

/ 05. ()			
05.00.001		, (16)	18600
05.00.002		()	4000
/ 06. / ()			
06.00.001		(19)	4000
06.00.002		/ (99,9 %) (20)	9900
06.00.003		/ / (99,9 %) (19)30	16000
06.00.004		/ (99,9 %) (20)30	9900
/ 07.			
07.00.001		(/ /) (99,9%) (16)30	19350
07.00.002		(/) (50-99,9%) (28)	27870
07.00.003		(/) (17 Y-)	15500
07.00.004		(/) (50-99,9%) (28)	27870
07.00.005		(/) (17 Y-)	14870
07.00.006		(/) (17 Y-)	14870
/ 08. -			
08.00.001			8000
08.00.002		ATP7B (1 .)	36500
/ 09.			
09.00.001		(4-5), (5), (2), (2), ,	5000
/ 10.			
10.00.001	19.03.002	10	4000
10.00.002	19.03.002	1	450
10.00.003	19.04.001.003	" " 1	660
10.00.004	19.04.001.003	" " 10	6600
10.00.005	19.04.001.003	" " 20	11000
10.00.006	19.04.001.003	" " 10	6600
10.00.007	19.04.001.003	" " 20	11000
10.00.008	19.04.001.003	" " 10	6600
10.00.009	19.04.001.003	" " 20	11000
10.00.010	19.04.001.003	"HUR" 10 c	2000
10.00.011	19.23.003	(,) SP 1000	200
10.00.012	19.004.001.023	"HUR" (60)	200
10.00.014	19.03.002	3 (8)-(60 -1)	1600
10.00.015	19.03.002	(" 12 ")-(60 -1)	4800
10.00.016	19.03.002	(" 8 ")-(60 -1)	3200
10.00.017	19.03.002	55 8 5 ()-	900
10.00.018	19.30.006	-Trainer (30)	300
10.00.019	19.30.006	-Trainer (60)	550
10.00.020		()	400
10.00.021	19.23.005		300
/ 11.			
11.00.025		- , 1,0 -	250
11.00.026		- , -	1000
11.00.027	21.01.001		2200

11.00.028		"VIBRAMATIC" ()	450
11.00.029		"VIBRAMATIC" ()	450
11.00.030		"VIBRAMATIC"	250
11.00.031		"VIBRAMATIC" (, ,)	800
11.00.032	21.01.001	(1 -3)-	500
11.00.033	21.01.001	(3 -7)	700
11.00.034	21.01.001	(7 -12)	900
11.00.044	17.001.008		350
11.00.035	21.01.003	1	250
11.00.036	21.01.003.001	1,5	375
11.00.037		2	500
11.00.038		2,5	625
11.00.039		3	750
11.00.040		3,5	875
11.00.041		4	1000
11.00.042		4,5	1125
11.00.043		5	1250
/ 12.			
12.00.032	11.12.003.001	(500)	430
12.00.033	11.12.003.001	(1000)	570
12.00.038	11.12.003.001	(10,0)	630
12.00.039	11.12.003	(5,0)	380
12.00.040	11.12.003	(10,0)	550
12.00.041	11.12.003.001	(4)	480
12.00.034	11.04.004	(1,0)	1200
12.00.035	11.04.004	(2,0)	1400
12.00.036	11.01.002	(-2,2 + -2,2 + -2,2)	1500
12.00.037	11.01.002	(-2, 2+ -2,2 + -2,2)	1500
12.00.051	A11.12.003.001	(1)	610
12.00.042		(1,0)	393
12.00.043		(3000)	775
12.00.044		(5,0)	500
12.00.045		(10,0)	780
12.00.046		(5,0)	280
12.00.047		(5,0)	410
12.00.048		(10,0,0)	340
12.00.052		- , , 1,5	970
12.00.053		(20,0)	1200
12.00.049		, , 2,0	650
12.00.050		(2,0)	240
12.00.001	11.04.006	(10) , ,	600
12.00.002	11.04.006	(10) , , c	970
12.00.003	11.04.004	(10)	950
12.00.004	11.04.004	(10)	1300
12.00.005	11.01.002	(1)-(10)	770
12.00.006	11.01.002	(2)-(10)	970
12.00.007	11.01.002	-(10)	700
12.00.008	11.12.009	(10)	150
12.00.009	11.12.003	(10)	220
12.00.010	11.02.002	(10)	165

12.00.011	11.02.002	(400)-(10)	415
12.00.012	11.02.002	(2,0)-(10)	230
12.00.013	11.12.003.001	(40)	350
12.00.014	11.12.003.001	(2)-(40)	420
12.00.015	11.12.003.001	(100)-(40)	480
12.00.016	11.12.003.001	(200)-(40)	590
12.00.017	11.12.003.001	(400)-(60)	630
12.00.018	11.02.002	2,0 (10)	677
12.00.019	11.02.002	(10)	380
12.00.020	11.02.002	(2,0)-(10)	415
12.00.021	11.02.002	(2,0)-(10)	220
12.00.022	11.02.002	(1,0)-(10)	309
12.00.023	11.02.002	(5,0)-(10)	215
12.00.024	11.02.002	(2,0)-(10)	200
12.00.025	11.12.003.001	(50)-(40)	570
12.00.026	11.12.003.001	(10,0)-(40)	700
12.00.027	11.12.003.001	(10)-(40)	500
12.00.028	11.02.002	(1,5)-(10)	405
12.00.029	11.02.002	(8)-(10)	315
12.00.030	11.02.002	(2,0)-(10)	190
12.00.031	11.12.003.001	(400)	695
/ 13.			
13.00.001			535
13.00.002		3-	670
13.00.003		2-	575
13.00.004		2	600
13.00.005		2	515
13.00.006			365
13.00.007		2	515
13.00.008			535
13.00.009	1	() 2-	535
13.00.010		2-	515
13.00.011			385
13.00.012		-	530
13.00.013		2	515
13.00.014		2	515
13.00.015		2-	535
13.00.016		2	555
13.00.017		2-	600
13.00.018		()	385
13.00.019		1, 2	460
13.00.020			535
13.00.021			385
13.00.022		2	535
13.00.023		2-	575
13.00.024		2- +	940
/ 14.			
14.00.001	01.041.001	+	1000
14.00.002	17.01.002	- -	700
14.00.003	17.01.002	-	500
14.00.004	21.23.002		500
14.00.005	17.01.001		500

14.00.006	01.041.001	(30)	700
/ 15.			
15.00.054	17.30.040	- (15) 1	330
15.00.055	17.30.040	- (20) 2	380
15.00.056	17.30.040	- (25) 3	430
15.00.053	17.30.040	- (20)	780
15.00.045	17.30.014	-	300
15.00.046	17.30.014	-	400
15.00.047	17.23.014	- +	500
15.00.048	17.30.009	() (15)	350
15.00.049	17.30.009	() (20)	400
15.00.050	17.30.009	() (30)	450
15.00.051	17.30.009	1 (15)	300
15.00.052	17.30.009	1 (30)	350
15.00.057	A17.02.001	" " 1 (20)	300
15.00.058	A17.02.001	" " 2	350
15.00.059	A17.02.001	" " 3	400
15.00.060	A17.02.001	" " 1 (10)	250
15.00.001	17.30.004	- () " " 1	300
15.00.002	17.30.004	- () " " 2	350
15.00.003	17.30.024.002	" -1"	350
15.00.004	17.30.024.002	" -1 "	350
15.00.005	17.30.003	" -1	330
15.00.006	17.30.005	" "	500
15.00.007	17.30.008	- " " 1	450
15.00.008	17.30.008	- " " 2	500
15.00.009	22.01.005	1 " "	370
15.00.010	22.01.005	2 " "	400
15.00.011	22.01.005	3 " "	550
15.00.012	17.01.009	" "	400
15.00.013	17.30.031	" "	165
15.00.014	17.30.031	" " 1	300
15.00.015	17.30.031	" " 2	350
15.00.016	17.30.031	" "	330
15.00.017	17.30.018	" "	220
15.00.018	17.30.018	" "	330
15.00.019	17.30.044	-1,01	550
15.00.020	17.30.044	-1,01 1	400
15.00.021	17.30.044	-1,01 2	450
15.00.022	17.02.001	" "	330
15.00.023	17.02.001	" "	400
15.00.024	17.29.003	" -1"	350
15.00.025	17.30.024.001	" -1 "	220
15.00.026	17.30.024.002	" -1 "	400
15.00.027	17.29.003	" " -1"	440
15.00.028	17.29.003	" -1"	440
15.00.029	17.01.007	" "	250
15.00.030	17.29.003	" -1"	468
15.00.031	22.01.005	1 " "	370
15.00.032	22.01.005	2 " "	400

