Primary Non-Motor module (PNM, including bilateral
452 Crus I and Crus II), Secondary Motor module (SM, including bilateral Lobule VIIb,
453 Lobule VIIIa and Lobule VIIIb) and Secondary Non-Motor module (SNM, including
454 bilateral lobule IX, with additional bilateral Lobule X for functional networks).

455

456 **Alterations in cerebellar module-based cortical thickness**

457 Significant group effects were found on the mean cortical thickness within the
458 cerebellar PMA, PMB, SM and SNM ($p < 0.05$, FDR corrected). Post hoc
459 comparisons revealed that the group effects were due to common cortical thickening
460 to the two patient groups ([MS = NMOSD] $>$ HCs): SM and SNM, MS-specific
461 cortical thickening (MS $>$ [NMOSD = HCs]: PMA), or NMOSD -specific cortical
462 atrophy (NMOSD $<$ [MS = HCs]: PMB) (Figure 4).

463

464 **Alterations in cerebellar morphological connectivity**

465 Differences in cerebellar morphological connectivity are shown in Figure 5 (top
466 panel).

467 ***Within-cerebellar morphological connectivity.*** Significant group effects were
468 found on the mean morphological connectivity within two cerebellar modules (PMB
469 and SM) and between four pairs of cerebellar modules (PMA - PMB, PMA - SM,
470 PMA - SNM and PMB - SNM) ($p < 0.05$, FDR corrected). Post hoc comparisons
471 revealed that the group effects were owing to MS-related decreases (MS < HCs:
472 PMB), MS-specific increases (MS > [NMOSD = HCs]: SM), common decreases
473 ([MS = NMOSD] < HCs: PMA - SNM) and increases ([MS = NMOSD] > HCs: PMA
474 - PMB and SNM - PMB) to the two patient groups, or NMOSD-related decreases
475 (NMOSD < HCs: PMA - SM).

476 ***Cerebello-cerebral morphological connectivity.*** In the context of the cerebrum^C,
477 significant group effects were found on the mean morphological connectivity for the
478 cerebellar SM and SNM with all cerebral cytoarchitectonic classes, cerebellar PMB
479 with cerebral PM, PS, LB and IC, and cerebellar PNM with cerebral AC2 ($p < 0.05$,
480 FDR corrected). Post hoc comparisons revealed that the group effects were because of
481 common decreases to the two patient groups ([MS = NMOSD] < HCs: SM - all
482 cerebral cytoarchitectonic classes, and SNM - AC1, AC2, PSS, PS and LB), NMOSD-
483 specific increases (NMOSD > [MS = HCs]: PMB - PM, PS and IC), NMOSD-related
484 increases (NMOSD > HCs: PMB - LB), or MS-related decreases (MS < HCs: PNM -
485 AC2, and SNM - PM and IC). When the cerebrum^F was used, significant group
486 effects were found on the mean morphological connectivity for the cerebellar SM and
487 SNM with all cerebral functional systems, cerebellar PMB with cerebral SMN, and
488 cerebellar PNM with cerebral DMN ($p < 0.05$, FDR corrected). The group effects
489 originated from common decreases to the two patient groups ([MS = NMOSD] <

490 HCs: SM and SNM - all cerebral functional systems), NMOSD-related decreases
491 (NMOSD < HCs: PMB - SMN), or MS-related decreases (MS < HCs: PNM - DMN).

492

493 **Alterations in cerebellar functional connectivity**

494 Differences in cerebellar functional connectivity are summarized in Figure 5 (bottom
495 panel).

496 ***Within-cerebellar functional connectivity.*** Significant group effects were found
497 on the functional connectivity only within the cerebellar SM ($p < 0.05$, FDR
498 corrected) due to MS-specific decreases (MS < [NMOSD = HCs]).

499 ***Cerebello-cerebral functional connectivity.*** In the context of the cerebrum^C,
500 significant group effects were found on the mean functional connectivity for the
501 cerebellar PMB with cerebral AC1 and AC2, cerebellar SM with cerebral AC1, and
502 cerebellar PMA with cerebral AC1 ($p < 0.05$, FDR corrected). Post hoc comparisons
503 revealed that the group effects were attributable to common decreases to the two
504 patient groups ([MS = NMOSD] < HCs: PMB - AC1 and AC2, and SM - AC1), or
505 NMOSD-related decreases (NMOSD < HCs: PMA - AC1). When the cerebrum^F was
506 used, significant group effects were found on the mean functional connectivity for
507 the cerebellar PMB with cerebral FPN and LN, cerebellar SM with cerebral FPN and
508 DMN, and cerebellar PMA with cerebral FPN ($p < 0.05$, FDR corrected). The group
509 effects were driven by common decreases to the two patient groups ([MS =
510 NMOSD] < HCs: PMB - FPN, and SM - FPN), or NMOSD-specific decreases
511 (NMOSD < [MS = HCs]: PMA - FPN, PMB - LN, and SM - DMN).

512

513 **Relationship between imaging features and other variables in the patients**

514 Only in the MS group, significant correlations were found characterized by positive

515 correlations between the mean cortical thickness of the cerebellar PMA and disease
516 duration ($r = 0.193, p = 0.006$), lesion volume ($r = 0.235, p < 0.001$) and EDSS ($r =$
517 $0.207, p = 0.003$), and between the mean cortical thickness of the cerebellar SNM and
518 lesion volume ($r = 0.230, p < 0.001$) (Figure 6).

