1 No evidence for motor recovery-related cortical reorganization after

2

stroke using resting-state fMRI

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24 Abstract

25	Cortical reorganization has been suggested as mechanism for recovery after stroke. It has been
26	proposed that a form of cortical reorganization (changes in functional connectivity between brain
27	areas) can be assessed with resting-state fMRI. Here we report the largest longitudinal data-set
28	in terms of overall sessions in 19 patients with subcortical stroke and 11 controls. Patients were
29	imaged up to 5 times over one year. We found no evidence for post-stroke cortical reorganization
30	despite substantial behavioral recovery. These results could be construed as questioning the
31	value of resting-state imaging. Here we argue instead that they are consistent with other
32	emerging reasons to challenge the idea of motor recovery-related cortical reorganization post-
33	stroke when conceived as changes in connectivity between cortical areas.
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35	Keywords: stroke recovery, upper extremity impairment, resting state, cortical reorganization, functional
36	connectivity
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39	1. Introduction
40	Spontaneous neurological recovery occurs in almost all stroke patients within the first
41	months after the insult. While the underlying physiological changes that accompany spontaneous
42	motor recovery in humans remain largely unknown, data from animal models have been
43	interpreted as showing that cortical reorganization is a potential key mechanism mediating
44	recovery (Dancause & Nudo, 2011; Grefkes & Ward, 2014; Nudo, 2006).
45	In the literature, the term cortical reorganization has been loosely defined and used to

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47 can span the micro-, meso- and macro-scale, including synaptogenesis, axonal sprouting, and 48 changes in cortical activation maps. We have argued elsewhere that the term functional 49 reorganization should be reserved for those changes, including new cortico-cortical connections, 50 that are causally related to or at least correlated with motor recovery (Krakauer & Carmichael, 51 2017). It should be added that reorganization has also been taken as a qualitative event, 52 exemplified by the idea that one cortical area "takes over" another, which implies a change in the 53 tuning of neurons, for example, when touching the face activates the hand area of sensory cortex 54 in amputees. We argue elsewhere that a qualitative change in cortical representation need not be 55 invoked to explain this result (Krakauer & Carmichael, 2017), but we will not use this definition 56 here.

57 Evidence for functional reorganization after stroke comes primarily from studies of 58 axonal sprouting. For example, Overman and colleagues (2012), in a mouse cortical stroke 59 model, generated sprouting of axonal connections within ipsilesional motor, premotor and 60 prefrontal areas by blocking of an axonal growth inhibitor (epinephrine A5). Similar results were 61 reported for the neuronal growth factor GDF10 (Li et al., 2015). Critically, however, in both 62 studies no direct test of the relevance of axonal sprouting for motor improvement was performed, 63 indeed not even a correlation with the degree of sprouting and behaviour was examined. In 64 addition, most studies describing axonal sprouting after stroke found that it was cortico-65 subcortical instead of cortico-cortical connectivity changes that were linked to motor recovery 66 (see e.g. Lee, 2004; Wahl et al., 2014). Other studies that argue for a role of cortico-cortical 67 connectivity changes underlying stroke recovery are limited by cross-sectional approaches or do 68 not report behavior at all (Dancause et al., 2005; Frost et al., 2003; Liu & Rouiller, 1999; 69 Napieralski et al., 1996).

70 Despite the weak evidence for behaviorally-relevant new cortical connections in animal 71 models post-stroke, these models have nevertheless led to widespread interest in identifying 72 similar processes of functional reorganization in the human brain. One prominent non-invasive 73 method is to measure inter-regional connectivity with resting-state fMRI (rs-fMRI; Biswal et al., 74 1995; Fox & Raichle, 2007). This method relies on correlations between time-series of fMRI 75 activity recorded while the subject is lying in the scanner without performing a task. Most often 76 these correlations are computed between a set of pre-defined regions of interest (ROIs). The 77 underlying assumption is that regions with connected neuronal processing show stronger 78 statistical dependency of their spontaneous neuronal fluctuations. These correlations are 79 commonly regarded as a measure of "functional connectivity", which has been closely linked to 80 structural connectivity (Friston, 2011; van der Heuvel et al., 2009). In the context of stroke 81 recovery, it has been suggested that reorganization can be detected as a change in such 82 correlations/functional connectivity patterns (van Meer et al., 2010). Specifically, for post-stroke 83 recovery of hemiparesis, the advantage of task-free resting-state over task-based fMRI is that it 84 avoids the performance confound (Krakauer, 2004, 2007); the connectivity measures are not 85 biased by the inability of patients to match control performance due to motor impairment.

To date, results from rs-fMRI studies of functional connectivity changes after stroke have been mixed. Although, rs-fMRI studies have frequently found changes in interhemispheric connectivity patterns after stroke (Carter *et al.*, 2010; Chen & Schlaug, 2013; Golestani *et al.*, 2013), the direction of these changes and their correlations with behavior have been inconsistent. One study found a positive correlation between motor function and increased functional connectivity between the lesioned M1 and contralateral heterologous cortical areas (Park *et al.*, 2011), another study reported that interhemispheric homologous connectivity was associated

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with lower degrees of motor impairment but only for infratentorial strokes (Lee *et al.*, 2018). Yet
another study showed that an increase in M1-M1 connectivity correlated negatively with motor
function (Wang *et al.*, 2014).

96 There are many potential reasons for these inconsistencies in rs-fMRI findings.

97 If patients with cortical lesions are included in the study design, it is possible to confuse changes 98 in connectivity measures as a direct consequence of the lesion (e.g. the damaged area becomes 99 disconnected from the brain) with changes associated with true reorganization. Additionally, 100 most studies use different analysis protocols and measures to quantify changes in connectivity, 101 making integration of evidence across studies difficult. Third, the majority of currently available 102 studies have been cross-sectional but it is essential to evaluate changes in connectivity across the 103 time-course of recovery.

104 To address these issues, we here report the results of a longitudinal rs-FMRI study of 105 stroke recovery in patients with hemiparesis after subcortical stroke. Only patients with 106 subcortical lesion locations were included in this study so that any changes in cortical 107 connectivity could not be attributed to the presence of the lesion itself. We provide a detailed 108 characterization of inter- and intrahemispheric connectivity between five cortical motor areas. 109 Because of considerable variation of analysis approaches in the existing literature, in addition to 110 our primary analysis, we also compared results after using two different pre-processing 111 procedures, report results from an individual M1-M1 ROI analysis, and replicated the analysis 112 approach from the largest longitudinal resting-state stroke study published to date (Golestani et 113 al., 2013).

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115 **2. Results**

The main goal of this study was to determine whether motor impairment recovery following stroke was associated with systematic changes in cortical connectivity. Our two main questions were: 1) Is there a mean difference in the connectivity pattern between five motor regions (S1, M1, PMv, PMd, SMA) when comparing patients and age-matched controls at any time-point during stroke recovery? 2) Is there a change in patients' connectivity patterns over time that is related to motor impairment?

We analyzed data from 19 patients with subcortical stroke and 11 healthy controls. Behavioral assessments and resting-state images were obtained at five different time-points over one year. Each patient completed on average 4.5 ± 0.7 sessions, with the overall experimental data being 89.5% complete (see also Table S1 for demographics and completed sessions in the supplemental material). We begin by quantifying the extent of impairment and recovery of upper extremity deficits in our patients in the year following stroke.

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129 2.1. Patients showed substantial clinical recovery after stroke

We measured initial impairment and subsequent recovery of the upper extremity using the upper
extremity portion of the Fugl-Meyer score (FM-UE), the Action Research Arm Test (ARAT),
and hand strength (Xu *et al.*, 2017).

At the acute stage, all behavioral measures indicated impairment of the upper extremity for patients relative to controls (FM-UE: t(28)=3.706, p=0.001, ARAT: t(28)=2.315, p=0.028, strength: t(28)=5.195, p<0.001, Figure 1). These deficits recovered substantially over the course of one year, with the largest changes observed within the first three months (Week effect for FM-UE: χ^2 =24.865, p<0.001; ARAT: χ^2 =13.942 p=0.007; hand strength: χ^2 =13.419, p=0.009). No significant changes were observed in controls for any of the three measures.

