

05.03.2024 .

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

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

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

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

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

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

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

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

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

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

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

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

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

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

/

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

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

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



A11.01.001	( )		400
A16.01.003			2000
A16.30.032		1 -3 1	1300
A16.30.032		3 -5 1	1900
A16.01.018			-
( , , 3 -5 1 )			2500
A16.01.003	(2 )		2600
A16.01.023	( 0,5-1 1 )		1500
A16.01.023	( 1 -3 1 )		1900
A16.01.023	( 3 -5 1 )		2400
A16.01.030	(1 )		2400
A16.01.030	(2 )		3100
A16.30.069	( ),		1000

A16.30.007.001			19000
A16.30.043.002			14800
A16.30.043.003			26000
A16.14.018.003	,		11000
A16.30.004.003	( )		42300
			56400
A16.30.004.007			65000
A16.30.004.016			55000
A16.30.001.002	-		35000
A16.30.002.002			35000
A16.30.001	( ) ( )		23100
A16.30.002	( )		20000
A16.14.009.002			35000
A16.19.018	( )		15000
A16.30.004.004	( )		46000
A16.01.018			-
( , , 5 10 -1 )			9500
A16.01.018			-
( , , 10 -1 )			12000
A16.30.032	( 5 -1 )		9800
A16.30.032	( 10 -1 )		12800
A16.30.006			34200
A16.30.006			65000
A16.18.022			28000
A16.18.022.001			30000
A16.22.002			120000
A16.22.001			55000

A16.30.004.010	1		40000
A16.30.004.010	2		55000

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

	(	)	2420	
A08.30.046	-	(	)	2420

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

A04.22.001			600
			800
A04.22.001.001			1100
			800

A04.16.001	( , , , , , )		1100
	( , ) , , , , , ,		2000
A04.06.001			500
A04.15.001			600
A04.28.002.005			600
A04.14.001			800
A04.14.002			500
A04.14.002.001			700
A04.18.001	( )		800
A04.28.002.001			800
A04.22.002			500
A04.28.001			800
A04.28.002.001			700
A04.28.002.003			500
A04.28.003			800
A04.10.002	( + )		1800
A04.01.001	( )		750
A04.09.001			600
A04.28.002.003	( )		1100
A04.18.001			800
A04.06.002			800
A04.06.002			800
A04.06.002			800
A04.06.002	/		800
A04.06.002			800
A04.06.002			800
A04.06.002	, / , , , , ,		2000
A04.21.001			1000
	( )		1200
	( )+ +		1400
A04.21.001.001	+ +		1100
	+ +		1300
A04.20.002			1200
			1200
A04.30.010			1100

A04.30.001	(	1200
A04.30.001.001	)	1300
A04.30.001.007	( )	1800
A04.12.024.003	III	1300
A04.04.001	II-III )	700
A04.20.003	( )	700
A04.20.003	( )	400
A04.20.001.004	( - )	700
A04.30.001.002	4D ( 26 , , 1 )	2500
A04.30.001	II	1500
A04.30.001.002	3D	1700
A04.30.001.001	( )	2000
A04.12.024.003	( ) ( II-III )	2400
A04.30.001.006	-	2500
( II)		
A04.30.001.008	III	2500
A04.30.001.002	4D ( 26 , , 1 )	3500
A04.30.001.002	3D	2400

--	--	--

A11.06.001.001		900
A11.20.010.003		900
A11.22.002.001		900
A11.30.024.001		900

--	--	--

A04.04.001	( )1	900
A04.04.001	( )1	700
A04.04.001	( )1	800
A04.04.001	( )1	800
A04.04.001	( )1	800
A04.04.001	( )1	800
A04.04.001	( )1	600
A04.04.002	( )	600
A04.12.022		1400

--	--	--

A04.24.001	( )	1100
A04.24.001	( )	3200

--	--	--

B03.037.001		500
A05.10.006		500
A02.12.002.001		1200

A05.10.008		24		2000
+		24		2200
A12.10.001	(	+	,	750
	)			750
A05.23.001				1100
A12.10.001				750
B01.058.001	(	,	)	-
				1400
B01.058.002	(	,	)	-
				1200
B01.001.001	(	,	)	-
				1500
B01.001.002	(	,	)	-
				1300
B01.001.002	(		)	-
				800
B01.001.001	(	,	)	-
	)		-	-
			(	
				1500
A11.20.011				1500
A11.20.014				1500
A11.20.008				5400
A11.20.008.001				2900
A11.20.002				450
A11.20.003				800
A03.20.001				1000
A11.20.015				500
A16.20.036				2100
A08.20.004			)	(
				1400
A11.20.015				500
A14.20.002	,			
(			)	
				900
A16.20.036.001				7500
A16.20.038				39000
A16.20.016				39900
A16.20.017.001				34900
A16.20.004.001				30900
A16.20.003.001	-			36900
A16.20.041.001				29900
A16.20.001.001				34900
(			)	
A16.20.001.001				37900
(			)	
A16.20.003.001				30900
A16.20.010.001			(	)
				73900

A16.20.010.001	2	( )		75900
A16.20.010.003		( 1 )		74900
A16.20.010.003		( 2 )		84900
A16.20.010		( )		58900
A16.20.011.001		( )		83900
A16.20.011.001		( )		94900
A16.20.011.002		( )		76900
A16.20.011.002	2	( )		87900
A16.20.063				92900
A16.20.033				35900
A16.20.012				77900
A11.20.003				1100
A03.20.003				19000
A16.20.042.003		( )		60750
A16.20.061.001			(1	35000
A16.20.061.001			(2	40500
-				
A16.20.035		( )		58900
A16.20.035.001		( 1- )	(- )	39900
A16.20.035.001		( 2- )	(- )	61900
A16.20.028.002				20900
A16.20.028.003				20900
A16.20.083				38900
A16.20.023				31900
A16.20.065				21900
A16.20.063.018				21900
A03.20.003				17000
A03.20.003.001	1			26900
A03.20.003.001	2			33900
A03.20.003.001	3			39900
A16.20.066				4100
A11.20.018		( )		2900
A16.20.059				950
A16.20.059.001				3900
A16.20.036				5900
A16.20.091.001				2800
A11.20.008.001				2000
A11.20.008.002				2700
A16.20.006				42000
A16.20.007				28900

A11.20.008		5400
A11.20.008.001		2900
A16.20.098	(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))	14 :	385
18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68) Screening)	( ) 14 : 16,	990
(Neisseria gonorrhoeae, DNA, Scrape of Urogenital Epithelial Cells)*		275
(Neisseria gonorrhoeae, ), (GC, Neisseria gonorrhoeae Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		990
(Bacterial Vaginosis, BV)		1694
(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Scrape of Urogenital Epithelial Cells)*		275
A08.20.017.002 ( , ThinPrep®)		1320
A08.20.004		660
( ; ; ; - )*		2420
(INBIOFLOR-Comprehensive Study of Microflora Composition of Urogenital Tract (UGT))		3058
APS) ( ), (Antiphospholipid Syndrome,		3927
(Genitourinary Tract Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*		1276

( ) (Vaginal Biocenosis: Bacteriophage and Antimycotic Susceptibility Testing (Gram Stain, Bacterioscopic Examination of Smear))*		1650
(The Bethesda System ? TBS) (Cytological Examination of Cervical Epithelium with Description on The Bethesda System, TBS)		627
MICROBIOCENOSIS (PCR Panel Femoflor 16)	16. (UROGENITAL TRACT	2310
		495
( ) (Cytological Examination: Cervix, Pap-test)		1100
(Candidiasis, Screening and Typing)		1012
MICROBIOCENOSIS, Screening ( PCR Panel Femoflor Screen))	. (UROGENITAL TRACT	2035
MICROBIOCENOSIS (PCR Panel Femoflor 8)	8. (UROGENITAL TRACT	1628
B01.053.001 ( , ) -		1400
B01.053.002 ( , ) -		1200
B01.053.002 ( , ) ( )		800
A16.28.040 (1 )		2100
		10000
A16.28.058		1000
A16.28.052.001		2400
A16.28.072.001		2400
A11.28.008		2000
A11.28.006.001		500
A21.21.001		550
A12.21.003		650
A15.21.001 ( )		1000
A03.28.002		3500
A03.28.003 ( )		6000
A03.28.001		5000
A16.28.077		5000
A11.28.012		1000
A16.21.015		2300
A16.28.013.001		1000
A06.28.007		5000
A06.28.011		5000
( )		10000
A16.28.077		6000
A11.28.012		2000
A11.28.007 ( )		1000
A16.01.016		10000
A11.28.008		2000
A22.28.001		25000
A22.28.002		25000



B01.015.001	( , ) -		1400
B01.015.002	( , ) -		1200

B01.008.003	( )		1400
B01.008.004	( )		1200

A11.02.002	. (1 )		260
A11.02.002	. (1 )		290
A11.01.003	. 1		2999
A11.01.003	. 2		4999
A11.01.003 PRP-	1 (1 )		5000
A11.01.003	mesoline ( ) 2,5 .		2500
A11.01.003	mesoline ( ) 2,5 .		2500

A16.01.024	(PRX-t33)		3750
A16.01.024			2000
A16.01.024			2500
A14.01.008			3000
			2000

B01.028.001	( , ) -		1500
B01.028.002	( , ) -		1300
B01.028.002	( ) -		800

A11.01.014			200
A12.25.001			1300
A11.07.004	,		2000
A11.08.001			5000
A11.08.002			3500
A11.25.006			3000
			200
A11.08.019			600
A16.01.012	( )		4000
A16.08.054			3000
A11.08.007			1000
A03.25.003			300
A22.30.033			2500
A11.08.022	( - ) - (1 )		2500
A11.08.022	( - ) - (2 )		3500
A21.25.002			250
A11.07.022			300
A16.01.004			1000

A15.01.002				1000
A16.08.006.001	(1	)		2500
A11.08.021.001				850
A16.08.016				750
A16.25.007	(1	)		700
A11.08.004				1500
A16.08.023	(1	)		2000
A16.08.023	(2	)		3800
A11.25.003.001				800
A16.01.017.001	1	3	.	3500
A16.25.008				1500
A02.25.001	(	)		1200
A03.08.004				1500
A03.08.001	(	)		1400
B01.003.004.004				300
B01.003.004.005	(I	)		500
B01.003.004.005	(II	)		700
B01.003.004.005	(III	)		900
	(	)		200
	(	)		200
	(	)		200
A11.08.020	(	)		700
A03.25.001				650
A16.08.012				1000
A16.08.009.001				10000
-1	:	1		
A16.08.009.001				20000
-1	:	2		
A16.08.009.001				30000
-1	:	3		
A16.08.010.001			:	10000
			1	
A16.08.010.001			:	15000
			2	
A16.08.010.001			:	20000
			3	
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			6700
( 1 )			
A16.16.052			7900
( 1 2 )			
A16.16.052			9900
( ( 2 5 )			
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

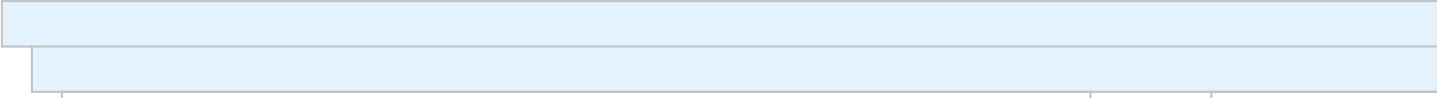
(60 )			2100
-------	--	--	------

Check-Up	45		12300
----------	----	--	-------

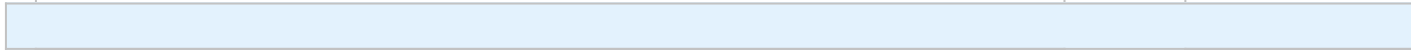
Check-Up	40		14500
Check-Up	45		17100
Check-Up	40		18700
Check-Up ( - )"	"	40	8600
Check-Up ( - )"	"	40	10700
"	" ( ;	;	2990
" )	" ( +		1600
+	+	+	7500



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
		2.0(1 )	30
		1.0(1 )	15
		1,5% 200 (1 )	200
		2 (1 )	25
		5 (1 )	150
	( )	(1 )	100
		50 / 2 (1 )	1115
		400 (1 )	450
		50 (1 )	5
		50 (1 )	5
		5% (1 )	5

