



22.02.2024 .

	(/)	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			1000
A06.20.004	2		1500
	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		(1-3 ,1)	1300
A16.01.017		(3-5 ,1)	1500
A16.01.017		(3-5 ,1)	1900
A16.01.017		(5 ,1)	2400
			500
A16.01.018 (, , 05-1 -1)		-	1300
A16.01.018 (, , 1-3 -1)		-	1500

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)			1900
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)			6500
A16.01.018 (, , 10 -1)			12000
A16.30.032		(5 10 -1)	6500
A16.30.032		(10 -1)	12000
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

A16.30.004.010	1		40000
A16.30.004.010	2		55000

A16.30.004.010		3		70000
	(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				900
				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			950
			1100
A04.30.010			950
A04.30.001	()		1000
A04.30.001.001	()		1000
A04.30.001.007		III	1500
A04.12.024.003		(1000
II-III)			
A04.04.001	()		700
A04.20.003		()	700
A04.20.003		()	400
A04.20.001.004	(-)		500

A04.30.001.002		4D (26 ,	2500
A04.30.001	II		1200
A04.30.001.002		3D	1350
A04.30.001.001		()	1500
A04.12.024.003		()	1500
A04.30.001.006		II-III	2000
A04.30.001.008		III	2000
A04.30.001.002		4D (26 , , 1)	3500
A04.30.001.002		3D	2000

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	()		87900
	2		
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	()		39900
	1-	()	
A16.20.035.001	()		61900
	2-	()	
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
A16.20.098	(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 (HPV DNA, Scrape of Urogenital Epithelial Cells, 2 Types (16, 18))	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))	385
	() 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) Screening)	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
	(), (Antiphospholipid Syndrome, APS)	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)	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
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 () 25 .		2500
A11.01.003	mesoline () 25 .		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			1500
A16.08.035	(1)	5000
A16.08.035	(2)	20000
A16.08.008.004			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

			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	(c		7000
OLGA/OLGIM)			
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
)	() 8 (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
)	() 10 (1	120
	20(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

()		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
A16.09.011	(1)	1000
A16.09.011	(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", ()	"A1"	"B1"	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	(,)	-	()	2500
	(60)			
A05.30.014				900
				150
				1837
(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-Saccharomyces Cerevisiae Antibodies, ASCA, IgA)	1232
IgG, IgA (Autoantibodies against Exocrine Pancreas, Pancreatic Antibodies, PAB)	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-Endomysial Antibodies, Anti-EMA, IgA, IgG, Total)	1276
IgA, IgG, IgM (Gastric Parietal Cell Antibodies, GPA, Anti-parietal cell antibodies, APCA, IgA, IgG, IgM, Total)	1518
IgG, IgA, GP2 (Anti-GP2)	1980
Ig (Anti-Deaminated Gliadin Peptide, Anti-DGP, IgA)	781
- (Birt-Hogg-Dube Syndrome, BHD, Gene FLCN, Mut.)	51348
- (Charcot-Marie-Tooth Disease Type 2A1, Gene MFN2, Freq. Mut.)	5148
4 TBP, . . .	3410
N1, . . .	3410
(Arthrogryposis Distal Type 2A, Gene MYH3, Freq. Mut.)	9999
(Charcot-Marie-Tooth Disease Type 1B, Genes NDRG1, SH3TC2, Mut.)	5148
ATXN8, . . . (Spinocerebellar Ataxia, Gene ATXN8, Freq. Mut.)	5148
(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
(3)	NS3, NS5A NS5B	12837
	RAB27A, . (Griscelli Syndrome, Gene RAB27A, Mut.)	21428
Angioedema Type I, Gene C1NH, Mut.)	C1NH, . (Hereditary	29975
	(22, . (Charcot-Marie-Tooth Disease Type 1B, Gene 22, Mut.)	17523
	HRAS, . (Costello Syndrome, Gene HRAS, Mut.)	7161
(Metaphyseal Chondrodysplasia, McKusick Type, Gene RMRP, Mut.)	RMRP, .	7161
Gene MEFV, Mut.)	MEFV, . (Familial Mediterranean Fever, FMF,	41525
(Shwachman-Diamond Syndrome, Gene SBDS1, Freq. Mut.)	SBDS1, . . .	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
Gene GLI3, Mut.)	(). GLI3, . (Greig Syndrome,	76989
		8910
	(EGR2, . (Charcot-Marie-Tooth Disease Type 1B, Gene EGR2, Mut.)	17160
Gene FGFR2, Mut.)	7 9 FGFR2, . (Crouzon Syndrome, Exons 7, 9	9999
- a - a ().	38522
ENG, . (Rendu-Osler-Weber Disease, Gene ENG, Mut.)		
Exudative Vitreoretinopathy, FEVR, Gene NDP, Mut.)	NDP, . (Familial	11264
Lipodystrophy 2, Gene LMNA, Mut.)	LMNA, . (Familial Partial	42801
(Nail-Patella Syndrome, NPS, Onychoosteodysplasia, Gene LMX1B, Mut.)	LMX1B, .	29975
Mut.)	SGCE, . (Myoclonic Dystonia, Gene SGCE,	51348
ATP7B, Freq. Mut.)	ATP7B, . . (Wilson Disease, Gene	10285
NGF, Mut.)	, NGF . (Hereditary Sensory and Autonomic olynuropathy, Gene	17160
syndrome, Gene ALMS1, Hot-Point Mut.)	« . » ALMS1, « . » . . (Alstrom	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
Primary Carnitine Deficiency, SPCD, Carnitine Deficiency Systemic Primary, CDSP, Gene SLC22A5, Mut.)	SLC22A5, . (Systemic	42801
Mut.)	FXN, . . (Friedrich A taxia, Gene FXN, Freq.	9020

EMG1, . (Bowen Conradi Syndrome, BCS, Gene EMG1, Mut.)	17160
(Leber Hereditary Optic Neuropathy, LHON, Mitochondrial DNA, 3 Freq. Mut.)	5148
Syndrome, Gene BCS1L, Mut.)	17523
BCS1L, . (Bjomstad	
TWIST1, Mut.)	14916
TWIST1, . (Saethre-Chotzen Syndrome, Gene	
GDAP, . (Charcot-Marie-Tooth Disease Type 2A1, Gene GDAP, Mut.)	25707
COMP, . . (Pseudoachondroplasia, Gene	
COMP, Freq. Mut.)	6820
Gene NPHS1, Mut.)	76989
NPHS1, . (Nephrotic Syndrome Type 1, NPHS1,	
Mut.)	12056
FGFR3, . . (Achondroplasia, Gene FGFR3, Freq.	
Syndrome, FCAS, Gene NLRP3, Mut.)	51348
NLRP3 . (Familial Cold Autoinflammatory	
O, . (Charcot-Marie-Tooth Disease Type 1B, Gene O, Mut.)	17160
DLL3, . (Spondylocostal Dysostosis, Gene	
DLL3, Mut.)	29975
NPHP1, . (Nephronophthisis 1, NPHP1, Gene NPHP1, Mut.)	17006
FHL1, . (Emery-Dreifuss	
Muscular Dystrophy, Gene FHL1, Mut.)	34254
ANKH, . (Chondrocalcinosis, Calcium Pyrophosphate	
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
ATXN7, . . (Spinocerebellar	
Ataxia, Gene ATXN7 Freq. Mut.)	5148
IGHMBP2, . (Distal Spinal Muscular Atrophy 1, DSMA1, Gene IGHMBP2, Mut.)	64163
RP2, . (Retinitis Pigmentosa, Gene	
RP2, Mut.)	21428
Syndrome, Type VI, Gene PLOD, Freq. Mut.)	10285
PLOD, . . (Ehlers-Danlos	
(Aarskog-Scott Syndrome, Faciodigitogenital Syndrome, Faciogenital Dysplasia, Gene FGD1, Mut.)	62722
FGD1, .	
Spinal and Bulbar Muscular Atrophy, Gene AR, Freq. Mut.)	5148
AR, . . (Kennedy	
PCSK9	11682
X	6369
/ , FMR1, . .	3410
GLI3, . (Pallister-Hall Syndrome, Gene GLI3,	
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
UNC13D, . .	
(Familial Hemophagocytic Lymphohistiocytosis, Gene UNC13D, Freq. Mut.)	5148
TNFRSF6, .	
(Autoimmune Lymphoproliferative Syndrome, ALPS, Gene TNFRSF6, Mut.)	34254
5, 8 RET, .	
(Familial Meddulary Thyroid Cancer, Exons 5, 8 Gene RET, Mut.)	9999
SBDS, . (Shwachman-Diamond	
Syndrome, Gene SBDS, Mut.)	21428
KCNJ2, . (Andersen-Tawil Syndrome, Gene KCNJ2,	
Mut.)	17523
ABCA4, . . (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 Nigrificans, CAN, Exon 10 Gene FGFR3, Mut.)	7161
Agammaglobulinemia, XLA, Gene BTK, Mut.)	76989
Progressive, Gene GRN, Mut.)	25707
	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.)	9999
Dominant, SCN1, Gene ELA2, Mut.)	21428
	4818
(Congenital Insensitivity To Pain With Anhidrosis, CIPA, Gene NTRK1, Mut.)	51348
Epiphyseal Dysplasia, MED, Gene COMP, Freq. Mut.)	6820
PHOX2B, . . (Congenital Central Hypoventilation Syndrome, CCHS, Gene PHOX2B, Freq. Mut.)	5148
(Genes CFTR, GJB2, PAH, SMN))	21142
Gene DMPK, Freq. Mut.)	5148
GJB1, . (Charcot-Marie-Tooth Disease Type 1B, Gene GJB1, Mut.)	9999
ongenital Ichthyosis, ARCI 1, All Known Mutations, Gene TGM1, Mut.)	42801
Motor Neuropathy, DHMN, Gene BSCL2, Mut.)	29975
Mut.)	64163
(Phosphoribosylpyrophosphate Synthetase Superactivity, PRS Superactivity, Gene PRPS1, Mut.)	29975
(Albinism oculocutaneous, Hermansky-Pudlak type, Gene HPS1, Freq. Mut.)	9999
	23540
(Klippel-Feil Syndrome, Gene GDF6, Mut.)	14916
. (Craniofacial Dysplasia, Gene ANKH, Hot-Point Mut.)	9999
(Nonbullous Congenital Ichthyosiform Erythroderma, NBCIE, Gene TGM1, Mut.)	42801
PRNP, Mut.)	14916
Gene CRYBA4, Mut.)	25707
	7590
(TNF-Receptor-Associated Periodic Syndrome, TRAPS, Gene TNFRSF1A, Mut.)	25707

