

579.81.21

**BACILLUS**

.. , .. , .. , ..

**Bacillus.** - . .

, .. , .. , .. -300

*Bacillus subtilis,*

*E.coli*

: , *Bacillus subtilis,* , , *E.coli.*

; *E-mail: safronova\_larisa@ukr.net* , 154, , 03680,

**Investigation of aerosil for development of complex probiotics on base of bacteria *Bacillus* genus.** – L.A. Safronova, A.I. Osadcha, V.M. Ilyach, V.O. Pryxodko. – Aerosil A-300 have been studied for their effect on growth and biological activity of probiotic strains *Bacillus subtilis*, some representatives obligatory microflora of intestines and pathogenic strains of *E.coli* under conditions of deep cultivation. It is shown that a erosil positively influences on probiotic strains of the family *Bacillus*, representatives of intestinal microflora and inhibits growth activity enteropathogenic strains of *E.coli*. The conducted researches have shown availability of use of the given sorbent for development complex probiotic preparations.

**Key words:** aerosil, *Bacillus subtilis*, probiotic, growth activity, obligatory microflora, *E.coli*.

**Address:** Zabolotnogo Institute of Microbiology and Virology, National Academy of Sciences of Ukraine , 154, Zabolotni str., Kyiv, 03680, Ukraine; *E-mail: safronova\_larisa@ukr.net*

[9].

-300,

[5].

[4,16].

-300

-300

[8].

[6].

(50 -100 / ),

[13].

[4, 6].

300-

1100 / (6,15).

*B. subtilis* *E. coli*  
*Lactobacillus*

( ) 1

[5, 13].

(°),

[3, 7, 11].

-300

*Bacillus subtilis*,

(Y) [1]:

*E. coli*

$$Y = N_{24}/N_0, (N_0 - N_{24} -$$

),

):

$$= Y / Y, Y Y$$

( )

*Bacillus subtilis* 39 51

(14),

*Lactobacillus* (*L. plantarum* 49, *L. fermentum* 71)-

*E. coli* 8 12 .

(*Staphylococcus*

*aureus* 209, *Salmonella typhi* 11, *Shigella flexneri* 36/39, *Escherichia coli* 028, *Proteus vulgaris* U-8 72, *Candida albicans* 690 *Pseudomonas aeruginosa* 4141),

-2.

37° 24  
*B. subtilis* *E. coli* -

*Lactobacillus*

[10].

Microsoft Excel 97.

-300,

4 40

0,05 3,0 %.

*B. subtilis*,

-300

$10^6-10^7$

( . 1).

1.

*Bacillus subtilis*

, %	<i>B. subtilis</i> 39		<i>B. subtilis</i> 51	
	( )**		( )	
0,05	0,65±0,11	7,0	0,26±0,05	6,2
0,1	0,91±0,07	6,95	0,70±0,03	6,4
0,3	1,53±0,41	6,75	0,95±0,07	5,9
0,5	1,85±0,71	6,8	1,33±0,14	6,0
1,0	1,70±0,53	7,0	1,0±0,10	6,0
3,0	0,50±0,10	7,4	0,50±0,15	6,8
( )*	1,0	7,2	1,0	7,2

\* -

\*\* - = Y / Y, Y Y

0,5 % 24 1,8-3,2·10<sup>9</sup> / ( . 2).

2. (0,5%) *Bacillus subtilis* ( 24 . )

	<i>B. subtilis</i> 39			<i>B. subtilis</i> 51		
	/	(Y) *		/	(Y)	
(0,5%)	3,2±0,61*10 <sup>9</sup>	152,8±27	6,8±0,2	1,78±0,15*10 <sup>9</sup>	148,3±12,5	6,0
( )	2,88±0,55*10 <sup>9</sup>	137,1±25	7,2±0,2	1,49±0,31*10 <sup>9</sup>	124,2±20	7,2

• -  $Y = N_{24}/N_0$ ,  $N_0$   $N_{24}$  - ( 24 )

*B. subtilis* 39 152,8±27, 11%  
*B. subtilis* 51 - (148,3 ±12,5) 19%

(0,5%)  
 1,85±0,71 *B. subtilis* 39 1,33±0,14  
 - *B. subtilis* 51. 1%  
 1% -

( . 3).

[15].

10%

[12].

*Lactobacillus*

0,5% ( . 4).

6,4-30,8%,

1

( . 4, 5).

