



10.04.2024 .

	(/)		400
A06.30.002.001			1000
	(, , ,)		200
	(, ,)		500
A06.30.002.002			1000
A06.30.002			500
B01.039.001	(,) -		1300
B01.039.002	(,) -		1100
A06.23.004	/		2600
A06.23.005.006	/		8700
A06.08.007	,		2600
A06.08.007.004			8700
A06.25.003			2600
A06.25.003.002			8700
A06.26.006			2600
A06.26.006.001			8700
A06.23.004.007			8700
A06.08.007.002			8700
A06.26.006.001	()		4000
			8700
A06.08.009.002			8700
A06.09.005			2600
A06.09.005.002			8700
A06.11.004			2600
A06.11.004.001			8700
A06.30.005			3500

A06.30.005.003						8700
A06.28.009						3500
A06.28.009.001						8700
A06.30.007						8700
A06.30.007.002						8700
A06.30.005.001						8700
-	()				
A06.03.021.001						2600
A06.03.021.002						8700
A06.03.021.001		-	2	(2		3600
,2	,2)				
A06.03.021.002		-	2	(2		9700
,2	,2)				
A06.03.036.001						2600
A06.03.036.002						8700
A06.03.058		()			2600
A06.03.058.003	()				8700
A06.03.062						2600
A06.04.020		-				2600
A06.04.017						2600
						2600
						8700
A06.12.052	-					8700
A06.12.053	-					8700
A06.12.050	-					8700
A06.12.054	-					8700
A06.12.055	-					8700
A06.12.057	-					8700
A06.12.058	-					8700
A06.12.001.001	-					8700
A06.12.001.002	-					8700
A06.23.004.007						8700
A06.10.009		()			3500
A06.10.006.001	-					12900
A06.10.006.001						12900
A05.23.009						3100
A05.23.009.001						7100
A05.12.004	()				3100

A05.12.005	()		3100
			7100
			7100
A05.22.002.001			6500
A05.26.008.001			7100
A05.08.001			3100
A05.26.008			3100
A05.22.002			3100
A05.04.001	-		5000
A05.30.008			3100
A05.23.009.004	-		3100
A05.23.009.008	-		3100

A05.04.001	-		3100
A05.30.008			3100
A05.03.002			3100
A05.03.002			3100
A05.03.002			3100
A05.03.002.001			7100
A05.30.008.001			7100
A05.03.002.001			7100
A05.03.002.001			7100
A05.23.009.010	()		3100
A05.23.009.011		()	7100
A05.23.009.013	-	()	3100
A05.23.009.016 ()	-		7100

		()	14000
A05.30.005			4000
A05.30.007			4000
A05.30.005.001			7100
A05.30.007.001			7100
A05.22.001	-		4000
A05.22.001.001	-		7100
A05.14.002			7100
A05.15.001			3500
A05.15.002			4000
A05.28.002			3100
A05.28.002.001	-		7100

A05.30.004.001			7100
A05.30.004			4000
A05.21.001			4000
	()+		7000
A05.21.001.001	c		7100

A05.30.004.001		7100
A05.30.004		4000
A05.30.004	() (, ,)	4000
A05.04.001	(,) , ,	3500
A05.30.011.002		3500
A05.30.012.002		3500
A05.30.010		4000
A05.01.002		4000
A05.01.001		7100
A05.30.010.001		7100
A05.08.002	-	4000
A05.08.004	-	4000
A05.03.002	+ +	7000
	+	5000
	+	5000
	+	5000
	: + +	7000
	+ +	7000
	+	4500
	+	4500
	+	5000
A05.03.002	+ +	4500
	+	4500
	+	7000
	+	4500
	+	4500
	+	5000
A06.30.008		1500
A06.08.003		1000
		1000
	2	1000
A06.08.003.002		1000
A06.08.003.002		1000
A06.03.005	,	1000
A06.03.056		1000
A06.04.001	-	1000

A06.26.001			1000
A06.07.008			1000
A06.07.009			1000
A06.25.002			1000
A06.25.002.001			1000
A06.03.010			1000
()	3		2000
A06.03.013			1000
A06.03.019			2000
A06.03.014			2000
A06.03.015		2	1000
		3	2000
A06.03.017			1000
A06.03.016			2000
A06.03.017.001			1000
A06.03.017.002			1000
A06.04.013	-		1000
	2		1000
A06.04.010			1000
A06.03.028			1000
A06.04.003			1000
A06.03.029			1000
A06.04.004			1000
A06.03.026			1000
A06.03.032			1000
	2		1000
A06.03.035	1		1000
A06.03.021			1000
A06.03.030			1000
A06.03.027			1000
A06.03.031			1000
A06.03.033			1000
A06.04.005			1000
	2	(1)	1000
	2		1000
A06.03.042			1000
	2		1000
A06.04.012			1000
A06.03.053	2		1000
			1000

A06.03.050				1000	
				1000	
		2		2000	
A06.03.036				1000	
A06.03.048				1000	
A06.03.054				1000	
A06.03.055	1			1000	
A06.04.011				1000	
A06.03.043				1000	
A06.03.045				1000	
A06.03.046				1000	
A06.03.049				1000	
A06.03.051				1000	
A06.03.052	1			1000	
A06.03.053.001				1000	
A06.30.004.001				1000	
				3000	
A06.28.001				1000	
	2			1000	
A06.28.002				3100	
		1	()	1000	
		2		1000	
A06.03.023		()		1000	
A06.03.024				1000	
A06.04.014		-		1000	
A06.03.022				1000	
A06.09.007				1000	
		-		1000	
				1000	
		1		1000	
		2		1000	
A06.20.004				1100	
A06.20.004	2			1900	
	c	(1	2-)	1500
	c	(2	2-)	2500
A06.20.004.002				1500	
A06.20.008				1000	
B01.023.001	(,)	-	1400

B01.023.002	(,) -		1200
B01.023.001	(,)		2500
B01.023.001	(,) - () 60		2300
B01.023.002	(,) - () 30		2100

A11.02.002	(1)		2500
A22.30.015	(1)		1000
A11.02.002			29900
A25.24.001.002			19500
A25.24.001.002			8900
A11.01.002	225 ()		24000
A11.24.001			2000
A11.02.002			1500
A11.02.002			1500
A11.02.002	()		1500

B01.047.001	(,) -		1400
B01.047.002	(,) -		1200
()			1200
()			2500

B01.031.001	()		1400
B01.031.002	()		1200
B01.031.001	()		2000
B01.031.002	()		2000
B01.031.001			1300
B01.031.001	()		1300
B01.031.001	-1 (,)		1500
B01.031.001	()		800
B01.031.001	/ ()		700
B01.031.001	(: ,)		800
B01.031.001	18 (095/)		800
- 095/			800
B01.031.001			800
B01.031.001	- (076/)		2000
B01.031.001	(079/)??		800

B01.029.001	(,) -		1400
B01.029.002	(,) -		1200

/

A02.26.015	-	(200
)			
A12.26.016			150
A03.26.010		(400
)			
A03.26.010		(400
)			
A02.26.015			200
A02.26.015	()	400
A02.26.015	()	200

B01.057.001	(,)	-	1400	
B01.057.002	(,)	-	1200	
B01.057.001	(,)	-	(800
)						
B01.057.002	(,)	()	800

A16.01.002	()	1500				
A16.01.012)	()()	-2	1900	
A16.01.011	()	1000				
A16.01.027			2600				
A16.30.060			1800				
A16.01.009	()	1500				
A16.30.069	()	500				
A16.01.001			1000				
A11.04.005			1000				
A11.30.024			1100				
A16.01.012)	()()	-1	1300	
A16.01.008.001			2100				
A16.01.004	()	700				
A16.01.004	()	2000				
A15.01.001			400				
A15.01.002			1000				
A11.01.001	()	400				
A16.30.076			1400				
A16.01.028	()	500				
A16.01.016	(05-1	,1)	1300			
A16.01.016	(1-3	,1)	1500			
A16.01.016	(3-5	,1)	1900			
A16.01.017	(05-1	,1)	1700			
A16.01.017	(1-3	,1)	1900			
A16.01.017	(3-5	,1)	2400			
A16.01.017	(5	,1)	2900		
			500				
A16.01.018	(,	05-1	-1)	-	1800
A16.01.018	(,	1-3	-1)	-	2100

A11.01.001	()	400
A16.01.003		2000
A16.30.032	1 -3 1	1300
A16.30.032	3 -5 1	1900
A16.01.018 (, , 3 -5 1)	-	2500
A16.01.003	(2)	2600
A16.01.023	(0,5-1 1)	1500
A16.01.023	(1 -3 1)	1900
A16.01.023	(3 -5 1)	2400
A16.01.030	(1)	2400
A16.01.030	(2)	3100
A16.30.069	(),	1000
A16.30.007.001		19000
A16.30.043.002		14800
A16.30.043.003		26000
A16.14.018.003	,	11000
A16.30.004.003	()	42300
		56400
A16.30.004.007		65000
A16.30.004.016		55000
A16.30.001.002	-	35000
A16.30.002.002		35000
A16.30.001	() ()	23100
A16.30.002	()	20000
A16.14.009.002		35000
A16.19.018	()	15000
A16.30.004.004)	(46000
A16.01.018 (, , 5 10 -1)	-	9500
A16.01.018 (, , 10 -1)	-	12000
A16.30.032	(5 -1)	9800
A16.30.032	(10 -1)	12800
A16.30.006		34200
A16.30.006		65000
A16.18.022		28000
A16.18.022.001		30000
A16.22.002		120000
A16.22.001		55000
A16.30.002 -	()	163000
A16.16.017.016		230000

A16.30.004.010		1	40000
A16.30.004.010		2	55000
A16.30.004.010		3	70000
A16.30.004.001 4	()1	22000
A16.30.004.001 4	()2	30000
A16.30.004.011	(1	4)	37000
A16.30.004.011	(2	4)	45000
A16.30.004.002	(5)	25000
A16.30.004.002	(5)	31000
A16.30.005.002			36000

	()	2420	
A08.30.046	-	()	2420

A04.12.005.003	()	1500
-	(+	2400
-	(+	2800
			1500
A04.12.001.004		-	800
A04.12.001.004			800
A04.12.001.001			1400
A04.12.002.002			1400
A04.12.002.001			2000
A04.12.002			1900
A04.12.002.003			1000
A04.12.001			1000
	()+	2000
A04.12.014			900
A04.12.001.006	()	1800
	()+	2800
A04.12.022			1400
A04.12.001.002			700
A04.12.003			1100
A04.12.003.001			1400
A04.12.003			1100
A04.12.014			900
A04.12.005.003			2000
A04.22.001			600

		800
A04.22.001.001		1100
		800
A04.16.001	(, , , , ,)	1100
	(,) , , , , , , , ,	2000
A04.06.001		500
A04.15.001		600
A04.28.002.005		600
A04.14.001		800
A04.14.002		500
A04.14.002.001		700
A04.18.001	()	800
A04.28.002.001		800
A04.22.002		500
A04.28.001		800
A04.28.002.001		700
A04.28.002.003		500
A04.28.003		800
A04.10.002	(+)	1800
A04.01.001	()	750
A04.09.001		600
A04.28.002.003	()	1100
A04.18.001		800
A04.06.002		800
A04.06.002		800
A04.06.002		800
A04.06.002	/	800
A04.06.002		800
A04.06.002		800
A04.06.002	, / , , , , ,	2000
A04.21.001		1000
	()	1200
	()+ +	1400
A04.21.001.001	+	1100
	+	1300
A04.20.002		1200

		1200
A04.30.010		1100
A04.30.001	(1200
A04.30.001.001)	1300
A04.30.001.007	() III	1800
A04.12.024.003	II-III)	1300
A04.04.001	()	700
A04.20.003	()	700
A04.20.003	()	400
A04.20.001.004	(-)	700
A04.30.001.002	, ,1) 4D (26 ,	2500
A04.30.001	II	1500
A04.30.001.002	3D	1700
A04.30.001.001	()	2000
A04.12.024.003) (II-III) (2400
A04.30.001.006	-	2500
(II)		
A04.30.001.008	III	2500
A04.30.001.002	(26 , ,1) 4D	3500
A04.30.001.002	3D	2400
A11.06.001.001		900
A11.20.010.003		900
A11.22.002.001		900
A11.30.024.001		900
A04.04.001	() 1	900
A04.04.001	() 1	700
A04.04.001	() 1	800
A04.04.001	() 1	800
A04.04.001	() 1	800
A04.04.001	() 1	800
A04.04.001	() 1	600
A04.04.002	()	600
A04.12.022		1400
A04.24.001	(1100
A04.24.001) (3200

B03.037.001			500			
A05.10.006			500			
A02.12.002.001			1200			
A05.10.008		24	2000			
+		24	2200			
A12.10.001	(+	750			
)	750			
A05.23.001			1100			
A12.10.001			750			
B01.058.001	()	-	1400		
B01.058.002	()	-	1200		
B01.001.001	()	-	1500		
B01.001.002	()	-	1300		
B01.001.002	()	-	800		
B01.001.001	()	-	-	(1500
A11.20.011			1500			
A11.20.014			1500			
A11.20.008			5400			
A11.20.008.001			2900			
A11.20.002			450			
A11.20.003			800			
A03.20.001			1000			
A11.20.015			500			
A16.20.036			2100			
A08.20.004)	(1400		
A11.20.015			500			
A14.20.002			900			
(
A16.20.036.001			7500			
A16.20.038			39000			
A16.20.016			39900			
A16.20.017.001			34900			
A16.20.004.001			30900			
A16.20.003.001	-		36900			
A16.20.041.001			29900			
A16.20.001.001	()	34900			

A16.20.001.001 ()		37900
A16.20.003.001		30900
A16.20.010.001 ()		73900
A16.20.010.001 2 ()		75900
A16.20.010.003 (1)		74900
A16.20.010.003 (2)		84900
A16.20.010 ()		58900
A16.20.011.001 ()		83900
A16.20.011.001 ()		94900
A16.20.011.002 ()		76900
A16.20.011.002 2 ()		87900
A16.20.063		92900
A16.20.033		35900
A16.20.012		77900
A11.20.003		1100
A03.20.003		19000
A16.20.042.003 ()		60750
A16.20.061.001 ()	(1	35000
A16.20.061.001 ()	(2	40500
-		
A16.20.035 (())		58900
A16.20.035.001 (1- (-))		39900
A16.20.035.001 (2- (-))		61900
A16.20.028.002		20900
A16.20.028.003		20900
A16.20.083		38900
A16.20.023		31900
A16.20.065		21900
A16.20.063.018		21900
A03.20.003		17000
A03.20.003.001 1		26900
A03.20.003.001 2		33900
A03.20.003.001 3		39900
-		
A16.20.066		4100
A11.20.018 ()	(2900
A16.20.059		950
A16.20.059.001		3900
A16.20.036		5900
A16.20.091.001		2800

A11.20.008.001		2000
A11.20.008.002		2700
A16.20.006		42000
A16.20.007		28900
A11.20.008		5400
A11.20.008.001		2900
A16.20.026.001		23000
A16.20.098	(1)	27770
A16.20.098 +)	(2)	41000
A16.20.083		46000
A16.20.029		27770
A16.20.024		63000
A16.20.028.005	(1)	49000
A16.20.028.005	(2)	60000
A16.20.019 +		120000
A11.01.013		10200
A11.01.013		14900
A11.01.013		26900
(7 +), (Identification of Sexually Transmitted Infections (STI) Pathogens, Scrape of Urogenital Epithelial Cells)*		1485
(4 +): Chlamydia trachomatis, Neisseria gonorrhoeae, Trichomonas vaginalis, Mycoplasma genitalium, (Identification of Sexually Transmitted Infections (STI) Pathogens, Chlamydia trachomatis DNA, Neisseria gonorrhoeae DNA, Trichomonas vaginalis DNA, Mycoplasma genitalium DNA, Human DNA)		1023
+ , o 16 18 Urogenital Epithelial Cells, 2 Types (16, 18) (HPV DNA, Scrape of		385
16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68))	14 :	385
18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68) Screening)	() 14 : 16,	990
(Neisseria gonorrhoeae, DNA, Scrape of Urogenital Epithelial Cells)*		275
(Neisseria gonorrhoeae,), (GC, Neisseria gonorrhoeae Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		990
(Bacterial Vaginosis, BV)		1694
(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Scrape of Urogenital Epithelial Cells)*		275
A08.20.017.002 (; ThinPrep®)		1320
A08.20.004		660
(; ; -)*		2420
(INBIOFLOR-Comprehensive Study of Microflora Composition of Urogenital Tract (UGT))		3058

APS)	(),	(Antiphospholipid Syndrome,	3927
	(Genitourinary Tract Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*		1276
	() (Vaginal Biocenosis: Bacteriophage and Antimycotic Susceptibility Testing (Gram Stain, Bacterioscopic Examination of Smear))*	1650
	(The Bethesda System ? TBS) (Cytological Examination of Cervical Epithelium with Description on The Bethesda System, TBS)		627
MICROBIOCENOSIS (PCR Panel Femoflor 16))		16. (UROGENITAL TRACT	2310
			495
-) (Cytological Examination: Cervix, Pap-test)	(1100
	(Candidiasis, Screening and Typing)		1012
MICROBIOCENOSIS, Screening (PCR Panel Femoflor Screen))		. (UROGENITAL TRACT	2035
MICROBIOCENOSIS (PCR Panel Femoflor 8))		8. (UROGENITAL TRACT	1628
B01.053.001	(,) -		1400
B01.053.002	(,) -		1200
B01.053.002	(,)	()	800
A16.28.040	(1)		2100
			10000
A16.28.058			1000
A16.28.052.001			2400
A16.28.072.001			2400
A11.28.008			2000
A11.28.006.001			500
A21.21.001			550
A12.21.003			650
A15.21.001		()	1000
A03.28.002			3500
A03.28.003	()		6000
A03.28.001			5000
A16.28.077		-	5000
A11.28.012			1000
A16.21.015			2300
A16.28.013.001			1000
A06.28.007			5000
A06.28.011			5000
	()		10000
A16.28.077			6000
A11.28.012			2000
A11.28.007	()		1000
A16.01.016			10000
A11.28.008			2000

A22.28.001			25000
A22.28.002			25000
B01.015.001	(,) -		1400
B01.015.002	(,) -		1200
B01.008.003	()		1400
B01.008.004	()		1200
A11.02.002	. (1)		260
A11.02.002	. (1)		290
A11.01.003	. 1		2999
A11.01.003	. 2		4999
A11.01.003 PRP-	1 (1)		5000
A11.01.003	mesoline () 2,5 .		2500
A11.01.003	mesoline () 2,5 .		2500
A16.01.024	(PRX-t33)		3750
A16.01.024			2000
A16.01.024			2500
A14.01.008			3000
			2000
B01.028.001	(,) -		1500
B01.028.002	(,) -		1300
B01.028.002	() -		800
A11.01.014			200
A12.25.001			1300
A11.07.004	,		2000
A11.08.001			5000
A11.08.002			3500
A11.25.006			3000
			200
A11.08.019			600
A16.01.012	()		4000
A16.08.054			3000
A11.08.007			1000
A03.25.003			300
A22.30.033			2500
A11.08.022	(-) - (1)		2500
A11.08.022	(-) - (2)		3500

A21.25.002				250
A11.07.022				300
A16.01.004				1000
A15.01.002				1000
A16.08.006.001		(1)		2500
A11.08.021.001				850
A16.08.016				750
A16.25.007		(1)		700
A11.08.004				1500
A16.08.023		(1)		2000
A16.08.023		(2)		3800
A11.25.003.001				800
A16.01.017.001	1 3 .			3500
A16.25.008				1500
A02.25.001		()		1200
A03.08.004				1500
A03.08.001		()		1400
B01.003.004.004				300
B01.003.004.005		(I)		500
B01.003.004.005		(II)		700
B01.003.004.005		(III)		900
		()		200
		()		200
		()		200
A11.08.020		()		700
A03.25.001				650
A16.08.012				1000
A16.08.009.001				10000
-1 :1				
A16.08.009.001				20000
-1 :2				
A16.08.009.001				30000
-1 :3				
A16.08.010.001			:1	10000
A16.08.010.001			:2	15000
A16.08.010.001			:3	20000
B01.003.004.001		()		1000
A16.07.087			:1	12000
A16.07.087			:2	15000
A16.07.087			:3	20000
				1000
A16.08.013			:1	18000
A16.08.013			:2	24000
A16.08.013			:3	29000
A16.08.001			1	35500
A16.08.014			1	10000
A15.03.003			()	3500

A16.08.017.001		1	30000
A16.08.017.001		2	37000
A16.08.017.001		3	42000
A16.08.035	(3)	35000
A16.08.002	1		32000
A16.08.002	2		38000
A16.08.002	3		42500
A16.08.001	2		37500
A16.08.001	3		42500
			5000
A16.08.064	1		18500
A16.08.064	2		23500
A16.08.064	3		28500
A16.08.014	2		15000
A16.08.014	3		25000
A11.25.006			3000
A17.30.021			6000
A11.08.022			2500
A11.07.022			300
A15.01.002			1000
A16.08.006.001			2500
A12.25.006			300
A11.25.003.001			800
A16.01.017.001	1		3300
A16.08.007			2500
A16.08.011			2500
A16.25.011			2000
A16.08.035	(1)	5000
A16.08.035	(2)	20000
A16.08.008.004	(1)	15000
A15.03.003		()	5000
A16.27.003.001		(1	38000
A16.27.003.001		(2	45000
A16.27.003.001		(3	50000
A16.25.042		(1)	3000
A16.25.042		(2)	5000
A16.25.042		(3)	7000
A16.08.008.004	(2)	35000
			781
			2420

A04.12.014		900
	()	1300
A04.12.003.001 ()		900
A04.12.002.002		1100
A04.12.002.001		2000
A04.12.002		1800
A04.12.002.003		1000
A04.12.001		1000
A04.12.001.001		1100
A04.12.001.002		600
	()	1500
A04.12.014		900
A04.22.001		500
		700
		800
A04.16.001 (, , , ,)		1100
	(,) , , , ,	1800
A04.14.001		700
A04.14.002		500
A04.14.002.001		650
A04.06.001		500
A04.15.001		500
A04.18.001		500
A04.28.002.001		800
A04.22.002		400
A04.28.001		700
A04.28.002.001		600
A04.28.002.005		600
A04.28.002.003		500
A04.28.003		700
A04.10.002 (+)		1650
A04.01.001 ()		750
A04.09.001		600
A04.06.003		600
A04.18.001		500
A04.06.002		800
A04.06.002		800

A04.06.002			600
A04.06.002	/		600
A04.06.002			600
A04.06.002			600
A04.06.002	, /	, , ,	1900
A04.21.001			700
A04.20.001	()		850
A04.20.002		()	850
A04.20.002.001 ()			950
A04.20.002	()		700
A04.23.001 (+)	()		900
A04.23.001	()	()	700
A04.03.001	1		700
A04.07.002			600
A04.08.001	(4-5)		500
A04.04.001.001	(1)		600
A04.03.001			600
A04.04.001	(- / -)		450
B01.059.001	(,) -		1400
B01.059.002	(,) -		1200
A03.08.004.002			6900
A16.30.074			4400
B01.059.001	(,) -	()	800
()			
A03.16.001	()		2300
A11.16.002	- ()		900
	IgA Helicobacter pylori)		850
A26.19.098	Helicobacter pylori ()		870
A11.16.002		/	1040
	Helicobacter pylori IgG (-)		3220
A11.16.002	()		400
A03.16.001.005	()		2900
			2000
A11.16.001	()		400
A11.16.003	()		400
A03.16.001.001			4900
A03.16.001.004			2400
A11.16.010			1100
A16.12.020.002			4900