(). Cerebelloparenchymal Disorder IV, CPD IV, Classic Joubert Syndrome, Joubert Syndrome type A, Joubert-Boltshauser Syndrome, Pure Joubert Syndrome, Gene NPHP1, Mut.)	NPHP1 (Joubert Syndrome,	17006
, 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
- congenital 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.)	SLC39A4, . (Acrodermatitis	34254
22, . (Hereditary Neuropathy with Liability to Pressure Palsies, HNPP, Gene 22, Mut.)	22, Mut.)	17523
. . (Osteopetrosis Autosomal Recessive 1, OPTB1, Gene TCIRG1, Freq. Mut.)	(). TCIRG1,	5148
SCN4A, . (Hypokalemic Periodic Paralysis Type 1, Exons 12, 18, 19 Gene SCN4A, Mut.)	12, 18 19 Gene SCN4A, Mut.)	14267
Mut.)	FLG, . (Ichthyosis Vulgaris, Gene FLG, Freq.	9999
	LDLR	12837
-IgD Mut.)	CD40LG, . (Hyper-IgD Syndrome, Gene CD40LG,	42801
IA, Gene TYR, Mut.)	1 . TYR, . (Albinism Oculocutaneous Type	21428
Dystrophy-Dystroglycanopathy, Gene FKR, Freq. Mut.)	FKRP, . . (Muscular	7876
Fundus Flavimaculatus Included, Gene ABCA4, Freq. Mut.)	ABCA4, . . (Stargardt Disease 1, STGD1,	12056
(Spondyloepiphyseal Dysplasia Tarda, SEDT, Gene TRAPPC2, Mut.)	(). TRAPPC2, .	17160
Pneumothorax, PSP, Gene FLCN, Mut.)	FLCN, . (Primary Spontaneous	51348
, NBN . . (Nijmegen Breakage Syndrome, NBS, Gene NBN, Freq. Mut.)		5148
PAX3, Mut.)	PAX3, . (Waardenburg Syndrome, WS, Gene	34254
Dystrophy Limb-Girdle Type 2A, Gene FKR, Mut.)	FKRP, . (Muscular	14916
	CTSK, . (Pyknodysostosis, PKND, Gene CTSK, Mut.)	25707
	PAH, . (Phenylketonuria, PKU, Gene PAH, Mut.)	51348
Syndrome, Gene DHCR7, Mut.)	(). DHCR7, . (Smith-Lemli-Opitz	38522
	CFTR, . . (Cystic Fibrosis, Gene CFTR, Freq. Mut.)	17006
		8800
, NOTCH3		12320
Muscular Dystrophy, X-Lyonization, Girls)	(Duchenne	8030
GDF6, Mut.)	GDF6, . (Microphthalmia Isolated 4, Gene	14916

		7590
SPM, Gene FHL1, Mut.)	FHL1, . (Scapulooperoneal Myopathy,	34254
Hemophagocytic Lymphohistiocytosis, Gene PRF1, Mut.)	PRF1, . (Familial	21428
Gene PRNP, Mut.)	PRNP, . (Fatal Familial Insomnia, FFI,	14916
	EXT1, . (Multiple Exostoses, Gene EXT1, Mut.)	51348
Myoclonic Epilepsy 1A Unverricht and Lundborg, Gene CSTB, Freq. Mut.)	CSTB, . . (Progressive	5148
ACVR1, « » . . (Fibrodysplasia Ossificans Progressiva, FOP, Gene ACVR1, Hot-Point Mut.)	« »	17160
	2, JPH3, . .	3410
(X-Linked Severe Combined Immunodeficiency, Gene IL2RG, Mut.)	IL2RG, .	17160
	NLRP3 . (Muckle-Wells Syndrome, MWS, Gene NLRP3, Mut.)	51359
Elasticum, Gene ABCC6, Freq. Mut.)	ABCC6, . . (Pseudoxanthoma	7161
	, CYB5R3 . (Methemoglobinemia, Gene CYB5R3, Mut.)	34254
Hypertension 1, PPH1, Gene BMPR2, Mut.)	BMPR2, . (Primary Pulmonary	64163
	FXN, . (Friedrich Ataxia, Gene FXN, Mut.)	21428
	NDP, . (Norrie Disease, Gene NDP, Mut.)	11264
(Nonbullous Congenital Ichthyosiform Erythroderma, NBCIE, Gene LOX12B, Mut.)	(). LOX12B, .	42801
Cardiomyopathy, Gene TNNT2, Mut.)	TNNT2, . (Familial Hypertrophic	51348
(Primary Open Angle Glaucoma 1A, POAG 1A, Gene CYP1B1, Mut.)	(). CYP1B1, .	17523
	FLT4, . (Lymphedema, Gene FLT4, Mut.)	111177
Thrombocytopenia, CAMT, Gene MPL, Mut.)	MPL, . (Congenital Amegakaryocytic	34254
Dystrophy Limb-Girdle Type 2A, Gene SGCB, Mut.)	SGCB, . (Muscular	25707
Hypodysplasia, Aplasia 1, Exons 10, 11, 13, 14, 15 Gene RET, Mut.)	10, 11, 13, 14, 15 RET, . (Renal	21428
Heteroplasia, POH, Gene GNAS, Mut.)	GNAS, . (Progressive Osseous	38522
	(), SOD1, .	7370
	1, TOR1A (DYT1), . .	3410
Syndrome, PPS, Gene IRF6, Mut.)	IRF6, . (Popliteal Pterygium	38522
PTEN, Mut.)	PTEN, . (Lhermitte-Duclos Syndrome, Gene	38522
Dystrophy-Dystroglycanopathy, Gene FKR1, Mut.)	FKR1, . (Muscular	14916
Syndrome, Gene PTEN, Mut.)	PTEN, . (Bannayan-Ruvalcaba-Riley	38522
VHL, . (Autosomal Recessive Erythrocytosis, Gene VHL, Mut.)	(c).	14916
disease type 1A (CMT1A))	, PMP22, (Charcot-Marie-Tooth	16577
Gene ANKH, Mut.)	ANKH, . (Craniometaphyseal Dysplasia,	51348
Disease, CGD, Gene CYBB, Mut.)	CYBB, . (Chronic Granulomatous	51348
Syndrome, Gene LMNA, Mut.)	LMNA, . (Hutchinson-Gilford Progeria	42801
(Crigler-Najjer Syndrome, Gene UGT1, Mut.)	(,). UGT1, .	21428
	MECP2, . (Retts Syndrome, Gene MECP2, Mut.)	21428
	HLA B51	3300

Analysis Gene SRY, Mut.)	SRY, . (Disorders Sex Determination,	5148
Dystrophy Limb-Girdle Type 2A, Gene SGCA, Mut.)	SGCA, . (Muscular	25707
NEFL, . (Charcot-Marie-Tooth Disease Type 2A1, Gene NEFL, Mut.)	NEFL, . (Charcot-Marie-Tooth Disease Type 2A1, Gene NEFL, Mut.)	25707
TNFRSF6, « . » . . (Autoimmune Lymphoproliferative Syndrome, ALPS, Gene TNFRSF6, Hot-Point Mut.)	TNFRSF6, « . » . . (Autoimmune Lymphoproliferative Syndrome, ALPS, Gene TNFRSF6, Hot-Point Mut.)	7161
-IgD Syndrome, Gene MVK, Hot-Point Mut.)	MVK, « . » . . (Hyper-IgD Syndrome, Gene MVK, Hot-Point Mut.)	9999
Fukuyama-Type, Gene FKTN, Mut.)	FKTN, . (Muscular Dystrophy	51348
ESC (.).	NR2E3, . (Enhanced S- one Syndrome, Goldmann-Favre Syndrome, Gene NR2E3, Mut.)	25707
Mut.)	IRF6, . (Van der Woude Syndrome, Gene IRF6, Mut.)	38522
Mut.)	RS1, . (Retinoschisis 1 X-Linked Juvenile, RS1, Gene RS1, Mut.)	25707
Hemophagocytic Lymphohistiocytosis, Gene UNC13D, Mut.)	UNC13D, . (Familial	76989
(Osteopetrosis Autosomal Recessive 1, OPTB1, Gene TCIRG1, Mut.)	TCIRG1, .	51348
1A Unverricht and Lundborg, Gene CSTB, Mut.)	CSTB, . (Progressive Myoclonic Epilepsy	14267
Mut.)	SH2D1A, . (X-Linked Lymphoproliferative Syndrome, XLP, Gene SH2D1A, Mut.)	17160
Dystrophy, All Known Mutations, Gene BEST1, Mut.)	BEST1, . (Best Vitelliform Macular	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, .	9999
Gene ABCC6, Mut.)	ABCC6, . (Pseudoxanthoma Elasticum,	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	10285
	(. - LDLR, APOB, PCSK9)	9218
Epiphysial Dysplasia, MED, Gene SLC26A2, Mut.)	SLC26A2, . (Multiple	25707
	PRPS1, . (Art's Syndrome, Gene PRPS1, Mut.)	29975
(Nonbullous Congenital Ichthyosiform Erythroderma, NBCIE, Gene ALOXE3, Mut.)	(.). ALOXE3, .	64163
	PAX3, . (Craniofacial-Deafness-Hand Syndrome, CDHS, Gene PAX3, Mut.)	34254
Muscular Dystrophy, Gene LMNA, Mut.)	LMNA, . (Emery-Dreifuss	42801
VHL, . (Autosomal Recessive Erythrocytosis, Gene VHL, Freq. Mut.)	(c .).	5148
(Keratitis-Ichthyosis-Deafness Syndrome, KID Syndrome, Gene GJB2, Mut.)	(.). GJB2, .	9999
« . » . . (Familial Partial Lipodystrophy 2, FPLD 2, Gene LMNA, Hot-Point Mut.)	« . » . . LMNA,	14267