15.00.033	22.01.005	3	" "	550
15.00.034	17.30.034		-1,01 ()	230
15.00.035	17.30.034		-1,01	270
15.00.036	17.30.024.002	- ()	" "	380
15.00.037	17.30.034		-1,01 -()	330
15.00.038	17.30.034	3000)	-1,01 (1 -	730
15.00.039	17.30.024		" -1" (1 -3000)	780
15.00.040	17.30.027		" " (1 -3000)	870
15.00.041	17.30.034		-1,01	260
15.00.042	17.30.034		-1,01	205
15.00.043	17.30.034		-1,01	205
15.00.044	17.30.034		-1,01	300
16. / 16.01.				
16.01.001	01.023.001			1300
16.01.002	01.023.002			950
16.01.003	01.023.001			1000
16.01.004	01.023.002			650
16.01.005	01.023.001			500
16. / 16.02.				
16.02.001	01.050.001			1300
16.02.002	01.050.001			2000
16.02.003	01.050.002			950
16. / 16.03.				
16.03.001	01.015.001			1300
16.03.002	01.015.002			950
16. / 16.04.				
16.04.001	01.040.001			1300
16.04.002	01.040.002			950
16. / 16.05.				
16.05.001	01.047.001			1300
16.05.002	01.047.002			950
16. / 16.06.				
16.06.001	01.004.001			1300
16.06.002	01.004.002			950
16. / 16.07.				
16.07.001	01.005.001			1300
16.07.002	01.005.002			950
16. / 16.08.				
16.08.001	01.020.001		+	1100
16.08.002	01.020.005			500
16.08.003	03.020.001			600
16.08.004	03.020.001			450
16. / 16.09.				
16.09.001				1300
16.09.002	01.026.002			950
16.09.003	01.070.010			600
16.09.004	01.070.009			1200
16.09.005	01.054.001			500
16.09.006	01.054.001			300
16. / 16.02.				
16.02.004				1300
16. / 16.01.				
16.01.006				1500

16.01.007			1000
16. / 16.09.			
16.09.007	01.026.001	(40)	800
16. / 16.03.			
16.03.003	01.026.001		500
16. / 16.04.			
16.04.003	01.026.001		500
16. / 16.05.			
16.05.003	01.026.001		500
16. / 16.06.			
16.06.003	01.026.001		500
16. / 16.08.			
16.08.005	01.020.001		800
16. / 16.09.			
16.09.008	05.069.007		800
16. / 16.10.			
16.10.001			800
16.10.002			1800
/ 17.			
17.00.003	03.070.004	(2)	2400
17.00.004	04.070.001	(30)	900
17.00.005	04.070.001	(60)	1100
17.00.006	03.070.004	(2)	2400
17.00.001	01.070.009		1300
17.00.002	01.070.010		800
21. (General Clinical Blood Tests) / 21.01.			
(General Clinical Blood Tests)			
21.01.001		: , , () (Clinical Blood Analysis: General Blood Analysis, Leucocyte Formula, ESR (with Microscopic Examination of Blood Smear if Presence of	250
21.01.002		: , , () (Clinical Blood Analysis: General Blood Analysis, Leucocyte Formula, ESR (with Manual Microscopic Examination of Blood Smear))	300
21.01.003		() () (General Blood Analysis, without White Blood Cell (WBC) Count and ESR)	70
21.01.004		() (Leucocyte Formula (Differential White Blood Cell Count) with Microscopic Examination of Blood Smear if Presence o	70
21.01.005		() (Leucocyte Formula (Differential White Blood Cell Count) with Manual Microscopic Examination of Blood Smear)*	150
21.01.006		(Reticulocytes)	170
21.01.007		() (Erythrocyte Sedimentation Rate, ESR)	100
22. (Immunoematology) / 22.01. (Immunoematology)			
22.01.001		(Blood Group, 0)	170
22.01.002		- (-) (Rh-factor, Rh)	170
22.01.003		Rh (C, E, c, e) Kell- (Rh C (E, c, e) Kell-Phenotyping)	450
22.01.004		, Rh- (Anti Rh)	400
23. (Assessment of Coagulation System) / 23.01.			
(Assessment of Coagulation System)			
23.01.001		() () () (Activated Partial Thromboplastin Time, APTT)	110

24.03.006		(Urine Protein Electrophoresis)	1700
24.03.007		(Bence-Jones Protein, Urine, Immunofixation, Quantification)	2000
24.03.008		(Bence-Jones Protein, Urine, Electrophoresis, Immunofixation, Kappa/Lambda Typing, Quantification)	2800
24.03.009		(Homocysteine)	650
24. (Blood Chemistry Tests) / 24.04. (Assessment of renal function)			
24.04.001		(Creatinine)	70
24.04.002		CKD-EPI – (Estimated Glomerular Filtration Rate, eGFR, CKD-EPI Creatinine Equation)	200
24.04.003		(Cystatin C)	700
24.04.004		CKD-EPI – (Estimated Glomerular Filtration Rate, eGFR, CKD-EPI Cystatin C Equation)	750
24.04.005		(Urea)	70
24.04.006		(Uric acid)	70
24. (Blood Chemistry Tests) / 24.05. (Pigments)			
24.05.001		(Bilirubin Total)	70
24.05.002		(Direct Bilirubin, DBIL, Conjugated Bilirubin)	70
24.05.003		(Indirect Bilirubin, IB, Free Blood Bilirubin)*	0
24. (Blood Chemistry Tests) / 24.06. (Enzymes)			
24.06.001		(Alanine Aminotransferase, ALT, Serum Glutamic Pyruvic Transaminase, SGPT)	70
24.06.002		(Aspartateaminotransferase, AST, Serum Glutamicoxaloacetic Transaminase, SGOT)	70
24.06.003		(?-Amylase, ?-Amylase)	80
24.06.004		(P-?-Amylase) (Pancreatic)	200
24.06.005		(Gamma-Glutamyl Transferase, GGT)	70
24.06.006		(Creatine Kinase, CK, Creatine Phosphokinase, CPK)	200
24.06.007		(Creatine Kinase-MB, CK-MB, Creatine Phosphokinase-MB, CPK-MB.)	250
24.06.008		(Lipase)	220
24.06.009		(Lactate Dehydrogenase, LDH)	120
24.06.010		(Lactate Dehydrogenase, Isoforms 1, Lactic Acid Dehydrogenase, LDH1, Alpha-Hydroxybutyrate Dehydrogenase, Alpha-HBDH)	200
24.06.011		(S-Cholinesterase, Pseudocholinesterase, PCHE)	200
24.06.012		(Acid Phosphatase, ACP)	180
24.06.013		(Alkaline Phosphatase, ALP)	70
24. (Blood Chemistry Tests) / 24.07. (Assessment of endocrine and secretory functions of gastrointestinal tract (GI Tract))			
24.07.001		(Gastrin)	600
24.07.002		I (Pepsinogen I)	400
24.07.003		II (Pepsinogen II)	400
24.07.004		(I/II) (Pepsinogen I/Pepsinogen II, PG1/PG2)	700
24.07.005		(GastroPanel)	3500

24.07.006		-17 () (GastroPanel, Helicobacter pylori IgG, Gastrin-17, G-17)*	1350
24. (Blood Chemistry Tests) / 24.08. (Bone Metabolism)			
24.08.001		(Gla) (steocalcin, N-Osteocalcin, Bone Gla Protein, BGP)	700
24.08.002		25-OH D (25-OH Vitamin D Total, 25(OH)D, 25-Hydroxycalciferol)	1700
24.08.003		12 (, Active-B12, Holotranscobalamin)	700
24.08.004		- I (-CrossLaps, - ,) (Carboxyterminal Cross-linking Telopeptide of Bone Collagen, Collagen Cross-linked C-Telopeptide, Beta-Cross Laps, ?-CrossLaps Serum, C-Telopeptide, Crosslaps, Type 1	850
24.08.005		N-1 (Procollagen Type 1 N-terminal Propeptide, P1NP, Total)	1000
24.08.006		() (Deoxypyridinoline, DPD, Urine)	900
24. (Blood Chemistry Tests) / 24.10. (Vitamins)			
24.10.001		12 (,) (Cobalamin)	500
24.10.002		(Folic Acid)	500
24.10.003		25-OH D (25-OH Vitamin D Total, 25(OH)D, 25-Hydroxycalciferol)	1700
24. (Blood Chemistry Tests) / 24.11. (Inorganic Matters)			
24.11.001		/ / (+/Potassium, Na+/Sodium, l-/Chloride, Serum)	220
24.11.002		(Ca) (Calcium Total)	70
24.11.003		(Ca ²⁺ , c) (Ionized Calcium, Free Calcium)	270
24.11.004		(g) (Magnesium (Mg), Serum)	70
24.11.005		(P) (Phosphorus (P))	70
24.11.006		(Fe) (Iron (Fe), Serum)	70
24.11.007		(,) (Unsaturated Iron Binding Capacity, UIBC)	90
24.11.008		() (Total Iron Binding Capacity, TIBC)*	0
24. (Blood Chemistry Tests) / 24.12. (Specific proteins)			
24.12.001		- (- ,) (Antistreptolysin-O, ASO)	250
24.12.002		- () (C-Reactive Protein, CRP)	150
24.12.003		() (Rheumatoid Factor, RF)	150
24.12.004		(Ceruloplasmin)	500
24.12.005		(Haptoglobin)	500
24.12.006		-2- (Alpha-2-Macroglobulin, ?2-Macroglobulin, A2M)	700
24.12.007		-1- (1), (Alpha-1-Antitrypsin, A1AT, AAT, Concentration)	1400
24.12.008		-1- (1), (Alpha-1-Antitrypsin, A1AT, AAT, Phenotyping)	2500
24.12.009		() (Transferrin)	250
24.12.010		(Ferritin)	550
24.12.011		(Myoglobin)	500
24.12.012		(-) N- (NT-proBNP, N-Terminal Pro-brain Natriuretic Peptide, Pro-B-Type Natriuretic Peptide)	2500
24.12.013		-I (Troponin-I)	500
24.12.014		- () (Carbohydrate-Deficient Transferrin, CDT)	2500
24. (Blood Chemistry Tests) / 24.13. (Tumor Markers)			
24.13.001		() (Prostate-Specific Antigen Total, PSA Total)	300
24.13.002		: (Male oncologic risk: prostate)	650