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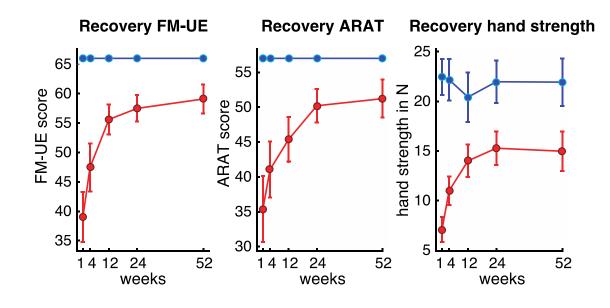


Figure 1: Recovery of upper extremity deficits after stroke over one year. For all behavioral assessments, the largest changes in recovery were seen within the first three months. Patients reached a plateau at 6 months and, on average, remained impaired compared to controls at all time-points. Note that patients had moderate to severe upper extremity impairment in the acute stage. Red lines = patients, blue lines = controls, FM-UE = Fugl-Meyer score Upper Extremity, ARAT = Arm Research Action Test.

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148 2.2. Connectivity patterns across sensorimotor areas were reliable and stable in controls

149 Next, we looked at changes in connectivity patterns (pattern of ROI-ROI connectivity 150 weights) between five key sensorimotor areas to determine if and how connectivity between 151 these sensorimotor areas changed alongside behavior during recovery. To determine the 152 connectivity patterns, we calculated pairwise correlations between the averaged time-series of 153 BOLD activities between all possible ROI pairs to get a 10×10 matrix of connectivity weights 154 (see Methods). An average connectivity pattern for patients and controls is shown in Figure 2a.

155 Connectivity patterns were highly reliable for both groups with high intrasession 156 reliabilities (*all connections*, controls: R=0.66, CI 0.62–0.71, patients: R=0.70, CI 0.66–0.74; see 157 supplementary material for inter- and intrahemispheric connections, Figure S2). An unbalanced

158 mixed-effects ANOVA (see Methods) showed that the intrasession reliability was not significantly different between groups ($\chi^2(1)=1.0782$, p=0.2991) and showed no changes over 159 time (controls: $\chi^2(4)=6.174$, p=0.187; patients: $\chi^2(4)=1.922$, p=0.75). 160 161 Furthermore, connectivity patterns for controls were stable, showing no significant 162 change over time (all connections: Δweek acute W4=0.841, confidence interval (CI) 0.597-163 0.687-2.821; 2.727; acute_W12=0.689, CI 0.582-2.412; acute_W24=1.079, CI 164 acute_W52=1.059, CI 0.611-2.531). Thus, for all subsequent analyses connectivity patterns for

165 controls were averaged over time-points.

166 We also confirmed that the connectivity pattern for controls reflected known anatomical 167 connectivity (Damoiseaux & Greicius, 2009). Within one hemisphere, the highest correlations 168 were found between S1-M1 (0.91 \pm 0.47, Fisher-Z transformed), while the weakest correlation 169 was found between M1-PMv (0.58 \pm 0.39). Between hemispheres, S1_{right}-S1_{left} demonstrated the 170 highest correlation (0.9 \pm 0.43), while M1_{right}-PmV_{left} showed a weaker correlation (0.59 \pm 0.37). 171 For correlations between hemispheres, homologous ROIs (e.g. M1-M1 or S1-S1) showed higher 172 correlations of the BOLD time series compared to heterologous ROI-ROI connectivity weights (e.g. M1_{right}-Pmv_{left} or S1_{left}-Pmd_{right}) as expected from interhemispheric neural-recordings 173 174 (Asanuma & Okamoto, 1959, see supplemental results and Figure S3 for comparison of 175 homologous versus heterologous interhemispheric connectivity).

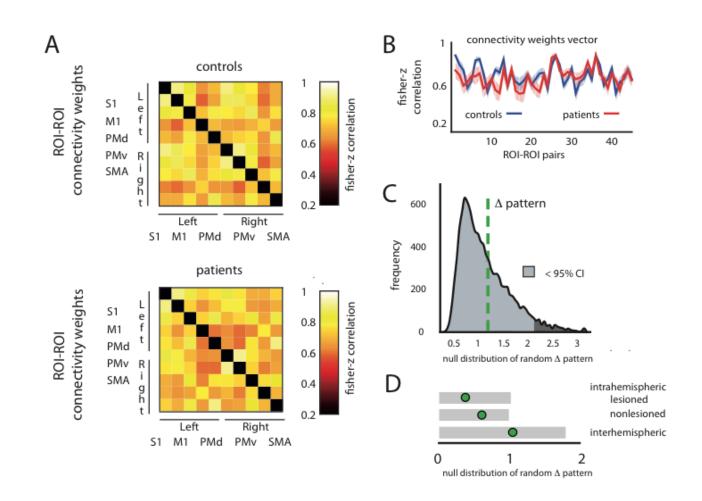
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177 2.3. There were no systematic changes in connectivity patterns in the acute recovery 178 period

179 If disruption of the cortical projections through subcortical stroke leads to an acute 180 reorganization of cortical circuits, one would expect that (on average) acute connectivity patterns

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181	of patients and controls would be different. Connectivity patterns for patients and controls were
182	highly correlated in the early period after stroke (acute stage: R=0.69, p=0.0002; see connectivity
183	matrices in Figure 2A and also Figure S4). To statistically test for significant differences
184	between connectivity patterns, we used the Euclidian distance between the two groups' mean
185	patterns and compared it to a null-distribution obtained by a permutation test (Figure 2c). We
186	found no systematic difference between patients and controls at the acute stage (Δ pattern=1.246,
187	CI 0.575-2.467). This was also true when only considering intrahemispheric connections of
188	either the lesioned (Δ pattern=0.367, CI 0.205–1.109) or non-lesioned side (Δ pattern=0.603, CI
189	0.196–1.1) or interhemispheric connections (Δ pattern=1.027, CI 0.394–1.968).
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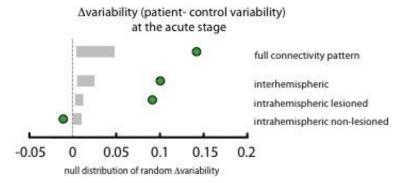
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192 Figure 2: No systematic differences in connectivity patterns of patients and controls in the acute 193 recovery period (week 1-2). (A) Heat map representation of average connectivity weights for 194 controls and patients at the acute stage after stroke. The y- and x-axis show the five ROIs (S1, 195 M1, Pmd, PMv, SMA) for the left and right hemisphere creating a connectivity matrix. One small square represents the connectivity weight for the respective ROI pairing. The diagonal (black) is 196 197 missing, as it is the correlation of a ROI with itself. (B) Vectorized upper triangular part of the 198 correlation matrix for the average full connectivity pattern of controls (blue line) and patients 199 (red line). (C) To quantify the differences between connectivity patterns for controls and 200 patients, we calculated the Euclidian distance between the two pattern vectors (*Apattern*, dashed 201 green line). The Euclidian distance is sensitive to differences of shape and scaling of patterns. 202 The measured distance was then tested against the expected distribution if there were no 203 differences between the two groups. To generate an empirical estimate of this distribution, we 204 randomly shuffled group assignments and repeatedly computed the Euclidean distance (x10.000 205 times, histogram with frequency on the y-axis and absolute value of the Euclidian distance on the 206 x-axis). For the acute stage after stroke, the Δ pattern lay within the lower 95% percentile (grey 207 shaded area) of the null-distribution. (D) The measured $\Delta patterns$ (green circle) for the 208 intrahemispheric lesioned, non-lesioned or interhemispheric ROIs also always fell within the 209 lower 95% range (grey boxes).