TAZ, . (Barth Syndrome, Gene TAZ, Mut.)	25707
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.)	16577
EXT2, . (Multiple Exostoses, Gene EXT2, Mut.)	64163
ANO5, SGCA CAPN3, FKRP,	12056
CYP1B1, . (Primary Congenital Glaucoma 3A, PCG 3A, Gene CYP1B1, Mut.)	17523
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.)	29975
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.)	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
EBP, . (Chondrodysplasia Punctata, CDP, Conradi-Hunermann Syndrome, Gene EBP, Mut.)	17160
STXBP2, . (Familial Hemophagocytic Lymphohistiocytosis, Gene STXBP2, Mut.)	51348
LPIN1, . (Myoglobinuria Acute Recurrent Autosomal Recessive, Gene LPIN1, Mut.)	94083
ADAMTSL2, . (Geleophysic Dysplasia 1, Gene ADAMTSL2, Mut.)	76989
PANK2, . . (Neurodegeneration With Brain Iron Accumulation 1, Gene PANK2, Freq. Mut.)	7161
GJB4, . (Erythrokeratodermia, Gene GJB4, Mut.)	9999
SLC26A2, . (Atelosteogenesis II, De la Chapelle Dysplasia, Gene SLC26A2, Mut.)	25707
RPS19, . (Diamond-Blackfan Anemia 1, DBA1, Gene RPS19, Mut.)	21428
NPHS2, . (Nephrotic Syndrome Type 1, NPHS1, Gene NPHS2, Mut.)	34254
IX B, . (Hemophilia B, Gene Factor IX, Mut.)	29975
TRPS1, . (Trichorhinophalangeal Syndrome, TRPS, Gene TRPS1, Mut.)	42801
IGHMBP2, . (Spinal Muscular Atrophy (SMA) with Diaphragmatic Paralysis, Gene IGHMBP2, Mut.)	64163
FGFR3, . . (Hypochondroplasia, Gene FGFR3, Freq. Mut.)	12056
NS3, NS5A NS5B (1 , 1b)	12837
13 24 SCN4A, . (Hyperkalemic Periodic Paralysis Type 2, Exons 13, 24 Gene SCN4A, Mut.)	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) (4290
(Hereditary Breast Cancer In Men: Cancer of Breast, Pancreatic, Prostate, Testicular Cancer (Genes BRCA1, BRCA2) (without Description))		
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	4785
Cancer In Men: Cancer of Breast, Pancreatic, Prostate, Testicular Cancer (Genes BRCA1, BRCA2))		
MGMT		11682
PIK3CA		11682
228 250 TERT		8327
(Hereditary Breast and/or Ovarian Cancer, HBOC (Genes BRCA1, BRCA2) (without Description))	BRCA1, BRCA2) (4290
IDH1		11682
A09.05.039 (Lactate Dehydrogenase, LDH)	(L- +)	165
G6PD		3278
A09.05.042 (Alanine Aminotransferase, ALT, Serum Glutamic Pyruvic Transaminase, SGPT)		165
A09.05.045 (?-Amylase)	(Ipha- milase, ?-Amylase)	231
A09.05.180 (Pancreatic ?-Amylase)	(P-)	275
(Cholinesterase, Pseudocholinesterase, PCHE)	(S- II, S-)	253
A09.05.046 (Alkaline Phosphatase, ALP)		165
A09.05.173 (Lipase)		319
(Acid Phosphatase, ACP)		220
GGT (Gamma-Glutamyl Transferase,		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

the fetus in the mother's blood)	Y-	(Y-chromosome of	4510
(Rh factor Definition)			6820
(Plasminogen)			
VIII () (Antihemophilic Globulin A, FVIII)	660
A09.05.051.001 D-		(D-Dimer)	1133
A09.05.050		(Fibrinogen, FG)	1133
A12.05.039		(231
() (Activated Partial Thromboplastin Time, APTT)	165
IX,		% (484
(Christmas Factor, anti-hemophilic globulin "B")		« ») Factor IX, Activity,%	
A12.05.027		(253
Ratio, INR)) (Prothrombin, rothrombin Time, PT, International Normalized	
A12.05.028		() (Thrombin Time, TT)	253
A09.05.029.001		() (Lupus Anticoagulant, LA)	803
C, %		(Protein C, % Activity)	1969
-		/ (1837
concentration, IU/ml)		, /), Anti-Xa activity, IU/ml (Heparin	
		, % (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
-		(2376
Serum Protein Electrophoresis (SPEP), Immunofixation with Polyvalent Antiserum, Quantification of M-Protein (without Typing))) (M-Gradient, Screening.	
		(Bence-Jones Protein, Urine, Electrophoresis, Immunofixation, Kappa/Lambda	3300
Typing, Quantification)			
A09.05.014		(Serum Protein Electrophoresis, SPE, SPE)*	275
-		(4389
(M-Gradient, Typing. Serum Protein Electrophoresis (SPEP), Immunofixation with Antisera (IgG, IgA, IgM, Kappa, Lambda), Quantification of M-Protein)		IgG, IgA, IgM,	
Concentration of Immunoglobulin Free Light Chains)		(Cerebrospinal Fluid	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
		(693
) (i6) IgE, ImmunoCAP	

(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
(wx1) 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
		(CMV DNA, Scrape of Conjunctiva Epithelial Cells)*	275
		(CMV DNA, Cerebrospinal Fluid)*	275
		(Copper, random urine; Cu)	1298
	(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), lood)	1298
(Zn)	(Zinc (Zn), lood)	1298
	(Iodine, serum)	275
(Au)	(Gold (Au), Serum)	275
(Cu)	(Copper (Cu), Serum)	275
	()	748
(Cu)	(Copper (Cu), lood)	1298
(Pb)	(Lead (Pb), lood)	1298
(Mn)	(Manganese (Mn), lood)	1298
(Zn)	(Zinc (Zn), Serum)	275
(Li)	(Lithium (Li), serum)	275
(Cd)	(Cadmium (Cd), lood)	1298
(Se)	(Selenium (Se), Serum)	275
(Se)	(Selenium (Se), lood)	1298
(Mn)	(Manganese (Mn), Serum)	275
(Co)	(Cobalt (Co), lood)	1298
	(3 , Reverse Triiodthyronine).	6567
(Tl)	(Thallium (Tl), Serum)	275
	(())	748
(Mo)	(Molybdenum (Mo), Serum)	275

ATM (FISH,) (Analysis of ATM gene rearrangements (FISH, quantitative))		10835
MPL, (Analysis of MPL gene mutations, PCR, qualitative)		4829
BCL- 6 (der(3)(q27)) (FISH,) (Analysis of BCL- 6 gene rearrangements (der(3)(q27)) on paraffin slides (FISH Histology, quantitative))		14927
MLL/AF4 -t(4;11) (,) (Analysis of chimeric gene MLL/AF4 -t(4;11) (PCR, qualitative))		2486
BCL- 6 (der(3)(q27)) (FISH,) (Analysis of BCL- 6 gene rearrangements (der(3)(q27) (FISH, quantitative))		10835
13 - (del(13), -13) (FISH,) (Analysis of chromosome 13 monosomy, deletion - (del(13), -13) (FISH, quantitative))		10835
53 (FISH,) (Analysis of 53 gene deletion (FISH, quantitative))		10835
t(2;5)(p23;q35) (FISH,) (Analysis of translocation t(2;5)(p23;q35) on paraffin slides (FISH Histology, quantitative))		14927
(FISH,) (Analysis of all specific aberrations on paraffin slides (FISH Histology, quantitative))		14927
t(11;14)(q13;q32) (FISH,) (Analysis of translocation t(11;14)(q13;q32) on paraffin slides (FISH Histology, quantitative))		14927
t(11;14)(q13;q32) (FISH,) (Analysis of translocation t(11;14)(q13;q32) (FISH, quantitative))		10835
PML/RAR? -t(15;17) (,) (Analysis of chimeric gene PML/RAR? -t(15;17) (PCR, qualitative))		2486
CBF?/MYH1- inv(16),t(16;16) (,) (Analysis of chimeric gene CBF?/MYH1- inv(16),t(16;16) (PCR, qualitative))		2486
12 (+12) (FISH,) (Analysis of chromosome 12 trisomy (FISH, quantitative))		10835
V617F 14 JAK2 (Qualitative assessment of presence of gene JAK2 617F somatic mutation)		1947
t(11;18)(q21;q21) (FISH,) (Analysis of translocation t(11;18)(q21;q21) (FISH, quantitative))		10835
(Karyotype, Hematologic Disorders, Peripheral Blood)		7722
53 (FISH,) (Analysis of 53 gene deletion (FISH, quantitative))		10835
t(14;16) (IGH/MAFB) (FISH,) (Analysis of translocation t(14;16) (IGH/MAFB) (FISH, quantitative))		10835