		40 / 5 (1 )		160
	( )			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	( , ) - ( ) ( 60 )	2500
A05.30.014	,	900
		150
	( )	
	- .	
	IgG4- (Diagnosis of Autoimmune Pancreatitis and other IgG4-Related Diseases)	1837
	Ig ( , ) (Anti- ndomysial ntibodies, Anti-EMA, Ig )	1430
	IgG ( ) (Anti-Sacch romyces Cerevisiae Antibodies, ASCA, IgG )	1232
	IgA IgG (Anti-Intestinal Goblet Cells Antibodies, GAB, IgA, IgG, Total)	1199
	Ig ( ) (Anti-Sacch romyces Cerevisiae Antibodies, ASCA, IgA )	1232
	, IgG IgA ( , 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- ndomysial ntibodies, Anti-EMA, IgA, IgG, Total)	1276
	IgA, IgG, IgM ( ), (Gastric Parietal Cell Antibodies, GPA, Anti- arietal cell antibodies, APCA, IgA, IgG, IgM, Total)	1518
	IgG IgA GP2 (Anti-GP2)	1980
	Ig (Anti-Deaminated Gliadin Peptide, Anti-DGP, IgA)	781
	- - ( ). FLCN, . (Birt-Hogg-Dube Syndrome, BHD, Gene FLCN, Mut.)	51348
	- ( - - ) II. MFN2, . . (Charcot-Marie-Tooth Disease Type 2A1, Gene MFN2, Freq. Mut.)	5148
	, 4 TBP, . .	3410
	, N1, . .	3410
	( - - ). MYH3, . . (Arthrogyposis Distal Type 2A, Gene MYH3, Freq. Mut.)	9999
	( - - ) I. NDRG1 SH3TC2, . . (Charcot-Marie-Tooth Disease Type 1B, Genes NDRG1, SH3TC2, Mut.)	5148
	ATXN8, . . (Spinocerebellar Ataxia, Gene ATXN8, Freq. Mut.)	5148
	( , , , ) TAZ, . (Left Ventricular Non-Compaction, LVNC, Gene TAZ, Mut.)	25707
	MVK, . (Mevalonic Aciduria, Gene MVK, Mut.)	42801
	8, 9 LMNA, . (Mandibuloacral Dysplasia, Exons 8, 9 Gene LMNA, Mut.)	7161

I, II, III, IV (Atrophy, SMA, Type I, II, III, IV (copy Number Variation SMN2))	SMN2) (Spinal Muscular	16577
CHRNG, . (Escobar Syndrome, Gene CHRNG, Mut.)		34254
BCL2, . (Silver Syndrome, Gene BCL2, Mut.)		29975
Hemophagocytic Lymphohistiocytosis, Gene STX11, Mut.)	STX11, . (Familial	14267
-1- , SERPINA1, . . .		2035
I, II, III, IV. ) (Spinal Muscular Atrophy, SMA, Type I, II, III, IV, Gene SMN1, Mut. (Only Presence One Gene Copy) )	SMN1, . (	34254
CINCA, NLRP3 . (Chronic Infantile Neurologic Cutaneous Articular, Gene NLRP3, Mut.)		51348
TBX3, . (Pallister W Syndrome, Gene TBX3, Mut.)		29975
(Oculopharyngeal Muscular Dystrophy, OPMD, Gene RABPN1, Freq. Mut.)	RABPN1, . . .	5148
(Gerstmann-Straussler Disease, Gene PRNP, Mut.)	PRNP, . .	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.)	( - - ) I.	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
( ) .	GLI3, . (Greig Syndrome,	76989
		8910
EGR2, . (Charcot-Marie-Tooth Disease Type 1B, Gene EGR2, Mut.)	( - - ) I.	17160
Gene FGFR2, Mut.)	7 9 FGFR2, . (Crouzon Syndrome, Exons 7, 9	9999
- a - a ( ) .		
ENG, . (Rendu-Osler-Weber Disease, Gene ENG, Mut.)		38522
Exudative Vitreoretinopathy, FEVR, Gene NDP, Mut.)	NDP, . (Familial	11264
Lipodystrophy 2, Gene LMNA, Mut.)	LMNA, . (Familial Partial	42801
( 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 polyneuropathy, 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 Ataxia, Gene FXN, Freq.	9020
EMG1, . (Bowen Conradi Syndrome, BCS, Gene EMG1, Mut.)	( , )	17160
(Leber Hereditary Optic Neuropathy, LHON, Mitochondrial DNA, 3 Freq. Mut.)	, 3 . .	5148
Syndrome, Gene BCS1L, Mut.)	( ) . BCS1L, . (Bjomstad	17523
TWIST1, Mut.)	TWIST1, . (Saethre-Chotzen Syndrome, Gene	14916
GDAP, . (Charcot-Marie-Tooth Disease Type 2A1, Gene GDAP, Mut.)	( - - ) II.	25707
COMP, Freq. Mut.)	COMP, . . (Pseudoachondroplasia, Gene	6820
Gene NPHS1, Mut.)	NPHS1, . (Nephrotic Syndrome Type 1, NPHS1,	76989
Mut.)	FGFR3, . . (Achondroplasia, Gene FGFR3, Freq.	12056
Syndrome, FCAS, Gene NLRP3, Mut.)	NLRP3 . (Familial Cold Autoinflammatory	51348
O, . (Charcot-Marie-Tooth Disease Type 1B, Gene O, Mut.)	( - - ) I.	17160
DLL3, Mut.)	DLL3, . (Spondylocostal Dysostosis, Gene	29975
NPHP1, . (Nephronophthisis 1, NPHP1, Gene NPHP1, Mut.)		17006
Muscular Dystrophy, Gene FHL1, Mut.)	FHL1, . (Emery-Dreifuss	34254
Dihydrate, CPPD, Gene ANKH, Mut.)	ANKH, . (Chondrocalcinosis, Calcium Pyrophosphate	51348
MULIBRAY. Mut.)	TRIM37, . (Muscle-Liver-Brain-Eye, Gene TRIM37,	9999
(Leber Hereditary Optic Neuropathy, LHON, Mitochondrial DNA, 12 Freq. Mut.)	, 12 . .	14267
Ataxia, Gene ATXN7 Freq. Mut.)	ATXN7, . . (Spinocerebellar	5148
IGHMBP2, . (Distal Spinal Muscular Atrophy 1, DSMA1, Gene IGHMBP2, Mut.)		64163
RP2, Mut.)	RP2, . (Retinitis Pigmentosa, Gene	21428
Syndrome, Type VI, Gene PLOD, Freq. Mut.)	VI. PLOD, . . (Ehlers-Danlos	10285
(Aarskog-Scott Syndrome, Faciodigitogenital Syndrome, Faciogenital Dysplasia, Gene FGD1, Mut.)	( ) . FGD1, .	62722
Spinal and Bulbar Muscular Atrophy, Gene AR, Freq. Mut.)	AR, . . (Kennedy	5148
PCSK9		11682
( X )		6369
/ , FMR1, . .		3410
Mut.)	GLI3, . (Pallister-Hall Syndrome, Gene GLI3,	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
( ) . FKRP, . (Walker-Warburg Syndrome, WWS, Gene FKRP, Mut.)	14916
10 FGFR3, . (Crouzon Syndrome with Acanthosis Nigrigan, CAN, Exon 10 Gene FGFR3, Mut.)	7161
BTK, . (X-Linked Agammaglobulinemia, XLA, Gene BTK, Mut.)	76989
D- ( ) . PHEX, . (Hypophosphatemic Vitamin D-Resistant Rickets, Gene PHEX, Mut. )	94083
GRN, . (Aphasia Primary Progressive, Gene GRN, Mut.)	25707
XK, . (McLeod Syndrome, Gene XK, Mut.)	17160
( ) . AR, . (Testicular Feminization Syndrome, Gene AR, Mut.)	42801
EDA, . (Anhidrotic Ectodermal Dysplasia, Gene EDA, Mut.)	34254
UPK3A, . (Renal Hypodysplasia, Aplasia 1, Gene UPK3A, Mut.)	25707
PAH, . . (Phenylketonuria, PKU, Gene PAH, Freq. Mut.)	17006
« » OPA1, (Optic Atrophy With Or Without Deafness, Ophthalmoplegia, Myopathy, A taxia And Neuropathy, Gene OPA1, Hot-Point Mut.)	9999
ELA2, . (Neutropenia Severe Congenital 1 Autosomal Dominant, SCN1, Gene ELA2, Mut.)	21428
APOB100	4818
NTRK1, . (Congenital Insensitivity To Pain With Anhidrosis, CIPA, Gene NTRK1, Mut.)	51348
COMP, . . (Multiple Epiphysial Dysplasia, MED, Gene COMP, Freq. Mut.)	6820
( ) . PHOX2B, . . (Congenital Central Hypoventilation Syndrome, CCHS, Gene PHOX2B, Freq. Mut.)	5148
( CFTR, GJB2, PAH, SMN) (Main Hereditary Diseases (Genes CFTR, GJB2, PAH, SMN))	21142
DMPK, . . (Myotonic Dystrophy 1, Gene DMPK, Freq. Mut.)	5148
( ) I. GJB1, . (Charcot-Marie-Tooth Disease Type 1B, Gene GJB1, Mut.)	9999
TGM1, . (Autosomal Recessive ongenital Ichthyosis, ARCI 1, All Known Mutations, Gene TGM1, Mut.)	42801
V. BSCL2, . (Distal Hereditary Motor Neuropathy, DHMN, Gene BSCL2, Mut.)	29975
ZEB2, . (Mowat-Wilson Syndrome, Gene ZEB2, Mut.)	64163
PRPS1, . (Phosphoribosylpyrophosphate Synthetase Superactivity, PRS Superactivity, Gene PRPS1, Mut.)	29975

(Albinism oculocutaneous, Hermansky-Pudlak type, Gene HPS1, Freq. Mut.)	HPS1, . . .	9999
/		23540
(Klippel-Feil Syndrome, Gene GDF6, Mut.)	GDF6, .	14916
. (Cranio metaphyseal Dysplasia, Gene ANKH, Hot-Point Mut.)	ANKH, « . » .	9999
(Nonbullous Congenital Ichthyosiform Erythroderma, NBCIE, Gene TGM1, Mut.)	TGM1, .	42801
PRNP, Mut.)	PRNP, . (Creutzfeldt-Jakob Disease, Gene	14916
Gene CRYBA4, Mut.)	CRYBA4, . (Microphthalmia with Cataract,	25707
:		7590
(TNF-Receptor-Associated Periodic Syndrome, TRAPS, Gene TNFRSF1A, Mut.)	TNFRSF1A, .	25707
( ) . NPHP1 (Joubert Syndrome, Cerebelloparenchymal Disorder IV, CPD IV, Classic Joubert Syndrome, Joubert Syndrome type A, Joubert-Boltshauser Syndrome, Pure Joubert Syndrome, Gene NPHP1, Mut.)		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	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 FKR1, Freq. Mut.)	FKR1, . . (Muscular	7876
Fundus Flavimaculatus Included, Gene ABCA4, Freq. Mut.)	ABCA4, . . (Stargardt Disease 1, STGD1,	12056
( ) .	TRAPPC2, .	17160
(Spondyloepiphyseal Dysplasia Tarda, SEDT, Gene TRAPPC2, Mut.)		
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 FKRP, Mut.)	FKRP, . (Muscular	14916
	CTSK, . (Pyknodysostosis, PKND, Gene CTSK, Mut.)	25707
	PAH, . (Phenylketonuria, PKU, Gene PAH, Mut.)	51348
Syndromе, 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, . (Scapuloperoneal 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, « . » Mut.)	(Fibrodysplasia Ossificans Progressiva, FOP, Gene ACVR1, Hot-Point	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
Syndromе, PPS, Gene IRF6, Mut.)	IRF6, . (Popliteal Pterygium	38522
PTEN, Mut.)	PTEN, . (Lhermitte-Duclos Syndrome, Gene	38522
Dystrophy-Dystroglycanopathy, Gene FKRP, Mut.)	FKRP, . (Muscular	14916

PTEN, . (Bannayan-Ruvalcaba-Riley Syndrome, Gene PTEN, Mut.)	38522
VHL, . (Autosomal Recessive Erythrocytosis, Gene VHL, Mut.)	14916
, PMP22, (Charcot-Marie-Tooth disease type 1A (CMT1A))	16577
ANKH, . (Craniometaphyseal Dysplasia, Gene ANKH, Mut.)	51348
CYBB, . (Chronic Granulomatous Disease, CGD, Gene CYBB, Mut.)	51348
LMNA, . (Hutchinson-Gilford Progeria Syndrome, Gene LMNA, Mut.)	42801
UGT1, . (Crigler-Najjar Syndrome, Gene UGT1, Mut.)	21428
MECP2, . (Retts Syndrome, Gene MECP2, Mut.)	21428
HLA B51	3300
SRY, . (Disorders Sex Determination, Analysis Gene SRY, Mut.)	5148
SGCA, . (Muscular Dystrophy Limb-Girdle Type 2A, Gene SGCA, Mut.)	25707
NEFL, . (Charcot-Marie-Tooth Disease Type 2A1, Gene NEFL, Mut.)	25707
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
FKTN, . (Muscular Dystrophy Fukuyama-Type, Gene FKTN, Mut.)	51348
ESC ( . ) NR2E3, . (Enhanced S- one Syndrome, Goldmann-Favre Syndrome, Gene NR2E3, Mut.)	25707
IRF6, . (Van der Woude Syndrome, Gene IRF6, Mut.)	38522
RS1, . (Retinoschisis 1 X-Linked Juvenile, RS1, Gene RS1, Mut.)	25707
UNC13D, . (Familial Hemophagocytic Lymphohistiocytosis, Gene UNC13D, Mut.)	76989
TCIRG1, . (Osteopetrosis Autosomal Recessive 1, OPTB1, Gene TCIRG1, Mut.)	51348
CSTB, . (Progressive Myoclonic Epilepsy 1A Unverricht and Lundborg, Gene CSTB, Mut.)	14267
SH2D1A, . (X-Linked Lymphoproliferative Syndrome, XLP, Gene SH2D1A, Mut.)	17160
BEST1, . (Best Vitelliform Macular Dystrophy, All Known Mutations, Gene BEST1, Mut.)	42801
TWIST1, . (Craniosynostosis Type 2, Gene TWIST1, Mut.)	14916
MSX2, . (Craniosynostosis Type 2, Gene MSX2, Mut.)	11264
PRNP, . (Spongiform Encephalopathy with Neuropsychiatric Features, Gene PRNP, Mut.)	14916
KRT2, . (Ichthyosis Bullosa Of Siemens, Gene KRT2, Mut.)	29975
RAB23, . (Carpenter Syndrome, Gene RAB23, Mut.)	29975
9 FGFR2 7A FGFR1, . (Jackson-Weiss Syndrome, JWS, Exon 9 Gene FGFR2, Exon 7A Gene FGFR1, Mut.)	9999
ABCC6, . (Pseudoxanthoma Elasticum, Gene ABCC6, Mut.)	119724
4, SPAST (SPG4), . .	6930
VHL, . (Von Hippel-Lindau Syndrome, VHL, Von Hippel-Lindau Hereditary Cancer Syndrome, Gene VHL, Mut.)	14916
HLA-Cw6	3300
	5599
( . ) FGFR2, . . (Apert Syndrome, AS, Gene FGFR2, Freq. Mut.)	10285