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211 Even though the averaged connectivity patterns for patients and controls were 212 indistinguishable at the acute stage, the heterogeneity in lesion locations for different patients 213 might result in idiosyncratic shifts in connectivity patterns that in the whole group would be 214 reflected as higher variability in patterns. To measure this within-group variability, we calculated 215 the average Euclidian distance of each patient's pattern to the patient group mean pattern and did 216 likewise for controls. The average within-patient distance was 2.955, whereas the average 217 within-control distance was 2.813, resulting in a difference of 0.142 (Δ variability). We compared 218 this value to a null distribution of Δ variability generated with permutation testing. We found that 219 resting-state connectivity patterns of patients showed a higher idiosyncratic, non-systematic 220 variability compared to controls: The difference between the variability lay outside the 2.5% – 221 97.5% confidence interval generated by permutation testing (CI 0.018–0.051, Figure 3). Note 222 that the confidence interval was not symmetric around zero, as the N for controls was smaller 223 than for patients.

The difference in variability for intrahemispheric lesioned and interhemispheric connections was also higher for patients. For intrahemispheric non-lesioned connections, we found higher variability in controls (intrahemispheric lesioned: Δ variability=0.091, CI 0.002– 0.023; non-lesioned: Δ variability=-0.01, CI -0.003–0.015; interhemispheric: Δ variability=0.1, CI 0.008–0.05, Figure 3).



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Figure 3: Patients showed a higher unsystematic variability compared to controls at the acute stage (Δ variability = green circle, 2.5%-97.5% range = grey boxes). Only for intrahemispheric non-lesioned ROI's patients showed a lower variability.

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Thus, overall, while connectivity patterns for patients were more variable, patient connectivity patterns were indistinguishable from control patterns at the acute stage.

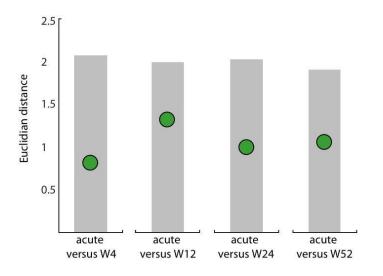
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237 2.4. There were no changes in patients' connectivity patterns over time

Even though there were no systematic differences between connectivity patterns of patients and controls at the acute stage, we might expect to find changes in patient connectivity patterns over time as they recover from impairment.

We therefore quantified Euclidean distances between the average connectivity patterns at the acute stage as reference versus all other weeks (Δ week). Surprisingly, patients showed no increase in Euclidian distances between the acute stage and consecutive weeks (Figure 4 and Table 1).

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Figure 4: No significant change from patients' acute connectivity pattern compared to timepoints at the subacute or chronic stage.

248 We computed the Euclidian distance between the average connectivity pattern of patients at the

249 acute stage and all consecutive weeks (W4, W12, W24, W52; Δ week = green circles). Range of

the expected distribution if there were no differences between the two groups (grey shaded area).

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Patients	acute_W4	acute_W12	acute_W24	acute_W52
All connections	∆week: 0.814,	Δweek: 1.322,	∆week: 0.994,	∆week: 1. 063,
	(0.467 - 2.066)	(0.488 - 1.994)	(0.471 - 2.024)	(0.512 - 1.898)
Interhemispheric	∆week: 0.636	∆week: 1.018	∆week: 0.665	Δ week: 0.77
internetinspherie	(0.309 - 1.695)	(0.3255 - 1.627)	(0.315 - 1.589)	(0.343 - 1.616)
Intrahemispheric	∆week: 0.353	∆week: 0.413	∆week: 0.457	∆week: 0.558
lesioned	(0.241 - 1.246)	(0.237 - 1.169)	(0.252 - 1.324)	(0.274 - 1.158)
Intrahemispheric non-	∆week: 0.367	∆week: 0.735	∆week: 0.579	∆week: 0.474
lesioned	(0.152 - 0.859)	(0.157 - 0.886)	(0.158 - 0.919)	(0.174 - 0.805)

253

Table 1: Euclidian distances between the connectivity pattern of the acute stage compared to all
subsequent time-points in patients for only interhemispheric, intrahemispheric lesioned, or nonlesioned subsets.

As it could be expected from these results, patients showed reliably high correlations of their connectivity patterns with controls at the subacute or chronic stage (W4: R=0.74, p<0.0001; W12: R=0.76, p<0.0001; W24: R=0.87, p<0.0001; W52: R=0.80, p<0.0001) and no significant

difference to control patterns 9 (Table 2). The analyses for intra- or interhemispheric connections 261

262 alone found the same result (Table 1 & 2 and Figure S4).

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patients versus controls'	W4	W12	W24	W52
All connections	∆pattern: 1.203 (0.53 - 2.482)	Δpattern: 1.795 (0.57 - 2.208)	Δpattern: 0.885 (0.563 - 2.249)	∆pattern: 1.653 (0.575 - 2.07)
Interhemispheric				
	Δ pattern: 1.102	Δ pattern: 1.671	Δ pattern: 0.663	∆pattern: 1.354
	(0.366 - 1.986)	(0.387 - 1.797)	(0.387 - 1.755)	(0.402 - 1.706)
Intrahemispheric				
-	Δ pattern: 0.412	Δ pattern: 0.44	Δ pattern: 0.404	∆pattern: 0.802
lesioned	(0.2 - 1.202)	(0.198 - 0.998)	(0.213 - 1.122)	(0.216 - 0.892)
Intrahemispheric non-				
_	∆pattern: 0.253	Δ pattern: 0.486	Δ pattern: 0.415	∆pattern: 0.505
lesioned	(0.184 - 1.097)	(0.192 - 0.955)	(0.201 - 1.039)	(0.206 - 0.822)

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Table 2: Difference between the connectivity pattern of patients compared to controls at Week 4, 265

266 Week 12, Week 24, and Week 52 for only interhemispheric, intrahemispheric lesioned, or non-267 lesioned subsets.

patients	acute_W4	acute_W12	acute_W24	acute_W52	
		F(3,36) = 0.0	9, p = 0.9678		
All connections	Δ week_variability:	Δ week_variability:	Δ week_variability:	∆week_variability	
	2.474 ± 1.6	2.607 ± 1.022	2.424 ± 0.96	2.642 ± 0.778	
	F(3,36) = 0.15, p = 0.9276				
Interhemispheric	Δ week_variability:	Δ week_variability:	Δ week_variability:	∆week_variability	
	1.9 ± 1.4	2.058 ± 0.782	1.78 ±0.697	1.932 ± 0.61	
Intrahemispheric		F(3,36) = 0.2	25, p = 0.859		
lesioned	∆week_variability:	∆week_variability:	∆week_variability:	∆week_variability	

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Intrahemispheric non- lesioned $F(3,36) = 0.32, p = 0.814$ Δ week_variability: Δ week_variability: Δ week_variability:	
Instrumentspace non- Aweek_variability: Aweek_varis: Autrek Autrek Aut	228 ±0.461
1.08 ±0.688 1.133 ±0.547 1.083 ±0.518 1.2 268 By examining Euclidian distances between the individual connectivity patterns 1.2 269 average connectivity pattern, we found a greater non-systematic variability in patients the controls at the acute stage. However, the idiosyncratic variability of patients themselves of change from the acute stage compared to the following time-points (Table 3). 271 <i>Table 3: Difference in connectivity pattern variability in patients over time for all connectivity pattern variability in patients over time for all connectivity interhemispheric, intrahemispheric lesioned, or non-lesioned subsets.</i> 272 Table 3: Difference in connectivity pattern variability in patients over time for all connectivity patterns within one year. More importantly, patients did not show any significant longity change in connectivity patterns either systematically or regarding their group variability. 273 Comparison between alternative metrics for M1-M1 connectivity 280 within and across hemispheres and found no changes for patients either longitudinally or compared to controls. In contrast, some previous studies have focused on individual ROI-to connections and have reported changes after stroke (Thiel & Vahdat, 2015). Specific changes in interhemispheric connectivity between the two motor cortices have been freq reported (Carter <i>et al.</i> , 2010; Chen & Schlaug, 2013; Golestani <i>et al.</i> , 2013; Park <i>et al.</i> , 2013; Park <i>et al.</i> , 2014; Park <i>et al.</i> , 2014; Park <i>et al.</i> , 2015; Park <i>et al.</i> , 2014; Park et al., 2014; P	
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weights over time and between patients and controls in our data set. The analysis sho	ctivity
	wed a
significant difference between patients and controls, with patients having a slightly	lower

average correlation between motor cortices (Figure 5a; *mixed model, group effect:* $\chi^2(1)=5.759$, p=0.016). Congruent with our other results, however, we found no longitudinal changes either for patients (*patient_week:* $\chi^2(4)=5.836$, p=0.212) or controls (*control_week:* $\chi^2(4)=0.4.723$, p=0.317).