519

520 **Classification results**

521 The cerebellar morphological and functional connectivity could distinguish the three
522 groups from each other with around 60% accuracy (MS vs HCs: 63.3%, $p < 0.001$;
523 NMOSD vs HCs: 64.4%, $p < 0.001$; MS vs NMOSD: 57.9%, $p = 0.001$). As shown in
524 Figure 7, the features contributing to the classification were mainly composed of
525 those showing between-group differences (MS vs HCs: 83.7%; NMOSD vs HCs:
526 80.0%; MS vs NMOSD: 55.6%). Interestingly, morphological connectivity
527 predominated in features contributing to the differentiation between the patients and
528 HCs (MS vs HCs: 75.5%; NMOSD vs HCs: 64.0%) while the classification between
529 the two diseases mainly benefited from functional connectivity (MS vs NMOSD:
530 88.9%). In the context of cerebellar modular architecture, the morphological
531 connectivity contributing to the classification between the patients and HCs were
532 mainly related to the PMB, SM and SNM (MS vs HCs: 91.9%; NMOSD vs HCs:
533 100%). For classifying the two patient groups, the functional connectivity
534 contributing to the classification were all involved in the three motor-related modules
535 (i.e., the PMA, PMB and SM; MS vs NMOSD: 100%).

536

537 **Discussion**

538 In this study, we investigated the cerebellar morphological and functional connectivity
539 in patients with MS and NMOSD. For cerebellar morphological connectivity,

540 common alterations of both increases and decreases in within-cerebellar coordination
541 and decreases in cerebello-cerebral communication were observed in the two patient
542 groups. The MS patients showed specific increase within the cerebellar SM module
543 while the NMOSD patients showed specific increases between cerebellar primary
544 motor module and cerebral areas involving motor and sensory domains. Relative to
545 morphological connectivity, fewer alterations were observed in cerebellar functional
546 connectivity in the two patient groups that were all characterized by disease-related
547 reductions and were mainly involved in cerebello-cerebral interactions between
548 cerebellar motor modules and cerebral association cortex and high-order networks.
549 The MS patients showed specific decrease within the cerebellar SM module while the
550 NMOSD patients exhibited specific decreases between cerebellar motor modules and
551 cerebral limbic system and default mode regions. These network-level cerebellar
552 dysfunctions provide novel insights into shared and unique pathophysiologic
553 mechanisms between MS and NMOSD.

554

555 **Cerebellar modular architecture**

556 Modular organization is one of the main organizational principles of the human brain
557 networks (Sporns and Betzel 2016). For the cerebellum, a hierarchical double-
558 motor/triple-non-motor organization has been well established previously based on
559 cerebellar representations of cerebral intrinsic connectivity networks (Buckner et al.
560 2011), cerebellar task-evoked activity (Guell, Gabrieli, and Schmahmann 2018),
561 cerebello-cerebral functional connectivity profiles (Guell, Gabrieli, and Schmahmann
562 2018) and gradients of within-cerebellar functional connectivity patterns (Guell et al.
563 2018). In contrast to single modality in these previous studies, here we utilized a
564 multiplex network model to merge both morphological and functional connectome

565 information within the cerebellum for cerebellar module detection. Five modules were
566 reliably identified with module composition largely comparable between
567 morphological and functional cerebellar networks. Moreover, the cerebellar modular
568 architecture well matches with the cerebellar double-motor/triple-non-motor
569 organization, and thus provide further, in particular new anatomical, evidence for the
570 cerebellar hierarchical organization. Specifically, the PMA and PMB correspond to
571 the first motor representation of the cerebellum and are mainly involved in leg and
572 foot movements (Marek et al. 2018;; Xue et al. 2021), and hand and face movements
573 (Guell, Gabrieli, and Schmahmann 2018; Marek et al. 2018; Xue et al. 2021),
574 respectively; The PNM is a mixture of the first and second non-motor representations
575 of the cerebellum and is primarily related to working memory, executive function,
576 language, social processing and emotion processing (E et al. 2014; Guell, Gabrieli,
577 and Schmahmann 2018). The SM mirrors the second motor representation of the
578 cerebellum and is mainly engaged in motor activities with demands of task focus,
579 visual process, attention and working memory (Brissenden et al. 2018; Guell et al.
580 2018); The SNM resembles the third non-motor representation of the cerebellum and
581 is predominantly engaged in compounded domains including working memory and
582 emotion processing, and sometimes additional visuo-spatial functions and balance
583 maintaining (Barmack and Pettorossi 2021; Guell, Gabrieli, and Schmahmann 2018;
584 Guell and Schmahmann 2020; Palesi et al. 2021).
585

586 **Alterations in cerebellar cortical thickness**

587 Our main finding of cortical thickness comparisons was cortical thickening in both
588 MS and NMOSD patients. This is in contrast to previous cerebellar studies based on
589 gray matter volume that consistently reported decreases in MS and NMOSD