24.13.003		(, , -2proPSA, phi)	5200
24.13.004		(Calcitonin)	700
24.13.005		- () (?-Fetoprotein, AFP)	250
24.13.006		- (,) (Carcinoembryonic Antigen, CEA)	350
24.13.007		-15-3 (15-3) (Carbohydrate Antigen -15-3, Cancer Antigen -15-3)	450
24.13.008		CA-72-4 (72-4) (Carbohydrate Antigen -72-4, Cancer Antigen CA-72-4)	800
24.13.009		-19-9 (19-9) (Carbohydrate Antigen -19-9, Cancer Antigen-GI)	420
24.13.010		(Cyfra 21-1, 19) (Cytokeratin 19 Fragments, C-terminus of Cytokeratin 19, CK19 Soluble Fragments, Cyfra 21-1)	800
24.13.011		-125 (125) (Carbohydrate Antigen -125, Cancer Antigen -125)	400
24.13.012		-125 + (Cancer Antigen -125 + CEA)	700
24.13.013		HE4 (4) (Human Epididymis Protein 4, HE4)	1200
24.13.014		ROMA (Risk of Ovarian Malignancy Algorithm,) (Risk of Ovarian Malignancy Algorithm, ROMA (Before Menopause))	1700
24.13.015		ROMA (Risk of Ovarian Malignancy Algorithm,) (Risk of Ovarian Malignancy Algorithm, ROMA (After Menopause))	1700
24.13.016		CA-242 (242, CA-242) (Carbohydrate Antigen -242, Tumor Marker CA-242)	1000
24.13.017		-2- (?-2-) (eta-2-Microglobulin, BMG, Serum)	800
24.13.018		() (Neuron-Specific Enolase, NSE)	1100
24.13.019		(Chromogranin A, CgA)	4500
24.13.020		S100 (S100 rotein)	2000
24.13.021		SCC () (Squamous Cell Carcinoma Antigen, SCCA, SCCAg)	2100
24.13.022		UBC (8 18 ,) (Urine Bladder Cancer Antigen, Urine Bladder Cancer, UBC)	2000
24.13.023		- , . , - () (M-Gradient, Screening. Serum Protein Electrophoresis (SPEP), Immunofixation with Polyvalent Antiserum, Quan	2000
24.13.024		- , . (IgG, IgA, IgM, , ,), - (M-Gradient, Typing. Serum Protein Electrophoresis (SPEP), Immunofixation with Antisera	3200
24. (Blood Chemistry Tests) / 24.10. (Vitamins)			
24.10.004		1 (, Vitamin K1, Phylloquinone, Serum)	1850
25. (Assessment of endocrine system function) / 25.01. (Assessment of Pituitary Function)			
25.01.001		(,) (Adrenocorticotropic Hormone, ACTH)	210
25.01.002		(,) (Thyroid Stimulating Hormone, TSH)	190
25.01.003		(,) (Growth Hormone, GH)	300
25.01.004		(1) (Somatomedin C, Insulin-like Growth Factor 1, IGF-1)	380
25.01.005		() (Follicle Stimulating Hormone, FSH)	200
25.01.006		() (Luteinizing Hormone, LH)	200
25.01.007		(Prolactin)	200

25. (Assessment of endocrine system function) / 25.02. (Assessment of Thyroid Function)			
25.02.001		(,) (Thyroid Stimulating Hormone, TSH)	220
25.02.002		(T4 ,) (Total Thyroxine, TT4)	220
25.02.003		(4) (Free Thyroxine, FT4)	220
25.02.004		(3) (Total Triiodthyronine, TT3)	220
25.02.005		(3) (Free Triiodthyronine, FT3)	220
25.02.006		(; ;) (Thyroid Uptake, T-Uptake, Thyroxine-Binding Capacity, TBC, Thyroxine-Binding Index, TBI, free T4 Index, fT4I)	450
25.02.007		() (Thyroglobulin, TG)	280
25.02.008		(-) (Anti- thyroglobulin Autoantibodies, Thyroglobulin Antibodies, Tg Autoantibodies, TgAb, Anti-Tg Ab, ATG)	280
25.02.009		(- ,) (Anti- thyroid peroxidase autoantibodies, Antimicrosomal Antibodies, TPO Antibodies, TPOAb, Anti-TPO)	280
25.02.010		() (Anti-Thyroid Microsomal Antibodies)	600
25.02.011		(, -) (Thyroid-Stimulating Hormone Receptor Antibodies, TSH Receptor Antibodies, TSHRABs, TSH binding inhibitor immunoglobulin, TBII)	1100
25. (Assessment of endocrine system function) / 25.03. (Assessment of adrenocortical function)			
25.03.001		() (Cortisol, Hydrocortisone)	230
25.03.002		(Free cortisol, Free Hydrocortisone, Urine)	500
25.03.003		, (Cortisol, Saliva)	600
25.03.004		(Aldosterone)	500
25.03.005		(,) (Direct Renin, Plasma)	600
25.03.006		- (Aldosterone-Renin Ratio, ARR)	1500
25. (Assessment of endocrine system function) / 25.04. (Assessment of Androgen Status)			
25.04.001		(Testosterone)	280
25.04.002		(Free Testosterone)	450
25.04.003		() (Dihydrotestosterone, DHT)	1000
25.04.004		(Androstenedione)	900
25.04.005		() (Androstanediol Glucuronide, 3?-Androstanediol Glucuronid, 3?-diol G)	1000
25.04.006		- (-S04,) (Dehydroepiandrosterone Sulfate, DHEA-S)	280
25.04.007		17- (17-) (17-Ketosteroids, Urine)	800
25.04.008		17- - (17-Hydroxyprogesterone, 17-OHP)	280
25.04.009		, () (Sex Hormone-Binding Globulin, SHBG)	280
25. (Assessment of endocrine system function) / 25.05. (Estrogens and Progestins)			
25.05.001		(2) (Estradiol, E2)	280
25.05.002		(Progesterone)	280
25. (Assessment of endocrine system function) / 25.06. (Nonsteroidal Gonadal Factors)			
25.06.001		() (Anti-Mullerian Hormone, AMH, Mullerian Inhibiting Substance, MIS)	1000
25.06.002		(Inhibin B)	1000
25.06.003		-1- () (Trophoblastic beta-1-Globulin, TBG)	400
25. (Assessment of endocrine system function) / 25.07. (Monitoring of Pregnancy, Biochemical Markers of Fetal Status)			