BCR/ABL - t(9;22), (Analysis of chimeric gene BCR-ABL - t(9;22), assessment of the BCR-ABL gene transcript type, PCR, qualitative)	BCR/ABL - (Analysis of chimeric gene BCR-ABL - t(9;22), assessment of the BCR-ABL gene transcript type, PCR, qualitative)	2486
12p (FISH,) (Analysis of 12p deletion (FISH, quantitative))	12p (FISH,) (Analysis of 12p deletion (FISH, quantitative))	10835
BCR-ABL (FISH,) (Analysis of chimeric gene BCR-ABL, FISH, quantitative)	BCR-ABL (FISH,) (Analysis of chimeric gene BCR-ABL, FISH, quantitative)	10835
BCL2 (FISH,) (Analysis of BCL2 gene rearrangements on paraffin slides (FISH Histology, quantitative))	BCL2 (FISH,) (Analysis of BCL2 gene rearrangements on paraffin slides (FISH Histology, quantitative))	14927
PDGFR?(FISH,) (Analysis of gene rearrangements PDGFR? (FISH, quantitative))	PDGFR?(FISH,) (Analysis of gene rearrangements PDGFR? (FISH, quantitative))	10835
FGFR1 (FISH,) (Analysis of gene rearrangements FGFR1 (FISH, quantitative))	FGFR1 (FISH,) (Analysis of gene rearrangements FGFR1 (FISH, quantitative))	10835
E2A/PBX1 - t(1;19) (,) (Analysis of chimeric gene E2A/PBX1 - t(1;19) (PCR, qualitative))	E2A/PBX1 - t(1;19) (,) (Analysis of chimeric gene E2A/PBX1 - t(1;19) (PCR, qualitative))	2486
BRAF (V600E) (,)	BRAF (V600E) (,)	8569
5 (FISH,) (Analysis of chromosome 5 rearrangements (FISH, quantitative))	5 (FISH,) (Analysis of chromosome 5 rearrangements (FISH, quantitative))	10835
t(4;14)(p16;q32) (FISH,) (Analysis of translocation t(4;14)(p16;q32) (FISH, quantitative))	t(4;14)(p16;q32) (FISH,) (Analysis of translocation t(4;14)(p16;q32) (FISH, quantitative))	10835
t(14;16) (IGH/MAFB) (FISH,) (Analysis of translocation t(14;16) (IGH/MAFB) (FISH,quantitative))	t(14;16) (IGH/MAFB) (FISH,) (Analysis of translocation t(14;16) (IGH/MAFB) (FISH,quantitative))	10835
BCR/ABL - RQ (,) (Analysis of the BCR/ABL relative expression, RQ-PCR, quantitative)	BCR/ABL - RQ (,) (Analysis of the BCR/ABL relative expression, RQ-PCR, quantitative)	4829
() (Cytogenetic analysis of bone marrow (karyotype))	() (Cytogenetic analysis of bone marrow (karyotype))	7722
FIP1L1/PDGFR?(FISH,) (Analysis of chimeric gene FIP1L1/PDGFR? (FISH, quantitative))	FIP1L1/PDGFR?(FISH,) (Analysis of chimeric gene FIP1L1/PDGFR? (FISH, quantitative))	10835
MLL (FISH,) (Analysis of MLL gene rearrangements (FISH, quantitative))	MLL (FISH,) (Analysis of MLL gene rearrangements (FISH, quantitative))	10835
7 (FISH,) (Analysis of chromosome 7 rearrangements (FISH, quantitative))	7 (FISH,) (Analysis of chromosome 7 rearrangements (FISH, quantitative))	10835
13 - (del(13), -13) (FISH,) (Analysis of chromosome 13 monosomy, deletion - (del(13), -13) (FISH,quantitative))	13 - (del(13), -13) (FISH,) (Analysis of chromosome 13 monosomy, deletion - (del(13), -13) (FISH,quantitative))	10835
12 JAK2 (,) (Analysis of JAK2 Exon 12 mutations (PCR qualitative))	12 JAK2 (,) (Analysis of JAK2 Exon 12 mutations (PCR qualitative))	4829
BCL2 t(14;18)(q32;q21),t(2;18)(p11;q21),t(18;22)(q21;q11) (FISH,) (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,) (Analysis of BCL2 gene rearrangements t(14;18)(q32;q21),t(2;18)(p11;q21),t(18;22)(q21;q11) (FISH, quantitative))	10835
BCR-ABL (,) (BCR-ABL1 Mutation Analysis using direct Sanger sequencing, qualitative)	BCR-ABL (,) (BCR-ABL1 Mutation Analysis using direct Sanger sequencing, qualitative)	9636
1 (FISH,)	1 (FISH,)	14267
3q (FISH,) (Analysis of 3q rearrangements (FISH, quantitative))	3q (FISH,) (Analysis of 3q rearrangements (FISH, quantitative))	10835
RUNX1/RUNX1T1 -t(8;21) (,) (Analysis of chimeric gene RUNX1/RUNX1T1 -t(8;21) (PCR, qualitative))	RUNX1/RUNX1T1 -t(8;21) (,) (Analysis of chimeric gene RUNX1/RUNX1T1 -t(8;21) (PCR, qualitative))	2486
20q (FISH,) (Analysis of 20q deletion (FISH, quantitative))	20q (FISH,) (Analysis of 20q deletion (FISH, quantitative))	10835
MYC (t(8;14)(q24;q32)-t(2;8)(p11;q24), t(8;22)(q24;q11)) (FISH,) (Analysis of MYC gene rearrangements (t(8;14)(q24;q32)-t(2;8)(p11;q24), t(8;22)(q24;q11) (FISH, quantitative))	MYC (t(8;14)(q24;q32)-t(2;8)(p11;q24), t(8;22)(q24;q11)) (FISH,) (Analysis of MYC gene rearrangements (t(8;14)(q24;q32)-t(2;8)(p11;q24), t(8;22)(q24;q11) (FISH, quantitative))	10835
CALR (,) (Analysis of CALR gene mutations, deletions, insertions, PCR, qualitative)	CALR (,) (Analysis of CALR gene mutations, deletions, insertions, PCR, qualitative)	4829
IGH (FISH,) (Analysis of IGH gene rearrangements (FISH, quantitative))	IGH (FISH,) (Analysis of IGH gene rearrangements (FISH, quantitative))	10835
t(2;5)(p23;q35) (FISH,) (Analysis of translocation t(2;5)(p23;q35) (FISH, quantitative))	t(2;5)(p23;q35) (FISH,) (Analysis of translocation t(2;5)(p23;q35) (FISH, quantitative))	10835
:		
		1320
() (Additional research 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))	19888
(Koprogramma, Stool)	374
	4059
	6127
(Fecal Calprotectin)	2618
	3542
-1- (Alpha-1-Antitrypsin, Feces)	1727
	1463
	253
() (PRO Stool, Helminth Eggs)	308
	1980
(), FOB Gold (Quantitative Immunochemical Fecal Occult Blood, Test FOB Gold)	715
1 (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: , IgE (FP50 (F84, F91, F92, F210), Food Panel: Kiwi Fruit, Mango, Banana, Pineapple, IgE)*	1045
, IgE (Pumpkin, IgE, F225)	484
, IgE (Hazelnut, IgE, F17)	484
, IgE (Orange, IgE, F33)	484
, (e3) IgE, ImmunoCAP (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 (Rb), 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 (La), 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
(ScI-70, ENP-A, CENP-B, RP11, RP155, NOR90, Th/To, PM-Sc100, PM-ScI75, Ku, PDGFR, Ro-52), (Scleroderma (Systemic Sclerosis) Antibody Panel: Anti-ScI-70, ENP-A, CENP-B, RP11, RP155, NOR90, Th/To, PM-ScI100, PM-ScI75, 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, ScI-70, PM-ScI, 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, ScI-70, PM-ScI, 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))	4048
	(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
	(Anti-HCV Total (IgG + IgM))*	396
	(Anti-HCV IgG, Immunoblot)	5621
	(HCV RNA, Serum, Qualitative)*	693
	(HCV RNA, Serum, Quantitative, PCR)*	3454
	(Hepatitis C Virus (HCV) RNA, Plasma, Genotyping, Subtypes (Types 1 (Subtypes 1a, 1b), 2, 3))*	924
	(HCV RNA, Plasma, Quantitative)*	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
(Herpes simplex virus, HSV-1, HSV-2)		
	(HSV-1, 2 DNA, Urine)*	275
	(HSV-1, 2 DNA, Serum, yping)*	660
	(HSV-1, 2 DNA, Scrape of Conjunctiva Epithelial Cells)*	275
	(HSV-1, 2 DNA, Scrape of Conjunctiva Epithelial Cells, yping)*	462
	(HSV-1, 2 DNA, Urine, yping)*	462
	(Anti-HSV-1, 2 IgG)	484