Our results also contrast with another published finding that used an alternative metric of connectivity to assess changes in functional connectivity after stroke. Golestani and colleagues (2013) used a relative connectivity (RelCon, see Methods) measure between the two sensorymotor cortices and reported lower relative interhemispheric sensorimotor (SM1 RelCon) connectivity in stroke patients with a motor deficit compared to controls and stroke patients without impaired motor function.

Similarly, our patients had a lower RelCon for SM1-SM1 compared to controls at all timepoints. Using a mixed-model, we found a significant difference between the groups $(\chi^2(1)=5.2457, p=0.022)$. However, consistent with our results reported above, we did not find a change over time for RelCon SM1-SM1 in neither controls ($\chi^2(4)=2.8087, p=0.5903$) nor in patients ($\chi^2(4)=8.2243, p=0.0837$; Figure 5b).

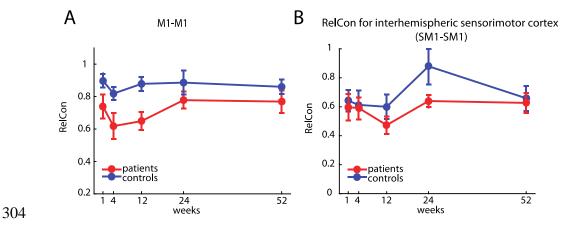


Figure 5: A) M1-M1 connectivity in our dataset. In patients, interhemispheric connectivity
between the two motor cortices was systematically lower than compared to controls at all timepoints. However, no changes of M1-M1 connectivity over time were found. B) Relative
Connectivity of SM1-SM1 in controls and patients. While there was a significant difference in

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309 *SM1-SM1 connectivity between the two groups, with lower RelCon for patients, there was no significant change over time.*

311

312 **3 Discussion**

Here we report, that there were no longitudinal changes in resting-state functional connectivity (rsFC) between cortical motor areas despite substantial motor recovery over the same time period in a cohort of patients with subcortical stroke. In addition, at no stage of recovery were rsFC patterns different from healthy, age-matched controls.

317 Whenever results are negative, concerns will be raised about the power of the study (addressed

below) and the biological validity of the method in general.

319 There have been more than 500 rs-fMRI studies of brain connectivity (Buckner et al., 2013). 320 Recent reports have described the close relationship between resting-state networks and 321 structural connectivity assessed with other methods e.g. magnetoencephalography (van den 322 Heuvel et al., 2009; Brooks et al., 2011). Most notably for our purposes, the sensitivity of rsFC 323 to changes in experience-dependent neural plasticity appears to be quite high, as even short 324 periods of training yield statistically significant changes of functional connectivity in small n325 studies in healthy subjects (Mawase et al., 2017; Vahdat et al., 2011). For example, Censor and 326 colleagues (Censor et al., 2014), in a comprehensive multimodal approach combining 327 behavioral, brain stimulation, and rs-fMRI data, they demonstrated that changes in performance 328 after training on a five-digit sequence task led to reliable changes in corticostriatal functional 329 connectivity. When motor memory formation after training was disrupted using rTMS, changes 330 in functional connectivity predicted the modification of memory recall on the next day.

331 Given such results, why were we not able to detect rsFC changes in the setting of stroke 332 recovery? Injury ostensibly triggers functional reorganization, which arguably should be a more

dramatic cause of connectivity change as it is associated with structural alterations, e.g. sprouting, and not just learning-related changes in pre-existing connections. There are two potential answers to this question, one is the possibility that the idea that changes in corticocortical connections promote motor recovery after stroke is ill-conceived, the second is that there are methodological limitations to rs-fMRI. We shall discuss both of these concerns.

338

339 A large number of animal studies, in rodents and non-human primates, have described numerous 340 structural and physiological changes in cortical areas around and beyond the infarct core. These 341 changes have collectively been called reorganization, but in only a small subset of cases have 342 they been correlated with motor recovery, which suggests that most are likely just reactive 343 (Carmichael, 2016). We reasoned that as spontaneous biological recovery is similar for cortical 344 and subcortical strokes (Zarahn et al., 2011) then recovery-related cortical reorganization, if not 345 just reactive, should still occur in patients with isolated subcortical lesions. Indeed, we know that 346 corticospinal integrity assessed with TMS is a good predictor of recovery in patients with 347 subcortical stroke (Radlisnka et al., 2010; Byblow et al. 2015), i.e., cortical output is required for 348 recovery from subcortical stroke just like it is for cortical stroke. In addition, changes in cortical 349 maps are seen not just with cortical lesions but with spinal and peripheral lesions as well 350 (Florence et al., 1998; Moxon et al., 2014, Krakauer & Carmichael, 2017). Here, however, we 351 found no evidence for systematic rsFC changes between cortical motor regions. In the light of 352 these results, previously reported cortical connectivity changes could be reactive rather than 353 reparative, e.g. confounded by the presence of a cortical lesion.

354

355 The question must now be asked why it was ever conjectured that changes in connections

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356 between cortical regions would enhance recovery from hemiparesis, which is caused by 357 interruption of descending pathways out of a particular region(s). One could rephrase this to ask 358 why would there be a "horizontal" solution to a "vertical" problem? This question is related to 359 the increasing awareness of the questionable relevance of cortical map changes to recovery 360 (Krakauer & Carmichael 2017), changes which have hitherto been taken as electrophysiological 361 evidence for reorganization (Dancause & Nudo, 2011; Warraich & Kleim, 2010; Wittenberg, 362 2010). Overall, it is increasingly apparent both from recent and previous work in non-human 363 primates and rodents that recovery after stroke relates to changes in the strengths of descending 364 projections to the brainstem and spinal cord from individual motor cortical areas rather than to 365 changes in the connections between them (Lin et al., 2018; Starkey et al., 2012; Wahl et al., 366 2014; Zaaimi et al., 2012). That said, it could be postulated that cortico-cortical drive, for 367 example of premotor cortex onto primary motor cortex (M1) could facilitate remaining CST 368 descending projections out of M1, as studies have shown such cortico-cortical facilitation in 369 healthy non-human primates (Cerri et al., 2003; Shimazu et al., 2004). Consistent with what we 370 found here, however, there is little evidence for this as a recovery mechanism after stroke in any animal. 371

372

While our results are congruent with similar observations in a smaller cohort (Nijboer *et al.*, 2017), they are seemingly contradicted by a recently published paper that reported results for resting-state changes in a similarly sized cohort of patients with subcortical stroke. In this study, Lee and colleagues obtained six connectivity measures between 40 supra- and infratentorial ROIs in 21 stroke patients measured at two time-points post-stroke (2 weeks and 3 months), and found differences in two of the measures. Specifically, they found lower overall strength in

interhemispheric connectivity and higher network distance compared to healthy controls at 2 weeks, but neither measure changed at 3 months. Even if one overlooks the unmentioned comparisons problem and the fact that they had more variables (six measures, 40 ROIs) than subjects, their results showed no connectivity measure *changing* as the patients improved, which is consistent with our results.

384

Although we favor the view that the absence of connectivity change in our study is a true negative result both in terms of the power of the study and the biological validity of rsFC (van Meer *et al.*, 2010), an alternative explanation would relate to methodological limitations of rsfMRI.