590 (Calabrese et al. 2010; Cocozza et al. 2018; Grothe et al. 2017; Sun et al. 2019). The
591 discrepancy is in line with the viewpoint that volumetric measures may overlook
592 specific morphological alterations in diseases since they reflect a composite of
593 cortical thickness, surface area and folding (Hutton et al. 2009). Interpretation of
594 cortical thickening is not straightforward. One speculative interpretation is attributable
595 to delays in normal physiological cleaning process caused by disease effects on
596 pruning redundancy (Hong et al. 2016). Another possible reason, which may be more
597 relevant to this study, is compensatory cortical reorganization to gain or maintain
598 function (Burge et al. 2016). We found that both the MS and NMOSD patients
599 exhibited cortical thickening in the SM and SNM modules. Given the functions of
600 these two modules as discussed above, the common cortical thickening might reflect
601 similar compensatory mechanisms in response to functional declines shared by the
602 two diseases in visual scanning, motor speed, attention, learning and working memory
603 (De Souza and Ashburn 1996; Guimarães and Sá 2012; Kawachi 2019; Oertel et al.
604 2019). In addition to the transdiagnostic cortical thickening, MS-specific cortical
605 thickening was observed in the PMA module, which might be due to adaptive
606 reorganization caused by leg restlessness, a frequent and unique sensory symptom in
607 MS patients (Manconi et al. 2007; Schürks and Bussfeld 2013). Moreover, the PMA
608 cortical thickness was positively correlated with EDSS scores and disease durations in
609 the MS patients, suggesting that cortical thickening may be a biomarker to monitor
610 clinical disability of MS. Finally, we found that the PMB module exhibited NMOSD-
611 specific cortical thinning. Several previous case studies have reported that NMOSD
612 patients presented choreoathetosis or pseudoathetosis, a series of involuntary and
613 irregular movements of the face, head, hand or limbs (Boddu and Shenker 2019;
614 Seok, Jang, and You 2018; Sugeno 2013). In addition, headaches and craniofacial pain

615 were reported in NMOSD patients (Nielsen et al. 2018). Since cortical thinning
616 typically reflects neuron loss and functional deficits, the thinner PMB is speculated to,
617 at least partly, be responsible for these symptoms in NMOSD patients.

618

619 **Alterations in cerebellar morphological connectivity**

620 Both the MS and NMOSD patients showed aberrant morphological connectivity
621 within the cerebellum. First, the patients with MS and NMOSD exhibited common
622 decreases in morphological connectivity between the PMA and SNM. As mentioned
623 above, the PMA is mainly involved in leg and foot movements, while the SNM is
624 predominately related to visuospatial function. Thus, the decreased morphological
625 coordination may contribute to impaired visual-motor integration in MS and NMOSD
626 (Harder et al. 2015; Julian et al. 2013; Nunan-Saah et al. 2015). Interestingly, both the
627 PMA and SNM exhibited increased connectivity with the PMB in the two patient
628 groups. The enhancement may reflect a common mechanism to the two diseases to
629 compensate for pathological disruptions in the morphological connectivity between
630 the PMA and SNM by constructing an indirect, alternative pathway. Since both the
631 patient groups exhibited decreased morphological connectivity within the PMB
632 (notably, the decrease did not reach significant in NMOSD after correcting for
633 multiple comparisons), the compensatory process may occur by adaptively altering
634 internal morphology of the PMB module in a bimodal way to enhance its
635 morphological homogeneity with the PMA and SNM, respectively, which finally
636 alleviates the loss of visual-motor integration in patients. Second, we found
637 morphological connectivity increase within the SM in the MS patients. This cannot be
638 simply interpreted as a derivative of MS-related overall cortical thickening in the SM
639 since cortical thickening but intact morphological connectivity was observed in the

640 NMOSD patients. Thus, the MS-specific morphological connectivity increase implies
641 the existence of unique mechanisms that make the SM towards more similar
642 morphology, presumably to compensate for motor-related dysfunction. The increase
643 provides possible physiological substrate of reversibility of neurological disability in
644 MS (Akaishi et al. 2020).

645 For cerebello-cerebral morphological connectivity, the two patient groups
646 showed extensive disruptions. Specifically, the MS and NMOSD patients showed
647 common decreases for the SM and SNM with almost all the cerebral components
648 regardless of the cytoarchitectonic classification or functional partition. The
649 cerebellum is closely linked with the cerebrum via complex cerebello-cerebral loops
650 to exchange information with each other and participate collectively in various motor
651 and non-motor activities. Thus, the widely disrupted cerebello-cerebral morphological
652 loops may be related to poor performance on a set of motor and non-motor activities
653 in MS and NMOSD. Notably, the disruptions were specific to the secondary rather
654 than primary modules. Previous studies have suggested that the secondary modules
655 are different from the primary modules in contributing to motor and non-motor
656 activities (Guell et al. 2018; Schmahmann et al. 2019). Compared with the primary
657 modules, the secondary modules are engaged in motor processes that require higher
658 task focus and sometimes visual process rather than pure movements, and are devoted
659 to processes involving multiple cognitive and affective activities rather than single
660 non-motor domain (Guell et al. 2018; Schmahmann et al. 2019). Taking these facts
661 into consideration, we could reasonably deduce that the common disruptions in
662 numerous cerebello-cerebral loops to both the patient groups that were selectively
663 involved the SM and SNM modules might account for the well described deficits in
664 those compounded motor and non-motor activities as reflected by worse performance

665 on BVMT, SDMT and CVLT scales in MS and NMOSD (Cao et al. 2020; Corfield
666 and Langdon 2018; Guo et al. 2019; Meng et al. 2017). Besides the common
667 morphological connectivity decreases, we found NMOSD-specific morphological
668 connectivity enhancement between the cerebellar PMB module and cerebral PM, PS,
669 IC and SMN, all of which play important roles in motor and sensory domains. The
670 morphological connectivity increases might be seen as a compensatory response to
671 NMOSD-specific cortical thinning in the PMB module to indemnify motor and
672 sensory functions in patients.