A16.16.041.001		6500
A16.16.041.003		5000
A16.16.048		5000
A03.08.003		1100
A16.16.052 (1)		6700
A16.16.052 (1 2)		7900
A16.16.052 ((2 5)		9900
A11.16.002 OLGA/OLGIM)	(c	7000

A03.18.001.001		2600
A03.19.002		1800
A11.18.001	()	400
A11.19.001 ()		400
A11.19.002)	(400
A16.18.019.001	(1)	6900
A16.18.019.001	(1 2-)	8100
A16.18.019.001	(2-)	10900
A03.18.002		11900
A03.19.004		3100

A03.09.001		2900
A03.08.001.001	()	2600
A11.07.016.001)	(400
A11.08.003.001 ()		400
A11.08.008.001 ()		400
A11.08.012.001 ()		400
A03.09.002		2900
A03.09.003		3300
A03.16.002		500
A03.30.006.001		2900
A11.09.008		400
A11.09.006		3000

	()	1400
	()	1200

B01.027.001	(,) -	1400
B01.027.002	(,) -	1200

(60)		2100
Check-Up 45		12300
Check-Up 40		14500
Check-Up 45		17100
Check-Up 40		18700
Check-Up (-)" " 40		8600
Check-Up (-)" " 40		10700
" " (; ;		2990
") "(+		1600
+ + +		7500
B01.004.001 (,) -		1400
B01.004.002 (,) -		1200
A11.12.003.001 ()		400
A11.12.003 () (1)		250
A11.01.002		170
A11.02.002 (1)		200
A11.16.010		1100
() 2 .(1)		200
) () 4 (1		100
) () 5 (1		100
) () 30 /1 (1		50
100 (1) () 5 /		450
5 / (1)		50
50 / (1)		50
() 10 (1)		230
() 5 (1)		300
2 .(1)		200
) () 5 (1		120
2.0(1)		30
1.0(1)		15
1,5% 200 (1)		200
2 (1)		25
5 (1)		150
() (1)		100
50 / 2 (1)		1115
400 (1)		450

		50 (1)		5
		50 (1)		5
			5% (1)	5
		40 / 5 (1)		160
		1000 /4 (1)		130
		2 (1)		65
		8 (1)		390
			1 (1)	260

	()			1400
	()			1200

B01.008.001	(,)	-		1400
B01.008.002	(,)	-		1200

B01.023.003		-	(1200
3-				
B01.058.006		-	(1200
3-				
B01.047.009		-	(1200
3-				
B01.001.007		-	-	1200
	3)		(
B01.057.005		-	(1200
	3)			

B01.003.001	()	-	-	1400
B01.003.002	()	-	-	950

B01.003.004.009		(30)		5500
B01.003.004.009		(1)		10000
B01.003.004.012			30	6500
B01.003.004.012			1	12000
B01.003.004.007		(1)		12000
B01.003.004.007		(2)		13000
B01.003.004.008	-		1	11900
B01.003.004.008	-		2-	16000
B01.003.004.010		(1)		13000
B01.003.004.010		(2)		15000
B01.003.004.009		(2-		13500
B01.003.004.009		(2-		15000
B01.003.004.007	(2)			14000
B01.003.004.006	(30)			8000

B01.003.004.006	(1)	12000
B01.003.004.006	(2)	13000
B01.003.004.006	(2)	15000
B01.003.004.008	- (30)	7000
B01.003.004.008	- (2)	15000
B01.003.004.010	(30)	7000
B01.003.004.010	(2-	17000
B01.003.004.012	(30)	6000
B01.003.004.012	(1)	12000
B01.003.004.012	(2)	13000
B01.003.004.012	(2)	16000
B01.003.004.011	(30)	6000
B01.003.004.011	(1)	11000
B01.003.004.011	(2)	15000
B01.003.004.011	(2)	16000
B01.003.004.009.001	(30)	4500
B01.003.004.009.001	(1)	7000
B01.003.004.009.001	(2-	8000
B01.003.004.009.001	(2-	10000
B01.003.004.009.001	()	3700
B01.003.004.009.001	()	4900
B01.003.004.009.001	(+)	7100
B01.003.004.007	(30)	5500
A 16.09.011	(1)	1000
A 16.09.011	(12)	7000
B01.003.004.001		1000

B01.003.003	- -	5000
		3000
B01.001.007	- - , ()	1500
B01.001.007	- - ,	5000
B01.057.005	-	5000
B01.028.003	-	5000
B01.047.009	-	5000
B02.001.001		1100
B02.003.001	,	2500
B02.003.002	,	2500
B02.003.004		3000
B02.003.005		3000
B02.004.001		1900
B02.007.001		3000
B02.008.001	,	3000

B02.015.002	-	3000
B02.018.001		1200
B02.018.001		1200
B02.029.001		1200
B02.057.001		1200
B02.070.001	,	3000
B03.003.006		1200

B01.018.001	(,) -	1400
B01.018.002	(,) -	1200

A16.19.003.001	(,1)	6500
A16.19.013.002	()1	3100
A11.30.005)1	(1500
A11.30.005)2	(2100
A16.19.018		15000
A16.19.033 (1)		2100
A16.19.024		5900
A16.19.034		15000
A03.19.002		1800
A03.19.001		1100
A16.19.040		600
		600
	/	700
A16.19.017		6100
A16.19.017		10000
A16.01.017	1	8600
A16.01.017	2	11600
A16.01.017	3	15100
A16.19.044	(1)	7000
A14.19.002		700
A16.19.034	- 1	15000
A16.19.034	- 2	20000
A16.19.034	- 3	29900
A15.19.001		700

A16.19.003.001)1	(,1	12100
A16.19.003.001)2	(,1	17100
A16.19.041	1	13900
A16.19.041	2	17900
A16.19.033 ()1		6300

A16.19.033 () 2		9300
A16.19.024 () 1		21000
A16.19.024 () 2		25000
A16.19.024 () 3		29000
A16.19.013 1		28500
A16.19.013 2		33900
A16.19.013 3		48100
A16.19.010 1		22100
A16.19.010 2		26100
A16.19.010 3		39100

"A" "M", "A1" "B1"		1300
"A" "M", ()		1700
"B" "BE", "B1" ()		1300
"B" "BE", "B1" () (-)		1700

B01.050.001 (,) - -		1400
B01.050.002 (,) - -		1200

A11.04.003		1100
A11.04.004		1200
A15.02.001		500
A15.03.001		500
A15.03.003		700
A15.03.003.002		600
A15.03.007		600
A15.03.009		700
A15.03.010 ()		500
A15.03.010.001		600
A15.04.001 ()		700
A16.02.007 ()		29900
A16.02.008 ()		29900
A16.02.009		26000
A16.03.034		2600
A16.04.003		23000
A16.04.018		3100
A16.30.032		1300
A16.30.032		1500

A16.30.032		1900
A16.30.032		2400

B01.013.001	(,) - (60)	2000
B01.013.002	(,) - (60)	1900
B01.013.002	(,) - (60)	2500

A05.30.014		900
------------	--	-----

		18411
24		13092
25		33715
I-III		61957
(,) -		1400
/ ()		1400
(PAPP-A)		1400
		200
I		2200
II		1500
		1000
III		2000

		150
--	--	-----

()

IgG4- (Diagnosis of Autoimmune Pancreatitis and other IgG4-Related Diseases)		1837
Ig () (Anti- ndomysial antibodies, Anti-EMA, Ig)		1430
IgG () (Anti-Sacch romyces Cerevisiae Antibodies, ASCA, IgG)		1232
IgA IgG (Anti-Intestinal Goblet Cells Antibodies, GAB, IgA, IgG, Total)		1199
Ig () (Anti-Sacch romyces Cerevisiae Antibodies, ASCA, IgA)		1232
, IgG IgA (Pancreatic Antibodies, PAB) , Autoantibodies against Exocrine Pancreas,		1199
IgG (Anti-Deaminated Gliadin Peptide, Anti-DGP, IgG)		781
, IgG		2189
Ig (Anti-Tissue Transglutaminase Antibodies, Anti-tTG, tTGA, IgA)		1122
IgG (Anti-Tissue Transglutaminase Antibodies, Anti-tTG, tTGA, IgG)		1122
IgA IgG (Anti-Reticulin Antibodies, ARA, IgA, IgG, Total)		1276
Ig () (Anti-Neutrophil Cytoplasmic Antibodies, ANCA, IgA)		1232
IgG (Anti-Intrinsic Factor, IFAb, Intrinsic Factor Antibodies, IgG)		1650
IgA IgG (Anti- ndomysial antibodies, Anti-EMA, IgA, IgG, Total)		1276

IgA, IgG, IgM (), (Gastric Parietal Cell Antibodies, GPA, Anti- arietal cell antibodies, APCA, IgA, IgG, IgM, Total)	1518
IgG IgA GP2 (Anti-GP2)	1980
Ig (Anti-Deaminated Gliadin Peptide, Anti-DGP, IgA)	781
() . FLCN, . (Birt-Hogg-Dube Syndrome, BHD, Gene FLCN, Mut.)	51348
() II. MFN2, . . (Charcot-Marie-Tooth Disease Type 2A1, Gene MFN2, Freq. Mut.)	5148
4 TBP, . .	3410
N1, . .	3410
() . MYH3, . (Arthrogryposis Distal Type 2A, Gene MYH3, Freq. Mut.)	9999
() I. NDRG1 SH3TC2, . . (Charcot-Marie-Tooth Disease Type 1B, Genes NDRG1, SH3TC2, Mut.)	5148
ATXN8, . . (Spinocerebellar Ataxia, Gene ATXN8, Freq. Mut.)	5148
() . TAZ, . (Left Ventricular Non-Compaction, LVNC, Gene TAZ, Mut.)	25707
MVK, . (Mevalonic Aciduria, Gene MVK, Mut.)	42801
8, 9 LMNA, . (Mandibuloacral Dysplasia, Exons 8, 9 Gene LMNA, Mut.)	7161
I, II, III, IV (SMN2) (Spinal Muscular Atrophy, SMA, Type I, II, III, IV (copy Number Variation SMN2))	16577
CHRNA9, . (Escobar Syndrome, Gene CHRNA9, Mut.)	34254
BSCL2, . (Silver Syndrome, Gene BSCL2, Mut.)	29975
STX11, . (Familial Hemophagocytic Lymphohistiocytosis, Gene STX11, Mut.)	14267
-1- , SERPINA1, . .	2035
I, II, III, IV. SMN1, . () (Spinal Muscular Atrophy, SMA, Type I, II, III, IV, Gene SMN1, Mut. (Only Presence One Gene Copy))	34254
CINCA, NLRP3 . (Chronic Infantile Neurologic Cutaneous Articular, Gene NLRP3, Mut.)	51348
TBX3, . (Pallister W Syndrome, Gene TBX3, Mut.)	29975
RABPN1, . . (Oculopharyngeal Muscular Dystrophy, OPMD, Gene RABPN1, Freq. Mut.)	5148
PRNP, . (Gerstmann-Straussler Disease, Gene PRNP, Mut.)	14916
() . IKBKG, . . (Bloch-Sulzberger Syndrome, Familial Incontinentia Pigmenti, Gene IKBKG, Freq. Mut.)	5148
GJB2	11264
NS3, NS5A NS5B (3)	12837
RAB27A, . (Griscelli Syndrome, Gene RAB27A, Mut.)	21428
C1NH, . (Hereditary Angioedema Type I, Gene C1NH, Mut.)	29975
() I. 22, . (Charcot-Marie-Tooth Disease Type 1B, Gene 22, Mut.)	17523
HRAS, . (Costello Syndrome, Gene HRAS, Mut.)	7161
RMRP, . (Metaphyseal Chondrodysplasia, McKusick Type, Gene RMRP, Mut.)	7161
MEFV, . (Familial Mediterranean Fever, FMF, Gene MEFV, Mut.)	41525
SBDS1, . . (Shwachman-Diamond Syndrome, Gene SBDS1, Freq. Mut.)	7161
HLA-A29	3300

ACVR1, « . » . . (Fibrodysplasia Ossificans Progressiva, FOP, Gene ACVR1, without Hot-Point Mut.)	29975
Gene EDNRB, Mut.) EDNRB, . (Waardenburg-Shah Syndrome,	29975
ERCC6, . (Cockayne Syndrome, Gene ERCC6, Mut.)	94083
SLC26A2, Mut.) SLC26A2, . (Diastrophic Dysplasia, Gene	25707
B1. ROR2, . (Brachydactyly Type B1, Gene ROR2, Mut.)	14267
(). GLI3, . (Greig Syndrome, Gene GLI3, Mut.)	76989
	8910
(- - -) I. EGR2, . (Charcot-Marie-Tooth Disease Type 1B, Gene EGR2, Mut.)	17160
7 9 FGFR2, . (Crouzon Syndrome, Exons 7, 9 Gene FGFR2, Mut.)	9999
- a - a (). ENG, . (Rendu-Osler-Weber Disease, Gene ENG, Mut.)	38522
Exudative Vitreoretinopathy, FEVR, Gene NDP, Mut.) NDP, . (Familial	11264
Lipodystrophy 2, Gene LMNA, Mut.) LMNA, . (Familial Partial	42801
(- - -) (- - -) LMX1B, . (Nail-Patella Syndrome, NPS, Onychoosteodysplasia, Gene LMX1B, Mut.)	29975
Mut.) SGCE, . (Myoclonic Dystonia, Gene SGCE,	51348
ATP7B, Freq. Mut.) ATP7B, . . (Wilson Disease, Gene	10285
, NGF . (Hereditary Sensory and Autonomic olynuropathy, Gene NGF, Mut.)	17160
« » ALMS1, « . » . . (Alstrom syndrome, Gene ALMS1, Hot-Point Mut.)	14267
(- - -) TCOF1, . (Treacher-Collins Syndrome, Franceschetti-Klein Syndrome, Mandibulofacial Dysostosis without Limb Anomalies, Gene TCOF1, Mut.)	94083
GJB3, . (Erythrokeratoderma, Gene GJB3, Mut.)	11264
, GLA, .	10120
« » TRPV4, « . » . . (Distal Spinal Muscular Atrophy Congenital Non-Progressive, Gene TRPV4, Hot-Point Mut.)	14267
SLC22A5, . (Systemic Primary Carnitine Deficiency, SPCD, Carnitine Deficiency Systemic Primary, CDSP, Gene SLC22A5, Mut.)	42801
Mut.) FXN, . . (Friedrich Ataxia, Gene FXN, Freq.	9020
(- - -) (- - -) EMG1, . (Bowen Conradi Syndrome, BCS, Gene EMG1, Mut.)	17160
(Leber Hereditary Optic Neuropathy, LHON, Mitochondrial DNA, 3 Freq. Mut.) , 3 . .	5148
(- - -) (- - -) BCS1L, . (Bjomstad Syndrome, Gene BCS1L, Mut.)	17523
TWIST1, Mut.) TWIST1, . (Saethre-Choitzen Syndrome, Gene	14916
(- - -) II. GDAP, . (Charcot-Marie-Tooth Disease Type 2A1, Gene GDAP, Mut.)	25707
COMP, Freq. Mut.) COMP, . . (Pseudoachondroplasia, Gene	6820
Gene NPHS1, Mut.) NPHS1, . (Nephrotic Syndrome Type 1, NPHS1,	76989
Mut.) FGFR3, . . (Achondroplasia, Gene FGFR3, Freq.	12056
Syndrome, FCAS, Gene NLRP3, Mut.) NLRP3 . (Familial Cold Autoinflammatory	51348
(- - -) I. O, . (Charcot-Marie-Tooth Disease Type 1B, Gene O, Mut.)	17160
DLL3, Mut.) DLL3, . (Spondylocostal Dysostosis, Gene	29975

NPHP1, . (Nephronophthisis 1, NPHP1, Gene NPHP1, Mut.)	17006
Muscular Dystrophy, Gene FHL1, Mut.)	34254
Dihydrate, CPPD, Gene ANKH, Mut.)	51348
MULIBRAY. TRIM37, . (Muscle-Liver-Brain-Eye, Gene TRIM37, Mut.)	9999
(Leber Hereditary Optic Neuropathy, LHON, Mitochondrial DNA, 12 Freq. Mut.)	14267
A taxia, Gene ATXN7 Freq. Mut.)	5148
IGHMBP2, . (Distal Spinal Muscular Atrophy 1, DSMA1, Gene IGHMBP2, Mut.)	64163
RP2, Mut.)	21428
Syndrome, Type VI, Gene PLOD, Freq. Mut.)	10285
(Aarskog-Scott Syndrome, Faciodigitogenital Syndrome, Faciogenital Dysplasia, Gene FGD1, Mut.)	62722
Spinal and Bulbar Muscular Atrophy, Gene AR, Freq. Mut.)	5148
PCSK9	11682
(X)	6369
/ , FMR1, . .	3410
Mut.)	76989
9 FGFR2, . (Antley-Bixler Syndrome, ABS, Exon 9 Gene FGFR2, Mut.)	7161
7, 9 FGFR2 7A FGFR1, . (Pfeiffer Syndrome, Exons 7, 9 Gene FGFR2, Exon 7A Gene FGFR1, Mut.)	14267
/	11803
(Familial Hemophagocytic Lymphohistiocytosis, Gene UNC13D, Freq. Mut.)	5148
(Autoimmune Lymphoproliferative Syndrome, ALPS, Gene TNFRSF6, Mut.)	34254
(Familial Meddular Thyroid Cancer, Exons 5, 8 Gene RET, Mut.)	9999
Syndrome, Gene SBDS, Mut.)	21428
Mut.)	17523
(Stargardt Disease 1, STGD1, Fundus Flavimaculatus Included, Gene ABCA4, Freq. Mut.)	12056
CHM, . (Choroideremia, CHM, Gene CHM, Mut.)	64163
	7590
WWS, Gene FKRP, Mut.)	14916
Syndrome with Acanthosis Nigrigan, CAN, Exon 10 Gene FGFR3, Mut.)	7161
Agammaglobulinemia, XLA, Gene BTK, Mut.)	76989
PHEX, . (Hypophosphatemic Vitamin D-Resistant Rickets, Gene PHEX, Mut.)	94083
Progressive, Gene GRN, Mut.)	25707
XK, . (McLeod Syndrome, Gene XK, Mut.)	17160
(Testicular Feminization Syndrome, Gene AR, Mut.)	42801
Ectodermal Dysplasia, Gene EDA, Mut.)	34254
UPK3A, Mut.)	25707
Mut.)	17006

« . . . » . . . (Optic Atrophy With Or Without Deafness, Ophthalmoplegia, Myopathy, Ataxia And Neuropathy, Gene OPA1, Hot-Point Mut.)	OPA1,	9999
Dominant, SCN1, Gene ELA2, Mut.)	ELA2, . (Neutropenia Severe Congenital 1 Autosomal	21428
, APOB100		4818
(Congenital Insensitivity To Pain With Anhidrosis, CIPA, Gene NTRK1, Mut.)	NTRK1, .	51348
Epiphysial Dysplasia, MED, Gene COMP, Freq. Mut.)	COMP, . . (Multiple	6820
PHOX2B, . . . (Congenital Central Hypoventilation Syndrome, CCHS, Gene PHOX2B, Freq. Mut.)	(.) .	5148
(Genes CFTR, GJB2, PAH, SMN))	(. CFTR, GJB2, PAH, SMN) (Main Hereditary Diseases	21142
Gene DMPK, Freq. Mut.)	DMPK, . . (Myotonic Dystrophy 1,	5148
GJB1, . (Charcot-Marie-Tooth Disease Type 1B, Gene GJB1, Mut.)	(. - -) l.	9999
ongenital Ichthyosis, ARCI 1, All Known Mutations, Gene TGM1, Mut.)	TGM1, . (Autosomal Recessive	42801
Motor Neuropathy, DHMN, Gene BSCL2, Mut.)	v. BSCL2, . (Distal Hereditary	29975
Mut.)	ZEB2, . (Mowat-Wilson Syndrome, Gene ZEB2,	64163
(Phosphoribosylpyrophosphate Synthetase Superactivity, PRS Superactivity, Gene PRPS1, Mut.)	PRPS1, .	29975
(Albinism oculocutaneous, Hermansky-Pudlak type, Gene HPS1, Freq. Mut.)	(. - -) . HPS1, . . .	9999
/ .		23540
(Klippel-Feil Syndrome, Gene GDF6, Mut.)	(.) . GDF6, .	14916
. (Cranio metaphyseal Dysplasia, Gene ANKH, Hot-Point Mut.)	« . » ANKH, « . » .	9999
(Nonbullous Congenital Ichthyosiform Erythroderma, NBCIE, Gene TGM1, Mut.)	(.) . TGM1, .	42801
PRNP, Mut.)	PRNP, . (Creutzfeldt-Jakob Disease, Gene	14916
Gene CRYBA4, Mut.)	CRYBA4, . (Microphthalmia with Cataract,	25707
:		7590
(TNF-Receptor-Associated Periodic Syndrome, TRAPS, Gene TNFRSF1A, Mut.)	TNFRSF1A, .	25707
(.) . NPHP1 (Joubert Syndrome, Cerebelloparenchymal Disorder IV, CPD IV, Classic Joubert Syndrome, Joubert Syndrome type A, Joubert-Boltshauser Syndrome, Pure Joubert Syndrome, Gene NPHP1, Mut.)	(.) . NPHP1 (Joubert Syndrome, Cerebelloparenchymal Disorder IV, CPD IV, Classic Joubert Syndrome, Joubert Syndrome type A, Joubert-Boltshauser Syndrome, Pure Joubert Syndrome, Gene NPHP1, Mut.)	17006
, CYB5R3 . . (Methemoglobinemia, Gene CYB5R3, Freq. Mut.)	, CYB5R3 . . (Methemoglobinemia, Gene CYB5R3, Freq. Mut.)	5148
Mut.)	IT15, . . (Chorea Huntington, Gene IT15, Freq.	5148
Syndrome, Type 1, SGBS1, Gene GPC3, Mut.)	GPC3, . (Simpson-Golabi-Behmel	34254
III, Gene OPA3, Mut.)	OPA3, . (3-Methylglutaconic Aciduria Type	11264
. (X-Linked Lymphoproliferative Syndrome, XLP, Gene XIAP, Mut.)	(.) , XIAP	34254
FMF, Gene MEFV, Freq. Mut.)	MEFV, . . (Familial Mediterranean Fever,	9977
Gene SRY, Mut.)	SRY, . (Disorders Sex Determination,	7161
Dysplasia, Gene GJB6, Mut.)	GJB6, . (Hidrotic Ectodermal	11264
ongenital 1, NYS1 X-Linked, Gene FRMD7, Mut.)	FRMD7, . (X-Linked Nystagmus	51348
Mut.)	ALX4, . (Parietal Foramina, PFM, Gene ALX4,	17160
WAS, Gene WAS, Mut.)	(.) . WAS, . (Wiskott-Aldrich Syndrome,	29975

