

SUPPLEMENTARY TABLE S3. REPRODUCIBILITY OF TRACT-DERIVED FEATURE (SD, PEAK HEIGHT, AND SKEWNESS) OF GFA AND MTR BETWEEN TP1 AND TP2 IN HEALTHY CONTROLS. PAIRED *t*-TEST ASSESSING THE NULL HYPOTHESIS: TP1 (SD, PEAK HEIGHT, AND SKEWNESS) = TP2 (SD, PEAK HEIGHT, AND SKEWNESS)

		<i>Controls (n = 10. tp1 vs. tp2). p-Value</i>																	
		<i>Intrahemispheric</i>						<i>Interhemispheric</i>											
<i>ROI_r-ROI_l</i>	<i>Features GFA/MTR</i>	<i>MI-SMA</i>		<i>PMd-MI</i>		<i>PMd-SMA</i>		<i>PMd-PMv</i>		<i>PMv-SMA</i>		<i>CC-m-PMd</i>		<i>CC-m-PMv</i>		<i>CC-m-MI</i>		<i>CC-m-SMA</i>	
		<i>GFA</i>	<i>MTR</i>	<i>GFA</i>	<i>MTR</i>	<i>GFA</i>	<i>MTR</i>	<i>GFA</i>	<i>MTR</i>	<i>GFA</i>	<i>MTR</i>	<i>GFA</i>	<i>MTR</i>	<i>GFA</i>	<i>MTR</i>	<i>GFA</i>	<i>MTR</i>	<i>GFA</i>	<i>MTR</i>
Mean		0.672	0.929	0.107	0.827	0.159	0.311	0.401	0.932	0.249	0.516	0.679	0.548	0.098	0.795	0.313	0.716	0.573	0.692
SD		0.807	0.521	0.732	0.286	0.314	0.332	0.650	0.767	0.390	0.674	0.669	0.439	0.561	0.543	0.346	0.337	0.209	0.912
Peak height		0.065	0.131	0.519	0.348	0.607	0.824	0.713	0.988	0.405	0.635	0.124	0.784	0.240	0.971	0.024	0.250	0.217	0.066
Skewness		0.004	0.611	0.645	0.624	0.389	0.203	0.159	0.449	0.782	0.086	0.881	0.595	0.977	0.845	0.467	0.501	0.662	0.432

p < 0.05 is considered significantly different.

CC-m, corpus callosum-midline; GFA, generalized fractional anisotropy; ROI, region of interest; SD, standard deviation.