673 Finally, we found MS-related cerebello-cerebral morphological connectivity
674 reductions for the cerebellar PNM with the cerebral AC2 and DMN. The cerebellar
675 PNM is constituted of the bilateral Crus I and Crus II, which are key cognitive lobules
676 in the cerebellum and play crucial roles in working memory, attention, language,
677 executive function and emotional self-experiences (S. H. A. Chen and Desmond 2005;
678 E et al. 2014; Guell, Gabrieli, and Schmahmann 2018; C. Stoodley and Schmahmann
679 2009). Previous studies have indicated that both cerebellar Crus I and Crus II have
680 close connection with cerebral association cortex (Sasaki et al. 1975; Xue et al. 2021),
681 and lesions affecting the Crus I and Crus II lead to cognitive impairments by
682 interrupting cerebellar regulation of cognitive loops with cerebral association cortex
683 (C. J. Stoodley and Schmahmann 2010). On the other hand, the cerebellar Crus I and
684 Crus II are also found to be major sites to interact with the cerebral DMN (Buckner et
685 al. 2011), and damages to cerebello-cerebral DMN connections are related to
686 cognitive impairments in MS patients (Savini et al. 2019). Accordingly, we speculate
687 that the disrupted cerebello-cerebral loops between the cerebellar PNM and the
688 cerebral AC2 and DMN may contribute to a variety of cognitive deficits in MS
689 patients. Notably, no alterations in cortical thickness and within-cerebellar

690 morphological connectivity were observed for the cerebellar PNM, suggesting a
691 susceptibility of MS pathology to long-range connectivity of this module with the
692 cerebrum. Using diffusion tensor images to estimate structural pathways in the
693 cerebrum, a previous study showed that long-range connections linking remote
694 regions were more severely disrupted in MS (Meijer et al. 2020). Expanding the
695 previous finding, our results suggest that the vulnerability of long-range connections
696 to MS holds for cerebellar connectivity and the susceptibility is salient for certain
697 regions or circuits. Moreover, our findings provide important clues to a previous
698 viewpoint that cerebellar inflammation amplifies other pathologic mechanisms in MS
699 via long-range connections, such as retrograde neurodegeneration of cortical regions
700 even far away from lesion sites in the cerebellum (Muthuraman et al. 2020). Deeper
701 understanding of these findings may benefit from future research by integrating
702 multidimensional data, animal models and computational modeling.

703

704 **Alterations in cerebellar functional connectivity**

705 Compared with morphological connectivity, much less functional connectivity was
706 detected to show alterations in the patients and the alterations exhibited an aggregated
707 pattern mainly involving cerebellar motor modules and cerebral association cortex or
708 high-order networks. Moreover, all the functional connectivity alterations were
709 characterized by decreases in the patients. Studies related to MS pathology have
710 suggested that moderate structural damage to the brain can be compensated by
711 adaptive functional connectome reconfiguration, but once the structural damage
712 exceeds a certain upper limit, the functional connectome collapses and clinical
713 symptoms and cognitive impairments ensue (Fleischer, Radetz, et al., 2019;
714 Schoonheim, Meijer, & Geurts, 2015). According to this theory, the observed

715 functional connectivity disruptions may reflect functional maladaptation or failure of
716 adaptive reorganization due to severe neurodegeneration and structural damage in the
717 patients. Specifically, for within-cerebellar functional connectivity, MS-specific
718 decreases were observed in the SM. According to the functions of the SM as
719 discussed above, the decreases may be responsible for degeneration of complex motor
720 functions and clinical disability in patients with MS.

721 With regard to cerebello-cerebral functional connectivity, both the patient groups
722 showed decreases for the cerebellar PMB and SM with the cerebral association cortex
723 and FPN. There are several lines of evidence indicating that the cerebellum is closely
724 related with both the cerebral association cortex and the FPN. From the perspective of
725 phylogenetic evolution, the newest parts of the cerebellum (e.g., the posterior lobe
726 that includes the SM) develops specifically in parallel with cerebral association cortex
727 rather than the cerebral cortex as a whole (Leiner, Leiner, and Dow 1986), facilitating
728 the coordination between the cerebellum and cerebral association areas to exchange
729 highly-processed multisensory information and cooperatively participate in high-level
730 functions (Leiner, Leiner, and Dow 1991). On the other hand, from the viewpoint of
731 functional network organization, the FPN is overrepresented in the cerebellum
732 compared with the cerebral cortex (Marek et al. 2018), which plays a major role in
733 task set initiation and task switching, modulating the integration of other association
734 and motor networks (Dosenbach et al. 2007; Marek and Dosenbach 2018). Moreover,
735 topography of the FPN in the cerebellum overlapped the cerebellar PMB and SM.
736 Together, the disrupted functional connectivity suggests failed or weakened
737 information exchange between the cerebellar motor modules and the cerebral
738 association cortex as well as FPN in supporting motor, multisensory and high-level
739 processes, which may further lead to poor performance of MS and NMOSD patients

740 in activities involving in motor and cognition.

741 In addition to the common decreases, the NMOSD patients exhibited additional
742 functional connectivity decrease for the cerebellar PMA with the cerebral association
743 cortex and FPN, cerebellar PMB module with cerebral LN, and cerebellar SM with
744 cerebral DMN. These findings suggest more serious disruptions in cerebello-cerebral
745 functional integration in NMOSD, which may be related unique pathology of the
746 disease and/or specific cognitive dysfunction of patients.

747

748 **Classification results**

749 Our classification results indicated that cerebellar connectivity had the potential to
750 help distinguish the patients from controls and the two diseases from each other
751 although the accuracies were relatively low. Clinical heterogeneity of the patients, a
752 critical concern for retrospective large-scale multisite studies, may be a main reason
753 for the low accuracies. In addition, the accuracies can be further improved for future
754 studies by using more sophisticated deep learning algorithms, such as convolutional
755 neural network. Interestingly, features contributing to the classification depended on
756 the connectivity style. This finding suggests unique susceptibility of different types of
757 connectivity to clinical research, possibly due to their poor cross-modal
758 correspondences as reported previously (Reid et al. 2016). Moreover, certain
759 cerebellar modules dominated the connectivity features in the classification.

760 Accordingly, future studies can improve the classification accuracies by exclusively
761 focusing on a specific set of regions or connections that are closely related to the
762 diseases.

763

764 **Limitations**

765 This study had several limitations. First, only a subset of participants in certain sites
766 completed a few neuropsychological assessments. This might explain why no
767 correlations were found for the cerebellar network alterations with neuropsychological
768 assessment. Second, there was a lack of standardized imaging protocols across
769 different sites due to the retrospective design of this multicenter study. Although we
770 utilized the Combat harmonization approach to mitigate site effects, it is still not clear
771 about the extent to which our findings are contaminated by residual site effects. Third,
772 MS and NMOSD are heterogeneous diseases both of which can be divided into
773 different phenotypes based on clinical evolution. Future studies are thus required to
774 explore subtype-specific cerebellar network alterations in MS and NMOSD. Finally,
775 from a methodological viewpoint, white matter fiber tractography should be adopted
776 in future studies which may provide additional insights beyond morphological and
777 functional connectivity into cerebellar network dysfunctions in MS and NMOSD.

778

779 **Competing interests**

780 The authors declare no competing interests.

781

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Table 1. Demographic, clinical and neuropsychological variables

	HCs (n = 228)	MS (n = 208)	NMO (n = 200)	p-value
Age (years)	37.0 (21.0)	36.0 (17.0)	41.0 (21.5)	0.003 ^{a,c}
Sex (female/male)	124/104	135/73	175/25	<0.001 ^{a,b,c}
Disease state (acute/chronic)*	-	48/149	48/147	0.954
Disease duration (months)	-	19.0 (54.0)	36.0 (52.0)	0.054
Lesion				
N (%)	-	208 (100%)	110 (55%)	<0.001
Volume (cm ³)	-	7.6 (16.0)	1.2 (4.0)	<0.001
EDSS	-	2.0 (2.5)	3.5 (3.0)	<0.001
CVLT [†]	52.0 (10.8)	47.5 (15.0)	47.0 (14.5)	<0.001 ^{b,c}
PASAT [†]	51.5 (14.0)	40.0 (17.0)	40.0 (18.0)	<0.001 ^{b,c}
BVMT [†]	28.0 (5.8)	27.0 (11.8)	22.0 (12.0)	0.003 ^c

Data are represented as median (interquartile range) unless stated otherwise. HC, healthy controls; MS, multiple sclerosis; NMO, neuromyelitis optica; EDSS, Expanded Disability Status Scale; CVLT, California Verbal Learning Test; PASAT, Paced Auditory Serial Addition Test; BVMT, Brief Visuospatial Memory Test.

*Data are missing for 16 patients.

†Data are available only for a subset of participants from three hospitals (see Materials and Methods for details).

^aSignificant differences between the two patient groups.

^bSignificant differences between the MS patients and HC.

^cSignificant differences between the NMO patients and HC.

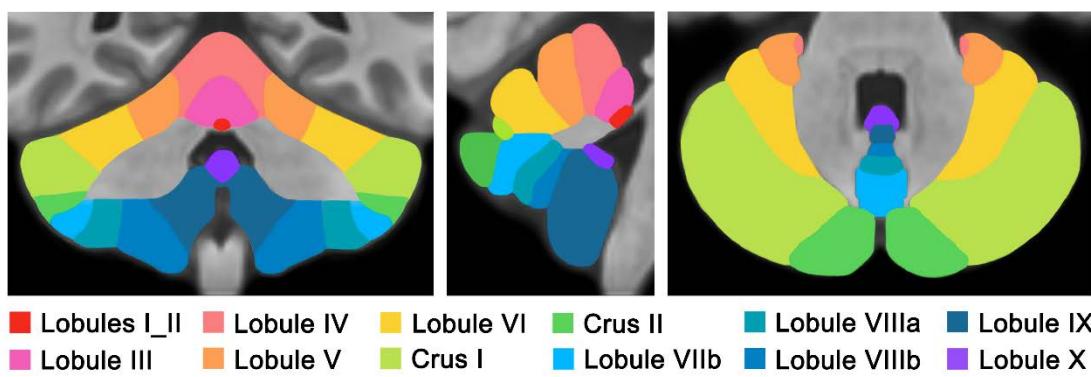


Figure 1. Illustrative representation of cerebellar lobules. According to the CERES partition, the cerebellum was segmented into 12 lobules in each hemisphere.