(Emery-Dreifuss Muscular Dystrophy, X-Linked Gene Emerine, Mut.)	14916
Enteropathica, Gene SLC39A4, Mut.)	34254
22, Mut.)	17523
(Osteopetrosis Autosomal Recessive 1, OPTB1, Gene TCIRG1, Freq. Mut.)	5148
SCN4A, Mut.)	14267
Mut.)	9999
LDLR	12837
-IgD Mut.)	42801
IA, Gene TYR, Mut.)	21428
Dystrophy-Dystroglycanopathy, Gene FKR, Freq. Mut.)	7876
Fundus Flavimaculatus Included, Gene ABCA4, Freq. Mut.)	12056
(Spondyloepiphyseal Dysplasia Tarda, SEDT, Gene TRAPPC2, Mut.)	17160
Pneumothorax, PSP, Gene FLCN, Mut.)	51348
, NBN . . (Nijmegen Breakage Syndrome, NBS, Gene NBN, Freq. Mut.)	5148
PAX3, Mut.)	34254
Dystrophy Limb-Girdle Type 2A, Gene FKR, Mut.)	14916
CTSK, Mut.)	25707
PAH, Mut.)	51348
Syndrome, Gene DHCR7, Mut.)	38522
CFTR, Mut.)	17006
	8800
NOTCH3	12320
Muscular Dystrophy, X-Lyonization, Girls)	8030
GDF6, Mut.)	14916
	7590
SPM, Gene FHL1, Mut.)	34254
Hemophagocytic Lymphohistiocytosis, Gene PRF1, Mut.)	21428
Gene PRNP, Mut.)	14916
EXT1, Mut.)	51348
Myoclonic Epilepsy 1A Unverricht and Lundborg, Gene CSTB, Freq. Mut.)	5148
ACVR1, Mut.)	17160
	3410
(X-Linked Severe Combined Immunodeficiency, Gene IL2RG, Mut.)	17160
NLRP3 Mut.)	51359
Elasticum, Gene ABCC6, Freq. Mut.)	7161
, CYB5R3 Mut.)	34254
Hypertension 1, PPH1, Gene BMPR2, Mut.)	64163

FXN, . (Friedrich Ataxia, Gene FXN, Mut.)	21428
NDP, . (Norrie Disease, Gene NDP, Mut.)	11264
(). LOX12B, . (Nonbullous Congenital Ichthyosiform Erythroderma, NBCIE, Gene LOX12B, Mut.)	42801
TNNT2, . (Familial Hypertrophic Cardiomyopathy, Gene TNNT2, Mut.)	51348
(). CYP1B1, . (Primary Open Angle Glaucoma 1A, POAG 1A, Gene CYP1B1, Mut.)	17523
FLT4, . (Lymphedema, Gene FLT4, Mut.)	111177
MPL, . (Congenital Amegakaryocytic Thrombocytopenia, CAMT, Gene MPL, Mut.)	34254
SGCB, . (Muscular Dystrophy Limb-Girdle Type 2A, Gene SGCB, Mut.)	25707
10, 11, 13, 14, 15 RET, . (Renal Hypodysplasia, Aplasia 1, Exons 10, 11, 13, 14, 15 Gene RET, Mut.)	21428
GNAS, . (Progressive Osseous Heteroplasia, POH, Gene GNAS, Mut.)	38522
(), SOD1, .	7370
1, TOR1A (DYT1), . .	3410
IRF6, . (Popliteal Pterygium Syndrome, PPS, Gene IRF6, Mut.)	38522
PTEN, . (Lhermitte-Duclos Syndrome, Gene PTEN, Mut.)	38522
FKRP, . (Muscular Dystrophy-Dystroglycanopathy, Gene FKRP, Mut.)	14916
PTEN, . (Bannayan-Ruvalcaba-Riley Syndrome, Gene PTEN, Mut.)	38522
(c). VHL, . (Autosomal Recessive Erythrocytosis, Gene VHL, Mut.)	14916
, PMP22, (Charcot-Marie-Tooth disease type 1A (CMT1A))	16577
ANKH, . (Craniometaphyseal Dysplasia, Gene ANKH, Mut.)	51348
CYBB, . (Chronic Granulomatous Disease, CGD, Gene CYBB, Mut.)	51348
LMNA, . (Hutchinson-Gilford Progeria Syndrome, Gene LMNA, Mut.)	42801
(, ,). UGT1, . (Crigler-Najjar Syndrome, Gene UGT1, Mut.)	21428
MECP2, . (Retts Syndrome, Gene MECP2, Mut.)	21428
HLA B51	3300
SRY, . (Disorders Sex Determination, Analysis Gene SRY, Mut.)	5148
SGCA, . (Muscular Dystrophy Limb-Girdle Type 2A, Gene SGCA, Mut.)	25707
(- -) II. NEFL, . (Charcot-Marie-Tooth Disease Type 2A1, Gene NEFL, Mut.)	25707
« » . (Autoimmune Lymphoproliferative Syndrome, ALPS, Gene TNFRSF6, Hot-Point Mut.)	7161
-IgD « » MVK, « » . (Hyper-IgD Syndrome, Gene MVK, Hot-Point Mut.)	9999
FKTN, . (Muscular Dystrophy Fukuyama-Type, Gene FKTN, Mut.)	51348
ESC (-). NR2E3, . (Enhanced S- one Syndrome, Goldmann-Favre Syndrome, Gene NR2E3, Mut.)	25707
IRF6, . (Van der Woude Syndrome, Gene IRF6, Mut.)	38522
RS1, . (Retinoschisis 1 X-Linked Juvenile, RS1, Gene RS1, Mut.)	25707
UNC13D, . (Familial Hemophagocytic Lymphohistiocytosis, Gene UNC13D, Mut.)	76989
(). TCIRG1, . (Osteopetrosis Autosomal Recessive 1, OPTB1, Gene TCIRG1, Mut.)	51348
CSTB, . (Progressive Myoclonic Epilepsy 1A Unverricht and Lundborg, Gene CSTB, Mut.)	14267

Mut)	SH2D1A, . (X-Linked Lymphoproliferative Syndrome, XLP, Gene SH2D1A, Mut.)	17160
Dystrophy, All Known Mutations, Gene BEST1, Mut.)	BEST1, . (Best Vitelliform Macular Dystrophy, All Known Mutations, Gene BEST1, Mut.)	42801
	TWIST1, . (Craniosynostosis Type 2, Gene TWIST1, Mut.)	14916
	MSX2, . (Craniosynostosis Type 2, Gene MSX2, Mut.)	11264
PRNP, . (Spongiform Encephalopathy with Neuropsychiatric Features, Gene PRNP, Mut.)	PRNP, . (Spongiform Encephalopathy with Neuropsychiatric Features, Gene PRNP, Mut.)	14916
Mut.)	KRT2, . (Ichthyosis Bullosa Of Siemens, Gene KRT2, Mut.)	29975
RAB23, . (Carpenter Syndrome, Gene RAB23, Mut.)	RAB23, . (Carpenter Syndrome, Gene RAB23, Mut.)	29975
(Jackson-Weiss Syndrome, JWS, Exon 9 Gene FGFR2, Exon 7A Gene FGFR1, Mut.)	9 FGFR2 7A FGFR1, . (Jackson-Weiss Syndrome, JWS, Exon 9 Gene FGFR2, Exon 7A Gene FGFR1, Mut.)	9999
Gene ABCC6, Mut.)	ABCC6, . (Pseudoxanthoma Elasticum, Gene ABCC6, Mut.)	119724
	4, SPAST (SPG4), . .	6930
VHL, . (Von Hippel-Lindau Syndrome, VHL, Von Hippel-Lindau Hereditary Cancer Syndrome, Gene VHL, Mut.)	VHL, . (Von Hippel-Lindau Syndrome, VHL, Von Hippel-Lindau Hereditary Cancer Syndrome, Gene VHL, Mut.)	14916
	HLA-Cw6	3300
		5599
Syndrome, AS, Gene FGFR2, Freq. Mut.)	FGFR2, . . (Apert Syndrome, AS, Gene FGFR2, Freq. Mut.)	10285
	LDLR, APOB, PCSK9)	9218
Epiphyseal Dysplasia, MED, Gene SLC26A2, Mut.)	SLC26A2, . (Multiple Epiphyseal Dysplasia, MED, Gene SLC26A2, Mut.)	25707
	PRPS1, . (Art's Syndrome, Gene PRPS1, Mut.)	29975
(Nonbullous Congenital Ichthyosiform Erythroderma, NBCIE, Gene ALOXE3, Mut.)	ALOXE3, . (Nonbullous Congenital Ichthyosiform Erythroderma, NBCIE, Gene ALOXE3, Mut.)	64163
	PAX3, . (Craniofacial-Deafness-Hand Syndrome, CDHS, Gene PAX3, Mut.)	34254
Muscular Dystrophy, Gene LMNA, Mut.)	LMNA, . (Emery-Dreifuss Muscular Dystrophy, Gene LMNA, Mut.)	42801
VHL, . . (Autosomal Recessive Erythrocytosis, Gene VHL, Freq. Mut.)	VHL, . . (Autosomal Recessive Erythrocytosis, Gene VHL, Freq. Mut.)	5148
(Keratitis-Ichthyosis-Deafness Syndrome, KID Syndrome, Gene GJB2, Mut.)	GJB2, . (Keratitis-Ichthyosis-Deafness Syndrome, KID Syndrome, Gene GJB2, Mut.)	9999
« . » . . (Familial Partial Lipodystrophy 2, FPLD 2, Gene LMNA, Hot-Point Mut.)	LMNA, « . » . . (Familial Partial Lipodystrophy 2, FPLD 2, Gene LMNA, Hot-Point Mut.)	14267
	TAZ, . (Barth Syndrome, Gene TAZ, Mut.)	25707
(Cerebrooculofacioskeletal Syndrome, COFS Syndrome, Gene ERCC6, Mut.)	ERCC6, . (Cerebrooculofacioskeletal Syndrome, COFS Syndrome, Gene ERCC6, Mut.)	94083
VHL, . (Von Hippel-Lindau Syndrome, VHL, Von Hippel-Lindau Hereditary Cancer Syndrome, Gene VHL, Copy Number Variation Gene VHL, Mut.)	VHL, . (Von Hippel-Lindau Syndrome, VHL, Von Hippel-Lindau Hereditary Cancer Syndrome, Gene VHL, Copy Number Variation Gene VHL, Mut.)	16577
	EXT2, . (Multiple Exostoses, Gene EXT2, Mut.)	64163
ANO5, SGCA	CAPN3, FKRP,	12056
3A, Gene CYP1B1, Mut.)	CYP1B1, . (Primary Congenital Glaucoma 3A, PCG 3A, Gene CYP1B1, Mut.)	17523
(Normokalemic Periodic Paralysis, Exon 13 Gene SCN4A, Mut.)	13 SCN4A, . (Normokalemic Periodic Paralysis, Exon 13 Gene SCN4A, Mut.)	7161
	ATP7B, PNPLA3, SERPINA1, . .	9240
	GLI3, . (Polydactyly, Gene GLI3, Mut.)	76989
-IgM Mut.)	CD40LG, . (Hyper-IgM Syndrome, Gene CD40LG, Mut.)	21428
HPGD, . (Hypertrophic Osteoarthropathy, Primary, Autosomal Recessive, 1, Gene HPGD, Mut.)	HPGD, . (Hypertrophic Osteoarthropathy, Primary, Autosomal Recessive, 1, Gene HPGD, Mut.)	29975
Gene CLCN1, Freq. Mut.)	CLCN1, . . (Myotonia Congenita, Gene CLCN1, Freq. Mut.)	10285
	, C9orf72, . .	4620

And Diabetes Syndrome, Gene HNF1B, Mut.)	HNF1B, . (Renal Cysts	38522
SH3TC2, FIG4, FGD4 GDAP1, . . (Charcot-Marie-Tooth Disease Type 1B, Gene GDAP1, Freq. Mut.)	(. . .) l.	10285
Gene RPS6KA3, Mut.)	RPS6KA3, . (Coffin-Lowry Syndrome,	94083
	SHH, . (Polydactyly, Gene SHH, Mut.)	9999
	, . PNPLA3, . .	3190
TAR. TAR-Syndrome, Gene RBM8A, Mut.)	RBM8A, . (Thrombocytopenia-Absent Radius Syndrome,	25707
	PTEN, . (Cowden Syndrome 1, Gene PTEN, Mut.)	38522
	, 2, CNBP (ZNF9), . .	3080
(Chondrodysplasia Punctata, CDP, Conradi-Hunermann Syndrome, Gene EBP, Mut.)	EBP, .	17160
Hemophagocytic Lymphohistiocytosis, Gene STXBP2, Mut.)	STXBP2, . (Familial	51348
(. . .). Autosomal Recessive, Gene LPIN1, Mut.)	LPIN1, . (Myoglobinuria Acute Recurrent	94083
Gene ADAMTSL2, Mut.)	ADAMTSL2, . (Geleophysic Dysplasia 1,	76989
(Neurodegeneration With Brain Iron Accumulation 1, Gene PANK2, Freq. Mut.)	PANK2, . .	7161
	GJB4, . (Erythrokeratoderma, Gene GJB4, Mut.)	9999
(. . .). II, De la Chapelle Dysplasia, Gene SLC26A2, Mut.)	SLC26A2, . (Atelosteogenesis	25707
DBA1, Gene RPS19, Mut.)	RPS19, . (Diamond-Blackfan Anemia 1,	21428
NPHS1, Gene NPHS2, Mut.)	NPHS2, . (Nephrotic Syndrome Type 1,	34254
IX, Mut.)	IX B, . (Hemophilia B, Gene Factor	29975
Syndrome, TRPS, Gene TRPS1, Mut.)	TRPS1, . (Trichorhinophalangeal	42801
Muscular Atrophy (SMA) with Diaphragmatic Paralysis, Gene IGHMBP2, Mut.)	IGHMBP2, . (Spinal	64163
Freq. Mut.)	FGFR3, . . (Hypochondroplasia, Gene FGFR3,	12056
(. . . 1 , 1b)	NS3, NS5A NS5B	12837
. (Hyperkalemic Periodic Paralysis Type 2, Exons 13, 24 Gene SCN4A, Mut.)	13 24 SCN4A,	15697
DFNB1	GJB2	6820
MET		12331
(Hereditary Breast and/or Ovarian Cancer, HBOC (Genes BRCA1, BRCA2)	/ (BRCA1, BRCA2)	4785
POLE		8327
1 /19q		11165
		10010
Neoplasia Type 2B (Gene RET))	2B (RET) (Multiple Endocrine	5148
BRCA- (BRCA1, BRCA2) (. . .) (Hereditary Breast Cancer In Men: Cancer of Breast, Pancreatic, Prostate, Testicular Cancer (Genes BRCA1, BRCA2) (without Description))	(BRCA1, BRCA2) (. . .)	4290
Medullary Thyroid Cancer (Exons 10, 11, 13, 14, 15 Gene RET))	(10, 11, 13, 14, 15 RET) (Familial	21406
Endocrine Neoplasia Type 2A (Exons 10, 11 Gene RET))	2A (10, 11 RET) (Multiple	9999
IDH2		11682
		23199

14	JAK2 (Quantification of wild-type and mutant allelic ratio of gene JAK2 617V/617F)	617V/617F	8800
	BRCA- (BRCA1, BRCA2) (Hereditary Breast Cancer In Men: Cancer of Breast, Pancreatic, Prostate, Testicular Cancer (Genes BRCA1, BRCA2))		4785
	MGMT		11682
	PIK3CA		11682
	228 250 TERT		8327
	(Genes BRCA1, BRCA2) (without Description)	(BRCA1, BRCA2) (Hereditary Breast and/or Ovarian Cancer, HBOC)	4290
	IDH1		11682
A09.05.039	(, L- , +) (Lactate Dehydrogenase, LDH)		165
	G6PD		3278
A09.05.042	() (Alanine Aminotransferase, ALT, Serum Glutamic Pyruvic Transaminase, SGPT)		165
A09.05.045	(?- ,) (Ipha- milase, ?-Amylase)		231
A09.05.180	(P-) (Pancreatic ?-Amylase)		275
	(S- , II, S-) (Cholinesterase, Pseudocholinesterase, PCHE)		253
A09.05.046	() (Alkaline Phosphatase, ALP)		165
A09.05.173	() (Lipase)		319
	() (Acid Phosphatase, ACP)		220
	(,) (Gamma-Glutamyl Transferase, GGT)		165
A09.05.043	(, ,) (Creatine Kinase, CK, Creatine Phosphokinase, CPK)		264
A09.05.041	(,) (Aspartateaminotransferase, AST, Serum Glutamicoxaloacetic Transaminase, SGOT)		165
	(, - , - , -) (Creatine Kinase-MB, CK-MB, Creatine Phosphokinase-MB, CPK-MB.)		341
HLA-			
	HLA II (DRB1, DQA1, DQB1) (System Human Leukocyte Antigen (HLA) Class II, Typing (Genes DRB1, DQA1, DQB1))		5841
	(Genotype of RH factor Definition (without Description))		9768
	Y- (Y-chromosome of the fetus in the mother's blood)		4510
	(Rh factor Definition)		6820
	(Plasminogen)		660
	VIII () (Antihemophilic Globulin A, FVIII)		1133
A09.05.051.001	D- (D-Dimer)		1133
A09.05.050	(Fibrinogen, FG)		231
A12.05.039	() (Activated Partial Thromboplastin Time, APTT)		165
	IX, , % (« » Factor IX, Activity, % (Christmas Factor, anti-hemophilic globulin "B"))		484
A12.05.027	(,) (Prothrombin, rothrombin Time, PT, International Normalized Ratio, INR)		253
A12.05.028	() (Thrombin Time, TT)		253
A09.05.029.001	() (Lupus Anticoagulant, LA)		803
	C, % (Protein C, % Activity)		1969
	(, /) (, /), Anti-Xa activity, IU/ml (Heparin concentration, IU/ml)		1837

	, % (Willebrand Factor, Antigen, %)	2233
A09.05.047	III, % (III, Antithrombin III, % Activity)	363
S	(Protein S, Free)	2442
	(Urine immunoglobulin free light chains (FLC) kappa and lambda)	1474
	(M-Gradient, Screening. Serum Protein Electrophoresis (SPEP), Immunofixation with Polyvalent Antiserum, Quantification of M-Protein (without Typing))	2376
	(Bence-Jones Protein, Urine, Electrophoresis, Immunofixation, Kappa/Lambda Typing, Quantification)	3300
A09.05.014	(Serum Protein Electrophoresis, SPE, SPE)*	275
	(M-Gradient, Typing. Serum Protein Electrophoresis (SPEP), Immunofixation with Antisera (IgG, IgA, IgM, Kappa, Lambda), Quantification of M-Protein)	4389
	(Cerebrospinal Fluid Concentration of Immunoglobulin Free Light Chains)	1738
	/	2156
	(Bence-Jones Protein, Urine, Immunofixation, Quantification)	2068
A09.05.011	(Albumin)	231
A09.05.010	(Protein Total)	165
A09.05.214	(Homocysteine)	1419
	(Urine Protein Electrophoresis)	1650
ImmunoCAP		
	(f216) IgE, ImmunoCAP	693
	(f33) IgE, ImmunoCAP	693
	, nArtv1 (w231) IgE, ImmunoCAP	2013
	(f9) IgE, ImmunoCAP	693
	(f260), IgE, ImmunoCAP (Broccoli, Brassica oleracea (f260), IgE, ImmunoCAP)	792
	(f26) IgE, ImmunoCAP	693
	() (i6) IgE, ImmunoCAP	693
	(f343), IgE, ImmunoCAP (Raspberry, Rubus idaeus, IgE, ImmunoCAP)	792
	(f35) IgE, ImmunoCAP	693
	, nGal d3 (f323) IgE, ImmunoCAP	913
	(Hollister-Stier) (hx2) IgE, ImmunoCAP	1375
	Candida albicans (m5) IgE, ImmunoCAP	693
	(i1) IgE, ImmunoCAP	693
	(i3) IgE, ImmunoCAP	693
	(c8) IgE, ImmunoCAP	693
	(f95) IgE, ImmunoCAP	693
	(f6) IgE, ImmunoCAP	792
	(Saccharomyces cerevisiae) (f45) IgE, ImmunoCAP	693
	, nGal d1 (f233) IgE, ImmunoCAP	913
	, rAra h 2 (f423) IgE, ImmunoCAP	2013
	(e81) IgE, ImmunoCAP	693
	/ D. pter nyssinus (d1) IgE, ImmunoCAP	693
	, rBet v1/PR-10 (t215) IgE, ImmunoCAP	2013

(f23) IgE, ImmunoCAP	693
, nGal d2 (f232) IgE, ImmunoCAP	913
- , (nBos d5) (f77) IgE, ImmunoCAP	913
- (nBos d4) (f76) IgE, ImmunoCAP	913
(f92) IgE, ImmunoCAP	693
(f91) IgE, ImmunoCAP	693
, (e6) IgE, ImmunoCAP	693
, ImmunoCAP	3025
, rGly m 4/PR-10 (f353) IgE, ImmunoCAP	2013
, (e213) IgE, ImmunoCAP	693
(f75) IgE, ImmunoCAP	693
, (e1) IgE, ImmunoCAP	693
, rAra h 1 (f422) IgE, ImmunoCAP	2013
(Hollister –Stier) (h2) IgE, ImmunoCAP	693
c, rAra h 9 LTP (f427) IgE, ImmunoCAP	2013
, rCan f 1 (e101) IgE, ImmunoCAP	2013
(f83) IgE, ImmunoCAP	693
(w5) IgE, ImmunoCAP	693
Malassezia spp. (m227) IgE, ImmunoCAP	693
, rCan f 2 (e102) IgE, ImmunoCAP	2013
/ (k80) IgE, ImmunoCAP	693
(f31) IgE, ImmunoCAP	693
(f14) IgE, ImmunoCAP	693
(Greer Labs.) (h1) IgE, ImmunoCAP	693
c (k82) IgE, ImmunoCAP	693
Cladosporium herbarum (m2) IgE, ImmunoCAP	693
(w8) IgE, ImmunoCAP	792
(mx2) IgE, ImmunoCAP	1375
(f24) IgE, ImmunoCAP	693
, rPen a1(f351) IgE, ImmunoCAP	2013
(f209) IgE, ImmunoCAP	693
(i71) IgE, ImmunoCAP	693
(f2) IgE, ImmunoCAP	693
() (f55) IgE, ImmunoCAP	693
(f210) IgE, ImmunoCAP	693
(f13) IgE, ImmunoCAP	693
, nArtv3 (w233) IgE, ImmunoCAP	2013
-5 , rTri a 19 (f416) IgE, ImmunoCAP	2013
(fx15) IgE, ImmunoCAP	1375
, (e5) IgE, ImmunoCAP	693
(fx73) IgE, ImmunoCAP	1375
(fx5) IgE, ImmunoCAP	1375
(w204) IgE, ImmunoCAP	792
(f49) IgE, ImmunoCAP	693
Penicillium notatum (P.chrysogenum) (m1) IgE, ImmunoCAP	693
(f25) IgE, ImmunoCAP	693

, rCyp c 1 (f355) IgE, ImmunoCAP	2013
(i75) IgE, ImmunoCAP	693
(gx1) IgE, ImmunoCAP	1375
, rBet v2, rBet v4 (t221) IgE, ImmunoCAP	2013
(f11), IgE, ImmunoCAP	693
(m80) IgE, ImmunoCAP	693
Phadiatop ImmunoCAP, IgE	1727
, rPhl p1, rPhl p5 (g213) IgE, ImmunoCAP	2013
(f302) IgE, ImmunoCAP	792
, (nBos d8) (f78) IgE, ImmunoCAP	913
(mx1) IgE, ImmunoCAP	1375
V (c2) IgE, ImmunoCAP	693
(f227) IgE, ImmunoCAP	792
(tx9) IgE, ImmunoCAP	1375
(f12), IgE, ImmunoCAP (Pea, Pisum sativum, IgE, ImmunoCAP)	792
(f20) IgE, ImmunoCAP	792
() (f212), IgE, ImmunoCAP (Mushrooms, Agaricus hortensis, IgE, ImmunoCAP)	792
/ D. farina (d2) IgE, ImmunoCAP	693
(f88) IgE, ImmunoCAP	693
(f94), IgE, ImmunoCAP (Pear, Pyrus communis, IgE, ImmunoCAP)	792
(f17) IgE, ImmunoCAP	693
(f4) IgE, ImmunoCAP	693
(f208) IgE, ImmunoCAP	693
, nBos d6 BSA (e204) IgE, ImmunoCAP	2013
(f262), IgE, ImmunoCAP (Eggplant, Solanum melongena, IgE, ImmunoCAP)	792
(f221), IgE, ImmunoCAP (Coffee, Coffea spp., IgE, ImmunoCAP)	792
(f222) IgE, ImmunoCAP	792
(vx1) IgE, ImmunoCAP	1375
(f322) IgE, ImmunoCAP	792
(f93) IgE, ImmunoCAP	693
, rPhl p7, rPhl p12 (g214) IgE, ImmunoCAP	2013
, rFel d1 (e94) IgE, ImmunoCAP	2013
(f27) IgE, ImmunoCAP	693
(f84) IgE, ImmunoCAP	693
, nGal d4 (k208) IgE, ImmunoCAP	913
, nAmb a1 (w230) IgE, ImmunoCAP	2013
(w6) IgE, ImmunoCAP	693
(f48), IgE, ImmunoCAP (Onion, Allium cepa, IgE, ImmunoCAP)	792
G (c1) IgE, ImmunoCAP	693
(g6) IgE, ImmunoCAP	693
(t3) IgE, ImmunoCAP	693
, rFel d2 (e220) IgE, ImmunoCAP	2013
B (m81) IgE, ImmunoCAP	693
(f237), IgE, ImmunoCAP (Apricot, Prunus armeniaca, IgE, ImmunoCAP)	792
(f242), IgE, ImmunoCAP (Cherry, Prunus avium, IgE, ImmunoCAP)	792
Ig E ImmunoCAP	759

