Supplementary Material

Anatomical landmarks for ROIs

Coordinates for left and right M1 were obtained from the 'Tapping' condition, with local maxima at the 'hand knob' formation of the precentral gyrus (Yousry et al., 1997). For the premotor ROIs of PMC and SMA, we computed a conjunction analysis across all four conditions, ensuring that the regions were not only activated in the higher motor control conditions, but also in the localizer task. The SMA was located on the mesial cortical surface anterior to the paracentral lobule, superior to the cingulate sulcus and posterior to the coronal plane running through the anterior commissure (y coordinate ≤ 0 , (Picard and Strick, 2001). Hence, we deliberately chose the caudal parts of SMA, i.e., SMA proper, instead of pre-SMA because of the strong BOLD activation in this region induced by our task, knowledge about structural connectivity from non-human primate studies (Rouiller et al., 1994) and experience in motor network connectivity based on previous studies using DCM (Boudrias et al., 2012; Grefkes et al., 2010; Pool et al., 2013). The PMC was located at the intersection of the superior branch of the precentral sulcus and the superior frontal sulcus (Picard and Strick, 2001). Hence, this region most likely reflects a subregion of the PMC with strong functional coupling with both medial IPS and PFC (see a recent meta-analysis by (Genon et al., 2016). This region is particularly important for both top-down/goaldriven processes and bottom-up integration of visuospatial information into motor plans (Abe and Hanakawa, 2009; Luppino et al., 2003). When referring to non-human primates, this ROI most likely reflects area F2vr (Boussaoud, 2001; Genon et al., 2016). Importantly, there was explicitly no overlap with the ROI used for primary motor cortex. For the IPS ROI we chose its anterior and medial portion, i.e., a region devoted to visuomotor integration and coordinate transformation, most likely referring to the MIP in non-human primates (Grefkes and Fink, 2005; Grefkes et al., 2004). This regions has strong structural and functional connectivity in monkeys and humans with premotor regions and is crucially involved in visuomotor transformation processes (Grefkes et al., 2004). Coordinates of all ROIs are provided in the Supplementary Table 1.

Supplementary Table 1

ROI coordinates for the DCM analysis

Mean MNI coordinates (x, y, z) for all ROIs across all subjects, n=24. PFC = prefrontal cortex, PMC = premotor cortex, SMA = supplementary motor area, M1 = primary motor cortex, IPS = intraparietal sulcus. $L = left$ -hemispheric ROIs, $R = right$ -hemispheric ROIs.

Model space Family 1 (including PFC-premotor modulation)

All models within Family 1 shared the same endogenous network connectivity matrix (DCM-A) and varied in complexity regarding condition-dependent modulations (DCM-B): (A) without interhemispheric PFC coupling, (B) without interhemispheric IPS coupling, (C) without interhemispheric PMC coupling, (D) without interhemispheric M1 coupling, (E) without any interhemispheric coupling, (F) without PFC-premotor coupling, (G) without IPS-premotor coupling, (H) without premotor-M1 coupling, (I) modulation of all endogenous connections.

Model space Family 2 (including PFC-IPS modulation)

All models within Family 2 shared the same endogenous network connectivity matrix (DCM-A) and varied in complexity regarding condition-dependent modulations (DCM-B): (A) without interhemispheric PFC coupling, (B) without interhemispheric IPS coupling, (C) without interhemispheric PMC coupling, (D) without interhemispheric M1 coupling, (E) without any interhemispheric coupling, (F) without PFC-IPS coupling, (G) without IPS-premotor coupling, (H) without premotor-M1 coupling, (I) modulation of all endogenous connections.

Bayesian model selection

Results of the random-effects Bayesian model selection. Upper panel: Results of the family comparison; Lower panel: The most likely model given the data (I, Family 1) is highlighted in red. Results are shown for the entire sample, i.e., n=24. Note that the model selection procedure revealed the same winning model when testing for each age group separately.

Between-group activity differences for ROI analysis

Note that all differences between groups remained significant when controlling for structural atrophy as informed by the VBM analysis. PFC = prefrontal cortex, PMC = premotor cortex, SMA = supplementary motor area, $M1$ = primary motor cortex, IPS = intraparietal sulcus. R = righthemispheric, $L = left$ -hemispheric. p<0.05, FDR-corrected for multiple comparisons. Error bars: SEM.

Subcortical BOLD activity

(I) Basal ganglia and thalamus activity for young and older subjects for each condition separately. (II) Significantly enhanced BOLD activity in old as compared to young individuals for each condition separately. All p<0.05, family-wise error (FWE) corrected at the cluster level. Note that at the subcortical level, strongest between group differences were found at the level of the thalamus, especially for the 'Extern' and 'Intern' condition.

Supplementary Table 2

'Free'

'Intern'

'Extern'

Local maxima of overactivation in old vs. young subjects (whole-brain analysis)

(P<0.05, FWE corrected at the cluster level), $L = left$, $R = right$

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