25.07.001		(, - , ?-) (Human Chorionic Gonadotropin, HCG)	280
25.07.002		?- (?-) (Free Human Chorionic Gonadotropin, Free HCG)	400
25.07.003		() (Placental Lactogen, PL, Human Placental Lactogen, hPL, Chorionic Somatomammotropin, CS, Human Chorionic Somatomammotropin, hCS)	600
25.07.004		- (Pregnancy-Associated Plasma Protein-A, PAPP-A)	400
25.07.005		(Estriol Free, 3)	390
25.07.006		- () (?-Fetoprotein, AFP)	280
25.07.007		: I (PRIS A-1) (Maternal Screen, First Trimester; Prenatal Screening I; PRIS A I (Prenatal Risk Calculation))	600
25.07.008		: II (PRIS A-2) (Maternal Screen, Second Trimester; Prenatal Screening II; PRISCA II (Prenatal Risk Calculation))	600
25. (Assessment of endocrine system function) / 25.08. (Assessment of Pancreatic Endocrine Function)			
25.08.001		(Insulin)	350
25.08.002		(Proinsulin)	600
25.08.003		- (C-Peptide)	300
25.08.004		: (), (), HOMA-IR (Insulin Resistance: Fasting Glucose/Insulin, Homeostasis Model Assessment of Insulin Resistance, HOMA-IR)	500
25. (Assessment of endocrine system function) / 25.09. (Biogenic Amines, BA)			
25.09.001		(, ,) - (Catecholamines: Epinephrine/Adrenaline, Norepinephrine/Noradrenaline, Dopamine, Plasma)	1700
25.09.002		(, ,) (Catecholamines: Epinephrine/Adrenaline, Norepinephrine/Noradrenaline, Dopamine, Urine)	1700
25.09.003		: (), (), 5- (5-) (Catecholamines and Serotonin Metabolites, 24 Hours-Urine: Vanillylmandelic Acid, V , Homovanillic Acid, V , 5-Hy	2200
25.09.004		() : (Metanephrines, Fractinated, 24 Hours-Urine – Metanephrine, Normetanephrine)	1600
25.09.005		(, ,) (Catecholamines: Epinephrine/Adrenaline, Norepinephrine/Noradrenaline, Dopamine, Urine)	1700
25.09.006		(Histamine, Plasma)	2500
25.09.007		(Serotonin, Serum)	2400
25.09.008		(Nephrines, Plasma)	2000
25. (Assessment of endocrine system function) / 25.10. (Calcium-regulating hormones)			
25.10.001		(Calcitonin)	650
25.10.002		(, ,) (Parathyroid Hormone, PTH)	300
25. (Assessment of endocrine system function) / 25.11. (Adipose Tissue Hormones)			
25.11.001		(Leptin)	700
25. (Assessment of endocrine system function) / 25.12. (Regulation of Erythropoiesis)			
25.12.001		(Erythropoetin)	900
26. (Microelements) / 26.01. (Microelements, Serum, Venous Blood)			
26.01.001		() (Essential Vital Elements, Essential Trace Elements, Serum)	1500
26.01.002		(e) (Toxic Trace Elements, Toxic Heavy Metals, Venous Blood)	1500
26.01.003		: (Trace Elements, Serum, Venous Blood: Screening)	3300
26.01.004		(Cd) (Cadmium (Cd), Serum)	850
26.01.005		(Co) (Cobalt (Co), Serum)	850

26.01.006	(Cu)	(Copper (Cu), Serum)	850
26.01.007	(Mn)	(Manganese (Mn), Serum)	850
26.01.008	(Se)	(Selenium (Se), Serum)	850
26.01.009	(Zn)	(Zinc (Zn), Serum)	850
26.01.010	(Ni)	(Nickel (Ni), Serum)	850
26.01.011	(Au)	(Gold (Au), Serum)	850
26.01.012	(Mo)	(Molybdenum (Mo), Serum)	850
26.01.013	(Tl)	(Thallium (Tl), Serum)	850
26.01.014	(As)	(Arsenic (As), Serum)	850
26.01.015		(Iodine, serum)	850
26.01.016	(Li)	(Lithium (Li), serum)	900
26.01.017	(Cd)	(Cadmium (Cd), lood)	850
26.01.018	(Co)	(Cobalt (Co), lood)	850
26.01.019	(Cu)	(Copper (Cu), lood)	850
26.01.020	(Mn)	(Manganese (Mn), lood)	850
26.01.021	(Ni)	(Nickel (Ni), lood)	850
26.01.022	(Se)	(Selenium (Se), lood)	850
26.01.023	(Zn)	(Zinc (Zn), lood)	850
26.01.024	(Pb)	(Lead (Pb), lood)	850
26.01.025	(Hg)	(Mercury (Hg), lood)	850
26. (Microelements) / 26.02. (Microelements, Urine)			
26.02.001		(Essential Vital Elements, Toxic Trace Elements, Urine)	2600
26.02.002	(Cd)	(Cadmium (Cd), Urine)	850
26.02.003	(Co)	(Cobalt (Co), Urine)	850
26.02.004		(Copper, random urine; Cu)	850
26.02.005	(Cu)	(Copper (Cu), 24-Hours Urine)	850
26.02.006	(Mn)	(Manganese (Mn), Urine)	850
26.02.007	(Se)	(Selenium (Se), Urine)	850
26.02.008	(Zn)	(Zinc (Zn), Urine)	850
26.02.009	(Ni)	(Nickel (Ni), Urine)	850
26.02.010	(Pb)	(Lead (Pb), Urine)	850
26.02.011	(Hg)	(Mercury (Hg), Urine)	850
26.02.012	(Tl)	(Thallium (Tl), Urine)	850
26.02.013	(As)	(Arsenic (As), Urine)	850
26.02.014	(Al)	(Aluminum (Al), Urine)	850
26.02.015	(Fe)	(Iron (Fe), Urine)	850
26.02.016	(I)	(Iodine (I), Urine)	850
26. (Microelements) / 26.03. (Microelements, Hair)			
26.03.001		(Toxic Trace Elements, Hair)	2800
26.03.002		(Toxic Trace Elements, Essential Vital Elements, Hair)	3000
26.03.003		(Elemental Composition of Hair: Screening)	5500
26.03.004	(Al)	(Aluminum (Al), air)	850
26.03.005	(Ba)	(Barium (Ba), air)	850
26.03.006	(Be)	(Beryllium (Be), air)	850
26.03.007	(B)	(Boron (B), air)	850
26.03.008	(V)	(Vanadium (V), air)	850
26.03.009	(Bi)	(Bismuth (Bi), air)	850
26.03.010	(W)	(Tungsten, Wolframium (W), air)	850
26.03.011	(Ga)	(Gallium (Ga), air)	850
26.03.012	(Ge)	(Germanium (Ge), air)	850
26.03.013	(Fe)	(Iron (Fe), air)	850
26.03.014	(Au)	(Gold (Au), air)	850
26.03.015	(I)	(Iodine (I), air)	850

26.03.016		(Cd) (Cadmium (Cd), air)	850
26.03.017		(K) (Potassium (K), air)	850
26.03.018		(Ca) (Calcium (Ca), air)	850
26.03.019		(Co) (Cobalt (Co), air)	850
26.03.020		(Si) (Silica (Si), air)	850
26.03.021		(La) (Lanthanum 9La), air)	850
26.03.022		(Li) (Lithium (Li), air)	850
26.03.023		(Mg) (Magnesium (Mg), air)	850
26.03.024		(Mn) (Manganese (Mn), air)	850
26.03.025		(Cu) (Copper (Cu), air)	850
26.03.026		(Mo) (Molybdenum (Mo), air)	850
26.03.027		(As) (Arsenic (As), air)	850
26.03.028		(Na) (Sodium (Na), air)	850
26.03.029		(Ni) (Nickel (Ni), air)	850
26.03.030		(Sn) (Tin (Sn), air)	850
26.03.031		(Pt) (Platinum (Pt), air)	850
26.03.032		(Hg) (Mercury (Hg), air)	850
26.03.033		(Rb) (Rubidium 9Rb), air)	850
26.03.034		(Pb) (Lead (Pb), air)	850
26.03.035		(Se) (Selenium (Se), air)	850
26.03.036		(Ag) (Silver (Ag), air)	850
26.03.037		(Sr) (Strontium (Sr), air)	850
26.03.038		(Sb) (Antimony (Sb), air)	850
26.03.039		(Tl) (Thallium (Tl), air)	850
26.03.040		(P) (Phosphorus (P), air)	850
26.03.041		(Cr) (Chromium (Cr), air)	850
26.03.042		(Zn) (Zinc (Zn), air)	850
26.03.043		(Zr) (Zirconium (Zr), air)	850
26.	(Microelements) / 26.04.	(Microelements, Nails)	
26.04.001		(Toxic Trace Elements, Nails)	1700
26.04.002		(Toxic Trace Elements, Essential Vital Elements, Nails)	3000
26.04.003		(Elemental Composition of Nails: Screening)	5000
26.04.004		(Al) (Aluminum (Al), Nails)	850
26.04.005		(Ba) (Barium (Ba), Nails)	850
26.04.006		(Be) (Beryllium (Be), Nails)	850
26.04.007		(B) (Boron (B), Nails)	850
26.04.008		(V) (Vanadium (V), Nails)	850
26.04.009		(Bi) (Bismuth (Bi), Nails)	850
26.04.010		(W) (Tungsten, Wolframium (W), Nails)	850
26.04.011		(Ga) (Gallium (Ga), Nails)	850
26.04.012		(Ge) (Germanium (Ge), Nails)	850
26.04.013		(Fe) (Iron (Fe), Nails)	850
26.04.014		(Au) (Gold (Au), Nails)	850
26.04.015		(I) (Iodine (I), Nails)	850
26.04.016		(Cd) (Cadmium (Cd), Nails)	850
26.04.017		(K) (Potassium (K), Nails)	850
26.04.018		(Ca) (Calcium (Ca), Nails)	850
26.04.019		(Co) (Cobalt (Co), Nails)	850
26.04.020		(Si) (Silica (Si), Nails)	850
26.04.021		(La) (Lanthanum (La), Nails)	850
26.04.022		(Li) (Lithium (Li), Nails)	850
26.04.023		(Mg) (Magnesium (Mg), Nails)	850
26.04.024		(Mn) (Manganese (Mn), Nails)	850