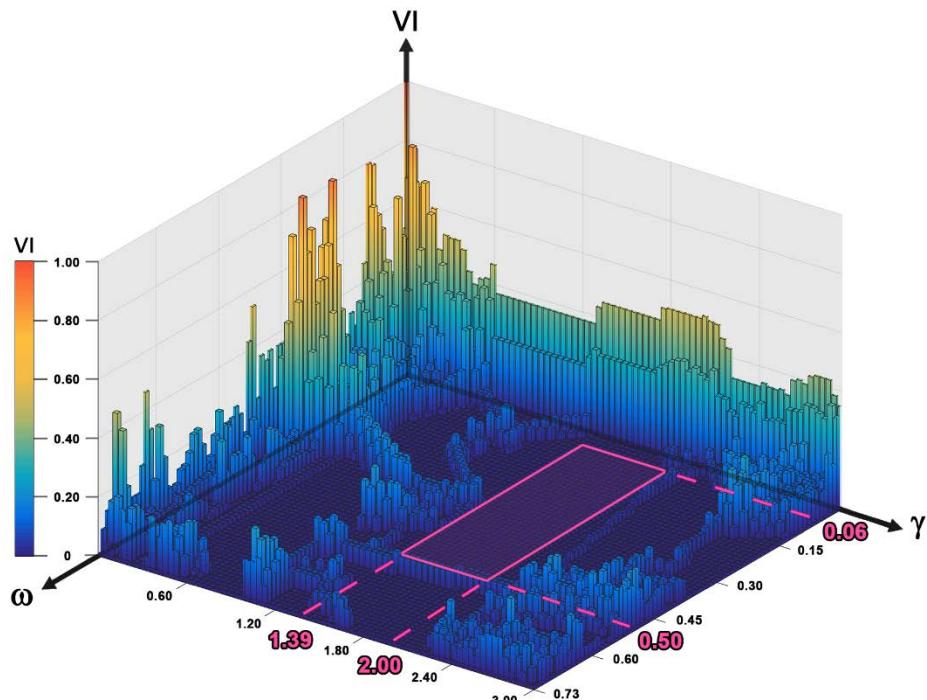


Figure 2. The stability of cerebellar modular architecture over different combinations of inter-layer connectivity strength (ω) and the module resolution (γ). The variation of information was used to evaluate the stability of cerebellar modular architecture. In the searched 2D parameter space ($\omega = [0.01 - 1]$; $\gamma = [0.01 - 3]$), a widest range ($\omega = 0.06 \sim 0.50$; $\gamma = 1.39 \sim 2.00$) was identified wherein cerebellar modular architecture maintained stable when the parameters fluctuated. VI, variation of information.

