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Note

Difference in cell surface hydrophobicity of *Halobacterium salinarium* strains

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The cell surface of halobacteria is remarkable for its ability to support the stressing conditions conferred by the high salt concentration of the surrounding medium. This is apparently due to an excess of surface charges that counteracts external ions¹⁻³ and to the internal accumulation of osmoregulators, which contributes to compensating for the high osmotic pressure from the exterior⁴⁻⁷.

The balance between the charged and non-polar groups of the cell surface in halobacteria may be important in determining its degree of halo-tolerance and/or halophilism. This property necessarily varies from one species to another, and its evaluation may be an interesting approach for classification. As shown in this paper, the chromatographic behaviour on hydrophobic matrices of three well known extreme halophile strains, regarded as belonging to the same species⁸, is distinct. It is concluded that hydrophobic interaction chromatography (HIC), a technique which takes advantage of the high salt concentration of the medium, is a useful alternative in the study of the cell surface hydrophobicity of halobacteria.

EXPERIMENTAL

The bacterial strains *Halobacterium halobium* (NRC 817), *H. cutirubrum* (CCM 2088) and *H. salinarium* (CCM 2148) were cultured in MH medium: 10 g/l MgSO₄ \cdot 7H₂O, 5 g/l KCl, 0.2 g/l CaCl₂ \cdot 6H₂O, 10 g/l yeast extract (DIFCO) and 2.5 g/l bacto tryptone (DIFCO), adjusted to 3.6 *M* NaCl at 37°C in a Lab-Line Orbit shaker at 120 rpm until the cell density reached a value of 0.45 optical units at 580 nm. Aliquots of 0.5 ml were reincubated in 50 ml as above and the bacteria from the mid-logarithmic phase were collected, washed twice with a saline solution (3.6 *M* sodium chloride) and kept at 4°C for chromatography. The adsorbent gels were obtained from Pharmacia (Uppsala, Sweden) and all other reagents from Sigma (St. Louis, MO, U.S.A.).

Hydrophobic adsorption

The hydrophobicity of the selected halobacteria was determined by adsorption on 1 ml of gel in test-tubes $(0.9 \times 11 \text{ cm})$ or on chromatographic columns $(33.5 \times 0.8 \text{ cm})$ bed volume). A 1-ml volume of cell suspension (about 0.45 optical units at 580 or 2.25 optical units at 280 nm, equivalent to $6.4 \cdot 10^6$ cells when compared with a standard plot of *Halobacterium halobium*) was either mixed with the gel in the test-tube or applied to the column at a flow-rate of 35 ml/h, collecting samples of 2 ml per tube. The absorbance of the supernatant or column effluent was determined at 280 nm in a Spectronic 2000 spectrophotometer. All experiments were run in triplicate and the mean values determined.

RESULTS AND DISCUSSION

A number of techniques have been tried for estimating the cell surface hydrophobicity of bacteria. One example is based on its degree of aggregation at different concentrations of ammonium sulphate⁹⁻¹¹; others utilize the distribution of microorganisms in a two-phase system of different polarity¹²⁻¹⁵ or the measurement of the contact angle between the bacteria and a given solid surface¹⁶; the binding of hydrophobic probes to the cell surface has also been an approach towards this end¹⁷.

Hydrophobic interaction chromatography $(\text{HIC})^{18-20}$ has been applied succesfully in the study of the hydrophobicity of many bacteria^{9,21-23} and for the separation of eukaryotic cells from various sources¹⁹. HIC is based on the interaction between an immobilized hydrophobic ligand and a hydrophobic group of the sample. The strength of interaction increases with increasing salt concentration and temperature; the nature of the hydrophobic ligand, its density on the matrix and its accesibility also determine the extent of adsorption. Hence it is possible to establish the hydrophobic character of a substance by examining its chromatographic behaviour, varying the ligand and the experimental conditions.

In general, the cell surface possesses exposed hydrophobic sites that are responsible for bacterial adsorption on hydrophobic gels²⁴. It has been shown, for instance, that enzymatic or chemical treatments modify the hydrophobicity of various kinds of microorganisms^{25–27}.