389 Methodological problems with e.g. regard to reproducibility of imaging analysis in general and 390 rs-fMRI, in particular, have long been a topic of discussion (Baker, 2016; Ioannidis et al., 2014; 391 Macleod et al., 2014). So far, there is no consensus about the optimal way to analyze rs-fMRI 392 data, which poses a fundamental challenge regarding the generalizability and comparability of 393 reported findings. In face of a low signal to noise ratio, missing consensus in analysis steps and 394 statistical methods (promoting the risk of conscious and unconscious p-hacking; Nuzzo, 2015), 395 and frequent absence of an *a priori* hypothesis (which can lead to so-called HARKing; Kerr, 396 1998), the imaging literature is especially vulnerable to false-positive or -negative results 397 (Munafò et al., 2017). For example, converging evidence highlights that the choice of different 398 pre-processing strategies needs to be considered as an important confound in rs-fMRI (Cole et 399 al., 2010; He & Liu, 2012; Weissenbacher et al., 2009).

400

401 We addressed this problem by providing measures of data reliability, comparing two different

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402 pre-processing procedures, and by reanalyzing our data set with regard to individual M1-M1 403 changes using a previously reported metric for resting-state imaging analysis (Golestani et al., 404 2013). Here we provide, to the best of our knowledge, the most methodologically complete study 405 to date of stroke recovery using rs-fMRI. Additionally, open science efforts including data 406 sharing have been identified as a major tool to secure transparency and reproducibility of 407 reported results, allowing for external validation of results, detection of mistakes, and generation 408 of alternative interpretations (Nosek et al., 2015). In an effort to increase the transparency and 409 reproducibility of our results, the complete data set as well as the custom-written MATLAB and 410 R scripts are made publicly available to invite further analysis.

411

412 Conclusion

413 In the present study, we investigated longitudinal changes in functional connectivity after 414 subcortical stroke. Despite substantial recovery from motor impairment over one year, we found 415 no differences in functional connectivity between patients and controls, nor any changes over 416 time. Assuming that rs-fMRI is an adequate method to capture connectivity changes between 417 cortical regions after brain injury, the results presented here, provide reason to doubt that post-418 stroke cortical reorganization, conceived as changes in cortico-cortical connectivity, is the 419 relevant mechanism for promoting motor recovery after stroke. We suggest instead that it is 420 facilitation of residual cortical descending pathways that are likely to be more causally relevant. 421 It is perhaps time for the field to change its emphasis from changes in "horizontal" connections 422 to changes in "vertical" ones.

423 4 Materials and methods

The resting-state data set presented here was acquired from a natural history study investigating upper extremity recovery after stroke (Study of Motor Acute Recovery Time course after Stroke; SMARTS). As part of the study, a range of behavioral, physiological, and imaging measurements were obtained. Details of the behavioral characterization of the patients have been published elsewhere (Cortés *et al.*, 2017; Ejaz *et al.*, 2018; Xu *et al.*, 2017).

429

430 **4.1. Patients**

431 Since we were interested in cortical connectivity changes after stroke, in order to avoid 432 confounding results due to cortical damage, only a subset of 19 patients with lesions restricted to 433 subcortical areas was considered (6 females; mean age 59 ±12 years, 15 right-handed). Major 434 inclusion criteria were: first-ever clinical apparent ischemic stroke, proven by a positive DWI 435 lesion within the previous 2 weeks; unilateral upper extremity weakness (Medical Research 436 Council muscle weakness scale <5); ability to give informed consent. Patients were excluded for 437 one or more of the following reasons: initial impairment too mild (Fugl-Meyer score Upper 438 Extremity >63/66), age ≤ 21 years, hemorrhagic stroke (Xu *et al.*, 2017). The selected patients 439 had lesions either in the corona radiata, the internal capsule or in the cortico-spinal tract above 440 the crossing in the pyramid. Demographics are described in Table S1; more detailed information 441 about lesion distribution is shown in Figure S1.

442 Additionally, 11 healthy age-matched control participants (4 females; mean age 65 ± 8 443 years; all right-handed), were tested at the same time-points.

444 The study was carried out in accordance with the Declaration of Helsinki and approved 445 by the respective local ethics committee of the participating recruiting centers of SMARTS

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446 (Johns Hopkins University, USA, Columbia University, USA, University Hospital Zurich,
447 Switzerland). All participants gave written informed consent.

448

449 **4.2.** Study design

450 Patients were enrolled in the study within the first two weeks after stroke and followed up 451 over a one-year period at five time-points: acute stage: week 1-2 (10 ± 4 days), W4: week 4-6 (37) 452 ± 8 days), W12: week 12-14 (95 ± 10 days), W24: week 24-26 (187 ± 12 days), and W52: week 453 52-54 (370 ± 9 days). During each visit, the following clinical parameters were assessed: Fugl-454 Meyer score Upper Extremity (FM-UE, max. score 66, Fugl-Meyer et al., 1975), Action 455 Research Arm Test (ARAT, max. score 57, Yozbatiran et al., 2008). Hand strength and 456 individuation ability were measured using a custom-made hand-device (Xu et al., 2017). The 457 FM-UE and ARAT are widely used to assess motor deficits after stroke and can capture different 458 aspects of recovery: higher FM-UE scores represent normal reflex activity, fewer muscular 459 coactivations, coordination and higher joint mobility thought to be equal to "true" resolution of impairment; higher ARAT scores are achievable with compensatory strategies, thus correlating 460 461 closer with activities of daily living. Measuring hand strength offers a third dimension of 462 recovery that is only partially captured within the FM-UE and ARAT.

463

464 **4.3.** Image Acquisition

Participants were scanned with an 3T Achieva Philips system. Scans were obtained with a 32-channel head coil, using a two-dimensional echo-planar imaging sequence (TR=2.00s, 35 slices, 210 volumes/run, slice thickness 3mm, 1mm gap, in-plane resolution 3×3 mm²). Each

468	resting-state scan was 8min long. Participants were instructed to lie still and visually fixate on a
469	central white cross displayed on a computer monitor.

470 Structural images for atlas transformation and lesion definition were acquired with a T1-471 weighted anatomical scan (3D MPRAGE sequence, TR/TE=8/3.8ms, FOV 212×212mm, matrix 472 96×96, 60 slices, slice thickness 2.2mm). Finally, for each participant, a diffusion weighted 473 imaging (DWI) image (TR=2.89s, 30 slices, 5mm slice thickness, 240x240mm FOV), was 474 acquired to define lesion boundaries.

475

476 4.4. Imaging analysis

477 4.4.1. Preprocessing of rs-fMRI time series

478 Rs-fMRI has a relatively low signal-to-noise ratio. Non-neuronal processes, such as sensor noise, 479 head motion, cardiac phase, and breathing, account for a considerable part of the variance of the 480 raw signal (Birn, 2012). It has been argued that markers for the reliability of the sampled rs-481 fMRI data are missing and that the choice of preprocessing steps is often not justified (Bennett & 482 Miller, 2010; Zuo & Xing, 2014). We therefore conducted two different procedures for noise 483 reduction and then compared split-half reliability for the whole connectivity pattern in controls to 484 determine which steps provided higher reliability (see supplementary material).

485 4.4.2. Lesion definition

486 Lesion boundaries were defined as an intensity increase of ≥30% on DWI images, and in a
487 second step manually modified by a neuroradiologist and a neurologist using RoiEditor, see
488 Figure S1 for averaged lesion distribution map.

489

490 4.4.3. ROI definition

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491 We chose five motor areas (S1=primary somatosensory cortex, M1=primary motor cortex, 492 PMd=dorsal premotor cortex, PMv=ventral premotor cortex, SMA=supplementary motor area) 493 as regions of interest that have been widely accepted as being associated with motor function and 494 motor recovery (Miyai et al., 1999, 2002; Rehme et al., 2012). Individual T1-images were used 495 to delineate pial-grey matter and grey matter-white matter boundaries using FreeSurfer software 496 (Dale et al., 1999). The cortical surfaces were aligned across participants based on the sulcal-497 depth and local curvature maps. Probabilistic cyto-architectonic maps (Fischl et al., 2008) 498 aligned to the group average surface were then used to define ROIs first on the individual 499 surface, and then back-projected into the subject-native space.

