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## MX-625 Report

We analysed cryo-cooled crystals (0.02 x 0.1 x 0.1 mm<sup>3</sup>) of the S-domain of SRP of *M.jannaschii* - consisting of 7S.S RNA, SRP19 and full-length SRP54, bound to the SRP receptor (SR) - on beam line ID29, ESRF, Grenoble in experiments conducted 29-30 November 2006. These crystals diffracted to 7.5 Å. Diffraction data were processed using the program MOSFLM. The crystals belong to space group C222 with cell parameters a = 141 Å, b = 272 Å, c = 164 Å. The structure of the complex comprising the 7S.S RNA bound to SRP19 alone is currently used in molecular replacement searches with the program CNS.