TSST (m226) IgE, ImmunoCAP	693
Alternaria alternata (m6) IgE, ImmunoCAP	693
Aspergillus fumigatus (m3) IgE, ImmunoCAP	693
(f47) IgE, ImmunoCAP	792
Phadiatop Infant ImmunoCAP, IgE	2189
, rAra h 3 (f424) IgE, ImmunoCAP	2013
(f1) IgE, ImmunoCAP	693
c (f7) IgE, ImmunoCAP	693
c, rAra h 8/PR-10 (f352) IgE, ImmunoCAP	2013
(f85) IgE, ImmunoCAP	693
, (f44) IgE, ImmunoCAP	693
(f225) IgE, ImmunoCAP	693
(i2) IgE, ImmunoCAP	693
(f300) IgE, ImmunoCAP	792
(f3) IgE, ImmunoCAP	693
, (e85) IgE, ImmunoCAP	693
, nCan f3 (e221) IgE, ImmunoCAP	2013
(w206) IgE, ImmunoCAP	792
Alternaria alternata, rAlt a 1 (m229) IgE, ImmunoCAP	2013
- HLA-B27 (Molecular Genetic Testing HLA-B27)	1705
() (Anti- ydic Citrullinated Peptide, anti-CCP)	1452
IgG () (Anti- eratin ntibodies, AKA, Anti-Filaggrin ntibodies, AFA, IgG)	2244
, IgA (IgA; Rheumatoid Factor, RF, IgA)	1232
() (Synovial Fluid Smear, Crystals)	1705
IgG (-M) (Anti-Mutated Citrullinated Vimentin Antibodies, Anti-MCV, Anti-Modified Citrullinated Vimentin Antibodies, Anti-Sa Antibodies, IgG)	1518
(CMV DNA, Scrape of Nasal Epithelial Cells)*	275
(CMV DNA, Scrape of Faucial Epithelial Cells)*	275
(CMV DNA, Blood)*	418
(CMV DNA, Exudate)*	275
(CMV DNA, Serum)*	418
A IgM (Anti-CMV IgM)	539
(CMV DNA, Prostatic Fluid, Semen)*	275
(CMV DNA, Scrape of Skin Epithelial Cells)*	275
(CMV DNA, Urine)*	275
(CMV DNA, Scrape of Urogenital Epithelial Cells)*	275
(CMV DNA, Saliva)*	275
(Cytomegalovirus, DNA)	407
Anti-CMV IgG	1144
A IgG (Anti-CMV IgG)	396

DNA, Scrape of Conjunctiva Epithelial Cells)*	(CMV	275
Fluid)*	(CMV DNA, Cerebrospinal	275
(Copper, random urine; Cu)		
(Cd)	(Cadmium (Cd), Urine)	1298
(Co)	(Cobalt (Co), Urine)	1298
(Tl)	(Thallium (Tl), Urine)	1298
(I)	(Iodine (I), Urine)	1298
(Mn)	(Manganese (Mn), Urine)	1298
(Fe)	(Iron (Fe), Urine)	1298
(Hg)	(Mercury (Hg), Urine)	1298
(Al)	(Aluminum (Al), Urine)	1298
()	()	748
(Zn)	(Zinc (Zn), Urine)	1298
(Cu)	(Copper (Cu), 24-Hours Urine)	1298
(Ni)	(Nickel (Ni), Urine)	1298
(Se)	(Selenium (Se), Urine)	1298
()	()	748
(Pb)	(Lead (Pb), Urine)	1298
(As)	(Arsenic (As), Urine)	1298
IgG (anti-Tetanus toxoid IgG)		
()	()	990
()		
(Ni)	(Nickel (Ni), Serum)	275
(Hg)	(Mercury (Hg), blood)	1298
()	()	748
(As)	(Arsenic (As), Serum)	275
(Cd)	(Cadmium (Cd), Serum)	275
(Co)	(Cobalt (Co), Serum)	275
(Ni)	(Nickel (Ni), blood)	1298
(Zn)	(Zinc (Zn), blood)	1298
(I)	(Iodine, serum)	275
(Au)	(Gold (Au), Serum)	275
(Cu)	(Copper (Cu), Serum)	275
()	()	748
(Cu)	(Copper (Cu), blood)	1298
(Pb)	(Lead (Pb), blood)	1298
(Mn)	(Manganese (Mn), blood)	1298
(Zn)	(Zinc (Zn), Serum)	275
(Li)	(Lithium (Li), serum)	275
(Cd)	(Cadmium (Cd), blood)	1298
(Se)	(Selenium (Se), Serum)	275
(Se)	(Selenium (Se), blood)	1298
(Mn)	(Manganese (Mn), Serum)	275
(Co)	(Cobalt (Co), blood)	1298

	(3 , Reverse Triiodthyronine).	6567
(Tl)	(Thallium (Tl), Serum)	275
	()	748
(Mo)	(Molybdenum (Mo), Serum)	275
quantitative))	ATM (FISH, .) (Analysis of ATM gene rearrangements (FISH,	10835
qualitative)	MPL, (Analysis of MPL gene mutations, PCR,	4829
(Analysis of BCL- 6 gene rearrangements (der(3)(q27)) on paraffin slides (FISH Histology, quantitative))	BCL- 6 (der(3)(q27)) (FISH, .)	14927
(PCR, qualitative))	MLL/AF4 -t(4;11) (, .) (Analysis of chimeric gene MLL/AF4 -t(4;11)	2486
rearrangements (der(3)(q27) (FISH, quantitative))	BCL- 6 (der(3)(q27)) (FISH, .) (Analysis of BCL- 6 gene	10835
chromosome 13 monosomy, deletion - (del(13), -13) (FISH, quantitative))	13 - (del(13), -13) (FISH, .) (Analysis of	10835
53 (FISH, .) (Analysis of 53 gene deletion (FISH, quantitative))	53 (FISH, .) (Analysis of 53 gene deletion (FISH, quantitative))	10835
translocation t(2;5)(p23;q35) on paraffin slides (FISH Histology, quantitative))	t(2;5)(p23;q35) (FISH, .) (Analysis of	14927
all specific aberrations on paraffin slides (FISH Histology, quantitative))	(FISH, .) (Analysis of	14927
translocation t(11;14)(q13;q32) on paraffin slides (FISH Histology, quantitative))	t(11;14)(q13;q32) (FISH, .) (Analysis of	14927
(FISH, quantitative))	t(11;14)(q13;q32) (FISH, .) (Analysis of translocation t(11;14)(q13;q32)	10835
-t(15;17) (PCR, qualitative))	PML/RAR? -t(15;17) (, .) (Analysis of chimeric gene PML/RAR?	2486
CBF?/MYH1- inv(16),t(16;16) (PCR, qualitative))	CBF?/MYH1- inv(16),t(16;16) (, .) (Analysis of chimeric gene	2486
12 (+12) (FISH, .) (Analysis of chromosome 12 trisomy (FISH, quantitative))	12 (+12) (FISH, .) (Analysis of chromosome 12 trisomy (FISH,	10835
assessment of presence of gene JAK2 617F somatic mutation)	V617F 14 JAK2 (Qualitative	1947
(FISH, quantitative))	t(11;18)(q21;q21) (FISH, .) (Analysis of translocation t(11;18)(q21;q21)	10835
	(Karyotype, Hematologic Disorders, Peripheral Blood)	7722
53 (FISH, .) (Analysis of 53 gene deletion (FISH, quantitative))	53 (FISH, .) (Analysis of 53 gene deletion (FISH, quantitative))	10835
(IGH/MAFB) (FISH, quantitative))	t(14;16) (IGH/MAFB) (FISH, .) (Analysis of translocation t(14;16)	10835
(Analysis of chimeric gene BCR-ABL - t(9;22), assessment of the BCR-ABL gene transcript type, PCR, qualitative)	BCR/ABL - t(9;22), BCR/ABL -	2486
12p (FISH, .) (Analysis of 12p deletion (FISH, quantitative))	12p (FISH, .) (Analysis of 12p deletion (FISH, quantitative))	10835
quantitative)	BCR-ABL (FISH, .) (Analysis of chimeric gene BCR-ABL, FISH,	10835
rearrangements on paraffin slides (FISH Histology, quantitative))	BCL2 (FISH, .) (Analysis of BCL2 gene	14927
quantitative))	PDGFR?(FISH, .) (Analysis of gene rearrangements PDGFR? (FISH,	10835
quantitative))	FGFR1 (FISH, .) (Analysis of gene rearrangements FGFR1 (FISH,	10835
t(1;19) (PCR, qualitative))	E2A/PBX1 - t(1;19) (, .) (Analysis of chimeric gene E2A/PBX1 -	2486
	BRAF (V600E) (, .)	8569
5 (FISH, .) (Analysis of chromosome 5 rearrangements (FISH, quantitative))	5 (FISH, .) (Analysis of chromosome 5 rearrangements (FISH,	10835
(FISH, quantitative))	t(4;14)(p16;q32) (FISH, .) (Analysis of translocation t(4;14)(p16;q32)	10835
(IGH/MAFB) (FISH, quantitative))	t(14;16) (IGH/MAFB) (FISH, .) (Analysis of translocation t(14;16)	10835
	BCR/ABL - RQ (4829
	.) (Analysis of the BCR/ABL relative expression, RQ-PCR, quantitative)	

marrow (karyotype)	() (Cytogenetic analysis of bone marrow)	7722
(FISH, quantitative)	FIP1L1/PDGFR? (FISH,) (Analysis of chimeric gene FIP1L1/PDGFR?)	10835
quantitative))	MLL (FISH,) (Analysis of MLL gene rearrangements (FISH, quantitative))	10835
quantitative))	7 (FISH,) (Analysis of chromosome 7 rearrangements (FISH, quantitative))	10835
chromosome 13 monosomy, deletion - (del(13), -13) (FISH, quantitative))	13 - (del(13), -13) (FISH, quantitative))	10835
qualitative))	12 JAK2 (,) (Analysis of JAK2 Exon 12 mutations (PCR qualitative))	4829
(Analysis of BCL2 gene rearrangements t(14;18)(q32;q21), t(2;18)(p11;q21), t(18;22)(q21;q11) (FISH, quantitative))	BCL2 t(14;18)(q32;q21), t(2;18)(p11;q21), t(18;22)(q21;q11) (FISH, quantitative))	10835
) (BCR-ABL1 Mutation Analysis using direct Sanger sequencing, qualitative)	BCR-ABL () (BCR-ABL1 Mutation Analysis using direct Sanger sequencing, qualitative)	9636
(FISH,)	1 (FISH,)	14267
(FISH, quantitative))	3q (FISH,) (Analysis of 3q rearrangements (FISH, quantitative))	10835
(PCR, qualitative))	RUNX1/RUNX1T1 -t(8;21) (,) (Analysis of chimeric gene RUNX1/RUNX1T1 -t(8;21) (PCR, qualitative))	2486
(FISH, quantitative))	20q (FISH,) (Analysis of 20q deletion (FISH, quantitative))	10835
(FISH, quantitative))	MYC (t(8;14)(q24;q32)-t(2;8)(p11;q24), t(8;22)(q24;q11) (FISH, quantitative)) (Analysis of MYC gene rearrangements (t(8;14)(q24;q32)-t(2;8)(p11;q24), t(8;22)(q24;q11) (FISH, quantitative))	10835
(Analysis of CALR gene mutations, deletions, insertions, PCR, qualitative)	CALR (,) (Analysis of CALR gene mutations, deletions, insertions, PCR, qualitative)	4829
(FISH, quantitative))	IGH (FISH,) (Analysis of IGH gene rearrangements (FISH, quantitative))	10835
(FISH, quantitative))	t(2;5)(p23;q35) (FISH,) (Analysis of translocation t(2;5)(p23;q35) (FISH, quantitative))	10835
:		
		1320
participant (child or mother or father))	() (Additional research participant (child or mother or father))	6303
(3) (Urgent Establishment of Biological Relationship for One Parent at Indisputable Relationship of Another (3 Persons))	(3) (Urgent Establishment of Biological Relationship for One Parent at Indisputable Relationship of Another (3 Persons))	41833
(2) (Urgent Establishment of Biological Relationship for One Parent in Absence of Another (2 Persons))	(2) (Urgent Establishment of Biological Relationship for One Parent in Absence of Another (2 Persons))	41833
(2) (Establishment of Biological Relationship for One Parent in Absence of Another (2 Persons))	(2) (Establishment of Biological Relationship for One Parent in Absence of Another (2 Persons))	17787
(3) (Establishment of Biological Relationship for One Parent at Indisputable Relationship of Another (3 Persons))	(3) (Establishment of Biological Relationship for One Parent at Indisputable Relationship of Another (3 Persons))	19888
:		
(Koprogramma, Stool)		374
		4059
		6127
(Fecal Calprotectin)		2618
		3542
-1- (Alpha-1-Antitrypsin, Feces)		1727
		1463
		253
() (PRO Stool, Helminth Eggs)		308
		1980
() (Quantitative Immunochemical Fecal Occult Blood, Test FOB Gold)		715
1 (), 1 (Elastase 1, E1)		2772

() (Stool Sugars, Reducing Substances, Fecal)		616
(PRO Stool)		308
(), (nterobiasis, Spatula)		286
(Stool osmotic gap)		1265
-10 (-10) (Interleukin 10, IL-10)		2035
-6 (-6) (Interleukin 6, IL-6)		2035
-? (-?) (Tumor Necrosis Factor Alpha, TNF-?, Cachectin)		2035
-1? (-1?) (Interleukin 1 Beta, IL-1)		2035
-8 (-8) (Interleukin 8, IL-8)		2035
IgE:		
, IgE (Food Allergy Panel, IgE)		4037
, IgE (Celery, IgE, F85)		484
, IgE (Milk, IgE, F2)		484
(f96), IgE, ImmunoCAP (Avocado, Persea americana, IgE, ImmunoCAP)		792
, IgE (Chicken Meat, IgE, F83)		484
(f244) IgE, ImmunoCAP		792
, IgE (Pineapple, IgE, F210)		484
, IgE (Grapefruit, IgE, F209)		484
- , IgE (Beta Lactoglobulin, IgE, F77)		484
, IgE (Egg Yolk, IgE, F75)		484
(g4) IgE, ImmunoCAP		792
, IgE (Shrimp, IgE, F24)		484
, IgE (Rice, IgE, F9)		484
, IgE (Strawberry, IgE, F44)		484
, IgE (Apple, IgE, F49)		484
, IgE (Pork, IgE, F26)		484
, IgE (Lamb, IgE, F88)		484
, IgE (Tomato, IgE, F25)		484
, IgE (Baker's Yeast, IgE, F45)		484
, IgE (Potato, IgE, F35)		484
, IgE (Lemon, IgE, F208)		484
, IgE (Peach, IgE, F95)		484
3: , IgE (FP73 (F26, F27, F83, F88), Food Panel: Pork, Beef, Chicken Meat, Lamb, IgE)*		1045
" 2"		1919,5
, IgE (Casein, IgE, F78)		484
, IgE (Kiwi Fruit, IgE, F84)		484
, IgE (Cabbage, IgE, F216)		484
, IgE (Codfish, IgE, F3)		484
, IgE (Wheat, IgE, F4)		484
1: , IgE (FP15 (F33, F49, F92, F95), Food Panel: Orange, Banana, Apple, Peach, IgE)*		1045
, IgE (Carrot, IgE, F31)		484
, IgE (Common Millet, IgE, F55)		484
, IgE (Banana, IgE, F92)		484
, IgE (Soybean, IgE, F14)		484

, IgE (Chocolate, IgE, F105)	484
, IgE (Egg White, IgE, F1)	484
2 Panel: Kiwi Fruit, Mango, Banana, Pineapple, IgE)* , IgE (FP50 (F84, F91, F92, F210), Food	1045
, IgE (Pumpkin, IgE, F225)	484
, IgE (Hazelnut, IgE, F17)	484
, IgE (Orange, IgE, F33)	484
, (e3) IgE, ImmunoCAP	792
(f329), IgE, ImmunoCAP (Watermelon, Citrullus lanatus, IgE, ImmunoCAP)	792
(f5) IgE, ImmunoCAP	792
, IgE (Beef, IgE, F27)	484
, IgE (Oat, IgE, F7)	484
" 1"	1919,5
, IgE (Buckwheat, IgE, F11)	484
, IgE (Brewer's Yeast, IgE, F403)	572
, IgE (Crab, IgE, F23)	484
, IgE (Peanut, IgE, F13)	484
, IgE (Mango, IgE, F91)	484

(Ca) (Calcium (Ca), air)	1298
(Co) (Cobalt (Co), air)	1298
(Zr) (Zirconium (Zr), air)	1298
(Mg) (Magnesium (Mg), air)	1298
(Be) (Beryllium (Be), air)	1298
(Pb) (Lead (Pb), air)	1298
(Mo) (Molybdenum (Mo), air)	1298
(Al) (Aluminum (Al), air)	1298
(Hg) (Mercury (Hg), air)	1298
(Mn) (Manganese (Mn), air)	1298
(Se) (Selenium (Se), air)	1298
(V) (Vanadium (V), air)	1298
(Si) (Silica (Si), air)	1298
(Rb) (Rubidium 9Rb), air)	1298
(Bi) (Bismuth (Bi), air)	1298
(P) (Phosphorus (P), air)	1298
()	748
(Cu) (Copper (Cu), air)	1298
(Sb) (Antimony (Sb), air)	1298
(I) (Iodine (I), air)	1298
(Ba) (Barium (Ba), air)	1298
(B) (Boron (B), air)	1298
(La) (Lanthanum 9La), air)	1298
(Fe) (Iron (Fe), air)	1298
(Ag) (Silver (Ag), air)	1298
(Cr) (Chromium (Cr), air)	1298
(Sr) (Strontium (Sr), air)	1298

(W)	(Tungsten, Wolframium (W), air)	1298
(Na)	(Sodium (Na), air)	1298
(Pt)	(Platinum (Pt), air)	1298
(Ge)	(Germanium (Ge), air)	1298
(As)	(Arsenic (As), air)	1298
(Cd)	(Cadmium (Cd), air)	1298
(K)	(Potassium (K), air)	1298
(Ni)	(Nickel (Ni), air)	1298
(Zn)	(Zinc (Zn), air)	1298
(Sn)	(Tin (Sn), air)	1298
(Li)	(Lithium (Li), air)	1298
(Au)	(Gold (Au), air)	1298
(Tl)	(Thallium (Tl), air)	1298
(Ga)	(Gallium (Ga), air)	1298

IgG () (Extractable Nuclear Antigen, ENA, Anti-Ribonucleoprotein Antibodies, Anti-RNP)		1232
Sc1-70, ENP-A, CENP-B, RP11, RP155, (, NOR90, Th/To, PM-Sc100, PM-Sc175, Ku, PDGFR, Ro-52), (Scleroderma (Systemic Sclerosis) Antibody Panel: Anti-Sc1-70, ENP-A, CENP-B, RP11, RP155, , NOR90, Th/To, PM-Sc100, PM-Sc175, Ku, PDGFR, Ro-52, Immunoblotting)		4411
(,), (Anti-Nuclear Antibodies, ANA, Screening)		517
, HEp-2 (, HEp-2-) (Antinuclear Antibodies, ANA, Hep-2 Substrate, ANA-Hep2, Fluorescent Anti-Nuclear Antibodies detection, FANA, iters)		1265
(Sm, RNP/Sm, SS-A (60), SS-A (52), SS-B, Scl-70, PM-Scl, PCNA, CENT-B, dsDNA, Histone, Nucleosome, Rib P, AMA-M2, Jo-1), (ANA: Anti-Sm, RNP/Sm, SS-A (60 kD), SS-A (52 kD), SS-B, Scl-70, PM-Scl, PCNA, CENT-B, dsDNA, Histone, Nucleosome, Rib P, AMA-M2, Anti-Jo-1, Immunoblotting)		3509
IgG (a), (Anti-Nuclear Antibodies, ANA, IgG, Screening)		1232
,		1089
IgG (- IgG, -) (Double-Stranded (Native) DNA IgG Antibodies, nti-dsDNA IgG)		627
, IgG		1727

B03.016.014 (Nechiporenko's Urine Test)		253
, () (Sulkowitch Urine Calcium Test)		143
B03.016.006 (Complete Urinalysis, Microscopic Examination)		253

(, Hepatitis C Virus, HCV)		
(Hepatitis C Virus (HCV) RNA, Quantitative PCR, Genotyping (Types 1, 2, 3)) (1, 2, 3)		4048
-28 (-28), (,) (Interleukin 28 Beta IL28B, Genotyping (Study of Genetic Markers Determining Effectiveness of Treatment of Chronic Hepatitis C in Interferon and Ribavirin))		792
(CITO), (HCV RNA, Plasma, Quantitative)*		23034
(), (Hepatitis C Virus (HCV) RNA, Ultrasensitive PCR)		3355
IgM IgG , (Anti-HCV Total (IgG + IgM))*		396
IgG C, (Anti-HCV IgG, Immunoblot)		5621
, (HCV RNA, Serum, Qualitative)*		693

RNA, Serum, Quantitative, PCR)*	(HCV	3454
1a 1b), 2, 3) (Hepatitis C Virus (HCV) RNA, Plasma, Genotyping, Subtypes (Types 1 (Subtypes 1a, 1b), 2, 3))*	(1 (924
Quantitative)*	(HCV RNA, Plasma,	11517
(, Staphylococcus aureus)		
(Staphylococcus aureus), (Staphylococcus aureus (Methicillin-Resistant Staphylococcus aureus – MRSA) Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)		1078
(Staphylococcus aureus), (Staphylococcus aureus (Methicillin-Resistant Staphylococcus aureus – MRSA) Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)		2145
(Staphylococcus aureus) (Staphylococcus aureus (Methicillin-Resistant Staphylococcus aureus – MRSA) Culture. Bacteria Identification)		715
(Staphylococcus aureus, (Staphylococcus aureus Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)		1078
(Staphylococcus aureus), (Staphylococcus aureus Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)		2145
(Staphylococcus aureus) (Staphylococcus aureus Culture. Bacteria Identification)		715
(Staphylococcus aureus), (Staphylococcus aureus Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		880
(Staphylococcus aureus), (Staphylococcus aureus (Methicillin-Resistant Staphylococcus aureus – MRSA) Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		880
(- 1 2 , Herpes simplex virus, HSV-1, HSV-2)		
1 2 , (HSV-1, 2 DNA, Urine)*		275
1 2 , Serum, yping)*	(HSV-1, 2 DNA,	660
1 2 , (HSV-1, 2 DNA, Scrape of Conjunctiva Epithelial Cells)*		275
1 2 , (HSV-1, 2 DNA, Scrape of Conjunctiva Epithelial Cells, yping)*		462
1 2 , (HSV-1, 2 DNA, Urine, yping)*		462
IgG 1 2 (Anti-HSV-1, 2 IgG)		484
1 2 , Blood, yping)*	(HSV-1, 2 DNA,	660
1 2 , 2 DNA, Cerebrospinal Fluid, yping)*	(HSV-1,	462
1 2 , (HSV-1, 2 DNA, Exudate)*		275
1 2 , (HSV-1, 2 DNA, Scrape of Faucial Epithelial Cells, yping)*		462
Ig 1 2 (Anti-HSV-1, 2 Ig)		506
1 2 , (HSV-1, 2 DNA, Scrape of Urogenital Epithelial Cells, yping)*		462
1 2 , yping)*	(HSV-1, 2 DNA, Saliva,	462
1 2 , yping)*	(HSV-1, 2 DNA, Exudate,	462
1 2 , (HSV-1, 2 DNA, Blood)*		418
1 2 , (HSV-1, 2 DNA, Saliva)*		275
1 2 , DNA, Scrape of Skin Epithelial Cells)*	(HSV-1, 2	275
1 2 , (HSV-1, 2 DNA, Scrape of Nasal Epithelial Cells, yping)*		462

