

Cover: The figure shows the ribbon and surface representation of the structural model of a beta-agarase from *Z. galactanivorans* (coloured in blue) in interaction with an agarose double helix (stick representation, the colour code is yellow for carbon atoms and red for oxygen atoms). The crystal structure of this beta-agarase in complex with agaro-octaoses reveals two parallel agarose binding sites, one located in the catalytic groove and a second on the molecular surface. Further details can be found in the article by W. Helbert, P. Nyvall-Collen, G. Michel, M. Czjzek on page 11.

