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**PRELIMINARY SURVEYS ON THE LACTIC ACID BACTERIA
OCCURENCE IN THE BALTIC SEA FISHES**

**WSTĘPNE BADANIA NAD WYSTĘPOWANIEM BAKTERII
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Qualitive and quantitative analysis proved lactic acid bacteria to be present in the alimentary tracks of the Baltic Sea fishes. Its numbers depended on feeding type and fishing period of the fish. Lactic acid bacteria were represented most numerously in the alimentary tracks of herrings, with the dominance of lactobacilli.

INTRODUCTION

Primarily treated as being indigenous to terrestrial environment, lactic acid bacteria were isolated, also, from marine environment far from populated areas (Schröder et al. 1980). Psychotrophic lactic acid bacteria of *L. plantarum* type were isolated from the alimentary tracks of some fish species caught in the North Atlantic, the Sea, the North Sea and the Baltic Sea as well as from krill (*Thysanoessa* sp.) from Norwegian fjord and from pacific cyster (*Grassostrea gigas*) Kraus, 1966; Preis and Lee, Schröder et al., 1980).

The aim of this work was to determine, wheather and to what extend Baltic Sea fishes harboured lactic acid bacteria.

MATERIAL AND METHODS

The subject of surveys were alimentary tracks of 64 fresh, Baltic Sea fishes, stored in ice, including:

33 herring (*Clupea harengus* L.)

Table 1

Lactic acid bacteria in the Baltic Sea fish

Fish species	No of individuals	Fishing period	CFU of lactic acid bacteria per 1g of alimentary track				Notice
			<2	2-99	100-999	> 10 ³	
herring (<i>Clupea harengus</i>)	10	01.88*	–	1	6	3	
	10	06.87	–	–	8	3	
	10	04.86	–	8	2	–	spawning
	3	07.85	–	–	3	–	
cod (<i>Gadus morhua</i>)	7	11.86	–	7	–	–	
	1	03.85	1	–	–	–	spawning
flounder (<i>Platichthys flesus</i>)	7	12.87	3	4	–	–	
	15	11.86	1	14	–	–	
	1	06.85	1	–	–	–	

* – data not confirmed

Table 2

Results of the confirmatory tests for strains isolated from alimentary tracks of the Baltic Sea fishes

Fish species	fishing period	method of estimation	No of isolated strains	No of regrown strains	G+; katalase-; OF+		others	Percent (%) of		
					G+, k-, OF+			G+, k-, OF+		others
					rods	cocci		rods	cocci	
								in No of regrown strains		
herring	01.88	immediate	—*							
	06.87		40	21	18	3	—	86	14	0
	04.86		50	27	6	3	18	22	14	64
	07.85		54	40	—	3	37	—	8	92
cod	11.86	"	45	43	39	2	2	90	5	5
	03.85		0	—	—	—	—	—	—	—
flounder	12.87	"	30	28	8	—	20	29	—	72
	11.86		4	4	2	2	—	50	50	0
	06.85		7	6	—	—	6	0	0	100
herring	01.88	after 48 h of incubation at 20°C with 1% glucose	52	50	46	4	—	92	8	0
	06.87		70	61	57	1	3	93	2	5
	04.86		48	22	17	1	4	77	5	18
	07.85		15	10	—	6	4	0	60	40
cod	11.86		30	15	15	—	—	100	0	0
flounder	12.87	"	20	20	18	2	—	90	10	0
	11.86		15	5	1	4	—	20	80	0
	06.85		—*							

* — not estimated

23 flounders (*Platichthys flesus* L.)
and 8 cods (*Gadus morhua* L.)

Microbiological analysis of samples was carried out within 24 to 48 hours after fish catching. Aseptically collected alimentary tracks were homogenized with 0,1% buffered peptone water (1:2) and then serially diluted.

In such prepared samples:

1. Total number of lactic acid bacteria was estimated. Material, from serial dilutions, transferred to Petri dishes was overlaid with MRS agar of pH 5,5 (2) and incubated at 20°C for 72 h. Grown up colonies were counted and selected, at random, for identification. Counted numbers of colonies were corrected after the preliminary identification of isolated strains.
2. Qualitative analysis was carried out.

Enriched, by addition of 1% glucose, samples were incubated for 48 h at 20°C prior to spreading on the plates. After incubation, samples were treated as the ones above.

In both cases preliminary identification of isolated strains was carried out. G-positive, katalase-negative, fermenting glucose rods and cocci were considered as lactic acid bacteria (Sharrpe, 1979).