In this study we first chose the chromatographic approach for analysing the

TABLE I

Gel	NaCl (M)														
	H. cutirubrum					H. halobium					H. salinarium				
	3.6	4.0	4.4	4.8	5.2	3.6	4.0	4.4	4.8	5.2	3.6	4.0	4.4	4.8	5.2
Control									_						
Agarose (4B CL)	55	49	37	41	46	80	71	76	71	71	87	89	85	86	83
Aminoalkyl															
Agarose-ethane	69	62	44	41	46	78	78	69	68	71	89	86	84	84	86
Agarose-butane	68	61	41	41	44	72	71	69	68	66	88	83	80	83	83
Agarose-hexane	72	52	45	36	44	84	80	76	73	66	88	81	81	79	81
Alkyl															
Agarose-phenyl	31	52	52	56	56	86	89	89	83	86	99	98	99	99	99
Agarose-octane	76	60	53	46	56	79	82	83	80	80	92	89	88	88	92

EFFECT OF THE NATURE OF THE HYDROPHOBIC LIGAND ON THE ADSORPTION OF HALOBACTERIA ON HYDROPHOBIC GELS AT DIFFERENT SALT CONCENTRATIONS Values correspond to mean percentages of three replicas. behaviour of extreme halophiles, which gave very low elution yields, that is, only a fraction of the cell population (about 15%) eluted from the column. This problem was first attributed to the physical entrapment of bacterial aggregates and therefore a batch-assay procedure was followed (Table I).

As shown in Table I, the percentage of adsorption of extreme halophiles (*H. halobium*, *H. salinarium* and *H. cutirubrum*) ranges from 30 to 99% according to the nature of the immobilized hydrophobic ligand and of the experimental conditions. *H. halobium*, *H. cutirubrum* and *H. salinarium* show important differences in their tendency to adsorb on the hydrophobic matrices employed in this study. This finding suggests that they posses different cell envelope properties and supports their designation as different strains of the *H. salinarium* species⁸. It is interesting to consider the cell surface hydrophobicity, which seems to be involved in the attachment of microorganisms to solid surfaces^{28–33}, for classification purposes; apparently, *H. cutirubrum* shows the weakest, *H. halobium* a moderate and *H. salinarium* the strongest hydrophobicity (Table I).

There is no clear pattern of the adsorption of H. salinarium strains on hydrophobic gels as a function of salt concentration (Table I). In this respect they do not follow the theoretical trend of most proteins and cell particles¹⁸⁻²⁰. In other words, the adsorption tendency does not increase with increasing salt concentration. This does not necessarily mean that another mechanism of adsorption (*i.e.*, avidity for the carbohydrate matrix) is involved. In fact, the strains employed here do not utilize sugars as a carbon source⁸ and their addition to the chromatographic buffer enhances. rather than diminishes, bacterial adsorption³⁵. Bacterial hydrophobicity is the sum of a number of cell surface characteristics and its expression may be extremely complex. As reported³⁴, the tendency of 23 strains of *Staphylococci* to adhere to hydrophobic materials may vary within the same species; this behaviour appears to be related to the presence of capsules. Our observations, derived from the chromatographic patterns of halobacteria³⁵, show that differences in hydrophobic character also exist among extreme halophiles. Hence it may necessary to consider the biological significance of this property in order to interpret correctly these findings and establish their importance in bacterial classification.