1 2 , , (HSV-1, 2 DNA, Blood, typing)*		660
1 2 , , (HSV-1, 2 DNA, Cerebrospinal Fluid, typing)*		462
1 2 , , (HSV-1, 2 DNA, Exudate)*		275
1 2 , , (HSV-1, 2 DNA, Scrape of Faucial Epithelial Cells, typing)*		462
Ig 1 2 (Anti-HSV-1, 2 Ig)		506
1 2 , , (HSV-1, 2 DNA, Scrape of Urogenital Epithelial Cells, typing)*		462
1 2 , , (HSV-1, 2 DNA, Saliva, typing)*		462
1 2 , , (HSV-1, 2 DNA, Exudate, typing)*		462
1 2 , , (HSV-1, 2 DNA, Blood)*		418
1 2 , , (HSV-1, 2 DNA, Saliva)*		275
1 2 , , (HSV-1, 2 DNA, Scrape of Skin Epithelial Cells)*		275
1 2 , , (HSV-1, 2 DNA, Scrape of Nasal Epithelial Cells, typing)*		462
1 2 , , (HSV-1, 2 DNA, Cerebrospinal Fluid)*		275
1 2 , , (HSV-1, 2 DNA, Prostatic Fluid, Semen)*		275
1 2 , , (HSV-1, 2 DNA, Prostatic Fluid, Semen, typing)*		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, typing)*		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-IgG- (N-, S-proteins) antibodies, IgG, qualitative	(SARS-CoV-2 (N-, S-), (Post-vaccination (EpiVacCorona Vector) SARS-CoV-2	2486
SARS-CoV-2, (Coronavirus SARS-CoV-2 RNA detection in nasopharyngeal and oropharyngeal smear)		2189
spike (S) protein (RBD), IgG, quantitative.	(S) (RBD) SARS-CoV-2, IgG (Anti-SARS-CoV-2,	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 (nucleocapsid protein), IgG, Abbott	(SARS-CoV-2 (), IgG, (Anti-SARS-CoV-2	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), pneumoniae, DNA, Plasma)*	(Mycoplasma pneumoniae)	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 (Anti-Mycoplasma hominis Ig)	462
	(Mycoplasma pneumoniae), pneumoniae, DNA, Sputum)*	(Mycoplasma pneumoniae)	649
	(Mycoplasma hominis), (Mycoplasma hominis, DNA, Prostatic Fluid, Semen)*		275
	IgG	Mycoplasma hominis (Anti-Mycoplasma hominis IgG)	462
	(Mycoplasma genitalium), (Mycoplasma genitalium, DNA, Prostatic Fluid, Semen)*		275
	(Mycoplasma pneumoniae), DNA, Saliva)*	(Mycoplasma pneumoniae)	231
	(Mycoplasma hominis), Urine)*	(Mycoplasma hominis, DNA)	275
	IgA	Mycoplasma hominis (Anti-Mycoplasma hominis IgA)	671
	(Mycoplasma genitalium), Urine)*	(Mycoplasma genitalium, DNA)	275
A09.05.127	(Mg)	(Magnesium (Mg), Serum)	253
A09.05.032	(Ca)	(Calcium Total)	209
	/	(Potassium, Na+ /Sodium, Cl-/Chloride, Serum)	275
	(Ca ²⁺ , c)	(Ionized Calcium, Free Calcium)	396
A09.05.033	(P)	(Phosphorus (P))	209
	(UIBC)	(Unsaturated Iron Binding Capacity, UIBC)	209
A09.05.007	(Fe)	(Iron (Fe), Serum)	209
	Helicobacter pylori (Associated Gastritis)*	(Helicobacter pylori)	4004
	PDGFRa		15070
	1	(PDGFRa):	1694
	PD-L1 clone SP263 (Ventana) antibodies.	PD-L1 expression in tumor tissue by IHC using PD-L1 clone SP263 (Ventana) antibodies.	18150
	KIT		12760
	Finished Histological Preparations (1 Glass + 1 Block)	(KIT + 1) (Consultation of)	1430
	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 clone SP142 (Ventana) antibodies.	PD-L1 expression in tumor tissue by IHC using PD-L1 clone SP142 (Ventana) antibodies.	18150
	(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 22 3 (Dako). (PD-L1 expression in tumor tissue by IHC using PD-L1 clone 22C3 (Dako) antibodies).	26400
PDL1	8140
() 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
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
(Trichomonas vaginalis, DNA, Prostatic Fluid, Semen)*	275
(Trichomonas vaginalis, DNA, Scrape of Urogenital Epithelial Cells)*	275
IgG Trichomonas vaginalis (nti-Trichomonas vaginalis IgG)	671
(Procedural services)	
(Blood serum derivation without further examinations)**	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
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(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, Sexually Transmitted Infections (STI) Pathogens, Chlamydia trachomatis DNA, Neisseria gonorrhoeae DNA, Trichomonas vaginalis DNA, Mycoplasma genitalium DNA, Human DNA)	(Identification of	1023
(Mobiluncus curtisii, DNA, Scrape of Urogenital Epithelial Cells)		231
(Mycoplasma hominis, Effectiveness Monitoring of Treatments)	(Mycoplasma hominis)	396
		1870
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

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

- , ()		
39, 45, 51, 52, 56, 58, 59, 66, 68 + , o	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
, o	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 + , o	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

Epithelial Cells, 3 Types (6, 11, 44)	3 : 6, 11, 44 + (HPV DNA, Scrape of Rectal	385
Epithelial Cells, 2 Types (16, 18)	16 18 + (HPV DNA, Scrape of Urogenital	385
(6, 11, 44)	3 : 6, 11, 44 (HPV DNA, Scrape of Faucial Epithelial Cells, 3 Types	385
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, 18,	990
Epithelial Cells, 3 Types (6, 11, 44)	3 : 6, 11, 44 + (HPV DNA, Scrape of Urogenital	385
39, 45, 51, 52, 56, 58, 59, 66, 68 + (HPV DNA, Scrape of Faucial Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68) Screening)*	14 : 16, 18, 31, 33, 35,	385
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))	21 : 6, 11, 16, 18, 26, 31, 33, 35, 39, 44, 45, 51,	2695
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, 18, 31, 33, 35,	385
(Candida albicans, DNA, Exudate)*		275
(Candidiasis, Typing)		737
(Candida, Cryptococcus) (Yeast Culture. Identification and Antimycotic Susceptibility testing)		759
(Candidiasis, Screening)		396
Semen)* (Candida albicans, DNA, Prostatic Fluid,		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
Scrape of Skin Epithelial Cells)* (Candida albicans, DNA,		275
albicans, DNA, Scrape of Urogenital Epithelial Cells)* (Candida		275
(Candida albicans, DNA, Saliva)*		275
()		1837
Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)* (Upper Respiratory		2178
(Eye Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*		1474
Enlarged Testing)* (Eye Culture. Bacteria Identification, Antibiotic Susceptibility,		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 Testing)*	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), Immunohromotographic Assay), (Adenovirus. One Step Rapid	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), Immunohromotographic Assay), (Helicobacter pylori. One Step Rapid	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
LG1 CASPR2 (), IgG, (VGKC-associated proteins LG1 and CASPR2 antibodies, serum)	6160
, IgG, (Neuronal antibodies, IgG, Indirect immunofluorescence (IIF))	3311
(Acetylcholine Receptor Antibodies, Anti-AChR, Total)	5731
IgG, NMDA, CASPR, LG1, AMPA1, AMPA2, GABAR1	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-Gangliosideantibodies, Ganglioside Antibodies Panel, Total)	5731

Crithidia luciliae, IgG, fluorescent test (CLIFT)	(Crithidia luciliae indirect	1265
IgG, IF)	(Anti-myelin antibody, IgG,	1474
IgA, IgG, IgM 4, (NMO) (Aquaporin-4 Receptor 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
IgG, NMDA, CASPR, LGI, AMPA1, AMPA2, GABAR1		13750
LG11 CASPR2 (VGKC-associated proteins LG11 and CASPR2 antibodies, CSF)	, IgG,	6160
IgG, CSF) GAD (, IgG, (Anti-GAD (glutamic acid decarboxylase),		2079
(ANNA2), (Anti-Neuronal Antibodies, Blot-Line (Hu (ANNA1), Yo-1 (PCA1), CV2, 2, Ri (ANNA2), Amphiphysin))	IgG (- : Hu (ANNA1), Yo-1 (PCA1), CV2, 2, Ri (ANNA2), Amphiphysin))	5731
IgG NMDA (N- -D-) (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
IgG (Anti-Rubella IgG, Immunoblot)		5621
Ig (Anti-Rubella Ig)		539
Anti-Rubella IgG		1067
(Rubella virus, RNA)		649
IgG (Anti-Rubella IgG)		396
B (Anti-HBc IgM, IgG, Antibodies to Hepatitis B Core Antigen; HBcAb, Total, HBV Core Total Antibodies (IgG + IgM))	B, (Anti-HBc IgM,	517
HBs- (HBs- (HBsAg, Hepatitis Surface Antigen, Quantitative)	B,	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
Quantitative)* B, (HBV DNA, Serum,		3751
Qualitative)* B, (HBV DNA, Serum,		418
HBs- (Anti-HBs, HBsAb)		627
HB - (Anti-HBe, HBeAb)		517
HBs- (HBs- (HBsAg, Hepatitis Surface Antigen, Qualitative)	B,	264
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		143
(Lactate)		572
(Fructosamine)		891

A09.05.083	HbA1 (HbA1 , Glycated Hemoglobin, GHB)	506
	() Oral Glucose Tolerance Test, Plasma, OGTT, Pregnancy	990
A12.22.005	2 (2-Hour Oral Glucose Tolerance Test, OGTT, Glucose Concentration (Fasting and 2 Hours after Load), Venous Blood)	671
A09.05.009	() (C-Reactive Protein, CRP)	341
	(- ,) (Antistreptolysin-O, ASO)	374
-1-	(1), (Alpha-1-Antitrypsin, A1AT, AAT, Phenotyping)	2596
	(-) N- (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
	ST2 (ST2, sST2,) (Soluble ST2 (Heart Failure's biomarker))	2739
A09.05.008	() (Transferrin)	484
	(Myoglobin)	572
-2-	(Alpha-2-Macroglobulin, ?2-Macroglobulin, A2M)	495
Pgp3 (IgG) 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 (Anti-Respiratory Syncytial Virus (RSV) IgG)	781