( - LDLR, APOB, PCSK9)	9218
Epiphyseal Dysplasia, MED, Gene SLC26A2, Mut.)	25707
PRPS1, . (Art's Syndrome, Gene PRPS1, Mut.)	29975
(Nonbullous Congenital Ichthyosiform Erythroderma, NBCIE, Gene ALOXE3, Mut.)	64163
PAX3, . (Craniofacial-Deafness-Hand Syndrome, CDHS, Gene PAX3, Mut.)	34254
Muscular Dystrophy, Gene LMNA, Mut.)	42801
VHL, . (Autosomal Recessive Erythrocytosis, Gene VHL, Freq. Mut.)	5148
(Keratitis-Ichthyosis-Deafness Syndrome, KID Syndrome, Gene GJB2, Mut.)	9999
« . » . (Familial Partial Lipodystrophy 2, FPLD 2, Gene LMNA, Hot-Point Mut.)	14267
TAZ, . (Barth Syndrome, Gene TAZ, Mut.)	25707
(Cerebrooculofacioskeletal Syndrome, COFS Syndrome, Gene ERCC6, Mut.)	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	12056
3A, Gene CYP1B1, Mut.)	17523
(Normokalemic Periodic Paralysis, Exon 13 Gene SCN4A, Mut.)	7161
ATP7B, PNPLA3, SERPINA1, . . .	9240
GLI3, . (Polydactyly, Gene GLI3, Mut.)	76989
-IgM Mut.)	21428
CD40LG, . (Hyper-IgM Syndrome, Gene CD40LG, Mut.)	29975
HPGD, . (Hypertrophic Osteoarthropathy, Primary, Autosomal Recessive, 1, Gene HPGD, Mut.)	10285
Gene CLCN1, Freq. Mut.)	4620
, C9orf72, . . .	38522
And Diabetes Syndrome, Gene HNF1B, Mut.)	10285
SH3TC2, FIG4, FGD4 GDAP1, . . . (Charcot-Marie-Tooth Disease Type 1B, Gene GDAP1, Freq. Mut.)	94083
Gene RPS6KA3, Mut.)	9999
RPS6KA3, . (Coffin-Lowry Syndrome, Mut.)	3190
SHH, . (Polydactyly, Gene SHH, Mut.)	25707
, . PNPLA3, . . .	38522
TAR. TAR-Syndrome, Gene RBM8A, Mut.)	3080
RBM8A, . (Thrombocytopenia-Absent Radius Syndrome, Mut.)	17160
PTEN, . (Cowden Syndrome 1, Gene PTEN, Mut.)	51348
, 2, CNBP (ZNF9), . . .	94083
EBP, . (Chondrodysplasia Punctata, CDP, Conradi-Hunermann Syndrome, Gene EBP, Mut.)	76989
STXBP2, . (Familial Hemophagocytic Lymphohistiocytosis, Gene STXBP2, Mut.)	7161
( ). LPIN1, . (Myoglobinuria Acute Recurrent Autosomal Recessive, Gene LPIN1, Mut.)	9999
ADAMTSL2, . (Geleophysic Dysplasia 1, Gene ADAMTSL2, Mut.)	
PANK2, . . . (Neurodegeneration With Brain Iron Accumulation 1, Gene PANK2, Freq. Mut.)	
GJB4, . (Erythrokeratoderma, Gene GJB4, Mut.)	

( ) II, De la Chapelle Dysplasia, Gene SLC26A2, Mut.)	SLC26A2, . (Atelosteogenesis	25707
DBA1, Gene RPS19, Mut.)	RPS19, . (Diamond-Blackfan Anemia 1,	21428
NPHS1, Gene NPHS2, Mut.)	NPHS2, . (Nephrotic Syndrome Type 1,	34254
IX, Mut.)	IX B, . (Hemophilia B, Gene Factor	29975
Syndrome, TRPS, Gene TRPS1, Mut.)	TRPS1, . (Trichorhinophalangeal	42801
Muscular Atrophy (SMA) with Diaphragmatic Paralysis, Gene IGHMBP2, Mut.)	IGHMBP2, . (Spinal	64163
Freq. Mut.)	FGFR3, . . (Hypochondroplasia, Gene FGFR3,	12056
( 1 , 1b)	NS3, NS5A NS5B	12837
. (Hyperkalemic Periodic Paralysis Type 2, Exons 13, 24 Gene SCN 4A, Mut.)	13 24 SCN 4A,	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- (Hereditary Breast Cancer In Men: Cancer of Breast, Pancreatic, Prostate, Testicular Cancer (Genes BRCA1, BRCA2) (without Description))	( BRCA1, BRCA2) (	4290
Medullary Thyroid Cancer (Exons 10, 11, 13, 14, 15 Gene RET))	( 10, 11, 13, 14, 15 RET) (Familial	21406
Endocrine Neoplasia Type 2A (Exons 10, 11 Gene RET))	2A ( 10, 11 RET) (Multiple	9999
IDH2		11682
		23199
14 JAK2 (Quantification of wild-type and mutant allelic ratio of gene JAK2 617V/617F)	617V/617F	8800
BRCA- Cancer In Men: Cancer of Breast, Pancreatic, Prostate, Testicular Cancer (Genes BRCA1, BRCA2))	( BRCA1, BRCA2) (Hereditary Breast	4785
MGMT		11682
PIK3CA		11682
228 250 TERT		8327
(Genes BRCA1, BRCA2) (without Description))	/ ( BRCA1, BRCA2) (	4290
IDH1		11682
A09.05.039 Dehydrogenase, LDH)	( , L- , + ) (Lactate	165
G6PD		3278
A09.05.042 ) (Alanine Aminotransferase, ALT, Serum Glutamic Pyruvic Transaminase, SGPT)	( , , -	165
A09.05.045 - (?- , ) ( lpha- milase, ?-Amylase)	( lpha- milase, ?-Amylase)	231
A09.05.180 - (P- ) (Pancreatic ?-Amylase)	(P- ) (Pancreatic ?-Amylase)	275
(S- (Cholinesterase, Pseudocholinesterase, PCHE)	II, S- , )	253

A09.05.046	( ) (Alkaline Phosphatase, ALP)	165
A09.05.173	( ) (Lipase)	319
	( ) (Acid Phosphatase, ACP)	220
GGT	( ) (Gamma-Glutamyl Transferase, GGT)	165
A09.05.043	( ) (Creatine Kinase, CK, Creatine Phosphokinase, CPK)	264
A09.05.041	( ) (Aspartateaminotransferase, AST, Serum Glutaminoxaloacetic Transaminase, SGOT)	165
	( ) (Creatine Kinase-MB, CK-MB, Creatine Phosphokinase-MB, CPK-MB.)	341
HLA-		
	HLA II ( DRB1, DQA1, DQB1) (System Human Leukocyte Antigen (HLA) Class II, Typing (Genes DRB1, DQA1, DQB1))	5841
	(Genotype of RH factor Definition (without Description))	9768
	Y- (Y-chromosome of the fetus in the mother's blood)	4510
	(Rh factor Definition)	6820
	(Plasminogen)	660
	VIII ( ) (Antihemophilic Globulin A, FVIII)	1133
A09.05.051.001	D- (D-Dimer)	1133
A09.05.050	(Fibrinogen, FG)	231
A12.05.039	( ) (Activated Partial Thromboplastin Time, APTT)	165
	IX, % ( « ») Factor IX, Activity, % (Christmas Factor, anti-hemophilic globulin "B")	484
A12.05.027	( ) (Prothrombin, rothrombin Time, PT, International Normalized Ratio, INR)	253
A12.05.028	( ) (Thrombin Time, TT)	253
A09.05.029.001	( ) (Lupus Anticoagulant, LA )	803
	C, % (Protein C, % Activity)	1969
	/ ( ), Anti-Xa activity, IU/ml (Heparin concentration, IU/ml)	1837
	, % (Willebrand Factor, Antigen, %)	2233
A09.05.047	III, % ( III, Antithrombin III, % Activity)	363
	S (Protein S, Free)	2442
	(Urine immunoglobulin free light chains (FLC) kappa and lambda)	1474
	(M-Gradient, Screening. Serum Protein Electrophoresis (SPEP), Immunofixation with Polyvalent Antiserum, Quantification of M-Protein (without Typing))	2376
	(Bence-Jones Protein, Urine, Electrophoresis, Immunofixation, Kappa/Lambda Typing, Quantification )	3300
A09.05.014	(Serum Protein Electrophoresis, SPE, SPE )*	275
	(M-Gradient, Typing. Serum Protein Electrophoresis (SPEP), Immunofixation with Antisera (IgG, IgA, IgM, Kappa, Lambda), Quantification of M-Protein)	4389
	(Cerebrospinal Fluid Concentration of Immunoglobulin Free Light Chains)	1738
	/	2156
	(Bence-Jones Protein, Urine, Immunofixation, Quantification )	2068

A09.05.011	(Albumin)	231
A09.05.010	(Protein Total)	165
A09.05.214	(Homocysteine)	1419
	(Urine Protein Electrophoresis)	1650
ImmunoCAP		
	(f216) IgE, ImmunoCAP	693
	(f33) IgE, ImmunoCAP	693
	, nArtv1 (w231) IgE, ImmunoCAP	2013
	(f9) IgE, ImmunoCAP	693
	(f260), IgE, ImmunoCAP (Broccoli, Brassica oleracea (f260), IgE, ImmunoCAP)	792
	(f26) IgE, ImmunoCAP	693
	( ) (i6) IgE, ImmunoCAP	693
	(f343), IgE, ImmunoCAP (Raspberry, Rubus idaeus, IgE, ImmunoCAP)	792
	(f35) IgE, ImmunoCAP	693
	, nGal d3 (f323) IgE, ImmunoCAP	913
	(Hollister-Stier) (hx2) IgE, ImmunoCAP	1375
	Candida albicans (m5) IgE, ImmunoCAP	693
	(i1) IgE, ImmunoCAP	693
	(i3) IgE, ImmunoCAP	693
	(c8) IgE, ImmunoCAP	693
	(f95) IgE, ImmunoCAP	693
	(f6) IgE, ImmunoCAP	792
	(Saccharomyces cerevisiae) (f45) IgE, ImmunoCAP	693
	, nGal d1 (f233) IgE, ImmunoCAP	913
	, rAra h 2 (f423) IgE, ImmunoCAP	2013
	, (e81) IgE, ImmunoCAP	693
	/ D. pter nyssinus (d1) IgE, ImmunoCAP	693
	, rBet v1/PR-10 (t215) IgE, ImmunoCAP	2013
	(f23) IgE, ImmunoCAP	693
	, nGal d2 (f232) IgE, ImmunoCAP	913
-	, (nBos d5) (f77) IgE, ImmunoCAP	913
-	(nBos d4) (f76) IgE, ImmunoCAP	913
	(f92) IgE, ImmunoCAP	693
	(f91) IgE, ImmunoCAP	693
	, (e6) IgE, ImmunoCAP	693
	, ImmunoCAP	3025
	, rGly m 4/PR-10 (f353) IgE, ImmunoCAP	2013
	, (e213) IgE, ImmunoCAP	693
	(f75) IgE, ImmunoCAP	693
	, (e1) IgE, ImmunoCAP	693
	, rAra h 1 (f422) IgE, ImmunoCAP	2013
	(Hollister -Stier) (h2) IgE, ImmunoCAP	693
	c, rAra h 9 LTP (f427) IgE, ImmunoCAP	2013
	, rCan f 1 (e101) IgE, ImmunoCAP	2013
	(f83) IgE, ImmunoCAP	693
	(w5) IgE, ImmunoCAP	693

