

Supplementary Information

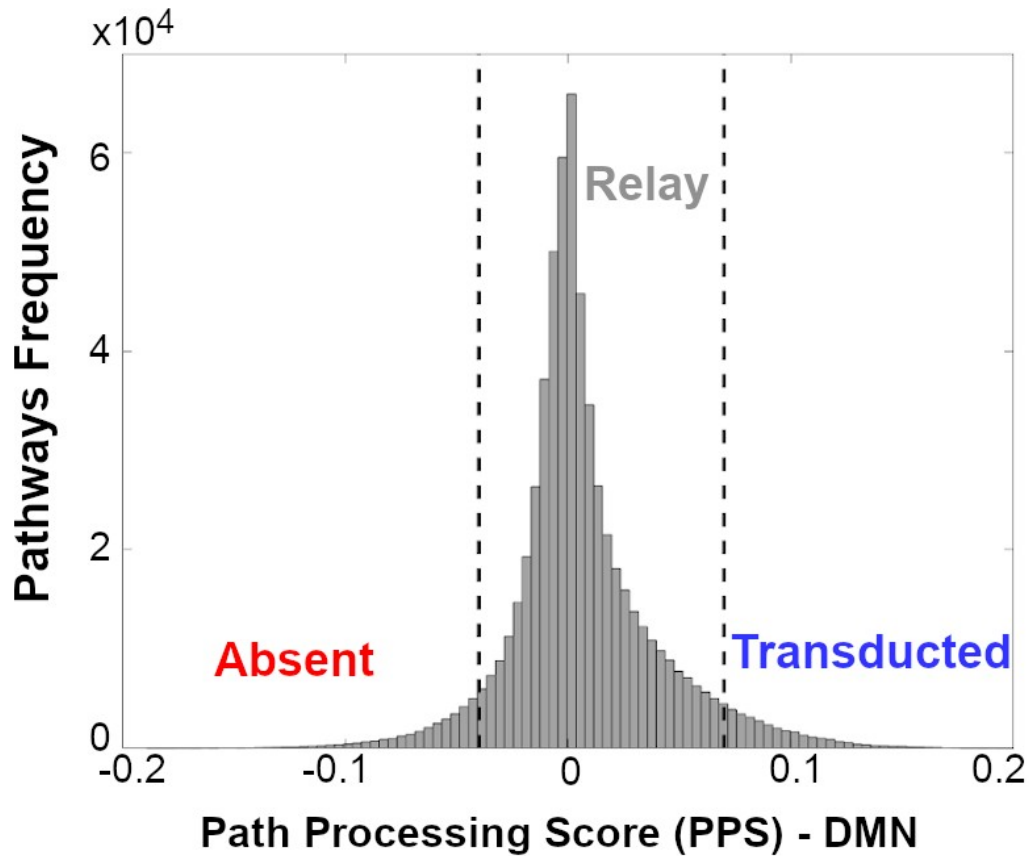


Figure S1. Relay baseline for path processing score (PPS) communication regimes. The DMN PPS distribution was obtained by considering the DMN-based shortest-paths and their corresponding (subject-level) PPS in resting state. Dashed vertical lines denote 5-95 percentiles respectively. Those percentiles were used as the range within which relay communications take place. Values below percentile 5 correspond to absent communication, whereas values above percentile 95 correspond to transducted communication.