A (RSV) IgM	IgM	-	(Anti-Respiratory Syncytial Virus)	781
(Genes F2, F5)	e	(F2, F5) (Risk of Oral Contraceptives, Ocs	3003
-	e	(F2, F5) (Risk of Oral Contraceptives, OCs (Genes F2, F5) (without Description))	2695
)	, 6	(AZF) (3916
MTHFR, MTRR, MTR, F2, F5)	(MTHFR, MTRR, MTR, F2, F5) (Preparation for Surgery (Genes		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
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
Food Panel: Pork, Beef, Chicken Meat, Lamb, IgG)*				1045
Food Profile, IgG				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
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	,	(HHV-6 DNA, Prostatic Fluid, Semen)*	275
6	,	(HHV-6 DNA, Scrape of Nasal Epithelial Cells)*	275
6	,	(HHV-6 DNA, Urine)*	275
6	,	(HHV-6 DNA, Cerebrospinal Fluid)*	275
6	,	(HHV-6 DNA, Exudate)*	275
6	,	(HHV-6 DNA, Blood)*	418
6	,	(HHV-6 DNA, Scrape of Faucial Epithelial Cells)*	275
6	,	(HHV-6 DNA, Serum)*	418
()			
(Chlamydia pneumoniae), DNA, Saliva)*		(Chlamydia pneumoniae, Chlamydia pneumoniae, DNA, Prostatic Fluid, Semen)*	429
IgG	()	Chlamydia trachomatis (Anti-cHSP60 IgG)	583
A	IgA	Chlamydia trachomatis (Anti-Chlamydia trachomatis IgA)	539
(Chlamydia pneumoniae), pneumoniae, DNA, Plasma)*		(Chlamydia pneumoniae, Chlamydia pneumoniae, DNA, Plasma)*	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)*		(Chlamydia trachomatis, DNA, Scrape of Rectal Epithelial Cells)*	275
(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Scrape of Urogenital Epithelial Cells)*		(Chlamydia trachomatis, DNA, Scrape of Urogenital Epithelial Cells)*	275
A	IgG	Chlamydia trachomatis (Anti-Chlamydia trachomatis IgG)	539
		(Helicobacter pylori, DNA, Biopsies of Gastric Mucosa and/or Duodenum, PCR)	2332
(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Scrape of Conjunctiva Epithelial Cells)*		(Chlamydia trachomatis, DNA, Scrape of Conjunctiva Epithelial Cells)*	275
A	IgA	Chlamydia pneumoniae (Anti-Chlamydia pneumoniae IgA)	671
(Chlamydia pneumoniae), DNA, Sputum)*		(Chlamydia pneumoniae, Chlamydia pneumoniae, DNA, Sputum)*	869
A	IgM	Chlamydia pneumoniae (Anti-Chlamydia pneumoniae IgM)	561
(Chlamydia trachomatis), Urine)*		(Chlamydia trachomatis, DNA, (Chlamydia trachomatis, DNA, Urine)*	275
(Chlamydia trachomatis), trachomatis, DNA, Cerebrospinal Fluid)*		(Chlamydia trachomatis, DNA, Cerebrospinal Fluid)*	275
(Chlamydia trachomatis), Exudate)*		(Chlamydia trachomatis, DNA, (Chlamydia trachomatis, DNA, Exudate)*	275
(Chlamydia trachomatis), trachomatis, DNA, Synovial Fluid)*		(Chlamydia trachomatis, DNA, Synovial Fluid)*	506

(Chlamydia pneumoniae), (Chlamydoghila 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, (Treponema pallidum, DNA, Scrape of Urogenital Epithelial Cells)*	275
(Treponema pallidum, DNA, Urine)*	275
DNA, Cerebrospinal Fluid)* (Treponema pallidum, (Treponema pallidum, DNA, Serum)*	407
(Treponema pallidum, DNA, Secretion)*	275
A IgM IgG Treponema pallidum, (Anti-Treponema pallidum IgM, IgG, Total)	418
pallidum, DNA, Scrape of Skin Epithelial Cells)* (Treponema (Treponema pallidum, DNA, Scrape of Faucial Epithelial Cells)*	275
A IgG Treponema pallidum, (Anti-Treponema pallidum IgG, Immunoblot)	1969
(Treponema pallidum, DNA, Scrape of Conjunctiva Epithelial Cells)*	275
A IgM Treponema pallidum (Anti-Treponema pallidum IgM)	891
RPR - (Syphilis RPR (Rapid Plasma Reagins), nticardioliipin est)	231
A IgM Treponema pallidum, (Anti-Treponema pallidum IgM, Immunoblot)	1969
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, 3?-Androstanediol Glucuronid, 3?-diol G)	1155
() (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
-			
	()		
	1 2 1 2 (HIV Ag/Ab Combo)		319
	-1, (HIV RNA, Plasma)*		14234
-			
	(Everolimus)		3630
	(Cyclosporine, Cyclosporine A, Sandimmune)		1045
	() (Teriflunomide, Leflunomide metabolite)		3630
	(Levetiracetam, Keppra®)		3740
	() (carbamazepine, Tegretol)		2882
	(Mitotane, o, p?-DDD, plasma)		3630
	() (Phenytoin)		1276
	() (FK506, Advagraf, Prograf, Protopic, Tacrosel)		1573
	(Lamotrigine)		3740
	() (Acidum Valproicum, Depakin, Convulexs)		891
	A09.05.035.002 () (Phenobarbitalum)		2882
-			
	()		
	(Neisseria gonorrhoeae, DNA, Prostatic Fluid, Semen)*		275
	(Neisseria gonorrhoeae, DNA, Scrape of Rectal Epithelial Cells)*		275
	(Neisseria gonorrhoeae, DNA, Synovial Fluid)*		506
	(Neisseria gonorrhoeae,), (GC, Neisseria gonorrhoeae Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		990
	(Neisseria gonorrhoeae, DNA, Scrape of Conjunctiva Epithelial Cells)*		275
	(Neisseria gonorrhoeae, DNA, Scrape of Faucial Epithelial Cells)*		275
	(Neisseria gonorrhoeae, DNA, Urine)*		275
	(Neisseria gonorrhoeae, DNA, Scrape of Urogenital Epithelial Cells)*		275
-			
	(Lactobacillus spp., DNA, Scrape of Urogenital Epithelial Cells)*		352
-			
	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 (Dermatophagoides farinae, IgG, D2)		572
/Greer (1), - IgG (House Dust – Greer, IgG, H1)		572
Cladosporium herbarum (2), - IgG (Cladosporium herbarum, IgG, M2)		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), - IgG (Dermatophagoides microceras, IgG, D3)		572
Dermatophagoides pteronyssinus (D1), - IgG (Dermatophagoides pteronyssinus, IgG, D1)		572
Alternaria tenuis (6), - IgG (Alternaria tenuis, IgG, M6)		572
: (AZF-) (-) (Impairment of Spermatogenesis: Full Panel (AZF-Region) (without Description))		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))		17270
, CYP21A2, . . . (Gene CYP21OHB, Freq. Mut.)		10769
: (F2, F5) (Thrombotic Tendency in Pregnancy: Minimum (Genes F2, F5))		3003
(AR, CFTR; AZF-) (Genetic Factors of Male Infertility (Genes AR, CFTR; AZF-Region))		18062
(MTHFR, MTRR, MTR) (Isolated Malformations in Fetus (Genes MTHFR, MTRR, MTR))		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))		19998
: (MTHFR, MTRR, MTR, F2, F5) (-) (Habitual Miscarriage, Thrombotic Tendency in Pregnancy: Extended Panel (Genes MTHFR, MTRR, MTR, F2, F5) (without Description))		7898
(F2, F5) (-) (Thrombotic Complications of Ovulation Induction (Genes F2, F5) (without Description))		2695
(MTHFR, MTRR, MTR) (-) (Isolated Malformations in Fetus (Genes MTHFR, MTRR, MTR) (without Description))		5203
(ACE, AGT, MTHFR, MTRR, MTR, F2, F5) (Gestosis and Placental Insufficiency (Genes ACE, AGT, MTHFR, MTRR, MTR, F2, F5))		11671
(F2, F5) (Thrombotic Complications of Ovulation Induction (Genes F2, F5))		3003
: (F2, F5) (-) (Thrombotic Tendency in Pregnancy: Minimum (Genes F2, F5) (without Description))		2695

(MTHFR, MTRR, MTR, F2, F5) (Habitual Miscarriage, Thrombotic Tendency in Pregnancy: Extended Panel (Genes MTHFR, MTRR, MTR, F2, F5))		8811
(ACE, AGT, MTHFR, MTRR, MTR, F2, F5) (MTHFR, MTRR, MTR, F2, F5) (without Description))	(Gestosis and Placental Insufficiency (Genes ACE, AGT, MTHFR, MTRR, MTR, F2, F5) (without Description))	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
Endoscopic Material: Presence of Helicobacter pylori)	Helicobacter pylori (Examination of	781
	(Cytological Examination of Material Obtained during Surgical Procedures and Other Urgent Research)	990
(ThinPrep ®)*		1320
(The Bethesda System ? TBS) (Cytological Examination of Cervical Epithelium with Description on The Bethesda System, TBS)		627
(Localization))	() (Cytological Examination: Scrapings (Smear) of Nasal Mucous Membrane (1	781
	(Examination of Breast Discharge)	528
Tissues)	(Examination of Punctates: Other Organs and	825
Device, IUD)	() (Examination of Imprint Intrauterine	561
	(Examination of Punctates: Breast)	649
of Scrapings and Prints Tumor and Tumor Like Formations)	(Examination	649
	(Examination of Urine)	528
	(The Bethesda System for Reporting Thyroid Cytopathology (TBSRTC), Fine-Needle Aspiration (FNA))	627
Scrapings and Prints of Skin and Mucous Membranes)	(Examination of	429
and Cervical Canal)	(Examination of Scrapings: Cervix	627
Ig	(nti-Measles IgM)	781
IgG	(nti-Measles IgG)	880
IgG)	IgG (Anti-Tick-borne Encephalitis Virus (TBEV)	528
	(Detection of pathogen DNA/RNA in ticks: Tick-borne encephalitis Virus (TBEV), Borrelia burgdorferi s. l., Anaplasma Phagocytophillum, Ehrlichia muris/chaffe nsis (RNA/DNA), PCR)	3696
IgM)	IgM (Anti-Tick-borne Encephalitis Virus (TBEV)	671
- -	() (HPLC-MS/MS Organic Acids (Succinylate))	3949