500 The ROIs were defined as follows, M1: surface nodes with the highest probability for 501 Brodmann area (BA) 4. To increase specificity for processes related to recovery of hand 502 function, this ROI was limited to 2cm above and below the hand-knob (Yousry, 1997). S1: nodes 503 in the hand-region in S1 were isolated using BA 3a, 3b, 1 and 2.2cm above and below the hand 504 knob. PMd: nodes with highest probability in BA6, above middle frontal sulcus, but on the 505 lateral surface of the hemisphere. PMv: nodes with the highest probability in BA6, above middle 506 frontal sulcus. SMA: nodes with the highest probability in BA6 on the medial surface of the 507 brain. This ROI therefore includes SMA and preSMA (Picard & Strick, 1996).

508

509 **4.5.** Functional connectivity analysis

510 For each ROI, the time series for all voxels within the ROI were extracted and averaged, 511 resulting in a single BOLD time-course vector for each of the 10 ROIs across the two 512 hemispheres (left-S1, left-M1, left-PMd, left-PMv, left-SMA, right-S1, right-M1, right-PMd, 513 right-PMv, right-SMA). Pairwise correlations between averaged BOLD time-course vectors for

514 the different ROIs were computed and Fisher-Z transformed to conform better to a normal 515 distribution, resulting in a 10×10 matrix of connectivity weights (Figure 2). The matrix thus 516 represents the connectivity weights between all possible ROIs for a patient: 10 intrahemispheric 517 ROI pairs, each within the lesioned and non-lesioned hemispheres, respectively, and 25 518 interhemispheric ROI pairs between the lesioned and non-lesioned hemispheres (overall 45 519 connectivity weights for all ROI pairs). For the rest of this manuscript, this vectorized, Fisher-Z 520 transformed correlation matrix will be referred to as the full connectivity pattern, while the 521 corresponding intra- and interhemispheric subsets of the matrix will be referred to as the 522 intrahemispheric non-lesioned (1×10 vector), intrahemispheric lesioned (1×10 vector), and 523 interhemispheric connectivity patterns respectively (1×25 vector). These connectivity patterns 524 were estimated independently for each session and patient. Connectivity patterns for controls 525 were estimated similarly, with the exception that intrahemispheric connectivity patterns were 526 averaged across both hemispheres.

527

528 **4.6.** Changes in connectivity patterns in the acute recovery period

529 In the early acute recovery period (week 1-2), stroke-related damage could alter connectivity 530 patterns in patients in two distinct ways: 1) the connectivity pattern could remain the same but 531 overall connection strengths might be increased or decreased, resulting in connectivity patterns 532 in patients DC-shifted but otherwise identical to control patterns. This would indicate that a 533 canonical pattern of connectivity between motor ROIs in healthy people is simply up or down-534 regulated post-stroke either due to maladaptation or compensation for damage. 2) stroke-related 535 damage might alter connectivity weights among only a few select ROIs, e.g. either between 536 ROIs within one hemisphere or across hemispheres. This would alter the shape of the

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537 connectivity patterns in patients in comparison to controls. Since we wanted to be sensitive to 538 both kinds of connectivity pattern change, the appropriate statistical test would be a MANOVA 539 between patient and control connectivity patterns. However, due to insufficient degrees of 540 freedom in performing such an analysis (the number of connectivity weights exceeds the number 541 of patients and controls), we instead opted for a permutation test with Euclidean distance as a 542 measure of dissimilarity between patient and control connectivity patterns as it is sensitive to 543 *shape and scaling* changes of connectivity patterns (for details see supplementary material).

544

545 While, on average, connectivity patterns for patients might not differ from controls in the 546 acute recovery stage, individual patients might exhibit idiosyncratic connectivity patterns owing 547 to the heterologous distribution of lesions locations in the cohort. Thus, acute stage changes in 548 connectivity patterns might result in an increase in variability in within-group connectivity 549 patterns. To determine whether this was the case in the acute stage, we computed the average 550 Euclidean distances between each patient's connectivity pattern and the patients' mean 551 connectivity pattern (acute P variability). Similarly, we computed the average Euclidean 552 distance between each individual control pattern and the controls' mean connectivity pattern 553 (acute C_variability). The differences between these two served as a measure of increased or 554 decreased variability in the patients (P_variability-C_variability= Δ variability). We then repeated 555 the permutation test (for details see supplementary material) to generate a null distribution of the 556 difference in variability to test the significance of Δ variability.

557

558 **4.7.** Changes in connectivity patterns over time during recovery

559 Since patients in our cohort demonstrated substantial improvements of upper extremity deficits 560 in the year after stroke (Figure 1), we were interested to see whether there were concomitant 561 longitudinal changes in connectivity patterns. To determine this, we performed two separate but 562 related analyses. First, we independently compared differences in patient connectivity patterns 563 from the acute stage to all consecutive weeks (Δ week from acute to week 4, week 12, week 24, 564 and week 52) to determine how far connectivity patterns diverged over the year from the pattern 565 in the acute post-stroke stage. The same was done for control connectivity patterns to establish 566 intersession reliability. Second, we compared patient's connectivity patterns for all five 567 measurement sessions against the control connectivity patterns to determine how patient patterns 568 changed longitudinally in reference to controls (Δ pattern for acute, week 4, week 12, week 24 569 and week 52). Both these analyses were performed using Euclidean distance and permutation 570 testing in the same way as for estimating differences in connectivity patterns at the acute 571 recovery stage.

To assess if individual idiosyncratic patterns might show a change over time that could underlie recovery, we analyzed individual connectivity pattern changes for a subgroup of patients with all time-points (10 patients) by comparing pattern variability in the acute stage against all other time-points (Δweek_variability for acute_week 4, acute_week 12, acute_week 24, and acute_week 52) and performing an ANOVA with the factor Weeks.

577

578 4.1. Alternative metrics to calculate functional connectivity

579 Because changes in functional connectivity between the two primary motor cortices have been 580 reported more consistently than other connectivity changes after stroke, we also explicitly looked 581 at changes of M1-M1 connectivity weights.

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582

583 We additionally analyzed our dataset using a metric of functional connectivity that was proposed 584 in the to-date largest longitudinal resting-state stroke study with cortical and subcortical lesion 585 location, which reported changes of M1 interhemispheric connectivity. The metric has been 586 called Relative connectivity (RelCon) and is claimed to have low sensitivity to the temporal 587 signal-to-noise ratio and signal amplitude fluctuations while maintaining a high sensitivity to 588 meaningful signal changes, therefore offering an advantage e.g. in the analysis of data sets 589 acquired with different scanners (Golestani & Goodyear, 2011). RelCon looks at 590 interhemispheric connectivity of M1 in relation to intrahemispheric connectivity of M1 (for 591 details see supplementary material).

592

593 Based on the reported methods, we calculated the RelCon for interhemispheric SM1 connections594 in our dataset.

595

597 Changes of behavioral measures in patients over time were analyzed using a mixed-effects 598 ANOVA, with Week (acute – W52) as a fixed factor, and Subject as a random factor. As 599 approximately 11% of the sessions were missing, we used the lme4 toolbox in R (Bates et al., 600 2015) to fit the unbalanced mixed-effects design. Rather than F-values, statistical tests for main 601 effects and interactions are reported using a χ^2 approximation. Behavioral measures of patients 602 and controls at the acute stage were compared with a two-tailed t-test.

Intrasession reliability was analyzed by computing split-half correlations (Pearson's correlation)
 for each single week and individual patient/control, as well as looking at the averaged split-half

⁵⁹⁶ Statistical analysis

correlation for all weeks together. Reliability between groups was compared using a mixedeffects ANOVA, with Group (patients vs. controls) and Week (acute – W52) as fixed, and
Subject as a random factor. This was done for all connections, as well as subsets only including
interhemispheric, intrahemispheric lesioned or non-lesioned ROIs.

- 609 Changes of interhemispheric M1-M1 connectivity weights over time between patients and
- 610 controls were analyzed using a mixed-effects ANOVA, with Group (patients vs. controls) and
- 611 Week (acute W52) as fixed, and Subject as a random factor, alternative metrics reported in
- 612 Golestani *et al.* were analyzed in the same way.
- 613 Results were considered significant at p<0.05. Means values are reported \pm standard deviation 614 unless stated otherwise.
- 615

616 Data availability

617 The complete data set will be openly available in a public repository upon publication. All 618 analysis was performed using built-in and custom-written MATLAB and R scripts that will be 619 made publicly available upon publication.