26.04.025	(Cu)	(Copper (Cu), Nails)	850
26.04.026	(Mo)	(Molybdenum (Mo), Nails)	850
26.04.027	(As)	(Arsenic (As), Nails)	850
26.04.028	(Na)	(Sodium (Na), Nails)	850
26.04.029	(Ni)	(Nickel (Ni), Nails)	850
26.04.030	(Sn)	(Tin (Sn), Nails)	850
26.04.031	(Pt)	(Platinum (Pt), Nails)	850
26.04.032	(Hg)	(Mercury (Hg), Nails)	850
26.04.033	(Rb)	(Rubidium (Rb), Nails)	850
26.04.034	(Pb)	(Lead (Pb), Nails)	850
26.04.035	(Se)	(Selenium (Se), Nails)	850
26.04.036	(Ag)	(Silver (Ag), Nails)	850
26.04.037	(Sr)	(Strontium (Sr), Nails)	850
26.04.038	(Sb)	(Antimony (Sb), Nails)	850
26.04.039	(Tl)	(Thallium (Tl), Nails)	850
26.04.040	(P)	(Phosphorus (P), Nails)	850
26.04.041	(Cr)	(Chromium (Cr), Nails)	850
26.04.042	(Zn)	(Zinc (Zn), Nails)	850
26.04.043	(Zr)	(Zirconium (Zr), Nails)	850
27. (Drug Monitoring) / 27.01. (Drug Monitoring)			
27.01.001	()	(Phenobarbitalum)	2100
27.01.002	(,)	(Phenytoin)	950
27.01.003	(,)	(Acidum Valproicum, Depakin, Convulexs)	700
27.01.004	(,)	(carbamazepine, Tegretol)	2100
27.01.005		(Cyclosporine, Cyclosporine A, Sandimmune)	900
27.01.006	(, ,)	(FK506, Advagraf, Prograf, Protopic, Tacrosel)	1400
27.01.007		(Lamotrigine)	3100
27.01.008		(Levetiracetam, Keppra®)	3200
27.01.009	(Li)	(Lithium (Li), serum)	1100
28. (Examinations of Urine) / 28.01. (Clinical Tests)			
28.01.001	()	(Complete Urinalysis, Microscopic Examination)	180
28.01.002		(Nechiporenko's Urine Test)	180
28.01.003	()	(Sulkowitch Urine Calcium Test)	200
28. (Examinations of Urine) / 28.02. () (Urine Biochemistry (Daily Excretion))			
28.02.001	(Albumin, Urine)		400
28.02.002	/ -)	(Urine Albumin Creatinine Ratio, UACR)	700
28.02.003	(Glucose, Urine)		100
28.02.004	(Creatinine, Urine)		100
28.02.005	()	(Glomerular Filtration Rate, GFR)*	100
28.02.006	(- ,)	(Amylase, 24-Hour or Timed Urine)	150
28.02.007	(Urea, Urine)		100
28.02.008	(Uric cid, Urine)		100
28.02.009	(Protein Total, Urine)		150
28.02.010	(Ca),	(Calcium (Ca), Urine)	100
28.02.011	(K), (Na)	(Potassium (K), Sodium (Na))	150
28.02.012	(P),	(Phosphorus (P), Urine)	100
28.02.013	(xalates, Urine)		1400

28.02.014		(Calcium-Creatinine Ratio, Random Urine)	400
28. (Examinations of Urine) / 28.03. (Psychoactive Substances, Urine)			
28.03.001		(Drugs and Psychotropic Substances Screening: Opiates, Amphetamines, Methamphetamine, Cocaine, Cannabinoids, Ca	3000
28.03.002		(Cannabinoids (Marijuana), Urine)*	1000
28.03.003		(Opiates (Morphine/Heroin), Urine)*	1000
28.03.004		(Barbiturates, Urine)*	1000
28.03.005		(Pernicious Habits: Nicotine, Drugs, Psychosti	3200
28. (Examinations of Urine) / 28.04. (Study of Kidney Stone)			
28.04.001		(Kidney Stone Analysis)	3000
29. (Examinations of Feces, Stool Examinations) / 29.01. (Examinations of Feces, Stool Examinations)			
29.01.001		(Koprogramma, Stool)	320
29.01.002		(PRO Stool, Helminth Eggs)	280
29.01.003		(PRO Stool, Helminth Eggs)	280
29.01.004		(PRO Stool)	280
29.01.005		(nterobiasis, Swab)	280
29.01.006		(Stool Sugars, Reducing Substances, Fecal)	490
29.01.007		(ccult Blood Feces nalysis)	200
29.01.008		(FOB Gold (Quantitative Immunochemical Fecal Occult Blood, Test FOB Gold)	600
29.01.009		1 (1), 1 (Elastase 1, E1)	1800
29.01.010		(Alpha-1-Antitrypsin, Feces)	2000
29.01.011		(Fecal Calprotectin)	2000
30. (Serology Diagnosis of Infectious Disease) / 30.01. (Adenovirus Infection)			
30.01.001		IgA (nti-Adenovirus IgA)	590
30.01.002		IgG (nti-Adenovirus IgG)	590
30. (Serology Diagnosis of Infectious Disease) / 30.02. (Ascariasis)			
30.02.001		IgG (Anti-Ascaris lumbricoides IgG)	600
30. (Serology Diagnosis of Infectious Disease) / 30.03. (Aspergillosis)			
30.03.001		Aspergillus fumigatus (3), - IgG (Aspergillus fumigatus, IgG, M3)	400
30. (Serology Diagnosis of Infectious Disease) / 30.04. (Bacterial flora (Bacteroides spp.))			
30.04.001		(Bacteroides spp., DNA, Scrape of Urogenital Epithelial Cells)*	300
30. (Serology Diagnosis of Infectious Disease) / 30.05. (Borreliosis, Lyme Borreliosis, Lyme Disease, LD)			
30.05.001		A IgG Borrelia burgdorferi (Anti-Borrelia burgdorferi IgG)	250
30.05.002		A IgM Borrelia burgdorferi (Anti-Borrelia burgdorferi IgM)	470
30.05.003		A IgG Borrelia burgdorferi, (Anti-Borrelia burgdorferi IgG, Immunoblot)	1250
30.05.004		A IgM Borrelia burgdorferi, (Anti-Borrelia burgdorferi IgM, Western Blot (WB))	1250
30.05.005		(orrelia burgdorferi, DNA)*	400
30.05.006		(orrelia burgdorferi, DNA, Synovial Fluid)*	500

30.05.007		(<i>Chlamydia trachomatis</i> , DNA, Cerebrospinal Fluid)*	500
30. (Serology Diagnosis of Infectious Disease) / 30.06. (Brucellosis)			
30.06.001		IgG (Anti-Varicella-Zoster Virus IgG, Anti-VZV IgG)	800
30.06.002		IgM (Anti-Varicella-Zoster Virus IgM, Anti-VZV IgM)	800
30. (Serology Diagnosis of Infectious Disease) / 30.07. (Human Immunodeficiency Virus, HIV)			
30.07.001		1 2 1 2 (HIV Ag/Ab Combo)	300
30.07.002		-1, (HIV RNA, Plasma)*	11000
30. (Serology Diagnosis of Infectious Disease) / 30.08. (Human papillomavirus, HPV, HPV Infection)			
30.08.001		16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68 + 14 (HPV DNA, Scrape of Rectal Epithelial Cells, 14 Types)	800
30.08.002		16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68 + 14 (HPV DNA, Scrape of Faucial Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68))	800
30.08.003		16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68 + 14 (HPV DNA, Scrape of Urogenital Epithelial Cells, 14 Types)	800
30.08.004		16 18 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 2 Types (16, 18))	500
30.08.005		14 : 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68 (HPV DNA, Scrape of Rectal Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68))	1100
30.08.006		14 : 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68 (HPV DNA, Scrape of Faucial Epithelial Cells, 14 Types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68))	1100
30.08.007		14 : 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 14 Types)	1100
30.08.008		: 6, 11, 44 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 3 Types (6, 11, 44))	500
30.08.009		: 6, 11, 44 + (HPV DNA, Scrape of Rectal Epithelial Cells, 3 Types (6, 11, 44))	500
30.08.010		: 6, 11, 44 (HPV DNA, Scrape of Faucial Epithelial Cells, 3 Types (6, 11, 44))	500
30.08.011		: 6, 11, 16, 18 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 4 Types (6, 11, 16, 18) Screening)	900
30.08.012		: 6, 11, 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 68 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 15 Types (6, 11, 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 68))	1000
30.08.013		: 6, 11, 16, 18, 26, 31, 33, 35, 39, 44, 45, 51, 52, 53, 56, 58, 59, 66, 68, 73, 82 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 21 Types (6, 11, 16, 18, 26, 31, 33, 35, 39, 44, 45, 51, 52, 53, 56, 58, 59, 66, 68, 73, 82))	2400
30. (Serology Diagnosis of Infectious Disease) / 30.09. (Gardnerella vaginalis)			
30.09.001		(<i>Gardnerella vaginalis</i> , DNA, Urine)*	250