*B. subtilis* -300.

3. *Bacillus subtilis*,

	<i>Staphylococcus aureus</i> 209	<i>Salmonella typhimurium</i> 11	<i>Es. herichia coli</i> 028	<i>Proteus vulgaris</i> U-8	<i>Proteus vulgaris</i> 72	<i>Candida albicans</i> 690	<i>Pseudomonas aeruginosa</i> 4141
<i>B. subtilis</i> 39	21	19	18	23	23	26	1
<i>B. subtilis</i> 51	13	10	12	25	25	27	13
( )							
<i>B. subtilis</i> 39	20	17	16	21	22	>25	0
<i>B. subtilis</i> 51	15	9	11	23	22	>25	11

4. *Lactobacillus* ( 24 )

	-	<i>Lactobacillus fermentum</i> 71				<i>Lactobacillus plantarum</i> 49			
		/	%	0	'	/	%	0	'
( - )	-	1,3±0,07* 10 <sup>8</sup>	100	105±20	4,2	3,1±0,30* 10 <sup>9</sup>	100	160±17	3,7-5,1
+	0,5	1,7±0,11* 10 <sup>8</sup>	130,8	110±30	4,6	3,3±0,09* 10 <sup>9</sup>	106,4	145±20	4,0-4,2

5.

		<i>Staphylococcus aureus</i> 209	<i>Salmonella typhimurium</i> 11	<i>Shigella flexneri</i> 36/39	<i>Es. herichia coli</i> 028	<i>Proteus vulgaris</i> U-8	<i>Proteus vulgaris</i> 72	<i>Candida albicans</i> 690	<i>Pseudomonas aeruginosa</i> 4141	<i>Pseudomonas aeruginosa</i> 20
		<i>Lactobacillus fermentum</i> 71		14	18	21	19	23	24	0
		15	20	25	17	24	26	2	23	-
<i>Lactobacillus plantarum</i> 49		22	24	21	-	18	19	1	23	30
		20	22	21	4	15	17	0	22	27

- coli 8 50,5-58,1 % E.  
 - coli 12 ( . 6). -

6. *Escherichia coli*

	<i>E. coli</i> 8			<i>E. coli</i> 12		
	CFU/g	%	CFU/g	CFU/g	%	CFU/g
(0,5 %)	$5,1 \pm 0,21 * 10^9$	50,5	4,2-4,5	$4,5 \pm 0,32 * 10^9$	58,1	4,0-4,2
	$1,01 \pm 0,9 * 10^9$	100	6,2-6,3	$7,75 \pm 0,53 * 10^9$	100	6,1-6,3

*Lactobacillus*

[17]. ( . 4).

[2].

[6].

[18].

*B. subtilis*

39 6,0 51 ( 6,8  
 7,2). *B. subtilis*.

1. Escherichia coli M-17 ( ) -2000.- 3.- .20-24.
2. -1996.-58, 2.- .22-28.
3. -1997.-59, 2.- .29-36.
4. -2003.- 5.- .24-25.
5. ( ) // .2 ( . ) : .55- .1999, .135-138.
6. / . . . . . /
7. . . . . . // 1991.- 4.- .31-35.

8. . . . .,1993, 22 . . . . .
9. . . . .,1998. . . . .
10. . . . .,1994. . . . .
11. . . . . Lactobacillus plan-  
tarum 8R-A3// . . . .-2003.-65, 3.- .39-46.
12. . . . . Azotobacter // . . . .-1994.-56,  
3.- .60-65.
13. // . . . .-1999.- 5.- .19-27.
14. . . . .,1993, 22 . . . . . 14569 . . . . . / . . . . .
15. . . . . 20.01.97 . . . . . 32145 61 31/00. / . . . . .
16. . . . . 7, 15.12.2000 . . . . . 2252956 , 7 12 N1/20, 23 1/165 / . . . . .
17. Harwood J.H., Pirt S.J. Quantitative aspect of growth of the methane – oxidizing bacterium Methylococcus capsulatus on methane in shake flask and continuous chemostate culture // J.Appl. Bacteriol.-1972.-55, 2.-P.597-607.
18. Iske U., Hubner K., Herold W. Investigation of the connection between the electrophoretic mobility (EFM) of microorganisms and their capability of metal uptake // Acta biotechnol. -1990.-10, 6.-P.541-549.

: 10 2007 .  
: 27 2007 .