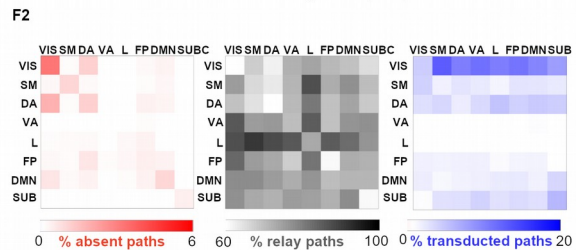
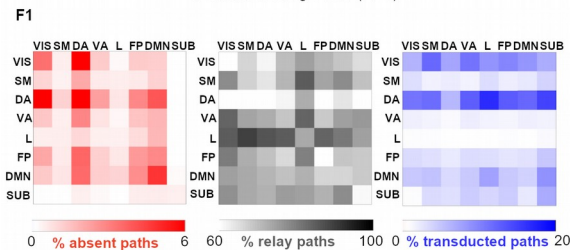
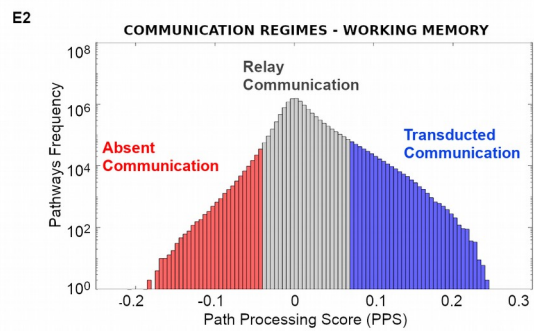
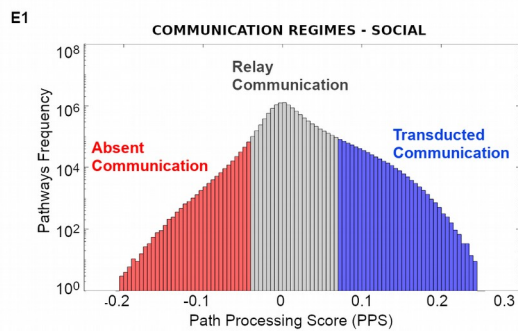
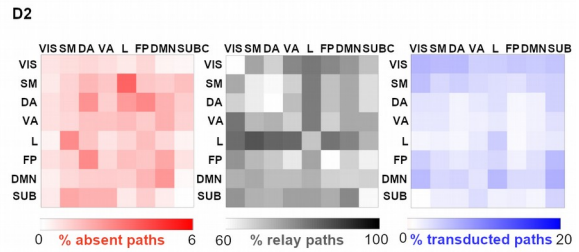
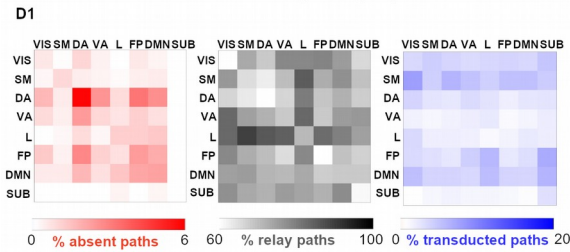
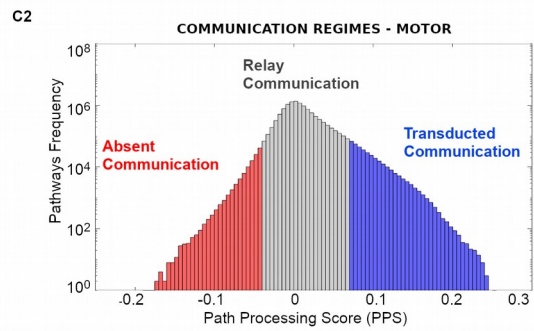
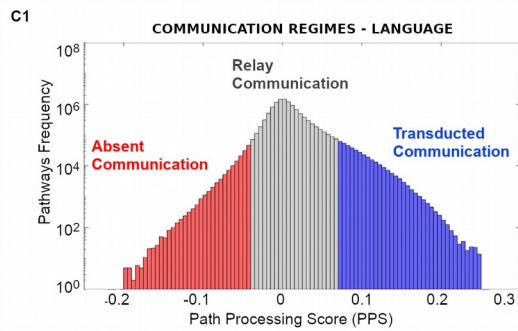
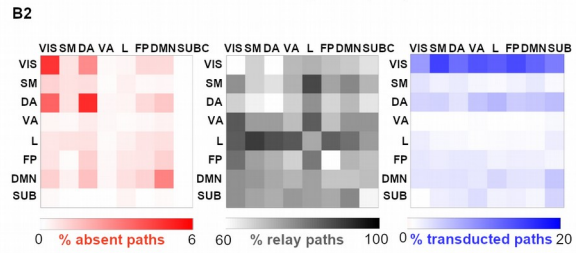
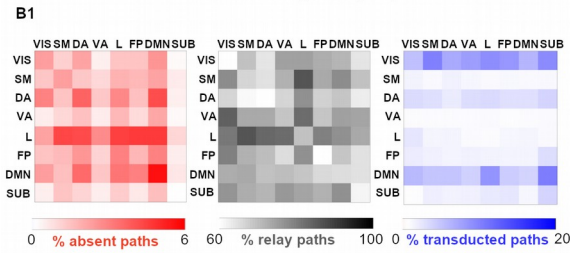
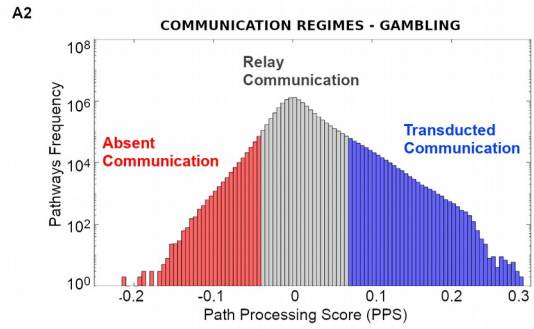
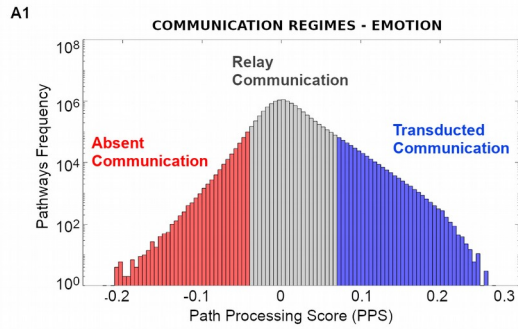


