Изпитвателен център ГЛОБАЛТЕСТ

иц глобалтест оод

1618 София, ул. "Крушовски връх" 31 теп./факс (02) 955 9862; 955 9837; 44 10 329 www.globaltest-bg.com - office@globaltest-bg.com



GLOBALTEST Testing Center

TC GLOBALTEST Ltd.

31, Krushovski vrah Str, 1618 Sofia, Bulgaria tel./fax (+359 2) 955 9862; 955 9837; 44 10 329 www.globaltest-bg.com - office@globaltest-bg.com

CERTIFICATE OF ANALYSIS

Number and date of issue: RN28159 / 15 Sept 2020

Application for testing №: RN28159 / 04 Sept 2020

Period of conducting the tests 11 Sept 2020 r. to 14 Sept 2020

Client:	ATA MERMER
Test object:	Toruko System, model Retail X, serial #20202800003, working under the Toruko technology with models: Dental, Jewelery, Hairdressing, Textile cabinet, Home and office set, I-box virus protection set, Portable set
Scope of tests:	Inoculation (contamination) of pre-sterilized samples of fabric, metal and plastic with a known number of microorganisms from the following strains: Staphylococcus aureus ATCC 6538, Pseudomonas aeruginosa ATCC 9027, Escherichia coli K 12, Candida albicans ATCC 10231, Aspergillus niger 16404. The inoculated samples of fabric, metal and plastic were placed in the Toruko system for an 8 minute cycle, after which the degree of reduction of microorganisms was reported. Expression of results in% for greater accuracy
Conclusion:	After the 8 minutes cycle in the closed Toruko system, model Retail X, serial #20202800003, where a pre-sterilized samples of fabric, metal and plastic were placed, inoculated with a known number of microorganisms from the following strains: Staphylococcus aureus ATCC 6538, Pseudomonas aeruginosa ATCC 9027, Escherichia coli K 12, Candida albicans ATCC 10231, Aspergillus niger ATCC 16404, a degree of reduction from 99.99562% to 100% inclusive is reported in the different samples, as shown in Table 1.
Study performed by:	Ms.Tanya Pancheva, Ph.D., microbiologist, Department manager "Microbiological analyses" at Testing center "GLOBALTEST"

Table 1

Degree of reduction %	Staphylococcus aureus ATCC 6538	Pseudomonas aeruginosa ATCC 9027	Escherichia coli ATCC 8739	Candida albicans ATCC 10231	Aspergillus niger ATCC 16404	
Fabric	99.99951	99.99998	99.99921	99.99936	99.99731	
Metal	100.00000	100.00000	99.99921	99.99962	99.99562	
Plastic	99.99960	99.99993	99.99921	NA	NA	

Lab manager:

Dimitar Tanev, Dipl.Eng

signature

Conducting the test:

Туре			Typical bact	eria	Yeast	Molds			
		S. aureus	Ps. aerug	E. coli	C. albicans	A.brasil			
Culture type			ATCC	ATCC	ATCC	ATCC	ATCC		
			6538	9027	8739	10231	16404		
Suspension solv	vent		NCI 9g/L; BP						
McFarland			0.5	0.7	0.5	4.0	5.0		
Dilution d 1:10		10-6	10 ⁻⁶	10-6	10-5	10-5			
			1/10 x	1/10 x	1/10 x	1/10 x	1/10 x		
			1/10 x	1/10 x	1/10 x	1/10 x	1/10 x		
			1/10 x	1/10 x	1/10 x	1/10 x	1/10 x		
Formulation			1/10 x	1/10 x	1/10 x	1/10 x	1/10 x		
			1/10 x	1/10 x	1/10 x	1/10	1/10		
			1/10	1/10	1/10				
ml, 1:10			1ml suspension + 9ml buffer						
Storing conditions of			2-8 °C	2-8 °C	2-8 °C	2-8 °C	2-8 °C		
microorganism suspensions		Up to 24 h	Up to 24 h	Up to 24 h	Up to 24 h	Up to 24 h			
		Determina	ation the conce	entration of tes	st suspension	S			
Start date: 11 Sept 2020									
End date: 13 Sept 2020									
Quantity of suspension tested on petri dishes, ml		0.1ml	0.1ml	0.1ml	0.1ml	0.1ml			
Incubation condi	itions		30-35 °C	30-35 °C	30-35 °C	20-25 °C	20-25 °C		
			24 h	24 h	24 h	3 d	3 d		
Media				CSA		SDA			
Distributed-fit		L							
Results cfu/petri		d	10-5	10-5	10-5	10 ⁻⁵	10-5		
petri1 max petri2 min			210	129	197	74	17		
			204	123	181	70	15		
X average cfu/pe			207.00	126.00	189.00	72.00	16.00		
TCoutput[cfu/ml]]= (X _{cp} x1/c	l _{1:10})/ml	207,000,000	126,000,000	189,000,000	72,000,000	16,000,000		
			2.07	1.26	1.89	7.20	1.60		
			×10 ⁸	x10 ⁸	x10 ⁸	x10 ⁷	x10 ⁷		