30.09.002		(Gardnerella vaginalis, DNA, Prostatic Fluid, Semen)*	250
30.09.003		(Gardnerella vaginalis, DNA, Scrape of Urogenital Epithelial Cells)*	250
30. (Serology Diagnosis of Infectious Disease) / 30.10. (Helminthiasis)			
30.10.001		() (PRO Stool, Helminth Eggs)	300
30.10.002		« » (Parasites: Opisthorchis, Echinococcosis, Toxocara, Trichinella)	400
30. (Serology Diagnosis of Infectious Disease) / 30.11. (Hepatitis A Virus, HAV) A			
30.11.001		IgG (Anti-HAV IgG)	500
30.11.002		IgM (Anti-HAV IgM)	600
30.11.003		(HAV RNA, Serum)*	500
30. (Serology Diagnosis of Infectious Disease) / 30.12. (Hepatitis B Virus, HBV) B			
30.12.001		HBs- (HBs- B, « »), (HBsAg, Hepatitis Surface Antigen, Qualitative)	150
30.12.002		HB - (Hepatitis Be Antigen, HBeAg)	400
30.12.003		IgM IgG HB-core B, (Anti-HBc IgM, IgG, Antibodies to Hepatitis B Core Antigen; HBcAb, Total, HBV Core Total Antibodies (IgG + IgM))	230
30.12.004		IgM HB-core B (Anti-HBc IgM Antibodies to Hepatitis B Core Antigen; HBV Core Antibodies IgM)	300
30.12.005		HB - (Anti-HBe, HBeAb)	400
30.12.006		HBs- (Anti-HBs, HBsAb)	400
30.12.007		HBs- (HBs- B, « »), (HBsAg, Hepatitis Surface Antigen, Quantitative)	1300
30.12.008		B, (HBV DNA, Serum, Qualitative)*	400
30.12.009		B, (HBV DNA, Serum, Quantitative)*	1800
30. (Serology Diagnosis of Infectious Disease) / 30.13. (Hepatitis C Virus, HCV)			
30.13.001		IgM IgG (Anti-HCV Total (IgG + IgM))*	240
30.13.002		IgG C, (Anti-HCV IgG, Immunoblot)	3000
30.13.003		-28 (-28), () (Interleukin 28 Beta IL28B, Genotyping (Study of Genetic Markers Determining Effectiveness))	1100
30.13.004		IgM (Anti-HCV IgM)	180
30.13.005		IgG core, NS3, NS4, NS5 (Anti-HCV core, NS3, NS4, NS5 IgG)	400
30.13.006		C (: 73, 78, 75, 76, 74, 77, 79, 1170) (Markers of Hepatitis B, C)	1800
30.13.007		(HCV RNA, Serum, Qualitative)*	600
30.13.008		(1 (1a 1b), 2, 3) (Hepatitis C Virus (HCV) RNA, Plasma, Genotyping, Subtypes (Types 1 (Subtypes 1a, 1b), 2, 3))*	1100
30.13.009		(1, 2, 3) (Hepatitis C Virus (HCV) RNA, Quantitative PCR, Genotyping (Types 1, 2, 3))	1100
30.13.010		(CITO), (HCV RNA, Plasma, Quantitative)*	17000
30.13.011		(HCV RNA, Serum, Quantitative, PCR)*	2000

30.13.012		(), (Hepatitis C Virus (HCV) RNA, Ultrasensitive PCR)	3000
30. (Serology Diagnosis of Infectious Disease) / 30.14. D (Hepatitis D Virus, HDV)			
30.14.001		IgM D (Anti-HDV IgM)	900
30.14.002		D, (Anti-HDV Total)	170
30.14.003		D, (HDV RNA, Serum)*	600
30. (Serology Diagnosis of Infectious Disease) / 30.15. G (Hepatitis E Virus, HEV)			
30.15.001		IgM E (Anti-HEV IgM)	800
30.15.002		IgG E (Anti-HEV IgG)	800
30. (Serology Diagnosis of Infectious Disease) / 30.16. G (Hepatitis G Virus, HGV)			
30.16.001		G, (HGV RNA, Serum)*	600
30. (Serology Diagnosis of Infectious Disease) / 30.17. D (Herpes simplex virus, HSV-1, HSV-2)			
30.17.001		IgG 1 2 (Anti-HSV-1, 2 IgG)	260
30.17.002		IgG 1 (Anti-HSV-1 IgG)	800
30.17.003		IgG 2 (Anti-HSV-2 IgG)	600
30.17.004		Ig 1 2 (Anti-HSV-1, 2 Ig)	280
30.17.005		Ig 1 2 IgG, (Anti-HSV-1, 2 IgM, IgG, Avidity)	600
30.17.006		1 2 , (HSV-1, 2 DNA, Exudate)*	250
30.17.007		1 2 , (HSV-1, 2 DNA, Scrape of Conjunctiva Epithelial Cells)*	250
30.17.008		1 2 , (HSV-1, 2 DNA, Scrape of Skin Epithelial Cells)*	250
30.17.009		1 2 , (HSV-1, 2 DNA, Blood)*	250
30.17.010		1 2 , (HSV-1, 2 DNA, Urine)*	250
30.17.011		1 2 , (HSV-1, 2 DNA, Scrape of Nasal Epithelial Cells)*	250
30.17.012		1 2 , (HSV-1, 2 DNA, Scrape of Faucial Epithelial Cells)*	250
30.17.013		1 2 , (HSV-1, 2 DNA, Serum)*	400
30.17.014		1 2 , (HSV-1, 2 DNA, Saliva)*	250
30.17.015		1 2 , (HSV-1, 2 DNA, Cerebrospinal Fluid)*	250
30.17.016		1 2 , (HSV-1, 2 DNA, Prostatic Fluid, Semen)*	250
30.17.017		1 2 , (HSV-1, 2 DNA, Scrape of Urogenital Epithelial Cells)*	250
30.17.018		1 2 , (HSV-1, 2 DNA, Exudate, typing)*	450
30.17.019		1 2 , (HSV-1, 2 DNA, Scrape of Conjunctiva Epithelial Cells, typing)*	450
30.17.020		1 2 , (HSV-1, 2 DNA, Scrape of Skin Epithelial Cells, typing)*	450
30.17.021		1 2 , (HSV-1, 2 DNA, Blood, typing)*	600
30.17.022		1 2 , (HSV-1, 2 DNA, Urine, typing)*	500
30.17.023		1 2 , (HSV-1, 2 DNA, Scrape of Nasal Epithelial Cells, typing)*	450

30.17.024		1 2 , (HSV-1, 2 DNA, Scrape of Faucial Epithelial Cells, typing)*	450
30.17.025		1 2 , (HSV-1, 2 DNA, Serum, typing)*	600
30.17.026		1 2 , (HSV-1, 2 DNA, Saliva, typing)*	450
30.17.027		1 2 , (HSV-1, 2 DNA, Cerebrospinal Fluid, typing)*	450
30.17.028		1 2 , (HSV-1, 2 DNA, Prostatic Fluid, Semen, typing)*	450
30.17.029		1 2 , (HSV-1, 2 DNA, Scrape of Urogenital Epithelial Cells, typing)*	450
30. (Serology Diagnosis of Infectious Disease) / 30.18.			
6 (Human herpesvirus 6, HHV-6)			
30.18.001		IgG 6 (Anti-HHV-6 IgG)	600
30.18.002		6 , (HSV-6 DNA, Exudate)*	250
30.18.003		6 , (HSV-6 DNA, Blood)*	500
30.18.004		6 , (HSV-6 DNA, Urine)*	250
30.18.005		6 , (HSV-6 DNA, Scrape of Nasal Epithelial Cells)*	250
30.18.006		6 , (HSV-6 DNA, Scrape of Faucial Epithelial Cells)*	250
30.18.007		6 , (HSV-6 DNA, Serum)*	500
30.18.008		6 , (HSV-6 DNA, Saliva)*	250
30.18.009		6 , (HSV-6 DNA, Cerebrospinal Fluid)*	250
30.18.010		6 , (HSV-6 DNA, Prostatic Fluid, Semen)*	250
30.18.011		6 , (HSV-6 DNA, Scrape of Urogenital Epithelial Cells)*	400
30.18.012		Ig IgG, 1 2 (Anti-HSV-1, 2 IgM, IgG, Avidity)	600
30.18.013		TORCH- IgG IgM, (ToRCH-Infection IgG, IgM, Avidity)	2000
30.18.014		TORCH- IgG IgM, (ToRCH-Infection IgG, IgM, without Avidity)	1900
30. (Serology Diagnosis of Infectious Disease) / 30.19.			
8 (Human herpesvirus 8, HHV-8)			
30.19.001		IgG 8 (Anti-HHV-8 IgG)	600
30. (Serology Diagnosis of Infectious Disease) / 30.20.			
() (Neisseria gonorrhoeae)			
30.20.001		, (Neisseria gonorrhoeae, DNA, Scrape of Conjunctiva Epithelial Cells)*	250
30.20.002		, (Neisseria gonorrhoeae, DNA, Urine)*	250
30.20.003		, (Neisseria gonorrhoeae, DNA, Scrape of Rectal Epithelial Cells)*	250
30.20.004		, (Neisseria gonorrhoeae, DNA, Scrape of Faucial Epithelial Cells)*	250
30.20.005		, (Neisseria gonorrhoeae, DNA, Synovial Fluid)*	500
30.20.006		, (Neisseria gonorrhoeae, DNA, Prostatic Fluid, Semen)*	250
30.20.007		, (Neisseria gonorrhoeae, DNA, Scrape of Urogenital Epithelial Cells)*	250
30.20.008		(Neisseria gonorrhoeae,), (GC, Neisseria gonorrhoeae Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	550