1 2 , Cerebrospinal Fluid)*	(HSV-1, 2 DNA,	275
1 2 , Prostatic Fluid, Semen)*	(HSV-1, 2 DNA,	275
1 2 , 2 DNA, Prostatic Fluid, Semen, yping)*	(HSV-1,	462
IgG 2 (Anti-HSV-2 IgG)		605
1 2 , (HSV-1, 2 DNA, Serum)*		418
1 2 , (HSV-1, 2 DNA, Scrape of Nasal Epithelial Cells)*		275
Anti-HSV IgG		770
1 2 , (HSV-1, 2 DNA, Scrape of Urogenital Epithelial Cells)*		275
1 2 , (HSV-1, 2 DNA, Scrape of Skin Epithelial Cells, yping)*		462
IgG 1 (Anti-HSV-1 IgG)		748
1 2 , (HSV-1, 2 DNA, Scrape of Faucial Epithelial Cells)*		275
COVID-19		
SARS-CoV-2, IgM (anti-SARS-CoV-2, IgM)		759
() SARS-CoV-2 (N-, S-), SARS-CoV-2-IgG- (Post-vaccination (EpiVacCorona Vector) SARS-CoV-2 (N-, S-proteins) antibodies, IgG, qualitative)		2486
SARS-CoV-2, (Coronavirus SARS-CoV-2 RNA detection in nasopharyngeal and oropharyngeal smear)		2189
(S) (RBD) SARS-CoV-2, IgG (Anti-SARS-CoV-2, spike (S) protein (RBD), IgG, quantitative).		1419
SARS-CoV-2, (Coronavirus SARS-CoV-2 RNA detection in nasopharyngeal and oropharyngeal smear)		1419
SARS CoV-2 (S- , RBD), IgG,		1419
(S) SARS-CoV-2, IgG, (anti-SARS-CoV-2 S (spike) protein antibody, IgG, qualitative. Assessment of immunity before and after vaccination)		979
SARS-CoV-2 (), IgG, (Anti-SARS-CoV-2 (nucleocapsid protein), IgG, Abbott)		759
SARS-CoV-2 (RBD), IgG (anti-SARS-CoV-2 (RBD) IgG avidity)		759
A		
IgM Mycoplasma pneumoniae (Anti-Mycoplasma pneumoniae IgM)		550
(Mycoplasma genitalium), (Mycoplasma genitalium, DNA, Scrape of Urogenital Epithelial Cells)*		275
(Mycoplasma hominis), (Mycoplasma hominis, DNA, Scrape of Urogenital Epithelial Cells)*		275
(Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Plasma)*		429
Ig Mycoplasma pneumoniae (Anti-Mycoplasma pneumoniae IgA)		671
(Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Scrape of Faucial Epithelial Cells)*		231
A IgG Mycoplasma pneumoniae (Anti-Mycoplasma pneumoniae IgG)		550
Ig Mycoplasma hominis (nti-Mycoplasma hominis Ig)		462
(Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Sputum)*		649
(Mycoplasma hominis), (Mycoplasma hominis, DNA, Prostatic Fluid, Semen)*		275
IgG Mycoplasma hominis (nti-Mycoplasma hominis IgG)		462
(Mycoplasma genitalium), (Mycoplasma genitalium, DNA, Prostatic Fluid, Semen)*		275
(Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Saliva)*		231
(Mycoplasma hominis), (Mycoplasma hominis, DNA, Urine)*		275
IgA Mycoplasma hominis (nti-Mycoplasma hominis IgA)		671

Urine)*	(Mycoplasma genitalium),	(Mycoplasma genitalium, DNA,	275
A09.05.127	(g)	(Magnesium (Mg), Serum)	253
A09.05.032	(Ca)	(Calcium Total)	209
/ /	(+ /Potassium, Na+ /Sodium, I- /Chloride, Serum)		275
	(Ca2+, c) (Ionized Calcium, Free Calcium)	396
A09.05.033	(P)	(Phosphorus (P))	209
(((209
(Unsaturated Iron Binding Capacity, UIBC)			209
A09.05.007	(Fe)	(Iron (Fe), Serum)	209
Associated Gastritis)*	Helicobacter pylori () (Helicobacter pylori	4004
	PDGFRa		15070
1	():	1694
PD-L1	PD-L1	c	18150
clone SP263 (Ventana) antibodies).	SP263 (Ventana). (PD-L1 expression in tumor tissue by IHC using PD-L1		
	KIT		12760
Finished Histological Preparations (1 Glass + 1 Block))	(1	+ 1) (Consultation of
	BRCA1, BRCA2		8140
	(PAS-)	319
	ROS1		8800
18,19,20, 21	EGFR		10890
2,3,4	NRAS		7700
15	BRAF		5280
2,3,4	KRAS		7700
PD-L1	PD-L1	c	18150
clone SP142 (Ventana) antibodies).	SP142 (Ventana). (PD-L1 expression in tumor tissue by IHC using PD-L1		
	(MSI)		6820
	HER2 ()	12760
S-100, Melan A (MART-1), HMB-45, SOX-10 (IHC verification of malignant melanoma using assessment of the expression S-100, Melan A (MART-1), HMB-45, SOX-10)*	()	19888
	() (Pathology of skin biopsies)*	2233
Histochemical Study)*	Helicobacter pylori () (Helicobacter pylori, Mucus,	2112
	ALK		8800
	()	2420
PD-L1	PD-L1		26400
clone 22C3 (Dako) antibodies).	22C3 (Dako). (PD-L1 expression in tumor tissue by IHC using PD-L1 clone		
	PDL1		8140
	()	19888
S-100, Melan A (MART-1), HMB-45, SOX-10 (IHC verification of malignant melanoma using assessment of the expression S-100, Melan A (MART-1), HMB-45, SOX-10)			
SCC () (Squamous Cell Carcinoma Antigen, SCCA, SCCAg)		2486
-19-9 (19-9) (Carbohydrate Antigen	-19-9, Cancer Antigen-GI)	682

(Chromogranin A, CgA)	5247
- () (Carcinoembryonic Antigen, CEA)	627
CA-242 (242, CA-242) (Carbohydrate Antigen -242, Tumor Marker CA-242)	924
ROMA1	77
() (Neuron-Specific Enolase, NSE)	1342
A09.05.130 () (Prostate-Specific Antigen Total, PSA Total)*	484
UBC (8 18) (Urine Bladder Cancer Antigen, Urine Bladder Cancer, UBC)	2046
(, , -2proPSA, phi)	3190
HE4 (4) (Human Epididymis Protein 4, HE4)	1122
-2- (?-2-) (eta-2-Microglobulin, BMG, Serum)	957
A09.05.130 () (Prostate-Specific Antigen Total, PSA Total)	484
ROMA2	77
- () (?-Fetoprotein, AFP)	385
-15-3 (15-3) (Carbohydrate Antigen -15-3, Cancer Antigen -15-3)	682
-125 (125) (Carbohydrate Antigen -125, Cancer Antigen -125)	638
A09.05.130.001 ()*	484
CA-72-4 (72-4) (Carbohydrate Antigen -72-4, Cancer Antigen CA-72-4)	957
(Cyfra 21-1, 19) (Cytokeratin 19 Fragments, C-terminus of Cytokeratin 19, CK19 Soluble Fragments, Cyfra 21-1)	946
-2- (?-2-) (Beta-2-Microglobulin, Urine)	957
S100 (S100 rotein)	2739
()	
(Trichomonas vaginalis, DNA, Urine)*	275
Prostatic Fluid, Semen)* (Trichomonas vaginalis, DNA,	275
(Trichomonas vaginalis, DNA, Scrape of Urogenital Epithelial Cells)*	275
IgG Trichomonas vaginalis (nti-Trichomonas vaginalis IgG)	671
(Procedural services)	
examinations)** (Blood serum derivation without further	320
	350
(Blood sampling without further examinations)**	190
(Buccal epithelium sampling, 2 persons)	180
(Cytological material sampling, PCR diagnosis material sampling, microbiology test material sampling)	350
(Saliva sampling for PCR)	50
A11.12.009	200
(Biomaterial sampling for the enterobiasis test)	77
(Buccal epithelium sampling, 3 persons)	270
() (capillary blood sampling)	190
(Buccal epithelium sampling)	90
C	
6 (1 ,1b,2,3 ,4,5 ,6)	2618
(Ureaplasma parvum, Effectiveness Monitoring of Treatments) (Ureaplasma parvum)	
	396

(INBIOFLOR ? Mycoplasma, Urogenital Screening)	()	528
(INBIOFLOR-Comprehensive Study of Microflora Composition of Urogenital Tract (UGT))		3058
(Bacterial Vaginosis, BV)		1694
MICROBIOCENOSIS (PCR Panel Femoflor 8)	8. (UROGENITAL TRACT	1628
(4 +): Chlamydia trachomatis, Neisseria gonorrhoeae, Trichomonas vaginalis, Mycoplasma genitalium, (Identification of Sexually Transmitted Infections (STI) Pathogens, Chlamydia trachomatis DNA, Neisseria gonorrhoeae DNA, Trichomonas vaginalis DNA, Mycoplasma genitalium DNA, Human DNA)		1023
(Mobiluncus curtisii, DNA, Scrape of Urogenital Epithelial Cells)		231
(Mycoplasma hominis, Effectiveness Monitoring of Treatments)	(Mycoplasma hominis)	396
		1870
(Ureaplasma ur alyticum) (Ureaplasma ur alyticum, Effectiveness Monitoring of Treatments)	(Ureaplasma	396
(7 +), (Identification of Sexually Transmitted Infections (STI) Pathogens, Scrape of Urogenital Epithelial Cells)*		1485
MICROBIOCENOSIS (PCR Panel Femoflor 16))	16. (UROGENITAL TRACT	2310
		2200
MICROBIOCENOSIS, Screening (PCR Panel Femoflor Screen))	. (UROGENITAL TRACT	2035
		1364
		4202

(As)	(Arsenic (As), Nails)	1298
(Pb)	(Lead (Pb), Nails)	1298
(Sb)	(Antimony (Sb), Nails)	1298
(Cr)	(Chromium (Cr), Nails)	1298
(Sn)	(Tin (Sn), Nails)	1298
(Hg)	(Mercury (Hg), Nails)	1298
(Fe)	(Iron (Fe), Nails)	1298
(Cd)	(Cadmium (Cd), Nails)	1298
(Ge)	(Germanium (Ge), Nails)	1298
(Mo)	(Molybdenum (Mo), Nails)	1298
(P)	(Phosphorus (P), Nails)	1298
()		748
(V)	(Vanadium (V), Nails)	1298
(Bi)	(Bismuth (Bi), Nails)	1298
(Ca)	(Calcium (Ca), Nails)	1298
(La)	(Lanthanum (La), Nails)	1298
(Cu)	(Copper (Cu), Nails)	1298
(Al)	(Aluminum (Al), Nails)	1298
(Au)	(Gold (Au), Nails)	1298
(Ga)	(Gallium (Ga), Nails)	1298
(Se)	(Selenium (Se), Nails)	1298
(Li)	(Lithium (Li), Nails)	1298
(Zn)	(Zinc (Zn), Nails)	1298
(B)	(Boron (B), Nails)	1298
(Ba)	(Barium (Ba), Nails)	1298

(I)	(Iodine (I), Nails)	1298
(Be)	(Beryllium (Be), Nails)	1298
(K)	(Potassium (K), Nails)	1298
(Co)	(Cobalt (Co), Nails)	1298
(W)	(Tungsten, Wolframium (W), Nails)	1298
(Tl)	(Thallium (Tl), Nails)	1298
(Si)	(Silica (Si), Nails)	1298
(Ni)	(Nickel (Ni), Nails)	1298
(Mg)	(Magnesium (Mg), Nails)	1298
(Sr)	(Strontium (Sr), Nails)	1298
(Rb)	(Rubidium (Rb), Nails)	1298
(Na)	(Sodium (Na), Nails)	1298
(Mn)	(Manganese (Mn), Nails)	1298
(Ag)	(Silver (Ag), Nails)	1298
(Zr)	(Zirconium (Zr), Nails)	1298
(Pt)	(Platinum (Pt), Nails)	1298

		6710
24-h urine	(Estrogens and progesterone metabolites,	2640
	(Melatonin, plasma)	1474

--	--	--

39, 45, 51, 52, 56, 58, 59, 66, 68 +	14 : 16, 18, 31, 33, 35, (HPV DNA, Scrape of Rectal Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68) Screening)*	385
4	: 6, 11, 16, 18 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 4 Types (6, 11, 16, 18) Screening)	605
18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68 +	14 : 16, (HPV DNA, Scrape of Urogenital Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68))	385
39, 45, 51, 52, 56, 58, 59, 66, 68 +	14 : 16, 18, 31, 33, 35, (HPV DNA, Scrape of Urogenital Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68) Screening)*	385
	3 : 6, 11, 44 + (HPV DNA, Scrape of Rectal Epithelial Cells, 3 Types (6, 11, 44))	385
	16 18 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 2 Types (16, 18))	385
	3 : 6, 11, 44 (HPV DNA, Scrape of Faucial Epithelial Cells, 3 Types (6, 11, 44))	385
31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68 +	() 14 : 16, 18, (HPV DNA, Scrape of Urogenital Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68) Screening)	990
	3 : 6, 11, 44 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 3 Types (6, 11, 44))	385
39, 45, 51, 52, 56, 58, 59, 66, 68 +	14 : 16, 18, 31, 33, 35, (HPV DNA, Scrape of Faucial Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68) Screening)*	385

o 21 : 6, 11, 16, 18, 26, 31, 33, 35, 39, 44, 45, 51, 52, 53, 56, 58, 59, 66, 68, 73, 82 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 21 Types (6, 11, 16, 18, 26, 31, 33, 35, 39, 44, 45, 51, 52, 53, 56, 58, 59, 66, 68, 73, 82))		2695
o 14 : 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68) Screening)*		385
(Candida albicans, DNA, Exudate)*		275
(Candidiasis, Typing)		737
(Candida, Cryptococcus) (Yeast Culture. Identification and Antimycotic Susceptibility testing)		759
(Candidiasis, Screening)		396
(Candida albicans, DNA, Prostatic Fluid, Semen)*		275
(Candida albicans, DNA, Scrape of Rectal Epithelial Cells)*		275
(Candida albicans, DNA, Scrape of Faucial Epithelial Cells)*		275
Candida albicans, IgG (M5) (M5 Candida albicans, IgG)		572
(Candidiasis, Screening and Typing)		1012
A IgG Candida albicans (Anti-Candida albicans IgG)		781
(Candida albicans, DNA, Urine)*		275
(Candida albicans, DNA, Scrape of Skin Epithelial Cells)*		275
(Candida albicans, DNA, Scrape of Urogenital Epithelial Cells)*		275
(Candida albicans, DNA, Saliva)*		275
()		1837
(Upper Respiratory Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*		2178
(Eye Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*		1474
(Eye Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*		2464
(Stool Culture with Bacteria Identification and Antibiotic+Bacteriophage Susceptibility Testing)		1606
(Stool Culture, Pathogenic Intestinal and Conditionally Pathogenic Microflora. Bacteria Identification and Antibiotic Susceptibility Testing)		1518
(Breast Milk Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*		1078
()		1727
(Ear Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Ttesting)*		2464
(Eye Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*		1276
(Breast Milk Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*		2145
(Punctate Fluid Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		913

(Wound/Pus/Aspirate/Tissue Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*	1078
(Bile Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	1144
(Anaerobic Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	1540
(Sputum and Tracheobronchial washings Culture. Bacteria Identification and Antibiotic Susceptibility Testing, Microscopy)*	1188
(Urine Culture. Bacteria Identification, Antibiotic susceptibility and Bacteriophage Efficiency Testing)*	1078
A12.20.001	495
(Ureaplasma spp. Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	748
(Breast Milk Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	880
(Genitourinary Tract Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	2464
(Wound/Pus/Aspirate/Tissue Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	880
(Mycoplasma hominis Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	847
(Genitourinary Tract Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	1276
(Ear Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*	1474
(Urine Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	2145
(Upper Respiratory Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*	1122
(Ear Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	1276
(Vaginal Biocenosis. Bacteriophage and Antimycotic Susceptibility Testing (Gram Stain, Bacterioscopic Examination of Smear))*	1650
(Wound/Pus/Aspirate/Tissue Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	2145
(Bile Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	2365
(Breast Milk Culture. Bacteria Identification)	715
(Genitourinary Tract Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*	1474
(Adenovirus), (Adenovirus. One Step Rapid Immun chromatographic Assay)	957
(Urine Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	880
(Sputum and Tracheobronchial washings Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing, Microscopy)*	2387
(Helicobacter pylori), (Helicobacter pylori. One Step Rapid Immun chromatographic Assay)	957
(Upper Respiratory Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	913

(Punctate Fluid Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	2178
(Stool Culture, Pathogenic Intestinal and Conditionally Pathogenic Microflora, Bacteria Identification)	1309
6 ()	2398
1 ()	2398
12 (,) (Cobalamin)	693
(-) (Vitamin E, alpha-Tocopherol, Serum)	2398
5 ()	2398
1,25-D3 (1,25-dihydroxivitamin D3)	2090
3 ()	2398
25(OH)D2 25(OH)D3, (- /)	6006
A09.05.080 (Folic Acid)	957
() (Vitamin A, Retinol, Serum)	2398
1 () (Vitamin K1, Phylloquinone, Serum)	2398
7, ()	2398
12 (, Active-B12, Holotranscobalamin)	1375
-	2398
2 ()	2398
()	2398
	2398
LGI1 CASPR2 (), IgG, (VGKC-associated proteins LGI1 and CASPR2 antibodies, serum)	6160
, IgG, (Neuronal antibodies, IgG, Indirect immunofluorescence (IIF))	3311
(,), o (Acetylcholine Receptor Antibodies, Anti-AChR, Total)	5731
NMDA, CASPR, LGI, AMPA1, AMPA2, GABAR1 IgG, ,	13607
IgG () (Anti-Skeletal Muscle Antibodies, AStMA, IgG)	1221
NMDA , IgG, (-NMDAR IgG, N-methyl-D-Aspartate Receptor Antibodies, CSF)	3036
- IgG (- : Mi-2, Ku, PM-Scl 100/75; Jo1 PL-7 PL-12 EJ OJ; SRP, SSA (Ro52)) (Myositis-Specific Panel)	4037
IgG IgM (- : GM1; GM2-GM3-GM4; GD1a, GD1b, GD2-GD3, GT1a, GT1b, GQ1b,), (Anti-GM1 Antibodies, Anti-GQ1b Antibodies, Anti-Ganglioside antibodies, Ganglioside Antibodies Panel, Total)	5731
Crithidia luciliae, IgG, (Crithidia luciliae indirect fluorescent test (CLIFT))	1265
IgG, (Anti-myelin antibody, IgG, IIF)	1474
IgA, IgG, IgM 4, (NMO) (Aquaporin-4Receptor Antibodies, anti-AQP4, Neuromyelitis Optica, NMO, IgA, IgG, IgM, Total)	2860
IgG () (Oligoclonal IgG, Cerebrospinal Fluid (CSF), Serum)	4389
(Muscle-specific tyrosine kinase (MuSK) antibody) (-MuSK)	5478
NMDA, CASPR, LGI, AMPA1, AMPA2, GABAR1 IgG, ,	13750
LGI1 CASPR2 (), IgG, (VGKC-associated proteins LGI1 and CASPR2 antibodies, CSF)	6160
GAD (), IgG, (Anti-GAD (glutamic acid decarboxylase), IgG, CSF)	2079

IgG (Anti-Neuronal Antibodies, Blot-Line (Hu (ANNA1), Yo-1 (PCA1), CV2, Ri (ANNA2), Amphiphysin))		5731
IgG NMDA (N-Methyl-D-Aspartate Receptor Antibodies IgG)		4389
Complement (CH50)		
(CH50) (Functionality Test of		1474
1- (C1-Esterase Inhibitor, 1-INH)		2156
4 (Complement Component C4)		396
3 (Complement Component C3)		396
Anti-Rubella IgG		
(Anti-Rubella IgG, Immunoblot)		5621
Ig (Anti-Rubella Ig)		539
Anti-Rubella IgG		1067
(Rubella virus, RNA)		649
IgG (Anti-Rubella IgG)		396
Hepatitis B Virus (HBV)		
IgM, IgG, HB-core B, (Anti-HBc IgM, IgG, Antibodies to Hepatitis B Core Antigen; HBcAb, Total, HBV Core Total Antibodies (IgG + IgM))		517
HBs- (HBsAg, Hepatitis Surface Antigen, Quantitative)		1452
HB - (Hepatitis Be Antigen, HBeAg)		561
IgM, HB-core B (Anti-HBc IgM Antibodies to Hepatitis B Core Antigen; HBV Core Antibodies IgM)		693
B, (HBV DNA, Serum, Quantitative)*		3751
B, (HBV DNA, Serum, Qualitative)*		418
HBs- (Anti-HBs, HBsAb)		627
HB - (Anti-HBe, HBeAb)		517
HBs- (HBsAg, Hepatitis Surface Antigen, Qualitative)		264
Oral Glucose Tolerance Test (OGTT)		
2 (2-Hour Oral Glucose Tolerance Test, OGTT, Glucose and C-Protein Concentration (Fasting and 2 Hours after Load), Venous Blood)		1529
A09.05.023 (Lactate)		143
(Fructosamine)		572
A09.05.083 HbA1 (HbA1, Glycated Hemoglobin, GHB)		891
(Oral Glucose Tolerance Test, Plasma, OGTT, Pregnancy)		506
A12.22.005 2 (2-Hour Oral Glucose Tolerance Test, OGTT, Glucose Concentration (Fasting and 2 Hours after Load), Venous Blood)		990
C-Reactive Protein (CRP)		
A09.05.009 (C-Reactive Protein, CRP)		341
(Antistreptolysin-O, ASO)		374
-1- (Alpha-1-Antitrypsin, A1AT, AAT, Phenotyping)		2596
(NT-proBNP, N-Terminal Pro-brain Natriuretic Peptide, Pro-B-Type Natriuretic Peptide)		2816
A09.05.076 (Ferritin)		506
(Carbohydrate-Deficient Transferrin with results on an electrophoregram (CDT))		3322