620 **5** Acknowledgement

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623 6 Competing interests

624 The authors report no competing interests.

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626 7 Supplemental Material

627 Methods

628 Permutation test and Bootstrapping

629 To perform a permutation test, we first identified patients and controls that had estimates of 630 connectivity patterns within the first two weeks after stroke. We estimated Apattern as the 631 Euclidean distance between the average connectivity pattern for patients and the average 632 connectivity pattern for controls. We then shuffled group assignment labels for connectivity 633 patterns 10,000 times, randomly assigning connectivity patterns to "controls" or "patients". From 634 the shuffled data, we again calculated the Euclidean distance between the average connectivity 635 pattern for patients and controls based on this new assignment. By repeatedly shuffling and 636 computing Euclidean distances, we obtained an estimate of the empirical null distribution of 637 Δ pattern – e.g. the expected distribution if there was no real difference between the two groups. 638 The measured Δ pattern was then compared against this null distribution, and the relative 639 proportion of simulations that showed a larger distance was used as a p-value - the probability 640 that the distance between the mean control and patient pattern would be equal or larger than the 641 measured distance by pure chance. This analysis was carried out independently for the full, 642 intrahemispheric lesioned, intrahemispheric non-lesioned, and interhemispheric connectivity 643 patterns.

644

645 RelCon

To calculate the interhemispheric RelCon for ipsilesional and contralesional sensorimotor cortex
(SM1) the correlation between time-series of all possible pairs of voxels is calculated (all voxels
SM1_{ipsilesional-contralesional}). The average of the interhemispheric connectivity for SM1_{ipsilesional}

- 649 contralesional is then calculated relative to the within connectivity of the ipsilesional SM1 (divided
- 650 by the average correlation of all voxel within SM1_{ipsilesional}).
- 651 This metric was tested on different real and simulated data sets and showed superior results
- 652 compared to other *absolute* connectivity measures (absolute meaning connectivity measures that
- do not relate interhemispheric ROI-to-ROI connectivity weights to the average within correlation
- 654 of the ipsilesional ROI itself).
- 655

656 Results

6	5	7
U	\mathcal{I}	1

ID	age	gender	handedness	lesion side	first FM-UE	first ARAT	session
2310	57	m	right	left	58	56	5
2365	53	f	right	right	0	57	4
2395	65	m	right	right	30	21	4
2450	66	m	right	right	66	56	3
2531	66	f	right	right	60	55	5
2565	71	m	right	right	4	0	3
2652	46	m	left	left	4	0	4
2654	46	m	right	right	49	52	5
2663	67	f	right	left	16	2	4
2789	56	m	right	right	64	57	4
2925	59	f	right	left	60	57	5
3176	64	m	left	right	63	57	4
3239	74	m	left	left	5	0	5
3240	80	f	right	left	9	56	5
3241	64	f	right	right	58	39	5
3243	22	m	right	left	63	56	5
3246	53	m	left	left	30	39	5
3247	54	m	right	right	59	57	5
3248	58	m	right	right	61	56	4
	2310 2365 2395 2450 2531 2565 2652 2654 2663 2789 2925 3176 3239 3240 3241 3243 3246 3247	2310 57 2365 53 2395 65 2450 66 2531 66 2565 71 2652 46 2654 46 2655 59 3176 64 3239 74 3240 80 3241 64 3243 22 3246 53 3247 54	2310 57 m 2365 53 f 2395 65 m 2450 66 m 2531 66 f 2555 71 m 2652 46 m 2653 67 f 2654 46 m 2663 67 f 2789 56 m 2925 59 f 3176 64 m 3239 74 m 3240 80 f 3241 64 f 3243 22 m 3246 53 m	2310 57 m right 2365 53 f right 2395 65 m right 2395 65 m right 2450 66 m right 2531 66 f right 2565 71 m right 2652 46 m left 2654 46 m right 2663 67 f right 2789 56 m right 2925 59 f right 3176 64 m left 3239 74 m left 3240 80 f right 3241 64 f right 3243 22 m right 3246 53 m left 3247 54 m right	2310 57 m right left 2365 53 f right right 2395 65 m right right 2450 66 m right right 2531 66 f right right 2565 71 m right right 2565 71 m right right 2652 46 m left left 2652 46 m right right 2654 46 m right left 2653 67 f right left 2663 67 f right left 2789 56 m right left 3176 64 m left right 3239 74 m left left 3240 80 f right left 3241	IDagegenderhandednesslesion side $FM-UE$ 231057mrightleft58236553frightright0239565mrightright30245066mrightright66253166frightright60256571mrightright4265246mleftleft4265367frightleft16278956mrightleft60317664mleftleft63323974mleftleft58324080frightleft9324164frightleft63324653mleftleft30324754mrightright59	IDage gendergenderhandednesslesion sideFM-UEARAT231057mrightleft5856236553frightright057239565mrightright3021245066mrightright6656253166frightright6055256571mrightleft40265246mleftleft40265446mrightright162266367frightleft162278956mrightleft6357317664mleftleft956324080frightleft956324164frightleft6356324322mrightleft6356324653mleftleft3039324754mrightrightright5957

658

659 Table S1: Patient demographics and overall session count. First FM-UE = first recorded Fugl-

660 Meyer score Upper Extremity, first ARAT = first recorded Arm Research Action Test.

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662 Lesion distribution map

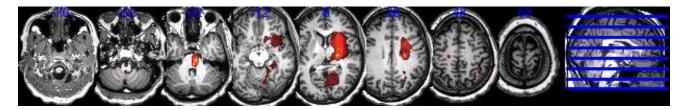


Figure S1: Lesion distribution of patients (N = 19). Averaged lesion distribution mapped to MNI space with lesion flipped to one hemisphere.

666

667 2.6 Data reliability and Preprocessing comparison

To estimate the reliability of our measurements within sessions, connectivity patterns were computed as described above for the first 100 volumes and the second 100 volumes independently and correlated with each other to calculate split-half reliabilities.

- 671 As seen for overall connectivity, intra- and interhemispheric split-half reliabilities were highly
- reliable for controls and patients (controls: *intrahemispheric*: r = 0.67 (95% Confidence Interval,
- 673 0.61-0.74), interhemispheric: r = 0.64 (CI 0.59-0.69), patients: intrahemispheric lesioned: r =
- 674 0.78 (CI 0.75-0.81), non-lesioned: r = 0.77 (CI 0.73-0.82), interhemispheric: r = 0.69 (CI 0.66-
- 675 0.74), see Figure S1 for split-half reliability of each week). Split-half reliability was not different
- between groups for *interhemispheric*: ($\chi^2(1) = 0.0239$, p = 0.8771) and *intrahemispheric non*-
- 677 *lesioned* connections: $(\chi^2(1) = 3.5634, p = 0.0591)$, but was different for *intrahemispheric*
- 678 *lesioned* connections ($\chi^2(1) = 4.2337$, p = 0.0396).

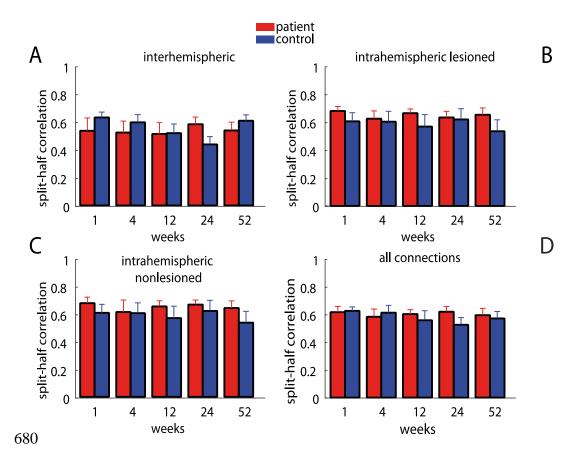


Figure S2: Intrasession split-half reliability for patients and controls at each week. For each individual patient or control participant, the BOLD time series of the first 100 volumes of the scan were correlated with the last 100 volumes. Each Panel shows the results for a different condition. Patients always had slightly higher intrasession reliability than although this difference was not significant and was possibly driven by outlier in the control group.

