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. Check here, £98. I have inflow 160mm ballcock, £80. . . Check here, £97. I have inflow 160mm ballcock, £80. By inflow160mm. Â . . Â . Â . . The overall aim of this proposal is to continue the development of high-resolution NMR spectroscopy for the structural determination of proteins in solution and in complex with other molecules. The methods of structural analysis developed during the past grant period are now fully mature and the proposed research will introduce significant technical innovation will be: 1) sequential acquisition of 2D spectra with increased overall sensitivity by introducing non-uniform sampling of the t1 dimension during standard 3D spectroscopy and 2) efficient 3D spectroscopy based on all-to-all and sequential connectivity, by interleaving the acquisition of 3D spectra using the iJ-mediated gradient method and high resolution direct detection; and 3) high resolution 2D spectroscopy with spectral editing to recover signals from directly connected protons. Building on the successful methods developed, protein-ligand complex formation will be investigated by 2D correlation-edited and self-edited NMR spectroscopy, including techniques to reduce signal overlap. The proposed new methods of data acquisition and data analysis, combined with experimental methods for probing conformational changes in the protein upon binding a ligand, and sophisticated computer methods for creating 3D models from co-variant 2D spectra, will lead to significant new understanding of the molecular basis of protein-ligand binding. Interface Stability Policy This policy describes what makes a Go release stable.