HADHA (3-Hydroxyacyl-CoA Dehydrogenase (LCHAD) Deficiency)		5511
e « » (Newborn Screening "HEEL")*		5489
Carboxylases Activity (Biotinidase Deficiency)		5511
GCDH (Glutaryl-CoA Dehydrogenase) Gene, Freq. Mut. (Glutaric Aciduria, Type 1)		5511
(/) (Analysis of the spectrum of organic urine acids by gas chromatography with mass spectrometry (GC / MS))		9130
ASS (ASS Gene, Freq. Mut. (Citrullinemia))		10483
GCDH (Glutaryl-CoA Dehydrogenase) Gene (Glutaric Aciduria, Type 1)		44440
ACADM (ACADM Gene, Freq. Mut. (Medium-Chain Acyl-CoA Dehydrogenase (MCAD) Deficiency))		5511
FAH (FAH Gene, Freq. Mut. (Tyrosinemia, Type 1))		9174
BTD (BTD (Biotinidase Deficiency) Gene, Freq. Mut.)		5511
FAH (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 chromatographic 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 chromatographic assay, antigen, urinae)		1584
(Respiratory Syncytial Virus, RSV, One step rapid immun chromatographic assay, antigen)		1023
(Hexagon Chlamydia, One step rapid immun chromatographic assay, antigen)		1122
(Neisseria gonorrhoeae test, One step rapid immun chromatographic assay)		946
(Campylobacter spp., One step rapid immun chromatographic assay, antigen, stool)		1144
(Norwalk virus) - I II (Norwalk virus GI, GII, One step rapid immun chromatographic assay, antigen, stool)		1925
(Streptococcus pneumoniae, One step rapid immun chromatographic assay, antigen, urinae)		1584
(Enterovirus, One step rapid immun chromatographic assay, antigen, stool)		1199
IgG -3 (Desmoglein 3, DSG3 Antibodies, IgG)		2376
IgG BP230 (Anti-Bp230 antibodies, Bullous Pemphigoid (230 kDa) Antibodies, Antibodies to BP Antigen 1, IgG)		2376

IgG BP180 (Anti-Bp180 antibodies, 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 pylori (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 pylori (Anti-Helicobacter pylori IgG)		539
1303HEL ?? - Helicobacter pylori (?? - , 13C-Urea Breath test, UBT).		2398
A IgM Helicobacter pylori (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
()		
Total) () (Circulating Immune Complexes (CIC)		1188
(Phagocytic Activity of Leucocytes)		1100
(Lymphocyte Activation Ability)		3795
CD4+ - , % (- , CD4+ T-cells, Percent and Absolute)		1507
(CD3+ HLA-DR+ , CD3-HLA DR+) (Activated Lymphocyte: CD3+ HLA-DR+ , CD3-HLA DR+)*		1507
- , % (CD19+ , B-cells, Percent and Absolute)		1507
() - CD3, CD4, CD8, CD19, CD16, CD56 (Lymphocyte Phenotyping: CD3, CD4, CD8, CD19, CD16, CD56)		3795
G (IgG1, IgG2, IgG3, IgG4)		13761
()		
IgM (Anti-Varicella-Zoster Virus IgM, Anti-VZV IgM)		858
Varicella-Zoster, (Varicella ZosterVirus, DNA, serum)		396
IgG (Anti-Varicella-Zoster Virus IgG, Anti-VZV IgG)		781
Varicella-Zoster, (Varicella Zoster Virus, DNA, scrape of faucial epithelial cells)		396
Varicella-Zoster, (VaricellaZosterVirus, DNA, saliva)		396
()		
(Stool Culture, Salmonella s p., Shigella s p. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)		1265

Shigella flexneri 1-5 (Shigella flexneri 1-5, IHA)	484
(Stool Culture (Salmonella spp., Shigella spp.). Bacteria Identification)	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
: - - - (P504S, AMACR), (34?E12), p63 (Prostate cancer – complex immunomorphological examination using assessment of the expression AMACR, high molecular weight cytokeratin (34?E12), p63)*	12540
() : () (Immunohistochemical diagnosis of lymphoproliferative diseases (Tissue Embedded in Paraffin Block))	27808
Ki-67 (MIB-1) , Ki-67 () (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
: p16INK4a () (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
Progesterone Receptors, Immunohistochemical Study)* (Estrogen and	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
: p16INK4a () (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 () (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- () (HER2/neu Expression, HER2 Status, Immunohistochemical Study (Fixed Biomaterial in Formalin Buffer))*	5489

Antibodies Pool, Glutamic Acid Decarboxylase-65, GAD and Insulinoma Antigen 2 (Tyrosine Phosphatase, IA2, ICA-512) Autoantibodies, Total)	GAD/IA-2, (Anti-GAD/IA2	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 nterocolitica :9 (Yersinia enterocolitica O:9, IHA)		484
IgG Yersinia nterocolitica (nti-Yersinia enterocolitica IgG)		539
IgA Yersinia nterocolitica (nti-Yersinia enterocolitica IgA)		539
Yersinia nterocolitica :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))		
(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)*		
(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 (Womwood, 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
		(Deoxy pyridinoline, DPD, Urine)	1375
		(osteocalcin, 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)	
		(Streptococcus group B, Streptococcus agalactiae), (Streptococcus agalactiae Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	968
		(Streptococcus group B, Streptococcus agalactiae) (Streptococcus agalactiae Culture. Bacteria Identification)	759
		(Streptococcus Group B. One Step Rapid Immunochromatographic Assay)	1155
		(Streptococcus Group A. One Step Rapid Immunochromatographic Assay)	957
		(Streptococcus group A, Streptococcus pyogenes), (Streptococcus pyogenes Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	891
		(Streptococcus spp., DNA, Saliva)*	429
		(Streptococcus group A, Streptococcus pyogenes) (Streptococcus pyogenes Culture. Bacteria Identification)	682
		(Streptococcus spp., DNA, Plasma)*	649
		(Streptococcus spp., DNA, Scrape of Faucial Epithelial Cells)*	429
		(Streptococcus spp., DNA, Sputum)*	869
A09.28.034.001		(Metanephrines fractionated, free and conjugated, 24-h urine)	2486
24-		(Catecholamines and Serotonin Metabolites, 24 Hours-Urine: Vanillylmandelic Acid, V, Homovanillic Acid, V, 5-Hydroxyindoleacetic Acid, 5- I)	2486
		(Serotonin, Serum)	2299
		(Catecholamines: Epinephrine/Adrenaline, Norepinephrine/Noradrenaline, Dopamine, Urine)	2266
		(Catecholamines: Epinephrine/Adrenaline, Norepinephrine/Noradrenaline, Dopamine, Plasma)	2299
		(Histamine, Plasma)	2662
		(Metanephrines fractionated, free + conjugated, random urine)	2013
		(Catecholamines: Epinephrine/Adrenaline, Norepinephrine/Noradrenaline, Dopamine, Urine)	2266
		(Opisthorchis felineus)	
		IgG (Anti-Opisthorchis felineus IgG)	869

(Giardia lamblia), Rapid Immunochromatographic Assay)	(Giardia lamblia. One Step	957
IgM, IgG, IgA IgA, Total)	(Anti-Giardia lamblia IgM, IgG,	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 acid, 24-Hour urine)		209
A09.28.003 (Protein Total, 24-Hour urine)		143
((Magnesium, 24 h urine excretion)		330
		2079
(Oxalates, 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 IgG)	(Anti-Complement 1q Antibodies, Anti-C1q,	1232
IgG, IgA, IgM (Anti-Endothelial Cell Antibodies, AECA, IgG, IgA, IgM, Total)	(HUVEC),	1705
IgG (- PO) (Myeloperoxidase Antibody, MPO)		1232
()		
1- 3- (Anti-Poliiovirus serotypes 1, 3, IgG)		1595
b, IgG (IgG (HiB), anti-PRP Haemophilus influenzae b 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 IgG Borrelia burgdorferi, (Anti-Borrelia burgdorferi IgG, Immunoblot)		2222

A	IgM Borrelia burgdorferi (Anti-Borrelia burgdorferi IgM)	627
(Barbiturates, Urine)*		
	() (Drugs and Psychotropic Substances Screening: Opiates, Amphetamines, Methamphetamine, Cocaine, Cannabinoids, Cannabinoid Metabolites, Urine)	1232
	() (Cannabinoids (Marijuana), Urine)*	3410
	() (Ethanol (Alcohol) Urine)*	1232
	« » () (Pernicious Habits: Nicotine, Drugs, Psychostimulants and Psychotropic Substances, Urine)*	1232
	(/) (Opiates (Morphine/Heroin), Urine)*	3608
- (-)		
	- , (EBV DNA, Exudate)*	275
	- , (EBV DNA, Cerebrospinal Fluid)*	275
	IgG - (nti-EBV Viral Capsid Antigens (VCA) IgG)	759
	IgG - (nti-EBV Early Antigen (EA) IgG)	627
	IgG - (nti-EBV Nuclear Antigen (EBNA) IgG)	528
	- , (EBV DNA, Blood)*	418
	- , (EBV DNA, Saliva)*	275
	- , (EBV DNA, Prostatic Fluid, Semen)*	275
	- , (EBV DNA, Scrape of Nasal Epithelial Cells)*	275
	- , (EBV DNA, Scrape of Urogenital Epithelial Cells)*	275
	Ig - (nti-EBV Viral Capsid Antigens (VCA) Ig)	528
	- , (EBV DNA, Scrape of Faucial Epithelial Cells)*	275
	- , (EBV DNA, Serum)*	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 Allergic components)	30030
	ALEX2, 300 IgE	29689
(MTHFR, MTRR, MTR) () (Folic Acid Metabolism (Genes MTHFR, MTRR, MTR) (without Description))		
	: D (VDR) () (Osteoporosis, Vitamin D Receptor (VDR) (Gene VDR) (without Description))	5203
	(MCM6) (Adult Lactase Deficiency (Gene MCM6))	1386
	, I (HFE) (Hemochromatosis Type 1 (Gene HFE))	1441
	: D (VDR) (Osteoporosis, Vitamin D Receptor (VDR) (Gene VDR))	2860
		1529