25 () (Hepcidin 25, bioactive)		6567
(Heart Failure's biomarker))	ST2 (ST2, sST2,) (Soluble ST2)	2739
A09.05.008 () (Transferrin)		484
(Myoglobin)		572
-2- (Alpha-2-Macroglobulin, ?2-Macroglobulin, A2M)		495
IgG Pgp3 () Chlamydia trachomatis	IgG	572
() (Rheumatoid Factor, RF)		374
-I (Troponin-I)		627
- ()		594
A09.05.077 (Ceruloplasmin)		649
(Haptoglobin)		638
-1- (1), (Alpha-1-Antitrypsin, A1AT, AAT, Concentration)		1386
- (Carbohydrate-Deficient Transferrin, CDT)		3091
(, Soluble Transferrin Receptor, sTfR)		1870
(Eosinophil Cationic Protein, ECP)		869
()		
(Ureaplasma parvum), (Ureaplasma parvum, DNA, Urine)*		275
(Ureaplasma arvum), (Ureaplasma parvum, DNA, Prostatic Fluid, Semen)*		275
IgG Ureaplasma urealyticum (nti-Ureaplasma urealyticum IgG)		671
(Ureaplasma urealyticum) (-960), (Ureaplasma urealyticum (T-960), DNA, Scrape of Urogenital Epithelial Cells)*		275
IgA Ureaplasma urealyticum (nti-Ureaplasma urealyticum IgA)		671
(Ureaplasma urealyticum + Ureaplasma arvum), (Ureaplasma urealyticum + Ureaplasma arvum, DNA, Prostatic Fluid, Semen)*		275
(Ureaplasma urealyticum + Ureaplasma arvum), (Ureaplasma urealyticum + Ureaplasma arvum, DNA, Urine)*		275
(Ureaplasma arvum), (Ureaplasma parvum, DNA, Scrape of Urogenital Epithelial Cells)*		275
(Ureaplasma urealyticum) (-960), (Ureaplasma urealyticum (T-960), DNA, Prostatic Fluid, Semen)*		275
(Ureaplasma urealyticum + Ureaplasma arvum), (Ureaplasma urealyticum + Ureaplasma arvum, DNA, Scrape of Urogenital Epithelial Cells)*		275
(Ureaplasma urealyticum) (-960), (Ureaplasma urealyticum (T-960), DNA, Urine)*		275
- (-)		
A IgG (RSV) IgG - (Anti-Respiratory Syncytial Virus)		781
A IgM (RSV) IgM - (Anti-Respiratory Syncytial Virus)		781
e (F2, F5) (Risk of Oral Contraceptives, Ocs (Genes F2, F5))		3003
e (F2, F5) ((Risk of Oral Contraceptives, Ocs (Genes F2, F5) (without Description))		2695
, 6 (AZF) ()		3916
(MTHFR, MTRR, MTR, F2, F5) (Preparation for Surgery (Genes MTHFR, MTRR, MTR, F2, F5))		8811
(MTHFR, MTRR, MTR, F2, F5) ((Preparation for Surgery (Genes MTHFR, MTRR, MTR, F2, F5) (without Description))		7898
IgG:		
(F24), - IgG (Shrimp, IgG, F24)		572

(F9),	-	IgG (Rice, IgG, F9)	572	
1: , , , , , IgG (FP15 (F33, F49, F92, F95), Food Panel: Orange, Banana, Apple, Peach, IgG)*			1045	
(F84),	-	IgG (Kiwi Fruit, IgG, F84)	572	
(F11),	-	IgG (Buckwheat, IgG, F11)	572	
(F3),	-	IgG (Codfish, IgG, F3)	572	
3: , , , , , IgG (FP73 (F26, F27, F83, F88), Food Panel: Pork, Beef, Chicken Meat, Lamb, IgG)*			1045	
Food Profile, IgG)		G (IgG)	(Basic	15840
(F44),	-	IgG (Strawberry, IgG, F44)	572	
(F7),	-	IgG (Oat, IgG, F7)	572	
(F105),	-	IgG (Chocolate, IgG, F105)	572	
(fx21) IgE, ImmunoCAP			1375	
(F4),	-	IgG (Wheat, IgG, F4)	572	
-		, IgG (Beta Lactoglobulin, IgG, F77)	572	
(F2),	-	IgG (Milk, IgG, F2)	572	
(F25),	-	IgG (Tomato, IgG, F25)	572	
(F55),	-	IgG (Common Millet, IgG, F55)	572	
		, IgG (Pineapple, IgG, F210)	572	
		, IgG (Lamb, IgG, F88)	572	
(F26),	-	IgG (Pork, IgG, F26)	572	
(F208),	-	IgG (Lemon, IgG, F208)	572	
		, IgG (Banana, IgG, F92)	572	
(F75),	-	IgG (Egg Yolk, IgG, F75)	572	
(F1),	-	IgG (Egg White, IgG, F1)	572	
(F83),	-	IgG (Chicken Meat, IgG, F83)	572	
(F35),	-	IgG (Potato, IgG, F35)	572	
		, IgG (Orange, IgG, F33)	572	
(F209),	-	IgG (Grapefruit, IgG, F209)	572	
(F91),	-	IgG (Mango, IgG, F91)	572	
		, IgG (Peanut, IgG, F13)	572	
2: , , , , , IgG (FP50 (F84, F91, F92, F210), Food Panel: Kiwi Fruit, Mango, Banana, Pineapple, IgG)*			1430	
(F225),	-	IgG (Pumpkin, IgG, F225)	572	
(F14),	-	IgG (Soybean, IgG, F14)	572	
(F31),	-	IgG (Carrot, IgG, F31)	572	
(F27),	-	IgG (Beef, IgG, F27)	572	
(F45),	-	IgG (Baker's Yeast, IgG, F45)	572	
(F216),	-	IgG (Cabbage, IgG, F216)	572	
(F403),	-	IgG (Brewer's Yeast, IgG, F403)	572	
(F17),	-	IgG (Hazelnut, IgG, F17)	572	
(F78),	-	IgG (Casein, IgG, F78)	572	
(F95),	-	IgG (Peach, IgG, F95)	572	
(F49),	-	IgG (Apple, IgG, F49)	572	
-	6			
	IgG	6 (Anti-HHV-6 IgG)	671	
6		(HHV-6 DNA, Saliva)*	275	

6 (HHV-6 DNA, Scrape of Urogenital Epithelial Cells)*		275
6 Fluid, Semen)*	(HHV-6 DNA, Prostatic	275
6 (HHV-6 DNA, Scrape of Nasal Epithelial Cells)*		275
6 Fluid)*	(HHV-6 DNA, Urine)*	275
6 Fluid)*	(HHV-6 DNA, Cerebrospinal	275
6 Fluid)*	(HHV-6 DNA, Exudate)*	275
6 Fluid)*	(HHV-6 DNA, Blood)*	418
6 DNA, Scrape of Faucial Epithelial Cells)*	(HHV-6	275
6 DNA, Scrape of Faucial Epithelial Cells)*	(HHV-6 DNA, Serum)*	418
()		
(Chlamydia pneumoniae), DNA, Saliva)*	(Chlamydia pneumoniae,	429
(Chlamydia trachomatis), trachomatis, DNA, Prostatic Fluid, Semen)*	(Chlamydia	275
IgG	() Chlamydia trachomatis (Anti-cHSP60 IgG)	583
A	IgA Chlamydia trachomatis (Anti-Chlamydia trachomatis IgA)	539
(Chlamydia pneumoniae), pneumoniae, DNA, Plasma)*	(Chlamydia	649
A	IgG Chlamydia pneumoniae (Anti-Chlamydia pneumoniae IgG)	561
IgM	Chlamydia trachomatis (Anti-Chlamydia trachomatis IgM)	561
(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Scrape of Rectal Epithelial Cells)*		275
(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Scrape of Urogenital Epithelial Cells)*		275
A	IgG Chlamydia trachomatis (Anti-Chlamydia trachomatis IgG)	539
Duodenum, PCR)	(Helicobacter pylori, DNA, Biopsies of Gastric Mucosa and/or	2332
(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Scrape of Conjunctiva Epithelial Cells)*		275
A	IgA Chlamydia pneumoniae (Anti-Chlamydia pneumoniae IgA)	671
(Chlamydia pneumoniae), DNA, Sputum)*	(Chlamydia pneumoniae,	869
A	IgM Chlamydia pneumoniae (Anti-Chlamydia pneumoniae IgM)	561
Urine)*	(Chlamydia trachomatis), (Chlamydia trachomatis, DNA,	275
(Chlamydia trachomatis), trachomatis, DNA, Cerebrospinal Fluid)*	(Chlamydia	275
(Chlamydia trachomatis), Exudate)*	(Chlamydia trachomatis, DNA,	275
(Chlamydia trachomatis), trachomatis, DNA, Synovial Fluid)*	(Chlamydia	506
(Chlamydia pneumoniae), (Chlamydia pneumoniae, DNA, Scrape of Faucial Epithelial Cells)*		429
(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Scrape of Faucial Epithelial Cells)*		275
()		
Prostatic Fluid, Semen)*	(Treponema pallidum, DNA,	275
(Treponema pallidum, DNA, Scrape of Urogenital Epithelial Cells)*		275
(Treponema pallidum, DNA, Urine)*		275
DNA, Cerebrospinal Fluid)*	(Treponema pallidum,	275
(Treponema pallidum, DNA, Serum)*		407
(Treponema pallidum, DNA, Secretion)*		275

A	IgM	IgG	Treponema pallidum,	(Anti-Treponema pallidum IgM,	418
IgG, Total)					
			pallidum, DNA, Scrape of Skin Epithelial Cells)*	(Treponema	275
			(Treponema pallidum, DNA, Scrape of Faucial Epithelial Cells)*		275
A	IgG		Treponema pallidum,		1969
(Anti-Treponema pallidum IgG, Immunoblot)					
			(Treponema pallidum, DNA, Scrape of Conjunctiva Epithelial Cells)*		275
A	IgM		Treponema pallidum (Anti-Treponema pallidum IgM)		891
	RPR -		(Syphilis RPR (Rapid Plasma Reagins),	nticardiolipin	231
est)					
A	IgM		Treponema pallidum,		1969
(Anti-Treponema pallidum IgM, Immunoblot)					
A09.05.078			(Testosterone)		385
	-		(-SO4, Dehydroepiandrosterone sulfate, DHEA-S)		385
17-	(17-)		(17-Ketosteroids, Urine)		2013
			() (Sex Hormone-Binding Globulin, SHBG)		396
			(Androstenedione)		1100
A09.05.139	17-	-	(17-Hydroxyprogesterone, 17-OHP)		539
A09.05.078.001			(Free Testosterone)		957
			() (Androstanediol Glucuronide,		1155
3?-Androstanediol Glucuronid, 3?-diol G)					
			() (Dih drotestosterone, DHT)		1430
			() (Protein, random urine, with creatinine and protein/creatinine ratio calculation)		286
			() (Magnesium, random urine, with creatinine and magnesium/creatinine ratio calculation)		583
					308
			() (Phosphorus, random urine, with creatinine and phosphorus/creatinine ratio calculation)		319
			() (Oxalates, random urine, with creatinine and oxalate/creatinine ratio calculation)		1485
			() (Albumin, random urine, with creatinine and albumin/creatinine ratio calculation, UACR)		495
			() (Calcium, random urine, with creatinine and calcium/creatinine ratio calculation)		220
			(Urine Creatinine)		55
IgE:					
			, IgE (Cockroach, IgE, I6)		484
			, IgE (Dog Epithelium, IgE, E2)		484
			, IgE (Sheep Epithelium, IgE, 81)		484
			, IgE (Budgerigar Feathers, IgE, 78)		484
			, IgE (EP70 (E6, E82, E84, E87, E88), Animal Panel: Guinea Pig Epithelium, Rabbit Epithelium, Hamster Epithelium, Rat, Mouse, IgE)*		1045
			, IgE (Guinea Pig Epithelium, IgE, 6)		484
			, IgE (Cat Dander-Epithelium, IgE, E1)		484
			, IgE (Chicken Feathers, IgE, 85)		484
			()		319
	1	2	1	2 (HIV Ag/Ab Combo)	319

-1,	(HIV RNA, Plasma)*	14234
(Everolimus)		
	(Cyclosporine, Cyclosporine A, Sandimmune)	3630
	(Teriflunomide, Leflunomide metabolite)	1045
	(Levetiracetam, Keppra®)	3630
	(carbamazepine, Tegretol)	3740
	(Mitotane, o, p?-DDD, plasma)	2882
	(Phenytoin)	3630
Tacrosel)	(FK506, Advagraf, Prograf, Protopic,	1276
	(Lamotrigine)	1573
	(Acidum Valproicum, Depakin, Convulexs)	3740
A09.05.035.002	(Phenobarbitalum)	891
()		
Prostatic Fluid, Semen)*	(Neisseria gonorrhoeae, DNA,	275
(Neisseria gonorrhoeae, DNA, Scrape of Rectal Epithelial Cells)*		275
Fluid)*	(Neisseria gonorrhoeae, DNA, Synovial	506
(Neisseria gonorrhoeae,), (GC, Neisseria gonorrhoeae Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		990
gonorrhoeae, DNA, Scrape of Conjunctiva Epithelial Cells)*	(Neisseria	275
gonorrhoeae, DNA, Scrape of Faucial Epithelial Cells)*	(Neisseria	275
(Neisseria gonorrhoeae, DNA, Urine)*		275
(Neisseria gonorrhoeae, DNA, Scrape of Urogenital Epithelial Cells)*		275
(Lactobacillus spp., DNA, Scrape of Urogenital Epithelial Cells)*		
Clostridium difficile		
(Toxin A and B Clostridium difficile. One step rapid immunochromatographic assay)		1320
(Clostridium difficile,) (Clostridium difficile Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		1375
?		
()		1100
() (Cytological Examination: Cervix, Pap-test)		1100
IgE:		
/H1-Greer, IgE (House Dust - Greer, IgE, H1)		484
Penicillium notatum, IgE (Penicillium notatum, IgE, M1)		484
Candida albicans, IgE (Candida albicans, IgE, M5)		484
Dermatophagoides pteronyssinus (D1), IgE (Dermatophagoides pteronyssinus, IgE, D1)		484
Aspergillus fumigatus, IgE (Aspergillus fumigatus, IgE, M3)		484
Alternaria tenuis, IgE (Alternaria tenuis, IgE, M6)		484
Dermatophagoides farinae (D2), IgE (Dermatophagoides farinae, IgE, D2)		484
: Penicillium notatum, Cladosporium herbarum, Aspergillus fumigatus, Candida albicans, Alternaria tenuis, IgE (MP1 (M1, M2, M3, M5, M6), Mold Panel 1: Penicillium notatum, Cladosporium herbarum, Aspergillus fumigatus, Candida albicans, Alternaria tenuis, IgE)*		1045
Cladosporium herbarum, IgE (Cladosporium herbarum, IgE, M2)		484

IgG:			
Dermatophagoides farinae (D2), IgG, D2)	-	IgG (Dermatophagoides farinae,	572
/Greer (1),	-	IgG (House Dust – Greer, IgG, H1)	572
Cladosporium herbarum (2), IgG, M2)	-	IgG (Cladosporium herbarum,	572
: Penicillium notatum, Cladosporium herbarum, Aspergillus fumigatus, Candida albicans, Alternaria tenuis, IgG (MP1 (M1, M2, M3, M5, M6), Mold Panel 1: Penicillium notatum, Cladosporium herbarum, Aspergillus fumigatus, Candida albicans, Alternaria tenuis, IgG)*			1045
Penicillium notatum (1),	-	IgG (Penicillium notatum, IgG, M1)	572
Dermatophagoides microceras (D3), microceras, IgG, D3)	-	IgG (Dermatophagoides	572
Dermatophagoides pteronyssinus (D1), pteronyssinus, IgG, D1)	-	IgG (Dermatophagoides	572
Alternaria tenuis (6),	-	IgG (Alternaria tenuis, IgG, M6)	572
-) (Impairment of Spermatogenesis: Full Panel (AZF-Region) (without Description))		(AZF-) (10219
: (F2, F5, MTHFR, MTRR, MTR, ACE, AGT, RHD) (Want to Become a Mother: Pregnancy Complications (Genes F2, F5, MTHFR, MTRR, MTR, ACE, AGT, RHD) (without Description))		(F2, F5, MTHFR, MTRR, MTR, ACE, AGT, RHD)	17270
, CYP21A2, . . . (Gene CYP21OHB, Freq. Mut.)			10769
Tendency in Pregnancy: Minimum (Genes F2, F5)		(F2, F5) (Thrombotic	3003
Male Infertility (Genes AR, CFTR; AZF-Region))		(AR, CFTR; AZF-) (Genetic Factors of	18062
Malformations in Fetus (Genes MTHFR, MTRR, MTR)		(MTHFR, MTRR, MTR) (Isolated	5808
: (F2, F5, MTHFR, MTRR, MTR, ACE, AGT, RHD) (Want to Become a Mother: Pregnancy Complications (Genes F2, F5, MTHFR, MTRR, MTR, ACE, AGT, RHD))		(F2, F5, MTHFR, MTRR, MTR, ACE, AGT, RHD)	19998
(MTHFR, MTRR, MTR, F2, F5) () (Habitual Miscarriage, Thrombotic Tendency in Pregnancy: Extended Panel (Genes MTHFR, MTRR, MTR, F2, F5) (without Description))		(MTHFR, MTRR, MTR, F2, F5) (7898
-) (Thrombotic Complications of Ovulation Induction (Genes F2, F5) (without Description))		(F2, F5) (2695
MTRR, MTR) (without Description))		(MTHFR, MTRR, MTR) (5203
(Gestosis and Placental Insufficiency (Genes ACE, AGT, MTHFR, MTRR, MTR, F2, F5))		(ACE, AGT, MTHFR, MTRR, MTR, F2, F5)	11671
Complications of Ovulation Induction (Genes F2, F5)		(F2, F5) (Thrombotic	3003
-) (Thrombotic Tendency in Pregnancy: Minimum (Genes F2, F5) (without Description))		(F2, F5) (2695
(MTHFR, MTRR, MTR, F2, F5) (Habitual Miscarriage, Thrombotic Tendency in Pregnancy: Extended Panel (Genes MTHFR, MTRR, MTR, F2, F5))		(MTHFR, MTRR, MTR, F2, F5) (8811
MTHFR, MTRR, MTR, F2, F5) (without Description))		(ACE, AGT, MTHFR, MTRR, MTR, F2, F5) (10384
(Hereditary Breast and/or Ovarian Cancer)	/	BRCA1, BRCA2, CHEK2, NBS1	10285
(Examination of Sputum)			825
Secrets)		(Examination of Transudates, Exudates,	528
Bronchial Washouts)		(Examination of	649
(Examination of Punctates: Skin)			649
(Examination of Endoscopic Material)			649

A08.20.004			660
	Helicobacter pylori (Examination of		781
Endoscopic Material: Presence of Helicobacter pylori)			
	(Cytological Examination of Material Obtained during Surgical		990
Procedures and Other Urgent Research)			
(ThinPrep ®)*		1320
	(The Bethesda System ? TBS) (Cytological Examination of Cervical Epithelium with		627
Description on The Bethesda System, TBS)			
	() (Cytological Examination: Scrapings (Smear) of Nasal Mucous Membrane (1		781
Localization))			
	(Examination of Breast Discharge)		528
	(Examination of Punctates: Other Organs and		825
Tissues)			
	() (Examination of Imprint Intrauterine		561
Device, IUD)			
	(Examination of Punctates: Breast)		649
	(Examination		649
of Scrapings and Prints Tumor and Tumor Like Formations)			
	(Examination of Urine)		528
	(The Bethesda System for Reporting Thyroid		627
Cytopathology (TBSRTC), Fine-Needle Aspiration (FNA))			
	(Examination of		429
Scrapings and Prints of Skin and Mucous Membranes)			
	(Examination of Scrapings: Cervix		627
and Cervical Canal)			
	Ig (Anti-Measles IgM)		781
	IgG (Anti-Measles IgG)		880
	IgG (Anti-Tick-borne Encephalitis Virus (TBEV)		528
IgG)			
	(Detection of pathogen DNA/RNA in ticks: Tick-borne		3696
encephalitis Virus (TBEV), Borrelia burgdorferi s. l., Anaplasma Phagocytophillum, Ehrlichia			
muris/chaffe nsis (RNA/DNA), PCR)			
	IgM (Anti-Tick-borne Encephalitis Virus (TBEV)		671
IgM)			
	/ « »		
	() (HPLC-MS/MS Organic Acids (Succinylate))		3949
	HADHA (5511
3- (HADHA Gene, Freq. Mut. (Long-Chain 3-Hydroxyacyl-Coa			
Dehydrogenase (LCHAD) Deficiency))			
	e « » (Newborn Screening "HEEL")*		5489
	(Biotin-Dependent		5511
Carboxylases Activity (Biotinidase Deficiency))			
	GCDH (1) (GCDH (Glutaryl-CoA Dehydrogenase)		5511
Gene, Freq. Mut. (Glutaric Aciduria, Type 1))			
	(/) (Analysis of the spectrum of organic urine acids by gas		9130
chromatography with mass spectrometry (GC / MS))			
	ASS () (ASS Gene, Freq. Mut. (Citrullinemia))		10483
	GCDH (1) (GCDH (Glutaryl-CoA Dehydrogenase)		44440
Gene (Glutaric Aciduria, Type 1))			
	ACADM (5511
MCAD) (ACADM Gene, Freq. Mut. (Medium-Chain Acyl-CoA Dehydrogenase (MCAD)			
Deficiency))			
	FAH () (FAH Gene, Freq. Mut. (Tyrosinemia, Type 1))		9174

Freq. Mut.)	BTD () (BTD (Biotinidase Deficiency) Gene,	5511
	FAH (I) (FAH Gene (Tyrosinemia, Type 1))	54890
	() (TC Gene (Ornithine Transcarbamylase (OTC) Deficiency))	41833
(, , Escherichia coli)		
	(Escherichia coli O157:H7,), (Escherichia coli O157:H7 Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)	1210
	(Escherichia coli O157:H7,), (Escherichia coli O157:H7 Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	1023
	(Escherichia coli O157:H7,), (Escherichia coli O157:H7. One Step Rapid Immun hromotographic Assay)	1023
	1, (Human immunodeficiency virus, quality, RNA)	2717
	19,	363
	- RHD (RHD gene of the fetus in the mother's blood)	6105
	19,	363
	19,	363
	(Legionella pneumophila, One step rapid immun hromotographic assay, antigen, urinae)	1584
	(Respiratory Syncytial Virus, RSV, One step rapid immun hromotographic assay, antigen)	1023
	(Hexagon Chlamydia, One step rapid immun hromotographic assay, antigen)	1122
	(Neisseria gonorrhoeae test, One step rapid immun hromotographic assay)	946
	(Campylobacter spp., One step rapid immun hromotographic assay, antigen, stool)	1144
	(Norwalk virus) - (Norwalk virus GI , GII, One step rapid immun hromotographic assay, antigen, stool)	1925
	(Streptococcus pneumoniae, One step rapid immun hromotographic assay, antigen, urinae)	1584
	(Enterovirus, One step rapid immun hromotographic assay, antigen, stool)	1199
	IgG -3 (Desmoglein 3, DSG3 Antibodies, IgG)	2376
	IgG BP230 (Anti-Bp230 ntibodies, Bullous Pemphigoid (230 kDa) Antibodies, Antibodies to BP Antigen 1, IgG)	2376
	IgG BP180 (Anti-Bp180 ntibodies, Bullous Pemphigoid (180 kDa) Antibodies, Antibodies to BP Antigen 2, IgG)	2376
	IgG (Desmoglein Antibodies, Desmoglein 1, DSG1 and Desmoglein 3, DSG3 Antibodies, IgG)	2376
	, IgG (Basement membrane zone antibodies, IgG)	2222
	IgG -1 (Desmoglein 1, DSG1 Antibodies, IgG)	2376
()		
A	IgA Helicobacter ylori (Anti-Helicobacter pylori IgA)	770
A	IgG Helicobacter pylori, (Anti-Helicobacter pylori IgG, Immunoblot)	3377
A	IgA Helicobacter pylori, (Anti-Helicobacter pylori IgA, Immunoblot)	3377
A	IgG Helicobacter ylori (Anti-Helicobacter pylori IgG)	539
	1303HEL ?? - (?? - , 13C-Urea Breath test, UBT). Helicobacter pylori	2398
A	IgM Helicobacter ylori (Anti-Helicobacter pylori IgM)	770