Malassezia spp. (m227) IgE, ImmunoCAP	693
, rCan f 2 (e102) IgE, ImmunoCAP	2013
/ (k80) IgE, ImmunoCAP	693
(f31) IgE, ImmunoCAP	693
(f14) IgE, ImmunoCAP	693
(Greer Labs.) (h1) IgE, ImmunoCAP	693
c (k82) IgE, ImmunoCAP	693
Cladosporium herbarum (m2) IgE, ImmunoCAP	693
(w8) IgE, ImmunoCAP	792
(mx2) IgE, ImmunoCAP	1375
(f24) IgE, ImmunoCAP	693
, rPen a1(f351) IgE, ImmunoCAP	2013
(f209) IgE, ImmunoCAP	693
(i71) IgE, ImmunoCAP	693
(f2) IgE, ImmunoCAP	693
( ) (f55) IgE, ImmunoCAP	693
(f210) IgE, ImmunoCAP	693
(f13) IgE, ImmunoCAP	693
, nArtv3 (w233) IgE, ImmunoCAP	2013
-5, rTri a 19 (f416) IgE, ImmunoCAP	2013
(fx15) IgE, ImmunoCAP	1375
, (e5) IgE, ImmunoCAP	693
(fx73) IgE, ImmunoCAP	1375
(fx5) IgE, ImmunoCAP	1375
(w204) IgE, ImmunoCAP	792
(f49) IgE, ImmunoCAP	693
Penicillium notatum (P.chrysogenum) (m1) IgE, ImmunoCAP	693
(f25) IgE, ImmunoCAP	693
, rCyp c 1 (f355) IgE, ImmunoCAP	2013
(i75) IgE, ImmunoCAP	693
(gx1) IgE, ImmunoCAP	1375
, rBet v2, rBet v4 (t221) IgE, ImmunoCAP	2013
(f11), IgE, ImmunoCAP	693
(m80) IgE, ImmunoCAP	693
Phadiatop ImmunoCAP, IgE	1727
, rPhl p1, rPhl p5 (g213) IgE, ImmunoCAP	2013
(f302) IgE, ImmunoCAP	792
, (nBos d8) (f78) IgE, ImmunoCAP	913
(mx1) IgE, ImmunoCAP	1375
V (c2) IgE, ImmunoCAP	693
(f227) IgE, ImmunoCAP	792
(tx9) IgE, ImmunoCAP	1375
(f12), IgE, ImmunoCAP (Pea, Pisum sativum, IgE, ImmunoCAP)	792
(f20) IgE, ImmunoCAP	792
( ) (f212), IgE, ImmunoCAP (Mushrooms, Agaricus hortensis, IgE, ImmunoCAP)	792
/ D. farina (d2) IgE, ImmunoCAP	693

(f88) IgE, ImmunoCAP	693
(f94), IgE, ImmunoCAP (Pear, <i>Pyrus communis</i> , 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, <i>Solanum melongena</i> , IgE, ImmunoCAP)	792
(f221), IgE, ImmunoCAP (Coffee, <i>Coffea</i> 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, <i>Allium cepa</i> , 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, <i>Prunus armeniaca</i> , IgE, ImmunoCAP)	792
(f242), IgE, ImmunoCAP (Cherry, <i>Prunus avium</i> , IgE, ImmunoCAP)	792
Ig E ImmunoCAP	759
TSST (m226) IgE, ImmunoCAP	693
<i>Alternaria alternata</i> (m6) IgE, ImmunoCAP	693
<i>Aspergillus fumigatus</i> (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
Peptide, anti-CCP)	( ) (Anti- ydic Citrullinated	1452
IgG (	) (Anti- eratin ntibodies, AKA, Anti-Filaggrin ntibodies, AFA, IgG)	2244
, IgA (	IgA; Rheumatoid Factor, RF, IgA)	1232
Fluid Smear, Crystals)	( ) (Synovial	1705
IgG (	-M )	1518
(Anti-Mutated Citrullinated Vimentin Antibodies, Anti-MCV, Anti-Modified Citrullinated Vimentin Antibodies, Anti-Sa Antibodies, IgG)		
DNA, Scrape of Nasal Epithelial Cells)*	(CMV	275
Scrape of Faucial Epithelial Cells)*	(CMV DNA,	275
,	(CMV DNA, Blood)*	418
,	(CMV DNA, Exudate)*	275
,	(CMV DNA, Serum)*	418
A IgM	(Anti-CMV IgM)	539
Semen)*	(CMV DNA, Prostatic Fluid,	275
of Skin Epithelial Cells)*	(CMV DNA, Scrape	275
,	(CMV DNA, Urine)*	275
(CMV DNA, Scrape of Urogenital Epithelial Cells)*		275
,	(CMV DNA, Saliva)*	275
,	(Cytomegalovirus, DNA)	407
Anti-CMV IgG		1144
A IgG	(Anti-CMV IgG)	396
DNA, Scrape of Conjunctiva Epithelial Cells)*	(CMV	275
Fluid)*	(CMV DNA, Cerebrospinal	275
,	(Copper, random urine; Cu)	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), blood)	1298
(Zn)	(Zinc (Zn), blood)	1298
(Iodine, serum)		275
(Au)	(Gold (Au), Serum)	275
(Cu)	(Copper (Cu), Serum )	275
( )	( )	748
(Cu)	(Copper (Cu), blood)	1298
(Pb)	(Lead (Pb), blood)	1298
(Mn)	(Manganese (Mn), blood)	1298
(Zn)	(Zinc (Zn), Serum)	275
(Li)	(Lithium (Li), serum)	275
(Cd)	(Cadmium (Cd), blood)	1298
(Se)	(Selenium (Se), Serum)	275
(Se)	(Selenium (Se), blood)	1298
(Mn)	(Manganese (Mn), Serum)	275
(Co)	(Cobalt (Co), blood)	1298
( )	( 3 , Reverse Triiodothyronine).	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, quantitative))	(. ) (Analysis of translocation t(11;14)(q13;q32)	10835
PML/RAR? -t(15;17) (FISH, quantitative))	(. ) (Analysis of chimeric gene PML/RAR?	2486
CBF?/MYH1- inv(16),t(16;16) (FISH, quantitative))	(. ) (Analysis of chimeric gene CBF?/MYH1- inv(16),t(16;16) (PCR, qualitative))	2486
12 (+12) (FISH, quantitative))	(. ) (Analysis of chromosome 12 trisomy (FISH, quantitative))	10835
V617F JAK2 (Qualitative assessment of presence of gene JAK2 617F somatic mutation)	14 (FISH, quantitative))	1947
t(11;18)(q21;q21) (FISH, quantitative))	(. ) (Analysis of translocation t(11;18)(q21;q21)	10835
(Karyotype, Hematologic Disorders, Peripheral Blood)		7722
53 (FISH, quantitative))	(. ) (Analysis of 53 gene deletion (FISH, quantitative))	10835
t(14;16) (IGH/MAFB) (FISH, quantitative))	(. ) (Analysis of translocation t(14;16) (IGH/MAFB) (FISH, quantitative))	10835
BCR/ABL - t(9;22), BCR/ABL - t(9;22), (Analysis of chimeric gene BCR-ABL - t(9;22), assessment of the BCR-ABL gene transcript type, PCR, qualitative)		2486
12p (FISH, quantitative))	(. ) (Analysis of 12p deletion (FISH, quantitative))	10835
BCR-ABL (FISH, quantitative))	(. ) (Analysis of chimeric gene BCR-ABL, FISH, quantitative))	10835
BCL2 (FISH, quantitative))	(. ) (Analysis of BCL2 gene rearrangements on paraffin slides (FISH Histology, quantitative))	14927
PDGFR?(FISH, quantitative))	(. ) (Analysis of gene rearrangements PDGFR? (FISH, quantitative))	10835
FGFR1 (FISH, quantitative))	(. ) (Analysis of gene rearrangements FGFR1 (FISH, quantitative))	10835
E2A/PBX1 - t(1;19) (FISH, quantitative))	(. ) (Analysis of chimeric gene E2A/PBX1 - t(1;19) (PCR, qualitative))	2486
BRAF (V600E) (FISH, quantitative))	(. )	8569
5 (FISH, quantitative))	(. ) (Analysis of chromosome 5 rearrangements (FISH, quantitative))	10835
t(4;14)(p16;q32) (FISH, quantitative))	(. ) (Analysis of translocation t(4;14)(p16;q32)	10835
t(14;16) (IGH/MAFB) (FISH, quantitative))	(. ) (Analysis of translocation t(14;16) (IGH/MAFB) (FISH, quantitative))	10835
BCR/ABL - RQ (FISH, quantitative))	(. ) (Analysis of the BCR/ABL relative expression, RQ-PCR, quantitative)	4829
(FISH, quantitative))	(. ) (Cytogenetic analysis of bone marrow (karyotype))	7722
FIP1L1/PDGFR?(FISH, quantitative))	(. ) (Analysis of chimeric gene FIP1L1/PDGFR?	10835
MLL (FISH, quantitative))	(. ) (Analysis of MLL gene rearrangements (FISH, quantitative))	10835
7 (FISH, quantitative))	(. ) (Analysis of chromosome 7 rearrangements (FISH, quantitative))	10835
13 - (del(13), -13) (FISH, quantitative))	(. ) (Analysis of chromosome 13 monosomy, deletion - (del(13), -13) (FISH, quantitative))	10835
12 JAK2 (FISH, quantitative))	(. ) (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, quantitative))	(. ) (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 (FISH, quantitative))	(. ) (BCR-ABL1 Mutation Analysis using direct Sanger sequencing, qualitative)	9636
1 (FISH, quantitative))	(. )	14267
3q (FISH, quantitative))	(. ) (Analysis of 3q rearrangements (FISH, quantitative))	10835
RUNX1/RUNX1T1 -t(8;21) (FISH, quantitative))	(. ) (Analysis of chimeric gene RUNX1/RUNX1T1 -t(8;21) (PCR, qualitative))	2486
20q (FISH, quantitative))	(. ) (Analysis of 20q deletion (FISH, quantitative))	10835
MYC (FISH, quantitative))	(. ) (Analysis of MYC gene rearrangements t(8;14)(q24;q32)-t(2;8)(p11;q24), t(8;22)(q24;q11) (FISH, quantitative))	10835
CALR (FISH, quantitative))	(. ) (Analysis of CALR gene mutations, deletions, insertions, PCR, qualitative)	4829

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))		10835
:		
		1320
participant (child or mother or father) ( ) (Additional research		6303
(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))		41833
(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 (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	792
(f329), IgE, ImmunoCAP (Watermelon, Citrullus lanatus, IgE, ImmunoCAP)	792
(f5) IgE, ImmunoCAP	792
, IgE (Beef, IgE, F27)	484
, IgE (Oat, IgE, F7)	484
" 1"	1919.5
, IgE (Buckwheat, IgE, F11)	484
, IgE (Brewer's Yeast, IgE, F403)	572
, IgE (Crab, IgE, F23)	484
, IgE (Peanut, IgE, F13)	484
, IgE (Mango, IgE, F91)	484
(Ca) (Calcium (Ca), air)	1298

(Co)	(Cobalt (Co), air)	1298
(Zr)	(Zirconium (Zr), air)	1298
(Mg)	(Magnesium (Mg), air)	1298
(Be)	(Beryllium (Be), air)	1298
(Pb)	(Lead (Pb), air)	1298
(Mo)	(Molybdenum (Mo), air)	1298
(Al)	(Aluminum (Al), air)	1298
(Hg)	(Mercury (Hg), air)	1298
(Mn)	(Manganese (Mn), air)	1298
(Se)	(Selenium (Se), air)	1298
(V)	(Vanadium (V), air)	1298
(Si)	(Silica (Si), air)	1298
(Rb)	(Rubidium (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 Antigen, ENA, Anti-Ribonucleoprotein Antibodies, Anti-RNP	( ) (Extractable Nuclear	1232
Sc1-70, ENP-A, CENP-B, RP11, RP155, PM-Sc175, Ku, PDGFR, Ro-52), Anti-Sc1-70, ENP-A, CENP-B, RP11, RP155, PDGFR, Ro-52, Immunoblotting)	( , NOR90, Th/To, PM-Sc100, (Scleroderma (Systemic Sclerosis) Antibody Panel: , NOR90, Th/To, PM-Sc100, PM-Sc175, Ku,	4411

(Anti-Nuclear Antibodies, ANA, Screening)	517
Hep-2 (Antinuclear Antibodies, ANA, Hep-2 Substrate, ANA-Hep2, Fluorescent Anti-Nuclear Antibodies detection, FANA, iters)	1265
(Sm, RNP/Sm, SS-A (60 ), SS-A (52 ), SS-B, Scl-70, PM-Scl, PCNA, CENT-B, dsDNA, Histone, Nucleosome, Rib P, AMA-M2, Jo-1 ), (ANA: Anti-Sm, RNP/Sm, SS-A (60 kD ), SS-A (52 kD ), SS-B, Scl-70, PM-Scl, PCNA, CENT-B, dsDNA, Histone, Nucleosome, Rib P, AMA-M2, Anti-Jo-1, Immunoblotting)	3509
IgG (a ), (Anti-Nuclear Antibodies, ANA, IgG, Screening)	1232
	1089
IgG ( - IgG, - ) (Double-Stranded (Native) DNA IgG Antibodies, nti-dsDNA IgG)	627
IgG	1727