Figure S2. Communication regimes in large-scale brain networks. A1) Emotion task. Path Processing Score (PPS) on indirect pathways allows to separate brain network communication in three different regimes: absent, relay communication and transduced communication). **B1) Emotion task.** The percentage of paths, for the three different communication regimes, corresponding to the within and between 7 functional networks source-target pairs, as specified by (Yeo et al., 2011). An eighth sub-cortical community was added for completeness. Analogously, results are shown for **Gambling task (A2-B2), Language task (C1-D1), Motor task (C2-D2), Social task (E1-F1) and Working Memory task (E2-F2).**

	MI bin-width 0.5	MI bin-width 0.75	MI bin-width 1	MI bin-width 2
MI bin-width 0.5	1	0.98	0.95	0.87
MI bin-width 0.75	0.98	1	0.98	0.90
MI bin-width 1	0.95	0.98	1	0.94
MI bin-width 2	0.87	0.90	0.93	1

Table S1. Effect of MI bin-width on MI-based functional connectomes. Table shows resting-state group average similarity between MI connectomes computed with different bin-widths (0.50, 0.75, 1 and 2, respectively). Note how the bin-width parameter minimally affects the MI connectome computation for the range of bin-widths explored.

	<i>REST</i>	<i>EMOT</i>	<i>GAMB</i>	<i>LANG</i>	<i>MOT</i>	<i>RELAT</i>	<i>SOC</i>	<i>WM</i>
<i>PPS r</i>	0.98	0.96	0.96	0.95	0.93	0.97	0.97	0.97
<i>PPS SD</i>	0.017	0.021	0.019	0.017	0.018	0.021	0.019	0.016
<i>PBS r</i>	0.99	0.99	0.99	0.99	0.99	0.99	0.99	0.99
<i>PBS SD</i>	0.010	0.010	0.010	0.010	0.009	0.010	0.009	0.010

Table S2. Stability of PBS and PPS between runs and across subjects. Table reports the PPS and PBS group-average similarity across HCP trials (LR and RL, see Methods for details).