()			
Sputum)*	(Mycobacterium tuberculosis, DNA,		649
tuberculosis, DNA, Synovial Fluid)*	(Mycobacterium		506
tuberculosis, DNA, Cerebrospinal Fluid)*	(Mycobacterium		275
Exudate)*	(Mycobacterium tuberculosis, DNA,		275
tuberculosis, DNA, Prostatic Fluid, Semen)*	(Mycobacterium		275
IgM, IgA, IgG Mycobacterium tuberculosis, (Anti-Mycobacterium tuberculosis IgM, IgA, IgG, total)			1793
Urine)*	(Mycobacterium tuberculosis, DNA,		275
DNA, Serum)*	(Mycobacterium tuberculosis,		429
tuberculosis, DNA, Menstrual Blood)*	(Mycobacterium		275
() (Circulating Immune Complexes (CIC)			
Total)			1188
	(Phagocytic Activity of Leucocytes)		1100
	(Lymphocyte Activation Ability)		3795
CD4+ - , % Absolute)	(- , CD4+ T-cells, Percent and		1507
CD3+ HLA-DR+, CD3-HLA DR+)*	(CD3+ HLA-DR+, CD3-HLA DR+) (Activated Lymphocyte:		1507
- , %	(CD19+ , B-cells, Percent and Absolute)		1507
(Lymphocyte Phenotyping: CD3, CD4, CD8, CD19, CD16, CD56)	(- CD3, CD4, CD8, CD19, CD16, CD56)		3795
G (IgG1, IgG2, IgG3, IgG4)			13761
()			
Virus IgM, Anti-VZV IgM)	(Anti-Varicella-Zoster		858
Varicella-Zoster,	(Varicella ZosterVirus, DNA, serum)		396
Virus IgG, Anti-VZV IgG)	(Anti-Varicella-Zoster		781
Varicella-Zoster,	(Varicella Zoster Virus, DNA, scrape of faucial epithelial cells)		396
Varicella-Zoster,	(VaricellaZosterVirus, DNA, saliva)		396
()			
Antibiotic Susceptibility and Bacteriophage Efficiency Testing)	(Stool Culture, Salmonella s p., Shigella s p. Bacteria Identification,		1265
Shigella flexneri 1-5 (Shigella flexneri 1-5, IHA)			484
Identification)	(Stool Culture (Salmonella spp., Shigella spp.). Bacteria		858
Shigella sonnei (Shigella sonnei, IHA)			484
(Stool Culture (Salmonella spp., Shigella spp.). Bacteria Identification and Antibiotic Susceptibility Testing)			1067
Shigella flexneri 6 (Shigella flexneri 6, IHA)			484
()			
HER2/neu , HER2-) (HER2/neu Expression, HER2 Status, Immunohistochemical Study (Fixed Biomaterial in Paraffin Block))			5489
: - - (34?E12), p63 (Prostate cancer – complex immunomorphological examination using assessment of the expression AMACR, high molecular weight cytokeratin (34?E12), p63)*	(P504S, AMACR),		12540

() () () (Immunohistochemical diagnosis of lymphoproliferative diseases (Tissue Embedded in Paraffin Block))	27808
Ki-67 (MIB-1) () () (Ki-67 (MIB-1) Expression, Assessment of Proliferative Activity by Expression Ki-67, Immunohistochemical Study (Fixed Biomaterial in Formalin Buffer))*	5599
(CD138) () (Chronic Endometritis, Identification of Plasma Cells CD138, Immunohistochemical Diagnosis (Fixed Biomaterial in Paraffin Block))	5500
() () (Immunohistochemical diagnosis of lymphoproliferative diseases (Fixed Biomaterial in Formalin Buffer))*	27808
() () (Estrogen and Progesterone Receptors, Immunohistochemical Study (Fixed Biomaterial in Paraffin Block))	7854
() () (Early Diagnosis Marker of Dysplasia with High Risk Malignancy: p16INK4a, Immunohistochemical Study (Fixed Biomaterial in Paraffin Block))	4873
() () () (Immunohistochemical diagnosis in cancer metastasis of unknown primary origin (Tissue Embedded in Paraffin Block))	27808
() () (Estrogen and Progesterone Receptors, Immunohistochemical Study)*	7051
() () (P504S, AMACR), (34?E12), p63 (Prostate cancer – complex immunomorphological examination using assessment of the expression AMACR, high molecular weight cytokeratin (34?E12), p63)	12540
() () (Early Diagnosis Marker of Dysplasia with High Risk Malignancy: p16INK4a, Immunohistochemical Study (Fixed Biomaterial in Formalin Buffer))*	4873
HER2 in situ (FISH) (Determination of HER2 Status of Tumor, Fluorescence In Situ Hybridization)	32626
Ki-67 (MIB-1) () () (Ki-67 (MIB-1) Expression, Assessment of Proliferative Activity by Expression Ki-67, Immunohistochemical Study (Fixed Biomaterial in Paraffin Block))	5599
(CD138) () (Chronic Endometritis, Identification of Plasma Cells CD138, Immunohistochemical Diagnosis (Fixed Biomaterial in Formalin Buffer))*	5500
HER2/neu () (HER2/neu Expression, HER2 Status, Immunohistochemical Study (Fixed Biomaterial in Formalin Buffer))*	5489
() () (Immunohistochemical diagnosis in cancer metastasis of unknown primary origin (Fixed Biomaterial in Formalin Buffer))*	27808
A09.05.065 ()	363
A09.05.064 (T4 () (Total Thyroxine, TT4))	385
A09.05.061 (3) (Free Triiodthyronine, FT3)	385
() (Anti-Thyroid Microsomal Antibodies)	550
A09.05.063 (4)	385
(-) (Anti-Thyroid Peroxidase Autoantibodies, Antimicrosomal Antibodies, TPO Antibodies, TPOAb, Anti-TPO)	429
A12.06.046.001 () (Thyroid-Stimulating Hormone Receptor Antibodies, TSH Receptor Antibodies, TSHRABs, TSH binding inhibitor immunoglobulin, TBII)	1540

	(-) (Anti- thyroglobulin Autoantibodies, Thyroglobulin Antibodies, Tg Autoantibodies, TgAb, Anti-Tg Ab, ATG)		484
A09.05.117	() (Thyroglobulin, TG)		693
	() (Thyroid Uptake, T-Uptake, Thyroxine-Binding Capacity, TBC, Thyroxine-Binding Index, TBI, free T4 Index, fT4I)		561
A09.05.060	(3) (Total Triiodothyronine, TT3)		385
	SARS-CoV-2, (Coronavirus SARS-CoV-2 RNA detection in nasopharyngeal and oropharyngeal smear)		1529
	(Soil: Agrochemical Evaluation)*		9988
	(Soil: Comprehensive Toxicological Evaluation)*		21934
A09.05.066	(,) (Growth Hormone, GH)		517
A09.05.067	(,) (Adrenocorticotrophic Hormone, ACTH)		682
	(1) (Somatomedin C, Insulin-like Growth Factor 1, IGF-1)		1089
A09.05.131	() (Luteinizing Hormone, LH)		385
	(Macroprolactin)*		1188
A09.05.087	(Prolactin)		385
A09.05.132	() (Follicle Stimulating Hormone, FSH)		385
	IgG Against Asialoglycoprotein Receptor, Anti-ASGPR, IgG) (-ASGPR) (Autoantibodies		1705
	IgA, IgG, IgM (Anti-Mitochondrial Antibodies, AMA, IgA, IgG, IgM, Total)		1518
	IgG (- 2, 2-3 , Sp100, PML, gp210, LKM-1, LC-1, SLA/LP, SSA/Ro-52), (Autoimmune Disease Liver Panel: AMA-M2, M2-3E (BPO), Sp100, PML, gp210, LKM-1, LC-1, SLA/LP, SSA/Ro-52, IgG, Immunoblotting)		3652
	IgA+IgG+IgM (anti-liver kidney microsomal antibody, anti-LKM, IgG+IgM+ IgA)		1573
	IgA, IgG, IgM (Smooth Muscle Antibodies, SMA, Anti-Smooth Muscle Antibodies, ASMA, IgA, IgG, IgM, Total)		1518
	1-		
	IgG (Insulin Autoantibodies, IAA, IgG)		671
	(IA-2) (Islet Antigen 2 Antibodies, Anti-IA2 antibodies, IA-2 Ab, Tyrosine Phosphatase Antibodies)		1705
	IgG - (Anti-Islet Cell Antibodies, Islet Cell Autoantibodies, ICA)		1518
	GAD/IA-2, (Anti-GAD/IA2 Antibodies Pool, Glutamic Acid Decarboxylase-65, GAD and Insulinoma Antigen 2 (Tyrosine Phosphatase, IA2, ICA-512) Autoantibodies, Total)		1705
	IgG (-GAD) (Anti-GAD Antibodies, Glutamate Decarboxylase Antibodies, AT-GAD, IgG)		1749
	(Yersinia enterocolitica, (Yersinia enterocolitica, Stool Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		1364
	Yersinia enterocolitica :9 (Yersinia enterocolitica O:9, IHA)		484
	IgG Yersinia enterocolitica (Anti-Yersinia enterocolitica IgG)		539
	IgA Yersinia enterocolitica (Anti-Yersinia enterocolitica IgA)		539
	Yersinia enterocolitica :3 (Yersinia enterocolitica O:3, IHA)		484
	Yersinia pseudotuberculosis (Yersinia pseudotuberculosis IHA)		484

(F2, F5, MTHFR, MTR, MTRR, F13, FGB, ITGA2, ITG 3, F7, PAI-1) (Extended Study of Hemostatic System (Genes F2, F5, MTHFR, MTR, MTRR, F13, FGB, ITGA2, ITG 3, F7, PAI-1))		15015
(ITGB3) ((Platelet Fibrinogen Receptor (Gene ITGB3) (without Description))		1386
(F2, F5, MTHFR, MTRR, MTR) ((Thrombosis: Advanced Panel (Genes F2, F5, MTHFR, MTRR, MTR) (without Description))		7898
ITGA2 .759 >T Hyperaggregation of platelets, gene polymorphism ITGA2 .759 >T		3036
(F2, F5) ((Thrombosis: Minimum (Genes F2, F5) (without Description))		2695
(MTHFR, MTRR, MTR) ((Hyperhomocysteinemia (Genes MTHFR, MTRR, MTR) (without Description))		5203
ITGA2 .759 >T ((Hyperaggregation of platelets, gene polymorphism ITGA2 .759 >T (without description))		2717
(F2, F5) (Thrombosis: Minimum (Genes F2, F5))		3003
(MTHFR, MTRR, MTR) (Hyperhomocysteinemia (Genes MTHFR, MTRR, MTR))		5808
(F2, F5, MTHFR, MTRR, MTR) (Thrombosis: Advanced Panel (Genes F2, F5, MTHFR, MTRR, MTR))		8811
(F2, F5, MTHFR, MTR, MTRR, F13, FGB, ITGA2, ITG 3, F7, PAI-1) ((Extended Study of Hemostatic System (Genes F2, F5, MTHFR, MTR, MTRR, F13, FGB, ITGA2, ITG 3, F7, PAI-1) (without Description))		10395
(Listeria monocytogenes, DNA, Cerebrospinal Fluid)*		253
(Listeria monocytogenes, DNA, Urine)*		253
(Listeria monocytogenes, DNA, Scrape of Nasal Epithelial Cells)*		231
(Listeria monocytogenes) (Listeria monocytogenes Culture. Bacteria Identification and Antibiotic Susceptibility testing)		759
(Listeria monocytogenes, DNA, Plasma)*		231
(Listeria monocytogenes, DNA, Scrape of Faucial Epithelial Cells)*		231
(Listeria monocytogenes, DNA, Synovial Fluid)*		506
IgE:		
IgE (GP3 (G1, G5, G6, G12, G13), Grass Panel: Sweet Vernal Grass, Perennial Rye Grass, Timothy Grass, Cultivated Rye Grass, Velvet Grass, IgE)*		1045
IgE (Cottonwood, IgE, T14)		484
IgE (Wormwood, IgE, W5)		484
IgE (Timothy Grass, IgE, G6)		484
IgE (WP1 (W1, W6, W9, W10, W11), Weed Panel: Common Ragweed, Mugwort, English Plantain, Lamb's Quarters, Russian Thistle, IgE)*		1045
IgE (Birch, IgE, 3)		484
IgE (Mugwort, IgE, W6)		484
IgE (TP9 (T2, T4, T12, T3, T7), Tree Panel: Alder, Hazelnut, Willow, Birch, Oak, IgE)*		1045
IgE (GP1 (G3, G4, G5, G6, G8), Grass Panel 1: Orchard Grass, Meadow Fescue, Perennial Rye Grass, Timothy Grass, June Grass (Kentucky Bluegrass), IgE)*		1045
25-OH D (25-OH Vitamin D Total, 25(OH)D, 25-Hydroxycalciferol)		2112
() (Deoxyypyridinoline, DPD, Urine)		1375
(Gla) (steocalcin, N-Osteocalcin, Bone Gla Protein, BGP)		737
N- P1NP, Total) 1 (Procollagen Type 1 N-terminal Propeptide,		1496

(Carboxyterminal Cross-linking Telopeptide of Bone Collagen, Collagen Cross-linked C-Telopeptide, Beta-Cross Laps, ?-CrossLaps Serum, C-Telopeptide, Crosslaps, Type 1 Collagen, b- Tx Serum)		957
(Human Cartilage Oligomeric Protein, COMP)		2695
(A B)		
agalactiae), (Streptococcus group B, Streptococcus (Streptococcus agalactiae Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		968
agalactiae) (Streptococcus agalactiae Culture. Bacteria Identification)		759
Immun chromatographic Assay) (Streptococcus Group B. One Step Rapid		1155
Step Rapid Immun chromatographic Assay) (Streptococcus Group A. One		957
pyogenes), (Streptococcus group A, Streptococcus (Streptococcus pyogenes Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		891
(Streptococcus spp., DNA, Saliva)*		429
pyogenes) (Streptococcus pyogenes Culture. Bacteria Identification)		682
(Streptococcus spp., DNA, Plasma)*		649
spp., DNA, Scrape of Faucial Epithelial Cells)* (Streptococcus		429
(Streptococcus spp., DNA, Sputum)*		869
(, Opisthorchis felineus)		
A09.28.034.001 (Metanephrines fractionated, free and conjugated , 24-h urine)		2486
and Serotonin Metabolites, 24 Hours-Urine: Vanillylmandelic Acid, V (5-), Homovanillic Acid, V (5-), 5-Hydroxyindoleacetic Acid, 5- I) (Catecholamines		2486
(Serotonin, Serum)		2299
Epinephrine/Adrenaline, Norepinephrine/Noradrenaline, Dopamine, Urine) (Catecholamines:		2266
(Catecholamines: Epinephrine/Adrenaline, Norepinephrine/Noradrenaline, Dopamine, Plasma)		2299
(Histamine, Plasma)		2662
(Metanephrines fractionated, free + conjugated, random urine) (+)		2013
Epinephrine/Adrenaline, Norepinephrine/Noradrenaline, Dopamine, Urine) (Catecholamines:		2266
(, Opisthorchis felineus)		
IgG (Anti-Opisthorchis felineus IgG)		869
(Giardia lamblia), (Giardia lamblia. One Step		
Rapid Immun chromatographic Assay) IgM, IgG, IgA (Anti-Giardia lamblia IgM, IgG,		957
IgA, Total)		627
(- ,)		
A09.28.027 (Amylase, 24-Hour or Timed Urine)		253
(Ca), (Calcium (Ca), 24-Hour urine)		220
(K), (Na), (Potassium (K), Sodium (Na), 24-Hour urine)		209
(Glomerular Filtration Rate, GFR)*		209
A09.28.011 (Glucose, 24-Hour urine)		165
A09.28.009 (Urea, 24-Hour urine)		165
A09.28.010 (Uric cid, 24-Hour urine)		209
A09.28.003 (Protein Total, 24-Hour urine)		143
(, (Magnesium, 24 h urine excretion)		330

		2079
	(xalates, 24-Hour urine)	1342
A09.28.003.001	(Albumin, 24-Hour urine)	352
A09.28.006	(Creatinine, 24-Hour urine)	165
(P),	(Phosphorus (P), 24-Hour urine)	220
A09.05.054.002	(Immunoglobulin A, IgA)	275
A09.05.054.003	(Immunoglobulin , Ig)	275
A09.05.054.004	G (Immunoglobulin G, IgG)	275
A09.05.054.001 (Immunoglobulin Total, IgE Total)	E (IgE,)	429
IgG Membrane antibodies, anti-GBM, IgG)	(-) (Glomerular Basement	1716
IgG antibodies, ANCA, IgG)	() (Anti-Neutrophil cytoplasmic	1342
IgG PR-3 ANCA, IgG)	-3 (-PR-3) (Anti- proteinase-3 antibodies, PR-3- antibodies,	1232
IgG, IgA, IgM (Anti-Phospholipase A2 Receptor Antibodies, Anti-PLA2R, IgG, IgA, IgM, Total)	2 (PLA2R),	2717
(Anti-Neutrophil Cytoplasmic Antibodies, ANCA, IgG, Panel)	(), IgG	3410
IgG C1q (Anti-Endothelial Cell Antibodies, AECA, IgG, IgA, IgM, Total)	(Anti-Complement 1q Antibodies, Anti-C1q,	1232
IgG (Anti-Endothelial Cell Antibodies, AECA, IgG, IgA, IgM, Total)	(HUVEC),	1705
IgG (Anti-Myeloperoxidase Antibody, MPO)	(- PO) (Myeloperoxidase Antibody, MPO)	1232
1- 3- (Anti-Poliovirus serotypes 1, 3, IgG)	(Anti-Poliovirus serotypes 1, 3, IgG)	1595
(HiB), anti-PRP Haemophilus influenzae b IgG)	b, IgG (IgG) (polyribosylribitolphosphate, PRP) (Haemophilus influenzae b	1991
()		
A IgG Borrelia burgdorferi (Anti-Borrelia burgdorferi IgG)		627
Cerebrospinal Fluid)*	(orrelia burgdorferi, DNA,	506
Fluid)*	(orrelia burgdorferi, DNA, Synovial	506
A IgM Borrelia burgdorferi, burgdorferi IgM, Western Blot (WB))	- (Anti-Borrelia	1980
A (Anti-Borrelia burgdorferi IgG, Immunoblot)		2222
A IgM Borrelia burgdorferi (Anti-Borrelia burgdorferi IgM)		627
(Barbiturates, Urine)*		1232
? (() (Drugs and Psychotropic	3410
Substances Screening: Opiates, Amphetamines, Methamphetamine, Cocaine, Cannabinoids, Cannabinoid Metabolites, Urine)		
()	(Cannabinoids (Marijuana), Urine)*	1232
()	(Ethanol (Alcohol) Urine)*	1232
« » ((;	3608
and Psychotropic Substances, Urine)*	() (Pernicious Habits: Nicotine, Drugs, Psychostimulants	
(/)	(Opiates (Morphine/Heroin), Urine)*	1232
-	(-)	
- ,	(EBV DNA, Exudate)*	275

Fluid)*	(EBV DNA, Cerebrospinal	275
Antigens (VCA) IgG)	(nti-EBV Viral Capsid	759
IgG)	(nti-EBV Early Antigen (EA	627
(EBNA) IgG)	(nti-EBV Nuclear Antigen	528
	(EBV DNA, Blood)*	418
	(EBV DNA, Saliva)*	275
Fluid, Semen)*	(EBV DNA, Prostatic	275
(EBV DNA, Scrape of Nasal Epithelial Cells)*		275
(EBV DNA, Scrape of Urogenital Epithelial Cells)*		275
Antigens (VCA) Ig)	(nti-EBV Viral Capsid	528
DNA, Scrape of Faucial Epithelial Cells)*	(EBV	275
Serum)*	(EBV DNA,	407
	(EBV DNA, Urine)*	275
	(EBV DNA, Serum)*	418
IgE:		
2		2024
, IgE (Pediatric Panel, IgE)		4037
1		2024
, IgE (Respiratory Panel, IgE)		4037
, IgE (Panel Different Allergens, IgE)		4037
ImmunoCAP ISAC, 112 (Allergochip ImmunoCAP ISAC, 112 llergic components)		30030
ALEX2, 300 IgE		29689
(MTHFR, MTRR, MTR) (5203
) (Folic Acid Metabolism (Genes MTHFR, MTRR, MTR) (without Description))		
: D (VDR) (1386
(Osteoporosis, Vitamin D Receptor (VDR) (Gene VDR) (without Description))		
(MCM6) (Adult Lactase Deficiency (Gene MCM6)		1441
, I (HFE) (Hemochromatosis Type 1 (Gene HFE))		2860
: D (VDR) (Osteoporosis, Vitamin D Receptor (VDR) (Gene VDR))		1529
(UGT1A1) (Gilbert's Syndrome (Gene UGT1A1))		4994
: (CALCR, COL1A1) (3993
) (Osteoporosis: Abridged Panel (Genes CALCR, COL1A1) (without Description))		
(MTHFR, MTRR, MTR) (Folic Acid Metabolism (Genes MTHFR, MTRR, MTR))		5808
: (CALCR, COL1A1) (Osteoporosis: Abridged Panel (Genes CALCR, COL1A1))		4455
: (CALCR, COL1A1, VDR) (5203
) (Osteoporosis: Full Panel (Genes CALCR, COL1A1, VDR) (without Description))		
HLA II (DRB1, DQA1, DQB1) (Hereditary Predisposition to Diabetes Type 1 (Insulin-Dependent Diabetes), HLA Class II (Genes DRB1, DQA1, DQB1))	1-	6864
: (CALCR, COL1A1, VDR) (Osteoporosis: Full Panel (Genes CALCR, COL1A1, VDR))		5808
123 / , 131 / , 141 / , 115 / , 124 / , 154 /) (Genetic Test Results: Description of the 2-nd Category Complexity)	2 (118 / , 121 / ,	1155

7014 , 125 / , 7207) (Genetic Test Results: Description of the 1-st Category Complexity)	1 (7201 , 7611 ,	583
120 / , 137 / , 138 / , 153 / , 110 / , 114 / , 140 / , 7661 , 7258 , 134 / , 135 / , 136 /) (Genetic Test Results: Description of the 3-rd Category Complexity)	3 (122 / , 129 / ,	2310
139 / , 145 / , 108 / , 19 /) (Genetic Test Results: Description of the 4-th Category Complexity)	4 (144 / , 143 / ,	5203
(4)		
(Isoprinosine)		539
(Immunomax)		539
(Immunal)		539
(Thymogen)		539
(Panavir)		539
(Tactivinum)		539
(Imunofan)		539
(Polyoxidonium)		539
(Galavit)		539
(Imunorix)		539
(Licopid)		539
()		
IgG oxoplasma gondii (nti-Toxoplasma gondii IgG)		396
Ig oxoplasma gondii (nti-Toxoplasma gondii Ig)		539
, (Toxoplasma gondii, DNA, Serum)*		418
Cerebrospinal Fluid)* (Toxoplasma gondii, DNA,		275
Anti-Toxopl gondii IgG		1045
, (Toxoplasma gondii, DNA, Exudate)*		275
(Streptococcus		
pneumoniae)		495
(Bordetella pertussis/parapertussis,		
/) (Bordetella pertussis/parapertussis, Nasopharyngeal Culture. Bacteria Identification)		1584
(2) (Estradiol, E2)		
(Progesterone)		385
(Cortisol, Hydrocortisone)		
A09.05.135 (Free ortisol, Free Hydrocortisone, 24-Hour urine)		385
A09.28.035		759
A09.05.069 (Aldosterone)		858
A09.05.121 (Direct Renin, Plasma)		913
, (Cortisol, Saliva)		616
(Cystatin C)		
A09.05.230		748
A09.05.017		165
A09.05.018 (Uric cid)		165
A09.05.020		165
- p16INK4a Ki-67		
		6050