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

(Hepatitis C Virus, HCV)	
(Hepatitis C Virus (HCV) RNA, Quantitative PCR, Genotyping (Types 1, 2, 3))	4048
-28 ( -28 ), ( ) (Interleukin 28 Beta IL28B, Genotyping (Study of Genetic Markers Determining Effectiveness of Treatment of Chronic Hepatitis C in Interferon and Ribavirin))	792
(CITO), (HCV RNA, Plasma, Quantitative)*	23034
( ), (Hepatitis C Virus (HCV) RNA, Ultrasensitive PCR)	3355
IgM IgG (Anti-HCV Total (IgG + IgM))*	396
IgG C, (Anti-HCV IgG, Immunoblot)	5621
Qualitative)* (HCV RNA, Serum,	693
RNA, Serum, Quantitative, PCR)* (HCV	3454
1a 1b), 2, 3) (Hepatitis C Virus (HCV) RNA, Plasma, Genotyping, Subtypes (Types 1 (Subtypes 1a, 1b), 2, 3))*	924
Quantitative)* (HCV RNA, Plasma,	11517

(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
Serum, (yping)* (HSV-1, 2 DNA, Scrape of Conjunctiva Epithelial Cells)*	(HSV-1, 2 DNA, (yping)*)	660
(HSV-1, 2 DNA, Scrape of Conjunctiva Epithelial Cells, (yping)*)		275
(HSV-1, 2 DNA, Scrape of Conjunctiva Epithelial Cells, (yping)*)		462
(HSV-1, 2 DNA, Urine, (yping)*)		462
IgG (Anti-HSV-1, 2 IgG)		484
Blood, (yping)* (HSV-1, 2 DNA, Cerebrospinal Fluid, (yping)*)	(HSV-1, 2 DNA, (yping)*)	660
(HSV-1, 2 DNA, Exudate)*		462
(HSV-1, 2 DNA, Scrape of Faucial Epithelial Cells, (yping)*)		275
Ig (Anti-HSV-1, 2 Ig )		462
(HSV-1, 2 DNA, Scrape of Urogenital Epithelial Cells, (yping)*)		506
(HSV-1, 2 DNA, Saliva, (yping)*)		462
(HSV-1, 2 DNA, Exudate, (yping)*)		462
(HSV-1, 2 DNA, Blood)*		418
(HSV-1, 2 DNA, Saliva)*		275
(HSV-1, 2 DNA, Scrape of Skin Epithelial Cells)*	(HSV-1, 2 DNA, (yping)*)	275
(HSV-1, 2 DNA, Scrape of Nasal Epithelial Cells, (yping)*)		462
(HSV-1, 2 DNA, Cerebrospinal Fluid)*	(HSV-1, 2 DNA, (yping)*)	275
(HSV-1, 2 DNA, Prostatic Fluid, Semen)*	(HSV-1, 2 DNA, (yping)*)	275
(HSV-1, 2 DNA, Prostatic Fluid, Semen, (yping)*)		462
IgG (Anti-HSV-2 IgG)		605
(HSV-1, 2 DNA, Serum)*		418
(HSV-1, 2 DNA, Scrape of Nasal Epithelial Cells)*		275
Anti-HSV IgG		770
(HSV-1, 2 DNA, Scrape of Urogenital Epithelial Cells)*		275
(HSV-1, 2 DNA, Scrape of Skin Epithelial Cells, (yping)*)		462
IgG (Anti-HSV-1 IgG)		748
(HSV-1, 2 DNA, Scrape of Faucial Epithelial Cells)*		275
COVID-19		
SARS-CoV-2, IgM (anti-SARS-CoV-2, IgM)		759
(SARS-CoV-2 (N-, S-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
(S) (RBD) SARS-CoV-2, IgG (Anti-SARS-CoV-2, spike (S) protein (RBD), IgG, quantitative).		1419



	SARS-CoV-2, (Coronavirus SARS-CoV-2 RNA detection in nasopharyngeal and oropharyngeal smear)		1419
	SARS CoV-2 (S- , RBD), IgG,		1419
	(S) SARS-CoV-2, IgG, (anti-SARS-CoV-2 S (spike) protein antibody, IgG, qualitative. Assessment of immunity before and after vaccination)		979
	SARS-CoV-2 ( , ), IgG, (Anti-SARS-CoV-2 (nucleocapsid protein), IgG, Abbott)		759
	SARS-CoV-2 (RBD), IgG (anti-SARS-CoV-2 (RBD) IgG avidity)		759
A	IgM Mycoplasma pneumoniae (Anti-Mycoplasma pneumoniae IgM) (Mycoplasma genitalium), (Mycoplasma genitalium, DNA, Scrape of Urogenital Epithelial Cells)*		550
	(Mycoplasma hominis), (Mycoplasma hominis, DNA, Scrape of Urogenital Epithelial Cells)*		275
	(Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Plasma)*		275
	Ig Mycoplasma pneumoniae (Anti-Mycoplasma pneumoniae IgA) (Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Scrape of Faucial Epithelial Cells)*		429
	Ig Mycoplasma pneumoniae (Anti-Mycoplasma pneumoniae IgG)		671
	Ig Mycoplasma hominis ( nti-Mycoplasma hominis Ig ) (Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Sputum)*		231
A	IgG Mycoplasma pneumoniae (Anti-Mycoplasma pneumoniae IgG)		550
	Ig Mycoplasma hominis ( nti-Mycoplasma hominis Ig ) (Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Prostatic Fluid, Semen)*		462
	IgG Mycoplasma hominis ( nti-Mycoplasma hominis IgG) (Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Prostatic Fluid, Semen)*		649
	(Mycoplasma genitalium), (Mycoplasma genitalium, DNA, Prostatic Fluid, Semen)*		275
	(Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Saliva)*		462
	(Mycoplasma hominis), (Mycoplasma hominis, DNA, Urine)*		275
	IgA Mycoplasma hominis ( nti-Mycoplasma hominis IgA) (Mycoplasma genitalium), (Mycoplasma genitalium, DNA, Urine)*		275
A09.05.127	( g) (Magnesium (Mg), Serum)		253
A09.05.032	(Ca) (Calcium Total)		209
	/ / ( + /Potassium, Na+ /Sodium, I- /Chloride, Serum)		275
	(Ca2+ , c ) (Ionized Calcium, Free Calcium)		396
A09.05.033	(P) (Phosphorus (P))		209
	( ) ( , ) (Unsaturated Iron Binding Capacity, UIBC)		209
A09.05.007	(Fe) (Iron (Fe), Serum)		209
	Helicobacter pylori ( ) (Helicobacter pylori Associated Gastritis)*		4004
	PDGFRa		15070
	(1 ): - 1 ( )		1694
	PD-L1 SP263 (Ventana). (PD-L1 expression in tumor tissue by IHC using PD-L1 clone SP263 (Ventana) antibodies). PD-L1 c		18150
	KIT		12760
	(1 + 1 ) (Consultation of Finished Histological Preparations (1 Glass + 1 Block))		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 c PD-L1 SP142 (Ventana). (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
Prostatic Fluid, Semen)*		(Trichomonas vaginalis, DNA,	275
		(Trichomonas vaginalis, DNA, Scrape of Urogenital Epithelial Cells)*	275
		IgG Trichomonas vaginalis ( nti-Trichomonas vaginalis IgG)	671
(Procedural services)			
examinations)**		(Blood serum derivation without further	320
			350
		(Blood sampling without further examinations)**	190
		(Buccal epithelium sampling, 2 persons)	180
		(Cytological material sampling, PCR diagnosis material sampling, microbiology test material sampling)	350
		(Saliva sampling for PCR)	50
A11.12.009			200
		(Biomaterial sampling for the enterobiasis test)	77
		(Buccal epithelium sampling, 3 persons)	270
		( ) (capillary blood sampling)	190
		(Buccal epithelium sampling)	90
C			
	6	(1 ,1b,2,3 ,4,5 ,6)	2618
(Ureaplasma parvum)			
(Ureaplasma parvum, Effectiveness Monitoring of Treatments)			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
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 )	(4 + ):	Chlamydia trachomatis, Neisseria	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
(Identification of Sexually Transmitted Infections (STI) Pathogens, Scrape of Urogenital Epithelial Cells)*	(7 + ):		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 +	( ) 14 : 16, 18, (HPV DNA, Scrape of Urogenital Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68) Screening )	990
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 + , o	14 : 16, 18, 31, 33, 35, (HPV DNA, Scrape of Faucial Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68) Screening )*	385
52, 53, 56, 58, 59, 66, 68, 73, 82 + , o	21 : 6, 11, 16, 18, 26, 31, 33, 35, 39, 44, 45, 51, (HPV DNA, Scrape of Urogenital Epithelial Cells, 21 Types (6, 11, 16, 18, 26, 31, 33, 35, 39, 44, 45, 51, 52, 53, 56, 58, 59, 66, 68, 73, 82))	2695
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
,	(Candida albicans, DNA, Exudate)*	275
,	(Candidiasis, Typing)	737
Susceptibility testing)	( Candida, Cryptococcus) (Yeast Culture. Identification and Antimycotic	759
,	(Candidiasis, Screening )	396
Semen)*	(Candida albicans, DNA, Prostatic Fluid,	275
(Candida albicans, DNA, Scrape of Rectal Epithelial Cells)*		275
DNA, Scrape of Faucial Epithelial Cells)*	(Candida albicans,	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
(Upper Respiratory Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	2178
(Eye Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*	1474
(Eye Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	2464
(Stool Culture with Bacteria Identification and Antibiotic+ Bacteriophage Susceptibility Testing)	1606
(Stool Culture, Pathogenic Intestinal and Conditionally Pathogenic Microflora. Bacteria Identification and Antibiotic Susceptibility Testing)	1518
(Breast Milk Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*	1078
( )	1727
(Ear Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged 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), (Adenovirus. One Step Rapid Immunohromotographic Assay)	957
(Urine Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	880
(Sputum and Tracheobronchial washings Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing, Microscopy)*	2387
(Helicobacter pylori), (Helicobacter pylori. One Step Rapid Immunohromotographic Assay)	957
(Upper Respiratory Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	913
(Punctate Fluid Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	2178
(Stool Culture, Pathogenic Intestinal and Conditionally Pathogenic Microflora, Bacteria Identification)	1309
6 ( )	2398
1 ( )	2398
12 ( ) (Cobalamin)	693
( - ) (Vitamin E, alpha-Tocopherol, Serum)	2398
5 ( )	2398
1,25-D3 (1,25-dihydroxivitamin D3)	2090
3 ( )	2398
25(OH)D2 25(OH)D3, ( - / )	6006
A09.05.080 (Folic Acid)	957
( ) (Vitamin A, Retinol, Serum)	2398
1 ( ) (Vitamin K1, Phylloquinone, Serum)	2398
7, ( )	2398
12 ( , Active-B12, Holotranscobalamin)	1375
-	2398
2 ( )	2398
( )	2398
	2398

LG11 CASPR2 ( ), IgG, (VGKC-associated proteins LG11 and CASPR2 antibodies, serum)	6160
, IgG, (Neuronal antibodies, IgG, Indirect immunofluorescence (IIF))	3311
(Acetylcholine Receptor Antibodies, Anti-AChR, Total)	5731
IgG, NMDA, CASPR, LGI, 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-Ganglioside antibodies, Ganglioside Antibodies Panel, Total)	5731
Critidia luciliae, IgG, (Critidia luciliae indirect fluorescent test (CLIFT))	1265
IgG, (Anti-myelin antibody, IgG, IIF)	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
- ( -MuSK) (Muscle-specific tyrosine kinase (MuSK) antibody)	5478
IgG, NMDA, CASPR, LGI, AMPA1, AMPA2, GABAR1	13750
LG11 CASPR2 ( ), IgG, (VGKC-associated proteins LG11 and CASPR2 antibodies, CSF)	6160
GAD ( ), IgG, (Anti-GAD (glutamic acid decarboxylase), IgG, CSF)	2079
IgG ( - : Hu (ANNA1), Yo-1 (PCA1), CV2, 2, Ri (ANNA2), ) (Anti-Neuronal Antibodies, Blot-Line (Hu (ANNA1), Yo-1 (PCA1), CV2, 2, Ri (ANNA2), Amphiphysin))	5731
IgG NMDA (N- -D- ) (N-Methyl-D-Aspartate Receptor Antibodies IgG)	4389
: (CH50) (Functionality Test of Complement (CH50))	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, (Anti-HBc IgM, IgG, Antibodies to Hepatitis B Core Antigen; HBcAb, Total, HBV Core Total Antibodies (IgG + IgM))	517
HBs- (HBs- , (HBsAg, Hepatitis Surface Antigen, « » ), 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
B, (HBV DNA, Serum, Quantitative)*	3751