30. (Serology Diagnosis of Infectious Disease) / 30.23.			
(Entamoeba histolytica)			
30.23.001		IgG (Anti-Entamoeba histolytica IgG)	600
30. (Serology Diagnosis of Infectious Disease) / 30.24.			
(Shigella flexneri, Shigella sonnei)			
30.24.001		Shigella flexneri 1-5 (Shigella flexneri 1-5, IHA)	550
30.24.002		Shigella flexneri 6 (Shigella flexneri 6, IHA)	550
30.24.003		Shigella sonnei (Shigella sonnei, IHA)	550
30.24.004		(Stool Culture (Salmonella spp., Shigella spp.). Bacteria Identification)	600
30.24.005		(Stool Culture (Salmonella spp., Shigella spp.). Bacteria Identification and Antibiotic Susceptibility Testing)	700
30.24.006		(Stool Culture, Salmonella s p., Shigella s p. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)	700
30. (Serology Diagnosis of Infectious Disease) / 30.25.			
(Corynebacterium diphtheriae)			
30.25.001		IgG (Anti-Diphtheria Toxoid IgG)	1100
30.25.002		(Corynebacterium diphtheriae Culture)	700
30. (Serology Diagnosis of Infectious Disease) / 30.26.			
(Yersinia enterocolitica)			
30.26.001		IgA Yersinia enterocolitica (Anti-Yersinia enterocolitica IgA)	550
30.26.002		IgG Yersinia enterocolitica (Anti-Yersinia enterocolitica IgG)	550
30.26.003		(Yersinia enterocolitica, (Yersinia enterocolitica, Stool Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	800
30.26.004		Yersinia pseudotuberculosis (Yersinia pseudotuberculosis IHA)	550
30.26.005		Yersinia enterocolitica :3 (Yersinia enterocolitica O:3, IHA)	550
30.26.006		Yersinia enterocolitica :9 (Yersinia enterocolitica O:9, IHA)	550
30. (Serology Diagnosis of Infectious Disease) / 30.27.			
(Campylobacter spp.)			
30.27.001		(Campylobacter s p.) (Campylobacter spp., Stool Culture. Bacterial Identification)	550
30. (Serology Diagnosis of Infectious Disease) / 30.28.			
(Candidiasis, Candida)			
30.28.001		A IgG Candida albicans (Anti-Candida albicans IgG)	240
30.28.002		Candida albicans, IgG (M5) (M5 Candida albicans, IgG)	400
30.28.003		(Candida albicans, DNA, Exudate)*	300
30.28.004		(Candida albicans, DNA, Scrape of Skin Epithelial Cells)*	250
30.28.005		(Candida albicans, DNA, Urine)*	250
30.28.006		(Candida albicans, DNA, Scrape of Rectal Epithelial Cells)*	250
30.28.007		(Candida albicans, DNA, Scrape of Faucial Epithelial Cells)*	250
30.28.008		(Candida albicans, DNA, Saliva)*	250
30.28.009		(Candida albicans, DNA, Prostatic Fluid, Semen)*	250
30.28.010		(Candida albicans, DNA, Scrape of Urogenital Epithelial Cells)*	250
30.28.011		(Candida albicans, Candida crusei, Candida glabrata), (Candida albicans, Candida crusei, Candida glabrata, DNA)*	550
30.28.012		(Candidiasis, Screening and Typing)	1000
30.28.013		(Candidiasis, Screening)	400
30.28.014		(Candidiasis, Typing)	700

30.28.015		(Candida, Cryptococcus) (Yeast Culture. Identification and Antimycotic Susceptibility testing)	550
30. (Serology Diagnosis of Infectious Disease) / 30.29. (Flavivirus, Tick-borne encephalitis virus)			
30.29.001		IgG (Anti-Tick-borne Encephalitis Virus (TBEV) IgG)	300
30.29.002		IgM (Anti-Tick-borne Encephalitis Virus (TBEV) IgM)	300
30. (Serology Diagnosis of Infectious Disease) / 30.30. (Clostridium difficile)			
30.30.001		(Clostridium difficile, (Clostridium difficile Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	800
30.30.002		Toxin A Clostridium difficile, (Toxin A Clostridium difficile. One Step Rapid Immun chromatographic Assay)	700
30. (Serology Diagnosis of Infectious Disease) / 30.31. (Bordetella pertussis, Whooping Cough)			
30.31.001		A IgA Bordetella pertussis (Anti-Bordetella pertussis IgA)	800
30.31.002		A IgG Bordetella pertussis (Anti-Bordetella pertussis IgG)	800
30.31.003		A IgM Bordetella pertussis (Anti-Bordetella pertussis IgM)	800
30.31.004		(Bordetella pertussis), (Bordetella pertussis, DNA)*	250
30.31.005		(Bordetella pertussis/parapertussis, /) (Bordetella pertussis/parapertussis, Nasopharyngeal Culture. Bacteria Identification)	900
30. (Serology Diagnosis of Infectious Disease) / 30.32. (Morbillivirus, Measles virus)			
30.32.001		IgG (Anti-Measles IgG)	1100
30.32.002		Ig (Anti-Measles IgM)	800
30. (Serology Diagnosis of Infectious Disease) / 30.33. (Rubella virus)			
30.33.001		IgG (Anti-Rubella IgG)	380
30.33.002		Ig (Anti-Rubella Ig)	280
30.33.003		IgG (Anti-Rubella IgG, Immunoblot)	2200
30.33.004		IgM IgG (Anti-Rubella IgM, IgG, Avidity)	900
30.33.005		(Rubella Virus, DNA, Serum)*	450
30. (Serology Diagnosis of Infectious Disease) / 30.34. (Cryptosporidium parvum)			
30.34.001		(Cryptosporidium parvum), (Cryptosporidium parvum. One Step Rapid Immun chromatographic Assay)	600
30. (Serology Diagnosis of Infectious Disease) / 30.35. (Lactobacillus spp.)			
30.35.001		(Lactobacillus spp., DNA, Scrape of Urogenital Epithelial Cells)*	260
30. (Serology Diagnosis of Infectious Disease) / 30.36. (Listeria monocytogenes, Listeriosis)			
30.36.001		(Listeria monocytogenes, DNA, Scrape of Nasal Epithelial Cells)*	350
30.36.002		(Listeria monocytogenes, DNA, Plasma)*	300
30.36.003		(Listeria monocytogenes, DNA, Scrape of Faucial Epithelial Cells)*	400
30.36.004		(Listeria monocytogenes, DNA, Synovial Fluid)*	450
30.36.005		(Listeria monocytogenes, DNA, Cerebrospinal Fluid)*	300
30.36.006		(Listeria monocytogenes,) (Listeria monocytogenes Culture. Bacteria Identification)	800

30. (Serology Diagnosis of Infectious Disease) / 30.37.			
(Giardia lamblia intestinalis)			
30.37.001		IgM, IgG, IgA (Anti-Giardia lamblia IgM, IgG, IgA, Total)	600
30.37.002		(Giardia lamblia), (Giardia lamblia. One Step Rapid Immunochromatographic Assay)	800
30.37.003		IgM (Anti-Giardia lamblia IgM)	300
30. (Serology Diagnosis of Infectious Disease) / 30.38.			
(Neisseria meningitidis)			
30.38.001		(Neisseria meningitidis Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	800
30. (Serology Diagnosis of Infectious Disease) / 30.39.			
(Skin and Nail Mycoses)			
30.39.001		(Fungal Infections of Nails)	1000
30.39.002		(Fungal Infections of Skin)	1000
30.39.003		(Parasitic Fungi, Microscopy and Culture (skin))	1700
30.39.004		(Parasitic Fungi, Microscopy and Culture (nails))	1700
30. (Serology Diagnosis of Infectious Disease) / 30.40.			
(Mycoplasma genitalium, Mycoplasma pneumoniae, Mycoplasma hominis)			
(Mycoplasmosis, Mycoplasma Infection)			
30.40.001		(Mycoplasma hominis), (Mycoplasma hominis, DNA, Urine)*	250
30.40.002		(Mycoplasma hominis), (Mycoplasma hominis, DNA, Prostatic Fluid, Semen)*	250
30.40.003		(Mycoplasma hominis), (Mycoplasma hominis, DNA, Scrape of Urogenital Epithelial Cells)*	250
30.40.004		(Mycoplasma genitalium), (Mycoplasma genitalium, DNA, Urine)*	250
30.40.005		(Mycoplasma genitalium), (Mycoplasma genitalium, DNA, Prostatic Fluid, Semen)*	250
30.40.006		(Mycoplasma genitalium), (Mycoplasma genitalium, DNA, Scrape of Urogenital Epithelial Cells)*	250
30.40.007		(Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Sputum)*	430
30.40.008		(Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Plasma)*	430
30.40.009		(Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Scrape of Faucial Epithelial Cells)*	380
30.40.010		(Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Saliva)*	380
30.40.011		IgM, IgG Mycoplasma hominis (Anti-Mycoplasma hominis IgM, IgG)	1100
30.40.012		IgM Mycoplasma hominis (Anti-Mycoplasma hominis IgM)	650
30.40.013		IgG Mycoplasma hominis (Anti-Mycoplasma hominis IgG)	450
30.40.014		A IgM, IgG Mycoplasma pneumoniae (Anti-Mycoplasma pneumoniae IgM, IgG)	1100
30.40.015		A IgM Mycoplasma pneumoniae (Anti-Mycoplasma pneumoniae IgM)	600
30.40.016		A IgG Mycoplasma pneumoniae (Anti-Mycoplasma pneumoniae IgG)	600
30.40.017		IgA Mycoplasma hominis (Anti-Mycoplasma hominis IgA)	400
30.40.018		IgG, IgA Mycoplasma hominis (Anti-Mycoplasma hominis IgG, IgA)	900
30. (Serology Diagnosis of Infectious Disease) / 30.41.			
(Urogenital Tract Microbiocenosis)			