	(PLGF)		3960
	(Inhibin B)		1232
MIS)	() (Anti-Mullerian Hormone, AMH, Mullerian Inhibiting Substance,		1232
	-1- () (Trophoblastic beta-1-Globulin, TBG)		484
	IgG V (Annexin V antibodies, aAnV, IgG)		1309
	Ig V (Annexin V antibodies, aAnV, Ig)		1309
Antibodies, anti-?-G 1, IgG, IgA, IgM, Total)	IgG, IgA, IgM -2- 1, (nti-?-Glycoprotein 1		1254
	IgM IgG (nti-Phospholipid Antibodies, APA, IgM, IgG)		803
	2 IgA		1199
(Anti- hosphatidylserine/ rothrombin ntibodies, Anti-PS/PT, IgG, IgM, Total)	IgG IgM -		1342
aCL, Screening)	IgA, IgM, IgG (ardiolipin Antibodies IgA, IgM, IgG,		1133
	IgG IgM (Anti- hosphatidylserine, IgG, IgM)		1782
	2 IgM		1309
	(Anti-Phospholipid Antibodies Panel)		9009
	IgA (Anticardiolipin IgA, aCL IgA)		836
	2 IgG		1199
	IgG (Anticardiolipin IgG, aCL IgG)		902
	IgM (Anticardiolipin IgM, aCL IgM)		1188
	(Bile Acids)		2651
	1 (1, 1) (Apolipoprotein A1, Apo A1)		572
Cholesterol)	(, , VLDL		407
	B (B,) (Apolipoprotein B, Apo B)		429
A09.05.025	() (Triglycerides)		209
A09.05.004			220
	(a), () (Lipoprotein (a), Lp (a))		869
A09.05.028			165
A09.05.026	() (Cholesterol Total)		209
	() Cholesterol LDL (direct)		253
	(4)		
	(Neovir)		539
	(Amixin)		539
	(Cycloferonum)		539
	(Kagocel)		539
Protein-A, PAPP-A)	(Pregnancy-Associated Plasma		693
A09.05.090 Gonadotropin, HCG)	(, - , ?-) (Human Chorionic		385
?- (?-) (Free Human Chorionic Gonadotropin, Free HCG)			539
PRISCA2			99
fms-	-1 (sFit-1)		3377
	(Estriol Free, 3)		484

PRISCA1		132
(Placental Lactogen, PL, Human Placental Lactogen, hPL, Chorionic Somatomammotropin, CS, Human Chorionic Somatomammotropin, hCS)		693
(Erythrocyte Sedimentation Rate, ESR)		143
(Leucocyte Formula (Differential White Blood Cell Count) with Manual Microscopic Examination of Blood Smear)*		330
(Leucocyte Formula (Differential White Blood Cell Count) with Microscopic Examination of Blood Smear if Presence of Pathologic Changes)*		209
(Platelets, Microscopy (Manual Platelet Count (PLT Count): Indirect Method by Fonio))*		275
(General Blood Analysis, without White Blood Cell (WBC) Count and ESR)		209
A12.05.123 (Reticulocytes)		264
		1980
(4)		
(Ingaron)		539
(Reaferonum)		539
Bordetella species: Bordetella pertussis (Bordetella bronchiseptica (Differentiated detection of DNA Bordetella spp.: Bordetella pertussis (pertussis pathogen) and Bordetella bronchiseptica (bronchosepticosis pathogen) in a scraping of the oropharynx and or nasopharynx)		924
A IgG Bordetella pertussis (Anti-Bordetella pertussis IgG)		891
A IgM Bordetella pertussis (Anti-Bordetella pertussis IgM)		891
A IgA Bordetella pertussis (Anti-Bordetella pertussis IgA)		891
-3 (Omega-3 Index)		4774
(Procalcitonin)		3003
, : -3,-6,-9, (Fatty acids panel, omega-3, -6, -9, plasma)		9174
Rh- (Anti Rh)		517
A12.05.005 (Blood Group, O)		253
Rh (C, E, c, e) Kell- (Rh C (E, c, e) Kell-Phenotyping)		660
A12.05.006 - (-) (Rh-factor, Rh)		253
: (Water: Complete and Comprehensive Quality Assessment)*		19998
: (Water: Suspicion Industrial Waste Contamination)*		14190
: (Water: Suspicion Products of Combustion and Emissions from Motorways Contamination)*		10483
: (Water: Suspicion Household Waste Contamination)*		8943
: (Water: Abridged Quality Assessment)*		9669
: (Water: Suspicion Excessive Use of Chemicals for Water Treatment)*		5159
(, Trichinella spiralis)		
IgG (Anti-Trichinella IgG)		517
: 20 (Water: Quality Assessment 20 Parameters)*		4103

Radionuclides)*	6	(Determination of Concentration 6	24673
?- (Radiological Drinking Water Study – Basic Test ?- and ?-Activity)*	-	?-	6787
:	30	(Water: Quality Assessment 30 Parameters)*	7766
Radionuclides)*	4	(Determination of Concentration 4	16434
Panel (Genes ACE, AGT, NOS3)	(ACE, AGT, NOS3) (Arterial Hypertension: Full	4554
Description))	(ACE, AGT, NOS3) (4081
ACE, AGT) (Arterial Hypertension, Renin-Angiotensin System Disorder (Genes ACE, AGT))	(3036
ACE, AGT) (-) (Arterial Hypertension,	2717
(NOS3) (-) (Arterial Hypertension, Endothelial	1386
NO-Synthase Disturbance (Gene NOS3) (without Description))		NO-	
()	(Compositional Analysis of Urine (Kidney) Stones, infrared spectrometry)	3872
()	(Compositional Analysis of Urine (Kidney) Stones, infrared spectrometry, -ray diffraction analysis)	3872
		(Alzheimer's Disease)	2860
Gastrin-17, G-17)	-	-17 (1210
I (Pepsinogen I)) (Gastrin-17 Stimulation Test,	968
A09.05.057 (Gastrin)			682
(GastroPanel)			4576
II (Pepsinogen II)			968
G	(G, Hepatitis G Virus, HGV)	
G,		(HGV RNA, Serum)*	660
(FibroTest)			11946
(FibroMax)			15268
(FibroTest)			13420
(SteatoS reen)			6336
(FibroMax)			17600
NASH-FibroTest			19305
NASH-FibroTest ()		18832
()		7920
()	(Karyotype)	7799
()		17787
		(Acute Intestinal Infections, PCR, Fecal)	1573
		(Enterovirus, RNA, Fecal)	517
Infections, PCR, Fecal)		(Acute Intestinal	1243

-	CYP2D6	(beta-Adrenergic Blockers, Gene CYP2D6)		8129
		ATII.		3036
(ACE)	(ACE Inhibitors, Fluvastatin, ATII Receptor Blockers.		5808
		(Methotrexatum. Genetic Markers of Increased Risk of Development of Adverse Reactions in Taking Methotrexate for Treatment of Rheumatoid Arthritis. Methotrexate Disrupts Metabolism		
A	IgG	(Anti-Mumps IgG)		781
A	IgM	(Anti-Mumps IgM)		781
()			
		(Streptococcus pneumoniae, DNA)		495
	(Calcitonin)			946
A09.05.058	() (Parathyroid Hormone, PTH)		671
()	(Echinococcus spp.)		
	IgG	(Anti-Echinococcus IgG)		869
	IgG	(Anti-Entamoeba histolytica IgG)		693
D	(D, Hepatitis D Virus, HDV)		
	IgM	IgG	D, o (Anti-HDV Total (IgG + IgM))	858
	D,		(HDV RNA, Serum)*	660
	IgM		D (nti-HDV IgM)	858
M3)	Aspergillus fumigatus (3),	-	IgG (Aspergillus fumigatus, IgG,	572
				231
(Bacteroides spp., DNA, Scrape of Urogenital Epithelial Cells)*				
A	()	(Hepatitis A Virus, HAV)	
	IgG		(Anti-HAV IgG)	561
			(HAV RNA, Serum)*	649
	IgM		(Anti-HAV IgM)	803
()			
				275
(Gardnerella vaginalis, DNA, Scrape of Urogenital Epithelial Cells)*				275
Prostatic Fluid, Semen)*			(Gardnerella vaginalis, DNA,	275
			(Gardnerella vaginalis, DNA, Urine)*	275
Converting Enzyme, ACE, Serum)	()	(Angiotensin	2420
()		(Neopterin, Serum)	1705
IgG			(),
(Anti-Heart Antibodies, IgG)				1342
HOMA-G	()		55

HOMA-IR		55
A09.05.056.001	(Proinsulin)	913
-	(C-Peptide)	429
A09.05.056	(Insulin)	539
Identification)	(Campylobacter s p.) (Campylobacter spp., Stool Culture. Bacterial	1364
(Rotavirus), Agglutination)	(Rotavirus Direct Detection by Latex	748
		4059
-	(Varicella-Zoster)	
Varicella-Zoster, Zoster Virus, DNA, scrape of skin epithelial cells)	(Varicella	396
	(Fungal Infections of Nails)	902
	(Fungal Infections of Skin)	902
IgG	(Platelet antibodies IgG, Indirect)	3278
(, Salmonella spp.)	
Salmonella gr.A (Salmonella gr.A, IHA)		484
Salmonella gr.B (Salmonella gr.B, IHA)		484
Salmonella gr.E, (Salmonella gr.E Antibodies, IHA)		484
Salmonella typhi, (Salmonella typhi Antibodies, IHA)		616
Salmonella O- (Salmonella O-antigens, IHA)		484
Salmonella gr.D (Salmonella gr.D, IHA)		484
Salmonella gr. (Salmonella gr.C, IHA)		484
IgG	(Anti-Strongyloides stercoralis IgG)	990
	(Rickettsia prowazekii, IHA)	484
A09.05.021		165
A09.05.022		165
	Anisakis IgG	814
IgG	(Anti-Diphtheria Toxoid IgG)	990
	(Corynebacterium diphtheriae Culture)	759
	(Anti-Spermatozoa Antibodies, ASA, Semen)	1419
	(Anti-Spermatozoa Antibodies, ASA, Serum)	1045
(E, Hepatitis E Virus, HEV)	
IgM	E (Anti-HEV IgM)	891
IgG	E (Anti-HEV IgG)	891
IgG:		
(2),	- IgG (Dog Epithelium, IgG, E2)	572

(1), -	IgG (Cat Dander-Epithelium, IgG, E1)		572
	(Streptococcus pneumoniae, DNA)		495
	IgG (Anti-Adenovirus IgG)		781
	IgA (Anti-Adenovirus IgA)		781
	(Streptococcus pneumoniae, DNA)		495
(, Toxocara canis)			
	IgG (Anti-Toxocara IgG)		517
	(Androflor® REAL-TIME PCR Detection Kit, the study of men's urogenital tract microbiocenosis in the epithelial scrapes from the balanus, urethra)		2838
	(Androflor® Screen REAL-TIME PCR Detection Kit, the study of men's urogenital tract microbiocenosis in the epithelial scrapes from the balanus, urethra)		1991
	(DLG5, NOD2, OCTN1, OCTN2) (Crohn's Disease (Genes DLG5, NOD2, OCTN1, OCTN2))		8899
	IgA, IgM, IgG (Anti-Ovarian Antibodies, AOA, IgA, IgM, IgG, Total)		1419
	IgA, IgM, IgG (Anti-Steroidal Cell Antibodies, StCAb, Steroidal Cell Autoantibodies, SCA, IgA, IgM, IgG, Total)		1232
	IgA, IgM, IgG (Anti-Testicular Steroid-ell Antibodies, Testicular Anti-Steroidal Cell Antibodies, Testicular StCAb, Steroidal Cell Autoantibodies, SCA against Testis, IgA, IgM, IgG, Total)		1705
	YP2D6 (YP2D6) (Cytochrome YP2D6 (Gene YP2D6))		8129
	(Neisseria meningitidis Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		935
	() (Total Antioxidant Status, TAS)		5247
	IgE: , IgE (Latex, IgG, K82)		484
	IgG (Anti-Ascaris lumbricoides IgG)		968
(, Demodex folliculorum, Demodex brevis)			
	(Demodex folliculorum, Demodex brevis)		374
	() (Consultation of Finished Cytological Preparations (1 Glass))		418
	Clonorchis sinensis IgG		1078
- 8			
	IgG 8 (Anti-HHV-8 IgG)		770
	()*(Provision of test findings in English (translation of test findings in English)).*		220

(Erythropoetin)	1056
(Leptin)	836
(Cryptosporidium parvum), (Cryptosporidium parvum. One Step Rapid Immunochromatographic Assay)	869
IgG - 1 2 (Anti-HTLV-1, 2 IgG)	858
(Study of Interferon Status)	2695
(Assessment of Androgen Status)	1606
(Female Hormonal Profile: Ovarian Dysfunction, Menstrual Irregularities)	3630
(Female Hormonal Profile: Ovarian Dysfunction, Menstrual Irregularities)	3762
D ()	2000
	3509
	3872
	7051
« »	8250
« »	9966
« »	17281
-	20911
() (Acute Respiratory Infections, ARI: Runny Nose, Cough, Sore Throat)	8129
A IgA IgG Chlamydia trachomatis, (Anti-Chlamydia trachomatis IgA, IgG)	1067
« : 6 () »	3465
(Comprehensive Study «Sex in City: 6 Infections (Blood Test)»)	
« : 6 () »	3597
(Comprehensive Study «Sex in City: 6 Infections (Blood Test)»)	
: (Joint Pain: Extended Survey)	7458
	7689
: (Want to Become a Mother: Pregnancy Planning, Comprehensive Survey)	7964
VIP- (VIP-Survey for Men)	16269
VIP- (VIP-Survey for Women)	17270
(Pediatric Infections: Immune Response)	5984
A IgM IgG Mycoplasma pneumoniae (Anti-Mycoplasma pneumoniae IgM, IgG)	1089
TORCH- (ToRCH-Infections)	3564
: I (1-13) (Pregnancy: First Trimester (1-13 Weeks))	8228
, , (HIV, Syphilis, Hepatitis B, C)	1540
« : 8 + »	2794
(Comprehensive Study «Sex in City: 8 Infections + Smear on Flora»)	
: III (29-30) (Pregnancy: Third Trimester (29-30 Weeks))	3993
« : 14 + »	4411
(Comprehensive Study «Sex in City: 14 Infections + Smear on Flora»)	
:	10923
() (Hemostasiogram (coagulogram), extended)	2761
: (Survey of Liver: Extended)	2893
(Hospitalization in Therapeutic Hospital)	3663

(Hospitalization in Surgical Hospital)	5060
Surgical Hospital: Extended Survey)	7392
« » (My Healthy Nurse)	7722
ROMA (Risk of Ovarian Malignancy Algorithm,) (Risk of Ovarian Malignancy Algorithm, ROMA (Before Menopause))	1815
ROMA (Risk of Ovarian Malignancy Algorithm,) (Risk of Ovarian Malignancy Algorithm, ROMA (After Menopause))	1815
-) (Breast Cancer, Immunohistochemistry, IHC (Formalin-Fixed Biomaterial))	17226
-) (Breast Cancer, Immunohistochemistry, IHC (Paraffin-Embedded Tissue Block))	17226
- : p16INK4a + Ki-67 () (Cervical Cancer - Study of Two Markers for Early Diagnosis Dysplasia with High Risk Malignancy: p16INK4a + Ki-67, Immunohistochemical Screening (Fixed Biomaterial in Paraffin Block))	8613
- : p16INK4a + Ki-67 () (Cervical Cancer - Study of Two Markers for Early Diagnosis Dysplasia with High Risk Malignancy: p16INK4a + Ki-67, Immunohistochemical Screening (Fixed Biomaterial in Formalin Buffer))*	8613
: (Metabolic bone and osteoporosis risk evaluation: comprehensive examination).	5896
: I (PRIS A-1) (Maternal Screen, First Trimester; Prenatal Screening I; PRIS A I (Prenatal Risk Calculation))	1287
: II (PRIS A-2) (Maternal Screen, Second Trimester; Prenatal Screening II; PRISCA II (Prenatal Risk Calculation))	1430
B03.005.006 (), (Coagulation, Gemostaziogram, Screening)	847
: (Miscarriage: Autoimmune Profile)	4004
(Immunological Survey Extended)	12078
: (Survey of Liver: Screening)	924
B03.016.004 :	1210
Supersport	2200
: (Serum Biochemistry: Minimum)	2222
Supersport	3410
: (Serum Biochemistry: Extended Profile)	3553
- : (40) (Healthy You ? Healthy Country: Annual Check-Up up to 40 Years of Age)	3861
(40) (Annual Check-Up after 40 Years of Age)	4928
: (Survey Before Diet: Additional)	5159
Supersport	7513
: (Survey of Kidneys: Extended)	4059
: (Diabetes Control: Extended)	2200
(Toxic Trace Elements, Hair)	3542
(Toxic Trace Elements, Essential Vital Elements, Hair)	1782
(Elemental Composition of Hair: Screening)	3399
() (Essential Vital Elements, Toxic Trace Elements, Urine)	5940
(Toxic Trace Elements, Nails)	2860
(Toxic Trace Elements, Essential Vital Elements, Nails)	1782
(Toxic Trace Elements, Essential Vital Elements, Nails)	3399

(Elemental Composition of Nails: Screening)	5940
	26455
	26455
	26455
	26455
	26455
	36300
(Testing for Kindergarten and School)	1738
: 0 14 (Healthy Child: for Children from 0 to 14 Years)	726
: (Survey of Kidneys: Screening)	957
	979
	20350
	19140
	19140
(Panel Chronic myelogenous leukemia, CML)	13090
	36300
	12210
	3465
	2431
	2376
	2376
	2475
	3179
	4103
IgG)	4345
	4367
: (Arthralgia: screening test)	5082
(Autoimmune Liver Disease: Screening)	7018
(Rheumatic arthritises)	1837
SARS-CoV-2, IgM () IgG	1859
(Anti-SARS-CoV-2, IgM/IgG)	4367
«)» (Bullous Dermatosi Diagnostics profile (antibodies to epidermis desmosomes, antibodies to skin basal membrane))	1419
SARS-CoV-2, IgM IgG (Abbott)	1859
(Anti-SARS-CoV-2, IgM/IgG)	1859
: (Thyroid Gland: Extended Survey)	1859
: (Thyroid Gland: Extended Survey)	1936
: (Thyroid Gland: Screening)	1111
: (Diabetes: Autoimmune Markers)	3916
(Rheumatoid arthritis).	3091
Ig IgG Mycoplasma hominis (nti-Mycoplasma hominis Ig , IgG)	913
	2904
	3003

4 « (), (- - IgG, 3)» (Systemic lupus erythematosus (SLE) profile, activity monitoring (anti-double-stranded DNA IgG, C3 and C4 complement components))	1342
, IgG; IgA (, IgA;	2167
Intolerance) : () (Coeliac Disease: Gluten	6226
, IgG, IgM	1980
APS) (), (Antiphospholipid Syndrome,	3927
cANCA, IgG; /ANCA, IgA; ASCA, IgG, IgA) () (/pANCA,	4774
(, IgA; IgA) (, IgA;	2717
IgA, IgG; IgA) (, IgA;	3553
steroid-producing cells Antibodies) (Reproductive tissue	2959
(Food Allergy)	8965
: (Lipid Profile: Extended)	2904
: (Lipid Profile: Extended)	2904
" / " IgE, ImmunoCAP	4961
" " IgE, ImmunoCAP	4961
(NOS3) (Arterial Hypertension, Endothelial NO-Synthase Disturbance (Gene NOS3))	1529
" / " IgE, ImmunoCAP	4961
)» (Comprehensive Study «Sex in City: 12 Infections (Urogenital Scraping)»)	2893
	1419
	1606
	4510
	12936
	792
(Diagnosis of Anemia)	3179
(Diagnosis of Anemia)	3520
Vessel Diseases) (Preventing Heart and Blood	3102
Diagnosis of Urogenital Tract Infection (UTI)) (Pregnancy Planning:	2486
: (Diabetes Control: Screening)	616
: (Survey Before Diet: Minimum)	1716
(Problems: Primary Survey)) (Weight	3223
(Healthy skin beauty)	1386
:	649
HOMA-IR (Insulin Resistance: Fasting Glucose/Insulin, Homeostasis Model Assessment of Insulin Resistance, HOMA-IR)	737
)" : (946
Problems: Metabolic Syndrome (Primary Identification, creening) (Weight	
(Diagnosis of Osteoporosis)	3047
: (Women's Oncorisk: Cervix)	1980
, , (Allergy to Animals, Dust, Mold)	6358
:	1650

(Strong hair and nails, velvet skin)	3465
: (Trace Elements, Serum, Venous Blood: Screening)	3377
(Mold Allergy)	2211
(Plant Allergy)	4818
(Immunological Survey, Screening)	7040
(AR, CFTR; AZF- ;) (Male Sterility (Genes AR, CFTR; AZF-Region; Karyotype))	23496
(F2, F5, MTHFR, MTRR, MTR, ACE, AGT, RHD, HLA II;) (Female Infertility, Pregnancy Complication (Genes F2, F5, MTHFR, MTRR, MTR, ACE, AGT, RHD, HLA II; Karyotype))	28534
(e) (Toxic Trace Elements, Toxic Heavy Metals, Venous Blood)	1375
()	2365
()	2244
	176
CKD-EPI – (Estimated Glomerular Filtration Rate, eGFR, CKD-EPI Creatinine Equation)	176
B03.016.003 ()	462
: II (14-28) (Pregnancy: Second Trimester (14-28 Weeks))	836
: (« ») (Clinical Blood Analysis: General Blood Analysis, Leucocyte Formula, ESR (with Manual Microscopic Examination of Blood Smear))	649
() (Essential Vital Elements, Essential Trace Elements, Serum)	1375
, 6 (AZF) (Spermatogenesis disorders (6 AZF))	5071
: (AZF-) (Impairment of Spermatogenesis: Full Panel (AZF-Region))	11407
: IgG ()	5830
(DQA1, DQB1) HLA II	6864
- (RH factor Genotype)	10901
(skin) () (Parasitic Fungi, Microscopy and Culture)	1727
(nails) () (Parasitic Fungi, Microscopy and Culture)	1727
I II (I/ II) (Pepsinogen I/Pepsinogen II, PG1/PG2)	2002
: sFit-1, PIGF, sFit-1/PIGF	6468
(Mycoplasma hominis Culture, Ureaplasma spp. Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	1485
: (Male oncologic risk: prostate)	957
A09.05.120.001 - (Aldosterone-Renin Ratio, ARR)	1408
3, 4 (Complement components C3, C4)	792
" , IgE, ImmunoCAP	2332
B03.016.005 : (Lipid Profile: Screening)	770
(ITGB3) (Platelet Fibrinogen Receptor (Gene ITGB3))	1529
: (Lipid Profile: Screening)	770
CKD-EPI – (Estimated Glomerular Filtration Rate, eGFR, CKD-EPI ystatin C Equation)	792
TREC KREC	5313

19.1	29	28.01.2021 . . .		10500
() . 5.1; . 5.2; . 19.1	29	28.01.2021 .		15000
(, 40) . 5.1; . 5.2; . 19.1	29	28.01.2021 .		12070
(, 40) . 5.1; . 5.2; . 19.1	29	28.01.2021 .		11570
/				2000
				4800
				6800