Figures and tables relevant to the main S&T results/foregrounds

Figure 1 & 2. Group comparisons between Remitters and Non-Remitters in the left hemisphere. Red indicates increased values and blue decreased values of cortical thickness for Remitters in comparison to Non-Remitters.

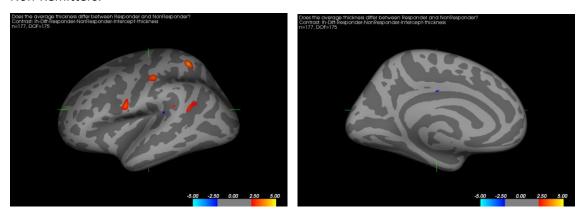


Figure 3 & 4. Group comparisons between Remitters and Non-Remitters in the right hemisphere. Red indicates larger values and blue smaller values of cortical thickness in Remitters in comparison to Non-Remitters.

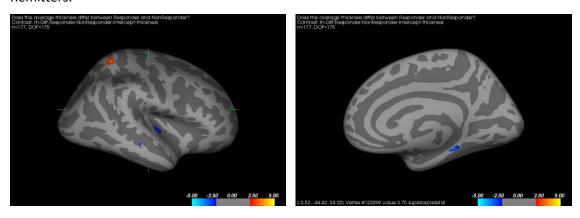


Figure 5. Group comparisons between Remitters and Non-Remitters. Red indicates smaller GM regions in Non-Remitters as compared to Remitters.

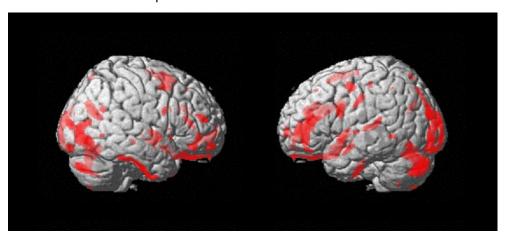
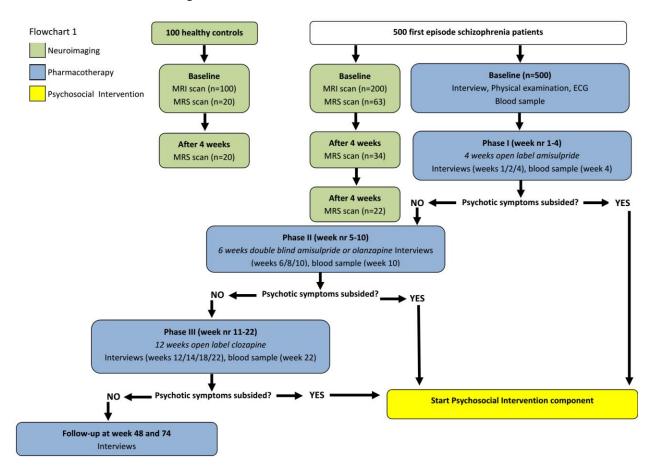


Table 1: Results from the Support Vector Machine treatment response prediction from white matter (top panel) and grey matter (bottom panel).

	All countries		Denmark		Israel		Spain		UK	
	Statistic	p-val	Statistic	p-val	Statistic	p-val	Statistic	p-val	Statistic	p-val
Accuracy	0.57	>0.05	0.57	>0.05	0.72	0.01	0.66	>0.05	0.52	-
Sensitivity	0.71	>0.05	0.48	>0.05	0.82	0.02	0.82	>0.05	0.85	>0.05
Specificity	0.37	>0.05	0.61	>0.05	0.58	0.01	0.35	>0.05	0.19	>0.05

	All countries		Denmark		Israel		Spain		UK	
	Statistic	p-val	Statistic	p-val	Statistic	p-val	Statistic	p-val	Statistic	p-val
Accuracy	0.55	>0.05	0.55	>0.05	0.571	>0.05	0.63	>0.05	0.65	>0.05
Sensitivity	0.67	>0.05	0.506	>0.05	0.735	>0.05	0.85	>0.05	0.92	>0.05
Specificity	0.37	>0.05	0.573	>0.05	0.327	>0.05	0.18	-	0.38	0.03

Flowchart 1: clinical trial design



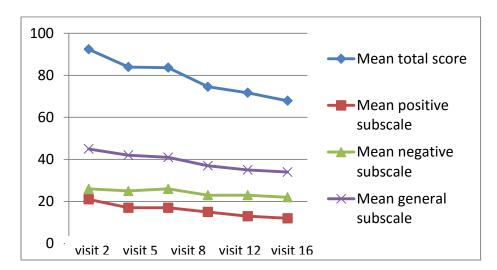


Figure 6: Positive and Negative Syndrome Scale (PANSS) scores of phase 3 patients, subdivided into total score and scores on the three subscales: positive, negative and general subscales.

Table 2: blood samples collected per study visit

	Visit 2	Visit 5	Visit 8	Visit 20
	(week 0)	(week 4)	(week 10)	(week 22)
Blood samples	403	314	55	17
collected				
DNA	377	296	53	17
RNA	391	306	52	17
Plasma	387	307	53	17
Serum	403	314	55	17
Proteomic plasma	390	303	54	17

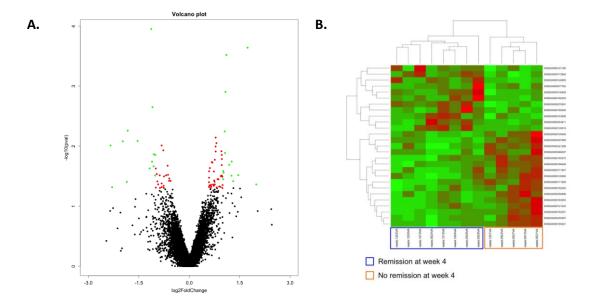


Figure 7: Genes differentially expressed between remitted and non-remitted patients. **A.** Volcano plot showing the difference in expression profile at inclusion between remitters and non-remitters to amisulpride. Up-regulated and down-regulated genes that are differentially expressed before and after treatment with amisulpride are shown in red and green, respectively. **B.** Heatmap of the cluster analysis of the 26 genes differentially expressed between remitters (blue) and non-remitters (orange) to amisulpride. Up-regulated and down-regulated genes are shown in red and green, respectively. Differentially expressed genes based cluster analysis showed a perfect discrimination between remitters and non-remitters.

Project logo:

