

Heritability of Fractional Anisotropy in Human White Matter: A Comparison of Human Connectome Project and ENIGMA-DTI Data

For submission to Neuroimage

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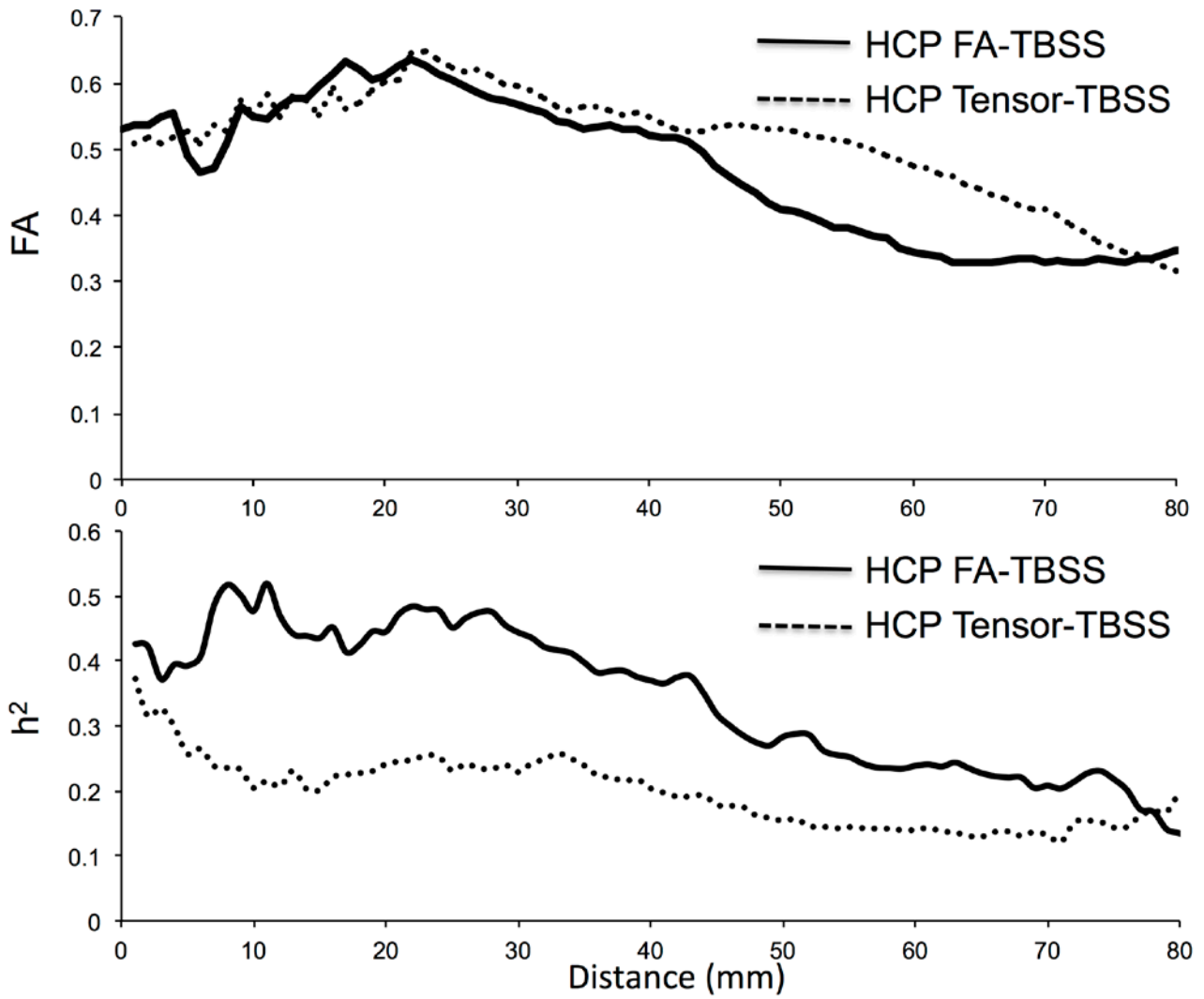


Figure 1S. Voxel-wise FA and heritability values were plotted for HCP data processed using standard TBSS and a modified TBSS approach that uses tensor-based alignment. The tensor-TBSS approach produced significantly higher FA values but lower heritability estimates.