Qualitative)*	B, (HBV DNA, Serum,	418
HBS-	(Anti-HBs, HBsAb)	627
HB -	(Anti-HBe, HBeAb)	517
HBS- « » ),	(HBsAg, Hepatitis Surface Antigen, Qualitative)	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	(Lactate)	143
	(Fructosamine)	572
A09.05.083	HbA1 (HbA1 , Glycated Hemoglobin, GHB)	891
	( ) Oral Glucose Tolerance Test, Plasma, OGTT, Pregnancy	506
A12.22.005	- 2 (2-Hour Oral Glucose Tolerance Test, OGTT, Glucose Concentration (Fasting and 2 Hours after Load), Venous Blood)	990
A09.05.009	- ( ) (C-Reactive Protein, CRP)	671
	- ( - , ) (Antistreptolysin-O, ASO)	341
-1-	( 1 ), (Alpha-1-Antitrypsin, A1AT, AAT, Phenotyping)	374
	( - ) N- (NT-proBNP, N-Terminal Pro-brain Natriuretic Peptide, Pro-B-Type Natriuretic Peptide)	2596
A09.05.076	(Ferritin)	2816
	(Carbohydrate-Deficient Transferrin with results on an electrophoregram (CDT))	506
	25 ( ) (Hepcidin 25, bioactive)	3322
	ST2 (ST2, sST2, ) (Soluble ST2 (Heart Failure's biomarker))	6567
A09.05.008	( ) (Transferrin)	2739
	(Myoglobin)	484
-2-	(Alpha-2-Macroglobulin, ?2-Macroglobulin, A2M)	572
Pgp3 (	IgG ) Chlamydia trachomatis IgG	495
	( ) (Rheumatoid Factor, RF)	572
	-I (Troponin-I)	374
	- ( )	627
A09.05.077	(Ceruloplasmin)	594
	(Haptoglobin)	649
-1-	( 1 ), (Alpha-1-Antitrypsin, A1AT, AAT, Concentration)	638
	(Carbohydrate-Deficient Transferrin, CDT)	1386
	( , Soluble Transferrin Receptor, sTfR)	3091
	(Eosinophil Cationic Protein, ECP)	1870
	(Ureaplasma parvum), (Ureaplasma parvum, DNA, Urine)*	869
	(Ureaplasma arvum), (Ureaplasma parvum, DNA, Prostatic Fluid, Semen)*	275
	IgG Ureaplasma urealyticum ( nti-Ureaplasma urealyticum IgG)	275
	(Ureaplasma urealyticum) ( -960), (Ureaplasma urealyticum (T-960), DNA, Scrape of Urogenital Epithelial Cells)*	671
	IgA Ureaplasma urealyticum ( nti-Ureaplasma urealyticum IgA)	275
		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 (RSV) IgG	(Anti-Respiratory Syncytial Virus)	781
A (RSV) IgM	(Anti-Respiratory Syncytial Virus)	781
-		
(Genes F2, F5)	( F2, F5) (Risk of Oral Contraceptives, Ocs)	3003
-	( F2, F5) (Risk of Oral Contraceptives, OCs (Genes F2, F5) (without Description))	2695
	( 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)*	1: , IgG (FP15 (F33, F49, F92, F95),	1045
(F84), -	IgG (Kiwi Fruit, IgG, F84)	572
(F11), -	IgG (Buckwheat, IgG, F11)	572
(F3), -	IgG (Codfish, IgG, F3)	572
F83, F88), Food Panel: Pork, Beef, Chicken Meat, Lamb, IgG)*	3: , IgG (FP73 (F26, F27, G (IgG) (Basic	15840
(F44), -	IgG (Strawberry, IgG, F44)	572
(F7), -	IgG (Oat, IgG, F7)	572
(F105), -	IgG (Chocolate, IgG, F105)	572
	(fx21) IgE, ImmunoCAP	1375
(F4), -	IgG (Wheat, IgG, F4)	572
-	, IgG (Beta Lactoglobulin, IgG, F77)	572
(F2), -	IgG (Milk, IgG, F2)	572
(F25), -	IgG (Tomato, IgG, F25)	572
(F55), -	IgG (Common Millet, IgG, F55)	572
	, IgG (Pineapple, IgG, F210)	572
	, IgG (Lamb, IgG, F88)	572
(F26), -	IgG (Pork, IgG, F26)	572
(F208), -	IgG (Lemon, IgG, F208)	572
	, IgG (Banana, IgG, F92)	572
(F75), -	IgG (Egg Yolk, IgG, F75)	572

(F1),	-	IgG (Egg White, IgG, F1)	572
(F83),	-	IgG (Chicken Meat, IgG, F83)	572
(F35),	-	IgG (Potato, IgG, F35)	572
, IgG (Orange, IgG, F33)			572
(F209),	-	IgG (Grapefruit, IgG, F209)	572
(F91),	-	IgG (Mango, IgG, F91)	572
, IgG (Peanut, IgG, F13)			572
2		, IgG (FP50 (F84, F91, F92, F210), Food Panel: Kiwi Fruit, Mango, Banana, Pineapple, IgG)*	1430
(F225),	-	IgG (Pumpkin, IgG, F225)	572
(F14),	-	IgG (Soybean, IgG, F14)	572
(F31),	-	IgG (Carrot, IgG, F31)	572
(F27),	-	IgG (Beef, IgG, F27)	572
(F45),	-	IgG (Baker's Yeast, IgG, F45)	572
(F216),	-	IgG (Cabbage, IgG, F216)	572
(F403),	-	IgG (Brewer's Yeast, IgG, F403)	572
(F17),	-	IgG (Hazelnut, IgG, F17)	572
(F78),	-	IgG (Casein, IgG, F78)	572
(F95),	-	IgG (Peach, IgG, F95)	572
(F49),	-	IgG (Apple, IgG, F49)	572
-	6		
IgG	6	(Anti-HHV-6 IgG)	671
6	,	(HHV-6 DNA, Saliva)*	275
6	,	(HHV-6 DNA, Scrape of Urogenital Epithelial Cells)*	275
6	,	(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
			275
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)*			275
(Chlamydia trachomatis), (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)*		275
A	IgA Chlamydia pneumoniae (Anti-Chlamydia pneumoniae IgA)		671
	(Chlamydia pneumoniae), (Chlamydia pneumoniae, DNA, Sputum)*		869
A	IgM Chlamydia pneumoniae (Anti-Chlamydia pneumoniae IgM)		561
	(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Urine)*		275
	(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Cerebrospinal Fluid)*		275
	(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Exudate)*		275
	(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Synovial Fluid)*		506
	(Chlamydia pneumoniae), (Chlamydia pneumoniae, DNA, Scrape of Faucial Epithelial Cells)*		429
	(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Scrape of Faucial Epithelial Cells)*		275
	(Treponema pallidum, DNA, Prostatic Fluid, Semen)*		275
	(Treponema pallidum, DNA, Scrape of Urogenital Epithelial Cells)*		275
	(Treponema pallidum, DNA, Urine)*		275
	(Treponema pallidum, DNA, Cerebrospinal Fluid)*		275
	(Treponema pallidum, DNA, Serum)*		407
	(Treponema pallidum, DNA, Secretion)*		275
A	IgM IgG Treponema pallidum, (Anti-Treponema pallidum IgM, IgG, Total)		418
	(Treponema pallidum, DNA, Scrape of Skin Epithelial Cells)*		275
	(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 Reagents), nticardiolipin 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
	( ) (Androstenediol Glucuronide, 3?-Androstenediol 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
<b>IgE:</b>	
, 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
<b>( )</b>	
1 2 1 2 (HIV Ag/Ab Combo)	319
-1, (HIV RNA, Plasma)*	14234
<b>(Everolimus)</b>	
(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
<b>( )</b>	
Prostatic Fluid, Semen)* (Neisseria gonorrhoeae, DNA,	275
(Neisseria gonorrhoeae, DNA, Scrape of Rectal Epithelial Cells)*	275
Fluid)* (Neisseria gonorrhoeae, DNA, Synovial	506
(Neisseria gonorrhoeae, ), (GC, Neisseria gonorrhoeae Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	990
(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) ( MTRR, MTR) (without Description)) (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
/ BRCA1, BRCA2, CHEK2, NBS1 (Hereditary Breast and/or Ovarian Cancer )	10285
(Examination of Sputum)	825
(Examination of Transudates, Exudates, Secrets)	528
( Examination of Bronchial Washouts)	649
(Examination of Punctates: Skin)	649
(Examination of Endoscopic Material)	649
A08.20.004	660
Helicobacter pylori (Examination of Endoscopic Material: Presence of Helicobacter pylori)	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
( ) (Cytological Examination: Scrapings (Smear) of Nasal Mucous Membrane (1 Localization))	781
(Examination of Breast Discharge)	528
(Examination of Punctates: Other Organs and Tissues)	825
( ) (Examination of Imprint Intrauterine Device, IUD)	561
(Examination of Punctates: Breast)	649
(Examination of Scrapings and Prints Tumor and Tumor Like Formations)	649
(Examination of Urine)	528
(The Bethesda System for Reporting Thyroid Cytopathology (TBSRTC), Fine-Needle Aspiration (FNA))	627
(Examination of Scrapings and Prints of Skin and Mucous Membranes)	429
(Examination of Scrapings: Cervix and Cervical Canal )	627
Ig ( nti-Measles IgM)	781

IgG	(Anti-Measles IgG)		880
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	(Anti-Tick-borne Encephalitis Virus (TBEV))		671
	(HPLC-MS/MS Organic Acids (Succinylate))		3949
	HADHA (HADHA Gene, Freq. Mut. (Long-Chain 3-Hydroxyacyl-Coa Dehydrogenase (LCHAD) Deficiency))		5511
	(Newborn Screening "HEEL")*		5489
	(Biotin-Dependent Carboxylases Activity (Biotinidase Deficiency))		5511
	GCDH (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 (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
	(OTC Gene (Ornithine Transcarbamylase (OTC) Deficiency))		41833
	(Escherichia coli O157:H7, Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)		1210
	(Escherichia coli O157:H7, Identification and Antibiotic Susceptibility Testing)		1023
	(Escherichia coli O157:H7, Assay) (Escherichia coli O157:H7. One Step Rapid Immun chromatographic		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 immunochromatographic assay)		946
(Campylobacter spp., One step rapid immunochromatographic assay, antigen, stool)		1144
(Norwalk virus) - (Norwalk virus GI, GII, One step rapid immunochromatographic assay, antigen, stool)	I II	1925
One step rapid immunochromatographic assay, antigen, urinae)	(Streptococcus pneumoniae,	1584
immunochromatographic assay, antigen, stool)	(Enterovirus, One step rapid	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) (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
	( ) (Immunohistochemical diagnosis in cancer metastasis of unknown primary origin (Fixed Biomaterial in Formalin Buffer))*		27808
A09.05.065	( )		363
A09.05.064	(T4, ) (Total Thyroxine, TT4)		385
A09.05.061	( 3 ) (Free Triiodthyronine, FT3)		385
	( ) (Anti-Thyroid Microsomal Antibodies)		550
A09.05.063	( 4)		385
	( - , ) (Anti- thyroid Peroxidase Autoantibodies, Antimicrosomal Antibodies, TPO Antibodies, TPOAb, Anti-TPO)		429
A12.06.046.001	( ) (Thyroid-Stimulating Hormone Receptor Antibodies, TSH Receptor Antibodies, TSHRabs, TSH binding inhibitor immunoglobulin, TBII)		1540
	( - ) (Anti- thyroglobulin Autoantibodies, Thyroglobulin Antibodies, Tg Autoantibodies, TgAb, Anti-Tg Ab, ATG)		484
A09.05.117	( ) (Thyroglobulin, TG)		693
	( ) (Thyroid Uptake, T-Uptake, Thyroxine-Binding Capacity, TBC, Thyroxine-Binding Index, TBI, free T4Index, FT4I)		561
A09.05.060	( 3 ) (Total Triiodthyronine, TT3)		385
	SARS-CoV-2, (Coronavirus SARS-CoV-2 RNA detection in nasopharyngeal and oropharyngeal smear)		1529
	: (Soil: Agrochemical Evaluation)*		9988
	: (Soil: Comprehensive Toxicological Evaluation)*		21934
A09.05.066	( , ) (Growth Hormone, GH)		517
A09.05.067	( , ) (Adrenocorticotrophic Hormone, ACTH)		682
	( 1) (Somatomedin C, Insulin-like Growth Factor 1, IGF-1)		1089
A09.05.131	( ) (Luteinizing Hormone, LH)		385
	(Macroprolactin)*		1188
A09.05.087	(Prolactin)		385
A09.05.132	( ) (Follide Stimulating Hormone, FSH)		385
	IgG ( -ASGPR) (Autoantibodies Against Asialoglycoprotein Receptor, Anti-ASGPR, IgG)		1705