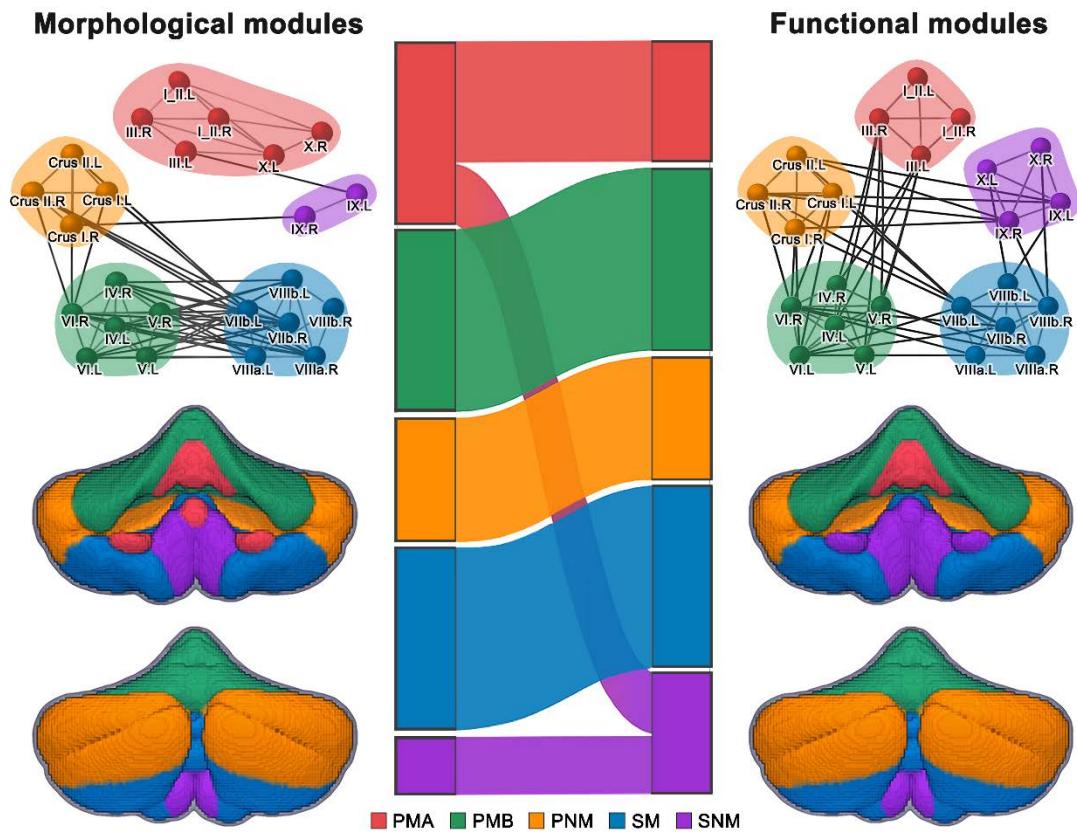


Figure 3. Modular architecture of the cerebellum. Five cerebellar modules were identified by applying a multilayer community detection algorithm to the group-level multiplex network of the HCs that integrated morphological and functional connectivity within the cerebellum. Module assignments of cerebellar lobules were largely comparable between morphological and functional networks. PMA, Primary Motor A; PMB, Primary Motor B; PNM, Primary Non-Motor; SM, Secondary Motor; SNM, Secondary Non-Motor.

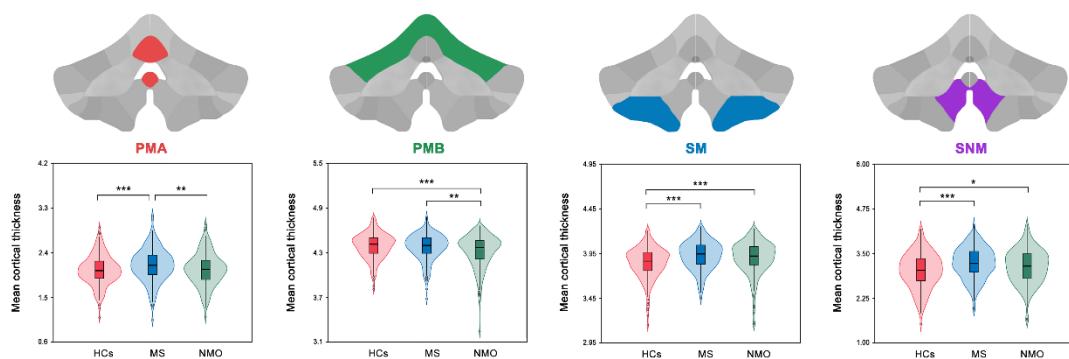


Figure 4. Alterations in cerebellar module-based cortical thickness. Significant group effects were found on mean cortical thickness within four cerebellar modules (PMA: MS-specific cortical thickening; PMB: NMO-specific cortical atrophy; SM and SNM: common cortical thickening to the two patient groups). PMA, Primary Motor A; PMB, Primary Motor B; SM, Secondary Motor; SNM, Secondary Non-Motor; *, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$.

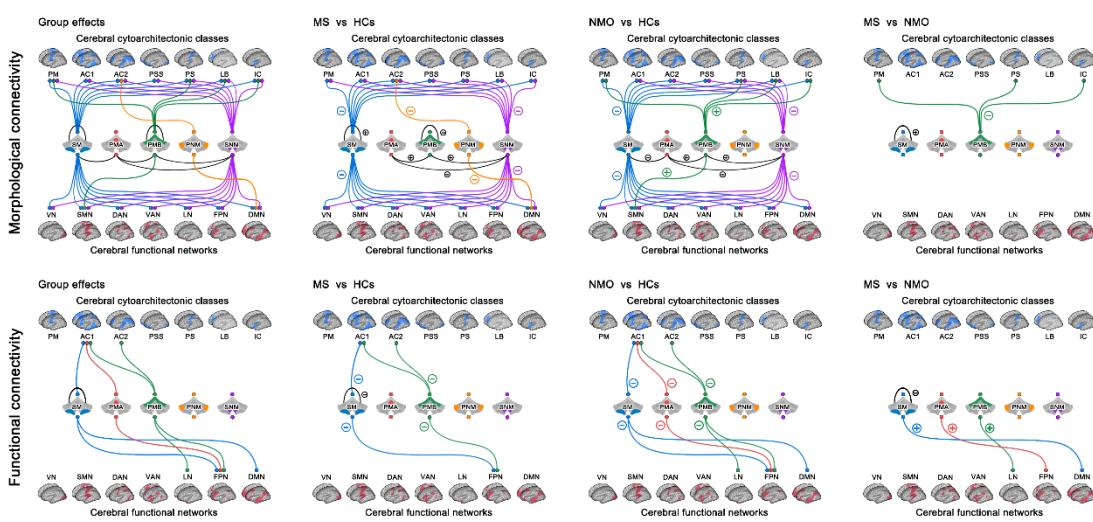


Figure 5. Alterations in cerebellar module-based morphological and functional connectivity. For morphological connectivity, numerous alterations of both increases and decreases were found in the two patient groups. Compared with morphological connectivity, fewer alterations were observed for functional connectivity in the two patient groups that were all characterized by disease-related decreases in particular for

connectivity between cerebellar motor modules and cerebral association cortex or high-order networks. PMA, Primary Motor A; PMB, Primary Motor B; PNM, Primary Non-Motor; SM, Secondary Motor; SNM, Secondary Non-Motor; PM, primary motor cortex; AC1, association cortex; AC2, association cortex; PSS, primary/secondary sensory; PS, primary sensory cortex; LB, limbic regions; IC, insular cortex; VN, visual network; SMN, somatomotor network; DAN, dorsal attention network; VAN, ventral attention network; LN, limbic network; FPN, fronto-parietal network; DMN, default mode network.