RESULTS AND DISCUSSION

Qualitative analysis of the alimentary tracks of the Baltic Sea fishes proved the number of lactic acid bacteria harboured there to be visibly different and dependant on the fish species and fishing period (Table 1).

Lactic acid bacteria were represented most numerously within the herring's alimentary tracks. Its numbers varied from $1,9 \times 10^1$, during spawning season, to $9,5 \times 10^3$ CFU*/g, when feeding intensively. At the same time number of lactic acid bacteria stated in the alimentary tracks of cods and flounders were much lower and ranged, respectively, from < 2 to $3,8 \times 10^1$ and < 2 to $4,5 \times 10^1$ CFU/g.

Number of lactic acid bacteria in the alimentary tracks of 1 cod and 5 flounders, lower than the level detectable by the applied method, did not signify lack of these bacteria in that environment. Qualitative analysis of the same samples, enriched in 1% glucose, resulted, in all cases, in isolation of G-positive, katalase-negative, fermenting glucose rods and cocci (Table 2).

The lactic acid bacteria's growth stimulation, by enrichment of natural environment with carbohydrates, applied by Schrøder et al. (1980), let them isolate psychrotrophic bacteria of *L. plantarum* type from alimentary tracks of saithe (*Pollachius virens*) and krill from Norwegian fjords as well as from saithe and capelin (*Mallotus villosus*) from the North Atlantic and the Barents Sea (Schrøder et al., 1980).

The most probable cause of sporadic isolation cases of lactic acid bacteria from marine fishes alimentary tracks could have been usually low number of those bacteria in that

environment; lower than the one, possible to be detected, by traditionally applied methods.

Herring seems to be the only exception to the above statement. Lactic acid bacteria were isolated from all the tested alimentary tracks of that, feeding on plancton, fish, no matter the isolation technic applied.

Variable due to fishing period, however, higher, than in the alimentary tracks of the other Baltic Sea fish species, numbers of lactic acid bacteria in herring, let, probably, Kraus, cited by Schrøder et al. (1980), isolate psychrotrophic bacteria of *L. plantarum* type from the Baltic Sea herring.

Preliminary differentiation of isolated strains proved G-positive lactobacilli to dominate in most of the tested samples.

Among the strains isolated from MRS medium, with directly spreaded, diluted material, other, than lactic acid bacteria, stated for 0 to 100% of isolates (Table 2). Share of another type of microorganisms, isolated from the enriched samples, was lower and ranged from 0 to 40%.

Low selectivity of the modified MRS medium, makes isolation of lactic acid bacteria from an environment, where being in minority, more difficult. Some authors also pointed out to an unsatisfactory selectivity of MRS medium, officially advised for isolation of lactic acid bacteria (Hitchener et al., 1982, Kitchell and Shaw, 1975).

Variable nutritional and environmental requirements of lactic acid bacteria make all, so far, offered mediums for general purposes, not fully satisfactory. Commonly used MRS medium, beyond its rather low selectivity, enables growth of most representatives of that group, which is the case when presence of lactic acid bacteria is to be stated.

CONCLUSION

1. Qualitative analysis of samples enriched with 1% glucose proved the G-positive, katalase-negative, fermenting sugar rods and cocci to be present in the alimentary tracks of all tested Baltic Sea fish species.
2. Number of lactic acid bacteria in the alimentary tracks of the Baltic Sea fishes depended on the feeding type and fishing period of the fish.
3. Lactic acid bacteria were represented most numerously in the alimentary track of herring exceeding, during intensive feeding of fish 10^3 CFU/g.
4. Lactobacilli were dominant group among the strains of lactic acid bacteria isolated from the alimentary tracks of the Baltic Sea fishes.

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WSTĘPNE BADANIA NAD WYSTĘPOWANIEM BAKTERII FERMENTACJI MLEKOWEJ W RYBACH BAŁTYCKICH

STRESZCZENIE

Przeprowadzone analizy ilościowe i jakościowe prób wykazały obecność bakterii fermentacji mlekowej w przewodach pokarmowych śledzi, dorszy i storni z Morza Bałtyckiego.

Liczba bakterii fermentacji mlekowej, stwierdzona w 1 g przewodu pokarmowego zależała od typu odżywiania się ryby i okresu jej połowu. Najliczniej G-dodatnie, katalazo-ujemne, fermentujące glukozę pałeczki i ziarniaki reprezentowane były w przewodach pokarmowych śledzi. Dominantę zaś, w większości prób, stanowiły pałeczki z rodzaju *Lactobacillus*.

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