(UGT1A1) (Gilbert's Syndrome (Gene UGT1A1))		4994
: (CALCR, COL1A1) (3993
-) (Osteoporosis: Abridged Panel (Genes CALCR, COL1A1) (without Description))		5808
(MTHFR, MTRR, MTR) (Folic Acid Metabolism (Genes MTHFR, MTRR, MTR))		4455
: (CALCR, COL1A1) (Osteoporosis: Abridged Panel (Genes CALCR, COL1A1))		5203
-) (Osteoporosis: Full Panel (Genes CALCR, COL1A1, VDR) (without Description))		6864
1- HLA II (DRB1, DQA1, DQB1) (Hereditary Predisposition to Diabetes Type 1 (Insulin-Dependent Diabetes), HLA Class II (Genes DRB1, DQA1, DQB1))		5808
: (CALCR, COL1A1, VDR) (Osteoporosis: Full Panel (Genes CALCR, COL1A1, VDR))		
-		
2 (118 / , 121 / , 123 / , 131 / , 141 / , 115 / , 124 / , 154 /) (Genetic Test Results: Description of the 2-nd Category Complexity)		1155
1 (7201 , 7611 , 7014 , 125 / , 7207) (Genetic Test Results: Description of the 1-st Category Complexity)		583
3 (122 / , 129 / , 120 / , 137 / , 138 / , 153 / , 110 / , 114 / , 140 / , 7661 , 7258 , 134 / , 135 / , 136 /) (Genetic Test Results: Description of the 3-rd Category Complexity)		2310
4 (144 / , 143 / , 139 / , 145 / , 108 / , 19 /) (Genetic Test Results: Description of the 4-th Category Complexity)		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
, (Toxoplasma gondii, DNA, Cerebrospinal Fluid)*		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)		385
(Progesterone)		385

A09.05.135	() (Cortisol, Hydrocortisone)	385
A09.28.035	(Free Cortisol, Free Hydrocortisone, 24-Hour urine)	759
A09.05.069	(Aldosterone)	858
A09.05.121	() (Direct Renin, Plasma)	913
	(Cortisol, Saliva)	616
A09.05.230	(Cystatin C)	748
A09.05.017		165
A09.05.018	(Uric acid)	165
A09.05.020		165
-	p16INK4a Ki-67	6050
	(PLGF)	3960
	(Inhibin B)	1232
MIS)	() (Anti-Mullerian Hormone, AMH, Mullerian Inhibiting Substance, MIS)	1232
	-1- () (Trophoblastic beta-1-Globulin, TBG)	484
	IgG V (Annexin V antibodies, aAnV, IgG)	1309
	Ig V (Annexin V antibodies, aAnV, Ig)	1309
	IgG, IgA, IgM -2- 1, (Anti-?2-Glycoprotein 1 Antibodies, anti-?2-G 1, IgG, IgA, IgM, Total)	1254
	IgM IgG (Anti-Phospholipid Antibodies, APA, IgM, IgG)	803
	2 IgA	1199
	IgG IgM (Anti- phosphatidylserine/ rothrombin antibodies, Anti-PS/PT, IgG, IgM, Total)	1342
	IgA, IgM, IgG (Cardiolipin Antibodies IgA, IgM, IgG, aCL, Screening)	1133
	IgG IgM (Anti- phosphatidylserine, IgG, IgM)	1782
	2 IgM	1309
	, IgG, IgM (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
	(, , 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 (sFlt-1)		3377
(Estriol Free, 3)		484
PRISCA1		132
Placental Lactogen, hPL, Chorionic Somatomammotropin, CS, Human Chorionic Somatomammotropin, hCS)	() (Placental Lactogen, PL, Human	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
Microscopy (Manual Platelet Count (PLT Count): Indirect Method by Fonio)*	(Platelets,	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
: Assessment)*	(Water: Complete and Comprehensive Quality	19998
: Contamination)*	(Water: Suspicion Industrial Waste	14190
: (Water: Suspicion Products of Combustion and Emissions from Motorways Contamination)*		10483
: Contamination)*	(Water: Suspicion Household Waste	8943
: (Water: Abridged Quality Assessment)*		9669
: Excessive Use of Chemicals for Water Treatment)*	(Water: Suspicion	5159
(, Trichinella spiralis)		
IgG	(nti-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, Renin-Angiotensin System Disorder (Genes ACE, AGT) (without Description))	(2717
(NOS3) (-) (Arterial Hypertension, Endothelial NO-Synthase Disturbance (Gene NOS3) (without Description))	NO-	1386
()	(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 () (Gastrin-17 Stimulation Test,	1210
I (Pepsinogen I)		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
(Acute Intestinal Infections, PCR, Fecal)		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
()		495
(Streptococcus pneumoniae, DNA)		
(Calcitonin)		946
A09.05.058 () (Parathyroid Hormone, PTH)		671
(, Echinococcus spp.)		869
IgG (Anti-Echinococcus IgG)		
IgG (Anti-Entamoeba histolytica IgG)		693
D (D, Hepatitis D Virus, HDV)		858
IgM IgG D, o (Anti-HDV Total (IgG + IgM))		660
D, (HDV RNA, Serum)*		858
IgM D (nti-HDV IgM)		
Aspergillus fumigatus (3), - IgG (Aspergillus fumigatus, IgG, M3)		572
(Bacteroides spp., DNA, Scrape of Urogenital Epithelial Cells)*		231

A (, Hepatitis A Virus, HAV)			
IgG	(Anti-HAV IgG)		561
	(HAV RNA, Serum)*		649
IgM	(Anti-HAV IgM)		803
()			
	(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	(),		1342
(Anti-Heart Antibodies, IgG)			
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 ntibodies 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)			
	(Anti-Spermatozoa Antibodies, ASA, Serum)		1419
			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
Cytological Preparations (1 Glass) () (Consultation of Finished	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
	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 + » (Comprehensive Study «Sex in City: 8 Infections + Smear on Flora»)	2794
: III (29-30) (Pregnancy: Third Trimester (29-30 Weeks))	3993
« : 14 + » (Comprehensive Study «Sex in City: 14 Infections + Smear on Flora»)	4411
:	10923
() (Hemostasiogram (coagulogram), extended	2761
: (Survey of Liver: Extended)	2893
(Hospitalization in Therapeutic Hospital)	3663
(Hospitalization in Surgical Hospital)	5060
: (Hospitalization in 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	3861
You ? Healthy Country: Annual Check-Up up to 40 Years of Age)	(40) (Annual Check-Up after 40 Years of	4928
Age)	(Survey Before Diet: Additional)	5159
Supersport		7513
		4059
	(Survey of Kidneys: Extended)	2200
	(Diabetes Control: Extended)	3542
	(Toxic Trace Elements, Hair)	1782
	(Toxic Trace Elements, Essential Vital	3399
Elements, Hair)	(Elemental Composition of Hair: Screening)	5940
	(Essential Vital	2860
Elements, Toxic Trace Elements, Urine)	(Toxic Trace Elements, Nails)	1782
	(Toxic Trace Elements, Essential Vital	3399
Elements, Nails)	(Elemental Composition of Nails: Screening)	5940
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		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
	(/pANCA, cANCA), IgG)	3179
	(, , IgG, IgM)	4103
IgG)	(, (), (/pANCA, cANCA),	4345
	(, IgG;	4367
	()	5082
	(Arthralgia: screening test)	7018
	(Autoimmune Liver Disease: Screening)	

(Rheumatic arthritises)	1837
SARS-CoV-2, IgM () IgG () (Anti-SARS-CoV-2, IgM/IgG)	1859
«)» (Bullous Dermatositis Diagnostics profile (antibodies to epidermis desmosomes, antibodies to skin basal membrane))	4367
SARS-CoV-2, IgM IgG (Abbott)	1419
SARS-CoV-2, IgM () IgG () (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
, (/pANCA, cANCA), IgG)	2904
;)	3003
4 «)» (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
((),)	2717
IgA, IgG; IgA) (, IgA;	3553
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: (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
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Vessel Diseases)	

Diagnosis of Urogenital Tract Infection (UTI))	(Pregnancy Planning:	2486
(Diabetes Control: Screening)		616
(Survey Before Diet: Minimum)		1716
(Problems: Primary Survey)) (Weight	3223
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HOMA-IR (Insulin Resistance: Fasting Glucose/Insulin, Homeostasis Model Assessment of Insulin Resistance, HOMA-IR)		649
HOMA-IR (Insulin Resistance: Fasting Glucose/Insulin, Homeostasis Model Assessment of Insulin Resistance, HOMA-IR)		737
(Problems: Metabolic Syndrome (Primary Identification, Screening)	(Weight	946
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(Women's Oncorisk: Cervix)		1980
(Allergy to Animals, Dust, Mold)		6358
(Strong hair and nails, velvet skin)		1650
(Trace Elements, Serum,		3465
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(e) (Toxic Trace Elements, Toxic		1375
Heavy Metals, Venous Blood)		2365
()		2244
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Filtration Rate, eGFR, CKD-EPI Creatinine Equation)		462
B03.016.003 ()		836
: II (14-28) (Pregnancy: Second Trimester (14-28 Weeks))		649
(Clinical Blood Analysis: General Blood Analysis, Leucocyte Formula, ESR (with Manual Microscopic Examination of Blood Smear))		1375
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, 6 (AZF) (Spermatogenesis		11407
disorders (6 AZF))	(AZF-) (Impairment of	5830
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IgG (,)		10901
()	HLA II	1727
(DQA1, DQB1)		
- (RH factor Genotype)		
() (Parasitic Fungi, Microscopy and Culture		
(skin))		

(nails)	() (Parasitic Fungi, Microscopy and Culture	1727
I II	(I/ II) (Pepsinogen	2002
I/Pepsinogen II, PG1/PG2)		6468
: sFlt-1, PlGF,	sFlt-1/PlGF	1485
(Mycoplasma hominis Culture, Ureaplasma spp. Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*		957
:	(Male oncologic risk: prostate)	1408
A09.05.120.001	- (Aldosterone-Renin Ratio, ARR)	792
	3, 4 (Complement components C3, C4)	2332
"	", IgE, ImmunoCAP	770
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	(ITGB3) (Platelet Fibrinogen Receptor (Gene ITGB3))	770
	: (Lipid Profile: Screening)	792
Filtration Rate, eGFR, CKD-EPI ystatin C Equation)	CKD-EPI – (Estimated Glomerular	
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