Precision:

Analysis:	Indo	or irradiatio	n	Suspensi	Suspension test - precision in execution between two petri							
Samples used: Fabric, Metal, Plast				tic								
Type MO	Cou	nt cfu/petri					Theoretical range of		Calculated range of variation			
							variation					
	a max	loga max	a min	loga min	average, Ā	log Ā	$\frac{1}{V\bar{A}}$ *100	2	$\frac{\log a \max - \log a \min}{\log A} * 1$			
Staphylococcus aureus	210	2.322219	204	2.309630	207.00	2.315925	6.950480	2	0.543590			
Pseudomonas aeruginosa	129	2.110590	123	2.089905	126.00	2.100247	8.908708	2	0.984865			
Escherichia coli	197	2.294466	181	2.257679	189.00	2.276072	7.273930	2	1.616278			
Candida albicans	74	1.869232	70	1.845098	72.00	1.857165	11.785113	2	1.299490			
Aspergillus brasiliensis	17	1.230449	15	1.176091	16.00	1.203270	25.000000	2	4.517495			

Materials:

Test	Inoculation with 0.1ml 10^8 10 ⁷ cfu test MO/ material
Materials used:	Fabric, Metal, Plastic – pre-sterilized

- 1. The surface of the material is contaminated with a control suspension, with a known concentration of the control microorganism
- 2. The material is irradiated in the device in one cycle
- 3. The material is immersed and washed with 100 ml of sterile solvent, leaving the remaining number of microorganisms in the solvent.
- 4. The residual number of microorganisms on the material was determined by a membrane filtration method, filtering the entire amount of solvent through a cellulose-nitrate filter with a pore size of 0,45 μ m
- 5. The filters were placed by aseptic technique on petri suitable for control microorganisms with sterile media, after which the petri were thermostated at suitable periods and temperatures suitable for the development of microorganisms.
- 6. The number of microorganisms on the used filters was determined

Dilution, d	Fabric cfu/ petri		Metal cfu/ petri		Plastica cfu/ petri	
Reported						
Staphylococcus aureus		101		0		82
Pseudomonas aeruginosa		3		0		9
Escherichia coli	>	150	>	150	>	150
Candida albicans		46		27		NA
Aspergillus brasiliensis		43		70		NA

Reduction:

Test MO	Inoculated cfu on material	Conte		nicroorganisms	Log reduction		
	(cfu/0.1 ml)	on the material			Fabric	Metal	Plastic
		Fabric	Metal	Plastic	log 10	log 10	log 10
Staphylococcus aureus	20,700,000	101	0	82	5.3	7.3	5.402156
Pseudomonas aeruginosa	12,600,000	3	0	9	6.6	7.1	6.146128
Escherichia coli	18,900,000	150	150	150	5.1	5.1	5.100371
Candida albicans	7,200,000	46	27	NA	5.2	5.4	-
Aspergillus brasiliensis	1,600,000	43	70	NA	4.6	4.4	-