REFERENCES

- 1 H. Hara and M. Masui, FEMS Microbiol. Ecol., 31 (1985) 279.
- 2 C. Pande, R. H. Callender, C.-H. Chang and T. G. Ebrey, Photochem. Photobiol., 42 (1985) 549.
- 3 F. Rodriguez-Valera, G. Juez and D. J. Kushner, System. Appl. Microbiol., 4 (1983) 369.
- 4 R. H. Vreeland, B. D. Mierau, C. D. Litchfield and E. L. Martin, Can. J. Microbiol., 29 (1983) 407.
- 5 S. M. Henrichs and R. Cuhel, Appl. Environ. Microbiol., 50 (1985) 543.
- 6 N. K. Birkeland and S. K. Ratkje, Membr. Biochem., 6 (1985) 1.
- 7 E. A. Galinski, H.-P. Pfeiffer and H. G. Truper, Eur. J. Biochem., 149 (1985) 135.
- 8 H. Larsen, in N. R. Krieg (Editor), Bergey's Manual of Systematic Bacteriology, Williams and Wilkins, Baltimore, London, 1984, p. 261.
- 9 T. J. Trust, W. W. Kay and E. E. Ishiguro, Curr. Microbiol., 9 (1983) 315.
- 10 A. Ljungh, A. Brown and T. Wadstrom, in J. Jeljaszewicz (Editor), *The Staphylococci, Zbl. Bakteriol.*, Suppl. 14 (1985) 157.
- 11 F. Rozgonyi, K. R. Szitha, S. Hjerten and T. Wadstrom, J. Appl. Bacteriol., 59 (1985) 451.
- 12 E. Weiss, M. Rosenberg, H. Judes and E. Rosenberg, Curr. Microbiol., 7 (1982) 125.
- 13 A. H. Hogt, J. Dankert, J. Feijen and J. A. DeVries, Antonie van Leeuwenhoek J. Microbiol. Serol., 48 (1982) 49.
- 14 M. Rosenberg, D. Gutnick and E. Rosenberg, FEMS Microbiol. Lett., 9 (1980) 29.

- 15 D. Lichtenberg, M. Rosenberg, N. Sharfman and I. Ofek, J. Microbiol. Methods, 4 (1985) 141.
- 16 S. Minagi, Y. Miyake, K. Inagaki, H. Tsuru and H. Suginaka, Infect. Immun., 47 (1985) 11.
- J. Brunner, A. J. Franzussof, B. Luscher, C. Zugliani and G. Semenza, *Biochemistry*, 24 (1985) 5422.
 J. L. Ochoa, *Biochimie*, 60 (1978) 1.
- 19 G. Halperin, M. Tauber-Finkelstein and S. Shaltiel, J. Chromatogr., 317 (1984) 103.
- 20 S. Shaltiel, Methods Enzymol., 104 (1984) 69.
- 21 T. Honda, M. M. A. Khan, Y. Takeda and T. Miwatani, FEMS Microbiol. Lett., 17 (1983) 273.
- 22 A. Faris, M. Lindhal and T. Wadstrom, Curr. Microbiol., 7 (1982) 357.
- 23 P. Johnsson and T. Wadstrom, Curr. Microbiol., 8 (1983) 347.
- 24 K. Pedersen, FEMS Microbiol. Lett., 12 (1981) 365.
- 25 J. H. Paul and W. H. Jeffrey, Appl. Environ. Microbiol., 50 (1985) 431.
- 26 H. F. Jenkinson, J. Gen. Microbiol., 8 (1983) 347.
- 27 F. Ascencio-Valle, A. Lopez-Cortes and J. L. Ochoa, Microbios Lett., (1988) in press.
- 28 E. E. Ishiguro, T. Ainsworth, T. J. Trust and W. W. Kay, J. Bacteriol., 164 (1985) 1233.
- 29 M. S. Hindhal and B. H. Iglewski, J. Bacteriol., 159 (1984) 107.
- 30 W. J. Peros, I. Etherden, R. J. Gibbons and Z. Skobe, J. Periodontal Res., 20 (1985) 24.
- 31 Y. Bar-or, M. Kessel and M. Shilo, J. Arch. Microbiol., 142 (1985) 21.
- 32 N. Garber, N. Sharon, D. Shohet, J. S. Lam and R. J. Doyle, Infect. Immun., 50 (1985) 336.
- 33 T. R. Tosteson, R. Revuelta, B. R. Zaidi, S. H. Imam and R. F. Bard, J. Colloid Interface Sci., 104(1985) 60.
- 34 A. H. Hogt, J. Dankert and J. Feijen, J. Gen. Microbiol., 131 (1985) 2485.
- 35 F. Ascencio-Valle and J. L. Ochoa, Rev. Latinoam. Microbiol., (1988) in press.