

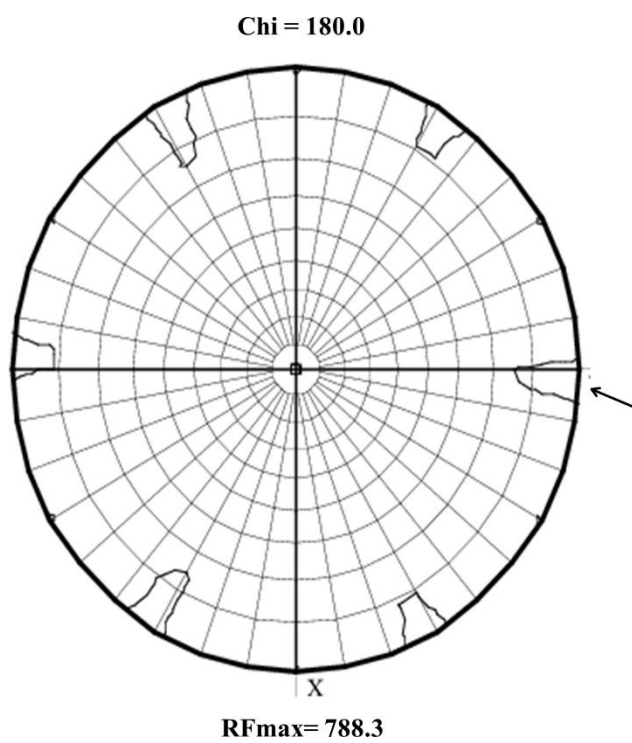


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**Supporting information for article:**

**Crystallization and X-ray crystallographic analysis of the novel  
thermostable cellulose-degrading enzyme DtCel5H from  
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**Figure S1** 180° section of the self-rotation function, computed with *MOLREP* (Vagin & Teplyakov, 2010) using a resolution range 10-4 Å and an integration radius of 30 Å.