IgA, IgG, IgM (Anti-Mitochondrial Antibodies, AMA, IgA, IgG, IgM, Total)		1518
IgG (Autoimmune Disease Liver Panel: AMA-M2, M2-3E (BPO), Sp100, PML, gp210, LKM-1, LC-1, SLA/LP, SSA/Ro-52), Immunoblotting)		3652
IgA+IgG+IgM (anti-liver kidney microsomal antibody, anti-LKM, IgG+IgM+ IgA)		1573
IgA, IgG, IgM (Smooth Muscle Antibodies, SMA, Anti-Smooth Muscle Antibodies, ASMA, IgA, IgG, IgM, Total)		1518
1-		
IgG (Insulin Autoantibodies, IAA, IgG)		671
(IA-2) (Islet Antigen 2 Antibodies, Anti-IA2 antibodies, IA-2 Ab, Tyrosine Phosphatase Antibodies)		1705
IgG (Anti-Islet Cell Antibodies, Islet Cell Autoantibodies, ICA)		1518
GAD/IA-2, (Anti-GAD/IA2 Antibodies Pool, Glutamic Acid Decarboxylase-65, GAD and Insulinoma Antigen 2 (Tyrosine Phosphatase, IA2, ICA-512) Autoantibodies, Total)		1705
IgG (-GAD) (Anti-GAD Antibodies, Glutamate Decarboxylase Antibodies, AT-GAD, IgG)		1749
( )		
(Yersinia enterocolitica, (Yersinia enterocolitica, Stool Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		1364
Yersinia enterocolitica :9 (Yersinia enterocolitica O:9, IHA)		484
IgG Yersinia enterocolitica (Anti-Yersinia enterocolitica IgG)		539
IgA Yersinia enterocolitica (Anti-Yersinia enterocolitica IgA)		539
Yersinia enterocolitica :3 (Yersinia enterocolitica O:3, IHA)		484
Yersinia pseudotuberculosis (Yersinia pseudotuberculosis IHA)		484
( F2, F5, MTHFR, MTR, MTRR, F13, FGB, ITGA2, ITG 3, F7, PAI-1) (Extended Study of Hemostatic System (Genes F2, F5, MTHFR, MTR, MTRR, F13, FGB, ITGA2, ITG 3, F7, PAI-1))		
( 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) (MTRR, MTR))		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

( <i>Listeria monocytogenes</i> ) ( <i>Listeria monocytogenes</i> Culture. Bacteria Identification and Antibiotic Susceptibility testing)		759
( <i>Listeria monocytogenes</i> , DNA, Plasma)*		231
( <i>Listeria monocytogenes</i> , DNA, Scrape of Faucial Epithelial Cells)*		231
( <i>Listeria monocytogenes</i> , DNA, Synovial Fluid)*		506
IgE:		
( <i>IgE</i> (GP3 (G1, G5, G6, G12, G13), Grass Panel: Sweet Vernal Grass, Perennial Rye Grass, Timothy Grass, Cultivated Rye Grass, Velvet Grass, <i>IgE</i> *)		1045
( <i>IgE</i> (Cottonwood, <i>IgE</i> , T14)		484
( <i>IgE</i> (Wormwood, <i>IgE</i> , W5)		484
( <i>IgE</i> (Timothy Grass, <i>IgE</i> , G6)		484
( <i>IgE</i> (WP1 (W1, W6, W9, W10, W11), Weed Panel: Common Ragweed, Mugwort, English Plantain, Lamb's Quarters, Russian Thistle, <i>IgE</i> *)		1045
( <i>IgE</i> (Birch, <i>IgE</i> , 3)		484
( <i>IgE</i> (Mugwort, <i>IgE</i> , W6)		484
( <i>IgE</i> (TP9 (T2, T4, T12, T3, T7), Tree Panel: Alder, Hazelnut, Willow, Birch, Oak, <i>IgE</i> *)		1045
( <i>IgE</i> (GP1 (G3, G4, G5, G6, G8), Grass Panel 1: Orchard Grass, Meadow Fescue, Perennial Rye Grass, Timothy Grass, June Grass (Kentucky Bluegrass), <i>IgE</i> *)		1045
(25-OH Vitamin D Total, 25(OH)D, 25-Hydroxycalciferol)		
(Deoxypyridinoline, DPD, Urine)		2112
(Osteocalcin, N-Osteocalcin, Bone Gla Protein, BGP)		1375
(Procollagen Type 1 N-terminal Propeptide, P1NP, Total)		737
(-CrossLaps, -CrossLaps Serum, C-Telopeptide, Crosslaps, Type 1 Collagen, b-Tx Serum)		1496
(Human Cartilage Oligomeric Protein, COMP)		957
(Streptococcus group B, Streptococcus agalactiae), (Streptococcus agalactiae Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		
(Streptococcus group B, Streptococcus agalactiae) (Streptococcus agalactiae Culture. Bacteria Identification)		968
(Streptococcus Group B. One Step Rapid Immunochromatographic Assay)		759
(Streptococcus Group A. One Step Rapid Immunochromatographic Assay)		1155
(Streptococcus group A, Streptococcus pyogenes), (Streptococcus pyogenes Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		957
(Streptococcus spp., DNA, Saliva)*		891
(Streptococcus group A, Streptococcus pyogenes) (Streptococcus pyogenes Culture. Bacteria Identification)		429
(Streptococcus spp., DNA, Plasma)*		682
(Streptococcus spp., DNA, Scrape of Faucial Epithelial Cells)*		649
(Streptococcus spp., DNA, Sputum)*		429
(Metanephrines fractionated, free and conjugated, 24-h urine)		869
A09.28.034.001 24-		2486

( ), 5- and Serotonin Metabolites, 24 Hours-Urine: Vanillylmandelic Acid, V (5- ) (Catecholamines, 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	(Anti-Giardia lamblia IgM, IgG, IgA, Total)	627
( )		
A09.28.027 (Amylase, 24-Hour or Timed Urine)	( - ,	253
(Ca), (Calcium (Ca), 24-Hour urine)		220
(K), (Na), (Potassium (K), Sodium (Na), 24-Hour urine)		209
(Glomerular Filtration Rate, GFR)*		209
A09.28.011 (Glucose, 24-Hour urine)		165
A09.28.009 (Urea, 24-Hour urine)		165
A09.28.010 (Uric acid, 24-Hour urine)		209
A09.28.003 (Protein Total, 24-Hour urine)		143
( ), (Magnesium, 24 h urine excretion)		330
( oxalates, 24-Hour urine)		2079
A09.28.003.001 (Albumin, 24-Hour urine)		1342
A09.28.006 (Creatinine, 24-Hour urine)		352
(P), (Phosphorus (P), 24-Hour urine)		165
		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 ( - ) (Glomerular Basement Membrane antibodies, anti-GBM, IgG)		1716
IgG ( ) (Anti-Neutrophil cytoplasmic antibodies, ANCA, IgG)		1342
IgG -3 ( -PR-3) (Anti- proteinase-3 antibodies, PR-3- antibodies, PR-3 ANCA, IgG)		1232
IgG, IgA, IgM (Anti-Phospholipase A2 Receptor Antibodies, Anti-PLA2R, IgG, IgA, IgM, Total)	2 (PLA2R),	2717
(Anti-Neutrophil Cytoplasmic Antibodies, ANCA, IgG, Panel)	( ), IgG	3410
IgG C1q (Anti-Complement 1q Antibodies, Anti-C1q, IgG)		1232

IgG, IgA, IgM (Anti-Endothelial Cell Antibodies, AECA, IgG, IgA, IgM, Total)	(HUVEC),	1705
IgG	( - PO) (Myeloperoxidase Antibody, MPO)	1232
1- 3-	(Anti-Poliavirus serotypes 1, 3, IgG)	1595
(HiB), anti-PRP Haemophilus influenzae b IgG)	(polyribosylribitolphosphate, PRP) (Haemophilus influenzae b IgG)	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, - (Anti-Borrelia burgdorferi IgM, Western Blot (WB))	1980
A	IgG Borrelia burgdorferi, (Anti-Borrelia burgdorferi IgG, Immunoblot )	2222
A	IgM Borrelia burgdorferi (Anti-Borrelia burgdorferi IgM)	627
	(Barbiturates, Urine)*	1232
Substances Screening: Opiates, Amphetamines, Methamphetamine, Cocaine, Cannabinoids, Cannabinoid Metabolites, Urine)	? ( ) (Drugs and Psychotropic	3410
( )	(Cannabinoids (Marijuana), Urine)*	1232
( )	(Ethanol (Alcohol) Urine)*	1232
« » ( ) ( ; ) (Pernicious Habits: Nicotine, Drugs, Psychostimulants and Psychotropic Substances, Urine)*		3608
( / )	(Opiates (Morphine/Heroin), Urine)*	1232
- ,	(EBV DNA, Exudate)*	275
Fluid)*	(EBV DNA, Cerebrospinal	275
Antigens (VCA) IgG )	- ( nti-EBV Viral Capsid	759
IgG )	- ( nti-EBV Early Antigen (EA)	627
(EBNA) IgG )	- ( nti-EBV Nuclear Antigen	528
- ,	(EBV DNA, Blood)*	418
- ,	(EBV DNA, Saliva)*	275
Fluid, Semen)*	(EBV DNA, Prostatic	275
(EBV DNA, Scrape of Nasal Epithelial Cells)*		275
(EBV DNA, Scrape of Urogenital Epithelial Cells)*		275
Ig Antigens (VCA) Ig )	- ( nti-EBV Viral Capsid	528
DNA, Scrape of Faucial Epithelial Cells)*	(EBV	275
Serum)*	(EBV DNA,	407
- ,	(EBV DNA, Urine)*	275
- ,	(EBV DNA, Serum)*	418
IgE:		2024
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))	5203
: D ( VDR) ( (Osteoporosis, Vitamin D Receptor (VDR) (Gene VDR) (without Description))	1386
( MCM6) (Adult Lactase Deficiency (Gene MCM6)	1441
, HFE) (Hemochromatosis Type 1 (Gene HFE))	2860
: D ( VDR) (Osteoporosis, Vitamin D Receptor (VDR) (Gene VDR))	1529
( UGT1A1) (Gilbert's Syndrome (Gene UGT1A1))	4994
: ( CALCR, COL1A1) ( - ) (Osteoporosis: Abridged Panel (Genes CALCR, COL1A1) (without Description))	3993
( MTHFR, MTRR, MTR) (Folic Acid Metabolism (Genes MTHFR, MTRR, MTR))	5808
: ( CALCR, COL1A1) (Osteoporosis: Abridged Panel (Genes CALCR, COL1A1))	4455
: ( CALCR, COL1A1, VDR) ( - ) (Osteoporosis: Full Panel (Genes CALCR, COL1A1, VDR) (without Description))	5203
HLA II ( DRB1, DQA1, DQB1) (Hereditary Predisposition to Diabetes Type 1 (Insulin-Dependent Diabetes), HLA Class II (Genes DRB1, DQA1, DQB1))	6864
: ( CALCR, COL1A1, VDR) (Osteoporosis: Full Panel (Genes CALCR, COL1A1, VDR))	5808
-	
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
Cerebrospinal Fluid)*	(Toxoplasma gondii, DNA,	275
	Anti-Toxopl gondii IgG	1045
	(Toxoplasma gondii, DNA, Exudate)*	275
	(Streptococcus	495
pneumoniae)		
	(Bordetella pertussis/parapertussis,	1584
/	) (Bordetella pertussis/parapertussis, Nasopharyngeal Culture. Bacteria	
Identification)		
	( 2) (Estradiol, E2)	385
	(Progesterone)	385
A09.05.135	( ) (Cortisol, Hydrocortisone)	385
A09.28.035	(Free cortisol, Free Hydrocortisone, 24-Hour	759
urine)		
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,	1232
	-1- ( ) (Trophoblastic beta-1-Globulin, TBG)	484
	IgG V (Annexin V antibodies, aAnV, IgG)	1309
	Ig V (Annexin V antibodies, aAnV, Ig )	1309
Antibodies, anti-?-G 1, IgG, IgA, IgM, Total)	-2- 1, ( nti-?-2-Glycoprotein 1	1254
	IgM IgG ( nti-Phospholipid Antibodies, APA, IgM, IgG)	803
2	IgA	1199
(Anti- hosphatidylserine/ rothrombin antibodies, Anti-PS/PT, IgG, IgM, Total)	-	1342
aCL, Screening)	IgA, IgM, IgG ( ardiolipin Antibodies IgA, IgM, IgG,	1133
	IgG IgM (Anti- hosphatidylserine, IgG, IgM)	1782
2	IgM	1309
	, IgG, IgM	9009
	(Anti-Phospholipid Antibodies Panel)	
	IgA (Anticardiolipin IgA, aCL IgA)	836
2	IgG	1199

IgG	(Anticardiolipin IgG, aCL IgG)	902
IgM	(Anticardiolipin IgM, aCL IgM)	1188
	(Bile Acids)	2651
1 (	1, 1) (Apolipoprotein A1, Apo A1)	572
Cholesterol)	( , , VLDL	407
B (	B, ) (Apolipoprotein B, Apo B)	429
A09.05.025	( ) (Triglycerides)	209
A09.05.004		220
(a), (	) (Lipoprotein (a), Lp (a))	869
A09.05.028		165
A09.05.026	( ) (Cholesterol Total)	209
(	) Cholesterol LDL (direct)	253
(4 )		
(Neovir)		539
(Amixin)		539
(Cycloferonum)		539
(KagoceI)		539
Protein-A, PAPP-A)	(Pregnancy-Associated Plasma	693
A09.05.090	( , - , ?- ) (Human Chorionic	385
Gonadotropin, HCG)		
?- (	?- ) (Free	539
Human Chorionic Gonadotropin, Free HCG)		
PRISCA2		99
fms-	-1 (sFit-1)	3377
	(Estril Free, 3)	484
PRISCA1		132
(	) (Placental Lactogen, PL, Human	693
Placental Lactogen, hPL, Chorionic Somatomammotropin, CS, Human Chorionic	Somatomammotropin, hCS)	
(	) (Erythrocyte Sedimentation Rate, ESR)	143
«	» (Leucocyte Formula (Differential White	330
Blood Cell Count) with Manual Microscopic Examination of Blood Smear)*		
(	)	209
(Differential White Blood Cell Count) with Microscopic Examination of Blood Smear if Presence of	Pathologic Changes)*	
(	) (Platelets,	275
Microscopy (Manual Platelet Count (PLT Count): Indirect Method by Fonio))*		
( ) (	) (General Blood Analysis,	209
without White Blood Cell (WBC) Count and ESR)		
A12.05.123	(Reticulocytes)	264
		1980
(4 )		
(Ingaron)		539
(Reaferonum)		539