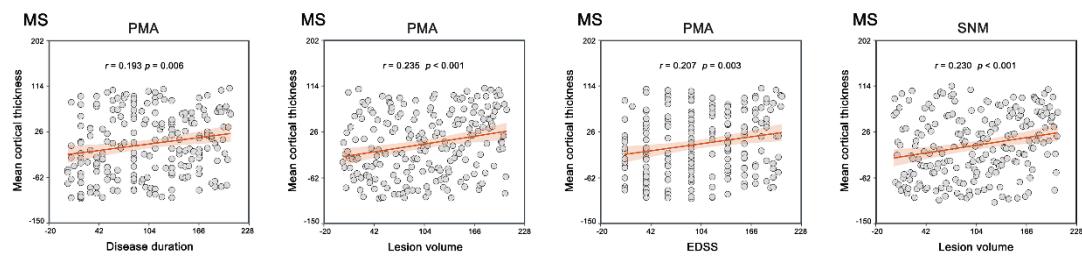


Figure 6. Relationships between cerebellar module-based cortical thickness and disease duration, lesion volume and EDSS in the MS patients. Among the patients, significantly positive correlations were found for mean cortical thickness within the PMA with disease duration, lesion volume and EDSS, and for mean cortical thickness within the SNM with lesion volume. MS, multiple sclerosis; PMA, Primary Motor A; SNM, Secondary Non-Motor; EDSS, expanded disability status scale.

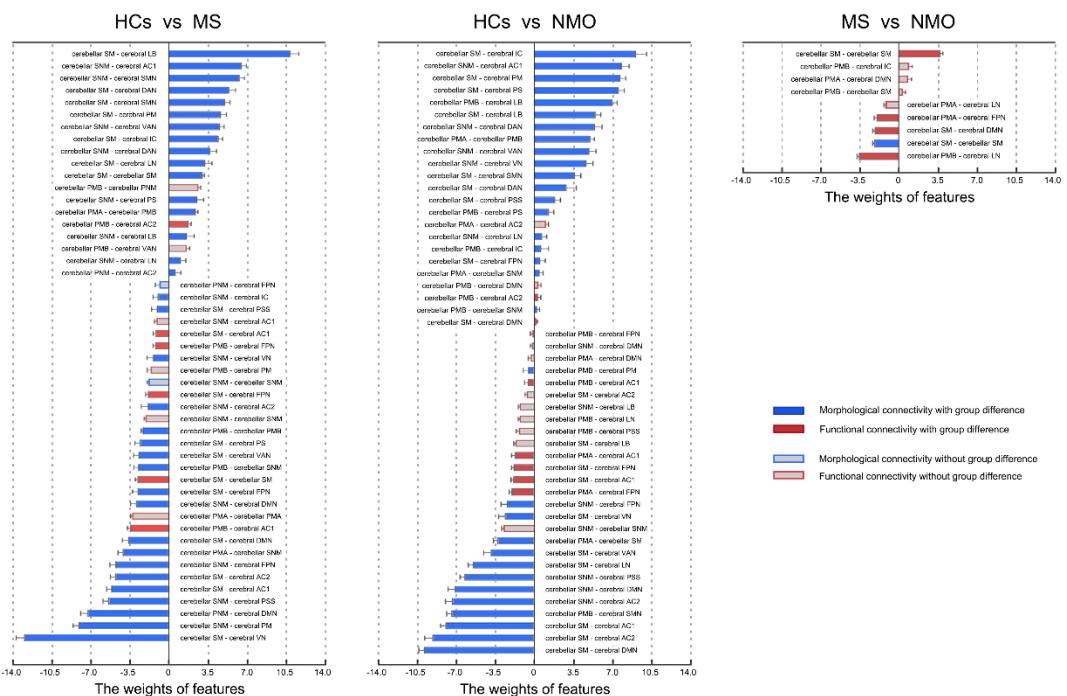


Figure 7. Features contributing to the classification between groups. The features contributing to the classification were mainly composed of connectivity that exhibited significant between-group differences. The classification between the patients and HCs mainly relied on morphological connectivity in particular those involving the PMB, SM and SNM, while the classification between the two diseases mainly benefited from functional connectivity that were all involved in the three motor-related modules (i.e., the PMA, PMB and SM). PMA, Primary Motor A; PMB, Primary Motor B; PNM, Primary Non-Motor; SM, Secondary Motor; SNM, Secondary Non-Motor; PM, primary motor cortex; AC1, association cortex; AC2, association cortex; PSS, primary/secondary sensory; PS, primary sensory cortex; LB, limbic regions; IC, insular cortex; VN, visual network; SMN, somatomotor network; DAN, dorsal attention network; VAN, ventral attention network; LN, limbic network; FPN, fronto-parietal network; DMN, default mode network.