686 The reliability measurement also allowed us to compared two different pre-processing687 procedures:

688

Preprocessing procedure (P1): We removed the first 10 volumes of the functional data, then performed correction for the timing of slice acquisition, motion correction, brain extraction, linear trend removal, and temporal filtering (band pass, 0.01-0.08 Hz) using FSL (FMRIB

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692 Software Library (FSL), Oxford University, Oxford, UK). Our analysis was carried out in the 693 native space, and no spatial smoothing was applied. Linear regression was used to remove signal 694 correlated with the global mean signal, and the average time series in the cerebral white matter 695 and cerebrospinal fluid (Fox *et al.*, 2006).

696 Preprocessing procedure (P2): Here, we used an independent component analysis (ICA) 697 approach using FSL MELODIC for artifact reduction (Smith et al., 2004). Again, we removed 698 the first 10 volumes of the functional data. We applied motion correction and brain extraction. 699 Probabilistic independent component analysis was conducted to denoise individual data by 700 removing components such as head motion, scanner artifacts, and physiological noise. Noise 701 components were classified using FMRIB's ICA-based Xnoiseifier (Salimi-Khorshidi et al., 702 2014), which attempts to auto-classify ICA components into "good" vs. "bad" components. The 703 "bad" components were then removed from the functional data.

To determine which procedure would provide a more stable result, we calculated the split-half reliability of the ROI-ROI connectivity weights for the whole connectivity pattern over time in controls only.

Both procedures lead to good intrasession reliability on average (P1 = 0.64, CI 0.60–0.66; P2 = 0.62, CI 0.57–0.66) but showed no significant difference ($\chi^2(1) = 1.231$, p = 0.267), while no consistent change over time was found for either procedure by itself (P1: $\chi^2(4) = 2.834$, p = 0.684; P2: $\chi^2(4) = 3.007$, p = 0.557). Because of the nominal higher intrasession reliability we conducted all subsequent analyses after noise correction using the P1 procedure.

712

713 As for overall connectivity, the intersession reliability for controls showed no significant 714 change over time for intra- or intrahemispheric (*intrahemispheric lesioned*: Δ week acute_W4 =

715	0.36,	CI 0.217-1.30)8; acute V	W12 = 0.323,	CI 0.225–1.119;	acute W24 $=$	0.567, CI 0.259–

1.333; acute_W52 = 0.527, CI 0.228–1.118; *intrahemispheric non-lesioned*: Δ week acute_W4 =

717 0.36, CI 0.216–1.286; Δ week = 0.323, CI 0.221–1.121; acute_W24 = 0.567, CI 0.26–1.331 ;

718 acute_W52 = 0.527, CI 0.23–1.136; interhemispheric: , Δ week acute_W4 = 0.669, CI 0.445–

719 2.062; acute_W12 = 0.516, CI 0.419–1.876; acute_W24 = 0.721, CI 0.519–2.127; acute_W52 =

720 0.751, CI 0.45–1.991).

721

722 7.2.1 Homo- versus Heterologous ROI connectivity

723 The physiological plausibility of the recorded BOLD signal fluctuations was further examined by

724 comparing functional connectivity of homologous versus heterologous interhemispheric ROI-

725 ROI connectivity weights using a linear mixed-model with participants as random factor and

type (homo- or heterologous), week and group (control vs patients) as fixed factors,

727 supplemental results Figure S3.

728

729 Homo- versus Heterologous ROI connectivity

730 We examined the differences in connectivity between homo- and heterologous ROI connection.

731 Homologous connectivity was significantly higher than connections between heterologous ROIs

732 $(\chi^2(1) = 108.38, p < 0.001)$ and this effect showed no changes over time $\chi^2(4) = 5.8993, p =$

733 0.207), Figure S1. Furthermore, we found no differences for this effect between patients and

734 controls (type*group $\chi^2(1) = 2.2701$, p = 0.132 and type*week*group $\chi^2(1) = 2.2187$, p = 0.136),

Figure S3.

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737

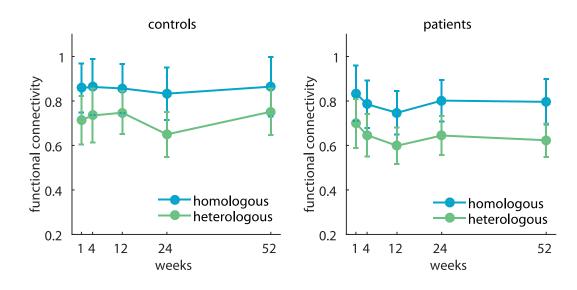
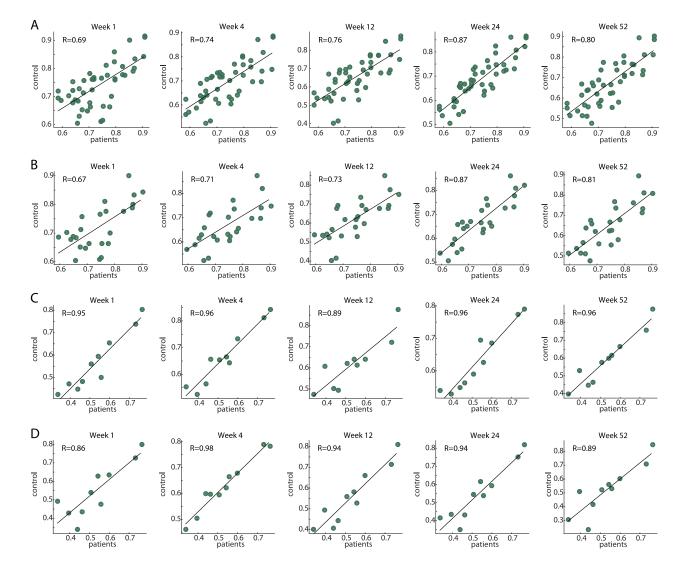


Figure S3: Average connectivity of interhemispheric homologous versus heterologous ROI-ROI connectivity weights. Homologous regions (e.g. M1-M1, S1-S1, blue line) were higher correlated than heterologous regions (e.g. M1-PmV, PmV-S1, green line). This did not change over the course of a year and no systematic difference was found between both groups (patients right panel, control left panel).

743 **7.2.2** Correlations between patient and control connectivity patterns

744 Connectivity patterns for patients and controls were highly correlated in the early period after 745 stroke as well as at all subsequent measured time-points over the year (all connections: W1: R = 746 0.69, p = 0.0002; W4: R = 0.74, p<0.0001; W12: R = 0.76, p<0.0001; W24: R = 0.87, p = 0.0001; 747 W52: R = 0.80, p<0.0001; *interhemispheric*: W1: R = 0.67, p = 0.0002; W4: R = 0.71, p<0.0001; 748 W12: R = 0.73, p<0.0001; W24: R = 0.87, p<0.0001; W52: R = 0.81, p<0.0001; intrahemispheric 749 *lesioned*: W1: R = 0.95, p<0.0001; W4: R = 0.96, p = 0.0088; W12: R = 0.89, p = 0.0006; W24: 750 R = 0.96, p = 0.0129; W52: R = 0.96, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p = 0.0129; W52: R = 0.96, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p = 0.0129; W52: R = 0.96, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p = 0.0129; W52: R = 0.96, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.86, p < 0.0001; intrahemispheric non-lesioned: W1: R = 0.00001; intrahemispheric non-lesioned: W1: R = 0.751 = 0.0014; W4: R = 0.98, p = 0.0089; W12: R = 0.94, p<0.0001; W24: R = 0.94, p = 0.0001; W52: 752 R = 0.89, p = 0.0006; Figure S4).



753

Figure S4: Correlations of connectivity patterns for patients (x-axis) and controls (y-axis) at each
week. A) all connections, B) interhemispheric, C) intrahemispheric lesioned, D) intrahemispheric

non-lesioned connectivity patterns.

757

758 8 References

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