Bordetella species: Bordetella pertussis ( ) Bordetella bronchiseptica ( ) (Differentiated detection of DNA Bordetella spp.: Bordetella pertussis (pertussis pathogen) and Bordetella bronchiseptica (bronchosepticosis pathogen) in a scraping of the oropharynx and or nasopharynx)		924
A	IgG Bordetella pertussis (Anti-Bordetella pertussis IgG)	891
A	IgM Bordetella pertussis (Anti-Bordetella pertussis IgM)	891
A	IgA Bordetella pertussis (Anti-Bordetella pertussis IgA )	891
-3	(Omega-3 Index)	4774
	(Procalcitonin)	3003
	, : -3,-6,-9, (Fatty acids panel, omega-3, -6, -9, plasma)	9174
	Rh- (Anti Rh)	517
A12.05.005	(Blood Group, O)	253
Rh (C, E, c, e) Kell-	(Rh C (E, c, e) Kell-Phenotyping)	660
A12.05.006	( - ) (Rh-factor, Rh)	253
:	(Water: Complete and Comprehensive Quality Assessment)*	19998
:	(Water: Suspicion Industrial Waste Contamination)*	14190
:	(Water: Suspicion Products of Combustion and Emissions from Motorways Contamination)*	10483
:	(Water: Suspicion Household Waste Contamination)*	8943
:	(Water: Abridged Quality Assessment)*	9669
:	(Water: Suspicion Excessive Use of Chemicals for Water Treatment)*	5159
	( , Trichinella spiralis)	
	IgG ( 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 (Gastrin-17 Stimulation Test, I (Pepsinogen I)		1210
A09.05.057 (Gastrin)		968
(GastroPanel)		682
II (Pepsinogen II)		4576
G (G, Hepatitis G Virus, HGV)		968
G, (HGV RNA, Serum)*		660
(FibroTest)		11946
(FibroMax)		15268
(FibroTest)		13420
(SteatoScreen)		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
ATI.		3036
( ACE) (ACE Inhibitors, Fluvastatin, ATI Receptor Blockers.		5808
(Methotrexatum. Genetic Markers of Increased Risk of Development of Adverse Reactions in Taking Methotrexate for Treatment of Rheumatoid Arthritis. Methotrexate Disrupts Metabolism		
A IgG (Anti-Mumps IgG)		781
A IgM (Anti-Mumps IgM)		781
( )		
(Streptococcus pneumoniae, DNA)		495
(Calcitonin)		946

A09.05.058	( ) (Parathyroid Hormone, PTH)		671
( , Echinococcus spp.)			
	IgG	(Anti-Echinococcus IgG)	869
	IgG	(Anti-Entamoeba histolytica IgG)	693
D ( D, Hepatitis D Virus, HDV)			
	IgM IgG	D, o (Anti-HDV Total (IgG + IgM))	858
	D,	(HDV RNA, Serum)*	660
	IgM	D ( nti-HDV IgM)	858
M3)	Aspergillus fumigatus ( 3),	- IgG (Aspergillus fumigatus, IgG,	572
	(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 antibodies IgG, Indirect)	3278
( , , , Salmonella spp.)			
	Salmonella gr.A	(Salmonella gr.A, IHA)	484
	Salmonella gr.B	(Salmonella gr.B, IHA)	484
	Salmonella gr.E,	(Salmonella gr.E Antibodies, IHA)	484
	Salmonella typhi,	(Salmonella typhi Antibodies, IHA)	616
	Salmonella O-	(Salmonella O-antigens, IHA)	484
	Salmonella gr.D	(Salmonella gr.D, IHA)	484
	Salmonella gr.	(Salmonella gr.C, IHA)	484
	IgG	(Anti-Strongyloides stercoralis IgG)	990
( )			
		(Rickettsia prowazekii, IHA)	484
A09.05.021			165
A09.05.022			165
		Anisakis IgG	814
( )			
	IgG	(Anti-Diphtheria Toxoid IgG)	990
		(Corynebacterium diphtheriae Culture)	759
		(Anti-Spermatozoa Antibodies, ASA, Semen)	1419
		(Anti-Spermatozoa Antibodies, ASA, Serum)	1045
( E, Hepatitis E Virus, HEV)			
	IgM	E (Anti-HEV IgM)	891
	IgG	E (Anti-HEV IgG)	891
IgG:			
	( 2),	- IgG (Dog Epithelium, IgG, E2)	572
	( 1),	- IgG (Cat Dander-Epithelium, IgG, E1)	572
		(Streptococcus pneumoniae, DNA)	495
	IgG	(Anti-Adenovirus IgG)	781
	IgA	(Anti-Adenovirus IgA)	781
		(Streptococcus pneumoniae, DNA)	495
( , Toxocara canis)			
	IgG	(Anti-Toxocara IgG)	517
		(Androflor® REAL-TIME PCR Detection Kit, the study of men's urogenital tract microbiocenosis in the epithelial scrapes from the balanus, urethra)	2838
		(Androflor® Screen REAL-TIME PCR Detection Kit, the study of men's urogenital tract microbiocenosis in the epithelial scrapes from the balanus, urethra)	1991

( DLG5, NOD2, OCTN1, OCTN2) (Crohn's Disease (Genes DLG5, NOD2, OCTN1, OCTN2))		8899
IgA, IgM, IgG (Anti-Ovarian Antibodies, AOA, IgA, IgM, IgG, Total)		1419
IgA, IgM, IgG (Anti-Steroidal Cell Antibodies, StCAb, Steroidal Cell Autoantibodies, SCA, IgA, IgM, IgG, Total)		1232
IgA, IgM, IgG (Anti-Testicular Steroid-ell Antibodies, Testicular Anti-Steroidal Cell Antibodies, Testicular StCAb, Steroidal Cell Autoantibodies, SCA against Testis, IgA, IgM, IgG, Total)		1705
YP2D6 ( YP2D6) (Cytochrome YP2D6 (Gene YP2D6))		8129
(Neisseria meningitidis Culture. Bacteria Identification and Antibiotic Susceptibility Testing)		935
(Total Antioxidant Status, TAS)		5247
IgE: , IgE (Latex, IgG, K82)		484
IgG (Anti-Ascaris lumbricoides IgG)		968
(Demodex folliculorum, Demodex brevis)		374
(Demodex folliculorum, Demodex brevis)		374
(Consultation of Finished Cytological Preparations (1 Glass))		418
Clonorchis sinensis IgG		1078
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 Immun hromotographic 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 ( )» (Comprehensive Study «Sex in City: 6 Infections (Blood Test)»)	3465
« : 6 ( )» (Comprehensive Study «Sex in City: 6 Infections (Blood Test)»)	3597
: (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



Ki-67 ( ) (Cervical Cancer – Study of Two Markers for Early Diagnosis Dysplasia with High Risk Malignancy: p16INK4a + Ki-67, Immunohistochemical Screening (Fixed Biomaterial in Formalin Buffer))*	8613
(Metabolic bone and osteoporosis risk evaluation: comprehensive examination).	5896
: I (PRIS A-1) (Maternal Screen, First Trimester; Prenatal Screening I; PRIS A I (Prenatal Risk Calculation))	1287
: II (PRIS A-2) (Maternal Screen, Second Trimester; Prenatal Screening II; PRISCA II (Prenatal Risk Calculation))	1430
B03.005.006 ( ), (Coagulation, Gemostaziogram, Screening)	847
: (Miscarriage: Autoimmune Profile)	4004
(Immunological Survey Extended)	12078
: (Survey of Liver: Screening)	924
B03.016.004 :	1210
Supersport	2200
: (Serum Biochemistry: Minimum)	2222
Supersport	3410
: (Serum Biochemistry: Extended Profile)	3553
- : ( 40 ) (Healthy You ? Healthy Country: Annual Check-Up up to 40 Years of Age)	3861
( 40 ) (Annual Check-Up after 40 Years of Age)	4928
: (Survey Before Diet: Additional )	5159
Supersport	7513
	4059
: (Survey of Kidneys: Extended )	2200
: (Diabetes Control: Extended)	3542
(Toxic Trace Elements, Hair)	1782
(Toxic Trace Elements, Essential Vital Elements, Hair)	3399
(Elemental Composition of Hair: Screening )	5940
( ) (Essential Vital Elements, Toxic Trace Elements, Urine)	2860
(Toxic Trace Elements, Nails)	1782
(Toxic Trace Elements, Essential Vital Elements, Nails)	3399
(Elemental Composition of Nails: Screening )	5940
	26455
	26455
	26455
	26455
	26455
	26455
	36300
(Testing for Kindergarten and School)	1738
: 0 14 (Healthy Child: for Children from 0 to 14 Years)	726
: (Survey of Kidneys: Screening)	957
	979
	20350
	19140
	19140
(Panel Chronic myelogenous leukemia, CML)	13090

	36300
	12210
	3465
	2431
	2376
	2376
	2475
	3179
	4103
	4345
	4367
	5082
	7018
	1837
	1859
	4367
	1419
	1859
	1859
	1936
	1111
	3916
	3091
	913
	2904
	3003
	1342
	2167
	6226
	1980
	3927
	4774
	2717
	3553
	2959
	8965
	2904
	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
(Preventing Heart and Blood	3102
Vessel Diseases )	
Diagnosis of Urogenital Tract Infection (UTI)) (Pregnancy Planning:	2486
: (Diabetes Control: Screening)	616
: (Survey Before Diet: Minimum )	1716
( ) (Weight	3223
Problems: Primary Survey)	
(Healthy skin beauty)	1386
:	649
HOMA-IR (Insulin Resistance: Fasting Glucose/Insulin, Homeostasis Model Assessment of Insulin Resistance, HOMA-IR)	737
)" : (	
Problems: Metabolic Syndrome (Primary Identification, creening) (Weight	946
(Diagnosis of Osteoporosis)	3047
: (Women's Oncorisk: Cervix )	1980
, , (Allergy to Animals, Dust, Mold)	6358
:	1650
, , (Strong hair and nails, velvet skin)	3465
: (Trace Elements, Serum,	3377
Venous Blood: Screening)	
(Mold Allergy)	2211
(Plant Allergy )	4818
(Immunological Survey, Screening)	7040
( AR, CFTR; AZF- ; ) (Male Sterility (Genes AR, CFTR; AZF-Region; Karyotype))	23496
( F2, F5, MTHFR, MTRR, MTR, ACE, AGT, RHD, HLA II; ) (Female Infertility, Pregnancy Complication (Genes F2, F5, MTHFR, MTRR, MTR, ACE, AGT, RHD, HLA II; Karyotype))	28534
( e ) (Toxic Trace Elements, Toxic	1375
Heavy Metals, Venous Blood)	
- , ( ,	2365
)	
( , , , , , )	2244
	176

CKD-EPI – (Estimated Glomerular Filtration Rate, eGFR, CKD-EPI Creatinine Equation)		176
B03.016.003 ( )		462
: II (14-28 ) (Pregnancy: Second Trimester (14-28 Weeks))		836
: ( « » ) (Clinical Blood Analysis: General Blood Analysis, Leucocyte Formula, ESR (with Manual Microscopic Examination of Blood Smear))		649
(Essential Vital Elements, Essential Trace Elements, Serum)		1375
, 6 ( AZF) (Spermatogenesis disorders (6 AZF))		5071
: (AZF- ) (Impairment of Spermatogenesis: Full Panel (AZF-Region))		11407
: IgG ( , ) ( )		5830
(DQA1, DQB1) HLA II		6864
- (RH factor Genotype)		10901
(skin) ( ) (Parasitic Fungi, Microscopy and Culture)		1727
(nails) ( ) (Parasitic Fungi, Microscopy and Culture)		1727
I II ( I/ II) (Pepsinogen I/Pepsinogen II, PG1/PG2)		2002
: sFlt-1, PlGF, sFlt-1/PlGF		6468
(Mycoplasma hominis Culture, Ureaplasma spp. Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*		1485
: (Male oncologic risk: prostate)		957
A09.05.120.001 - (Aldosterone-Renin Ratio, ARR)		1408
3, 4 (Complement components C3, C4)		792
" , IgE, ImmunoCAP		2332
B03.016.005 : (Lipid Profile: Screening)		770
( ITGB3) (Platelet Fibrinogen Receptor (Gene ITGB3))		1529
: (Lipid Profile: Screening)		770
Filtration Rate, eGFR, CKD-EPI ystatin C Equation) CKD-EPI – (Estimated Glomerular Filtration Rate, eGFR, CKD-EPI Creatinine Equation)		792
TREC KREC		
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