30.41.001		(Lactobacillus spp., DNA, Scrape of Urogenital Epithelial Cells)*	250
30.41.002		(Bacteroides spp., DNA, Scrape of Urogenital Epithelial Cells)*	250
30.41.003		(Mobiluncus curtisii, DNA, Scrape of Urogenital Epithelial Cells)	250
30.41.004		(13 +), (Microflora of Urogenital Tract, Scrape of Urogenital Epithelial Cells, Screening)*	1550
30.41.005		(7 +), (Identification of Sexually Transmitted Infections (STI) Pathogens, Scrape of Urogenital Epithelial Cells)*	950
30.41.006		-9 + , (FEMOFLO [®] 9, Scrape of Urogenital Epithelial Cells)*	1100
30.41.007		-17 + , (FEMOFLO [®] 17, Scrape of Urogenital Epithelial Cells)*	1760
30.41.008		? (INBIOFLO [®] ? Comprehensive Study of Microflora Composition of Urogenital Tract (UGT))	3000
30.41.009		(4 +): Chlamydia trachomatis, Neisseria gonorrhoeae, Trichomonas vaginalis, Mycoplasma genitalium, (Identification of Sexually Transmitted Infections (STI) Pathogens, Chlamydia trachomatis DNA,	1700
30.41.010		(Bacterial Vaginosis, BV)	1700
30.41.011		? - (INBIOFLO [®] ? Mycoplasma, Urogenital Screening)	550
30.41.012		, (Candidiasis, Screening and Typing)	1050
30.41.013		, (Candidiasis, Screening)	450
30.41.014		, (Candidiasis, Typing)	700
30.41.015		- , (Ureaplasma urealyticum) (Ureaplasma urealyticum, Effectiveness Monitoring of Treatments)	450
30.41.016		- , (Ureaplasma parvum) (Ureaplasma parvum, Effectiveness Monitoring of Treatments)	450
30.41.017		- , (Mycoplasma hominis) (Mycoplasma hominis, Effectiveness Monitoring of Treatments)	450
30. (Mobiluncus curtisii, M. mulieris) (Serology Diagnosis of Infectious Disease) / 30.42.			
30.42.001		(Mobiluncus curtisii, DNA, Scrape of Urogenital Epithelial Cells)	250
30. (, Opisthorchis felinus) (Serology Diagnosis of Infectious Disease) / 30.44.			
30.44.001		IgG (Anti-Opisthorchis felinus IgG)	300
30.44.002		IgM, IgG () (Anti-Opisthorchis felinus IgM, IgG, Circulating Immune Complexes (CIC))	300
30. (Acute Intestinal Infections) (Serology Diagnosis of Infectious Disease) / 30.45.			
30.45.001		, (Enterovirus, RNA, Fecal)	500
30.45.003		, - (Acute Intestinal Infections, PCR, Fecal)	1300
30.45.004		, - (Acute Intestinal Infections, PCR, Fecal)	1400
30. (Bordetella parapertussis) (Serology Diagnosis of Infectious Disease) / 30.46.			

30.46.001		(Bordetella pertussis/parapertussis, /) (Bordetella pertussis/parapertussis, Nasopharyngeal Culture. Bacteria Identification)	900
30. (Serology Diagnosis of Infectious Disease) / 30.47. (Parotitis epidemica, Mumps, Epidemic Parotitis)			
30.47.001	A	IgG (Anti-Mumps IgG)	700
30.47.002	A	IgM (Anti-Mumps IgM)	700
30. (Serology Diagnosis of Infectious Disease) / 30.48. (Entamoeba histolytica, Giardia intestinalis, Balantidium coli)			
30.48.001		(PRO Stool)	300
30. (Serology Diagnosis of Infectious Disease) / 30.49. (Respiratory Syncytial Virus, RSV)			
30.49.001	A	IgG - (Anti-Respiratory Syncytial Virus (RSV) IgG)	700
30.49.002	A	IgM - (Anti-Respiratory Syncytial Virus (RSV) IgM)	700
30. (Serology Diagnosis of Infectious Disease) / 30.50. (Typhus, Rickettsia prowazekii)			
30.50.001		(Rickettsia prowazekii, IHA)	550
30. (Serology Diagnosis of Infectious Disease) / 30.51. (Rotavirus)			
30.51.001		(Rotavirus), (Rotavirus Direct Detection by Latex Agglutination)	500
30. (Serology Diagnosis of Infectious Disease) / 30.52. (Salmonella spp.)			
30.52.001		Salmonella typhi, (Salmonella typhi Antibodies, IHA)	500
30.52.002		Salmonella gr.A (Salmonella gr.A, IHA)	500
30.52.003		Salmonella gr.B (Salmonella gr.B, IHA)	500
30.52.004		Salmonella gr. (Salmonella gr.C, IHA)	500
30.52.005		Salmonella gr.D (Salmonella gr.D, IHA)	500
30.52.006		Salmonella gr.E, (Salmonella gr.E Antibodies, IHA)	500
30.52.007		Salmonella O- (Salmonella O-antigens, IHA)	500
30.52.008		(Stool Culture (Salmonella spp., Shigella spp.). Bacteria Identification)	600
30.52.009		(Stool Culture (Salmonella spp., Shigella spp.). Bacteria Identification and Antibiotic Susceptibility Testing)	800
30.52.010		(Stool Culture, Salmonella s p., Shigella s p. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)	800
30. (Serology Diagnosis of Infectious Disease) / 30.53. (Treponema pallidum, Syphilis)			
30.53.001		RPR – (Syphilis RPR (Rapid Plasma Reagins), nticardiolipin est)	100
30.53.002	A	IgM IgG Treponema pallidum, (Anti-Treponema pallidum IgM, IgG, Total)	300
30.53.003	A	IgM Treponema pallidum (Anti-Treponema pallidum IgM)	600
30.53.004	A	IgM Treponema pallidum, (Anti-Treponema pallidum IgM, Immunoblot)	2600
30.53.005	A	IgG Treponema pallidum, (Anti-Treponema pallidum IgG, Immunoblot)	2100
30.53.006		(RW) ((RW) (Wassermann Reaction (RW), Wassermann Test)	900
30.53.007		(Treponema pallidum, DNA, Scrape of Conjunctiva Epithelial Cells)*	250
30.53.008		(Treponema pallidum, DNA, Scrape of Skin Epithelial Cells)*	250
30.53.009		(Treponema pallidum, DNA, Urine)*	250

30.53.010		(Treponema pallidum, DNA, Secretion)*	250
30.53.011		(Treponema pallidum, DNA, Scrape of Faucial Epithelial Cells)*	250
30.53.012		(Treponema pallidum, DNA, Serum)*	400
30.53.013		(Treponema pallidum, DNA, Cerebrospinal Fluid)*	250
30.53.014		(Treponema pallidum, DNA, Prostatic Fluid, Semen)*	250
30.53.015		(Treponema pallidum, DNA, Scrape of Urogenital Epithelial Cells)*	250
30. (Serology Diagnosis of Infectious Disease) / 30.54. (Staphylococcus aureus)			
30.54.001		(Staphylococcus aureus) (Staphylococcus aureus (Methicillin-Resistant Staphylococcus aureus – MRSA) Culture. Bacteria Identification)	500
30.54.002		(Staphylococcus aureus), (Staphylococcus aureus (Methicillin-Resistant Staphylococcus aureus – MRSA) Culture. Ba	500
30.54.003		(Staphylococcus aureus), (Staphylococcus aureus (Methicillin-Resistant Staphylococcus aureus	800
30.54.004		(Staphylococcus aureus), (Staphylococcus aureus (Methicillin-Resistant Staphylococcus aureus – M	760
30.54.005		(Staphylococcus aureus) (Staphylococcus aureus Culture. Bacteria Identification)	760
30.54.006		(Staphylococcus aureus), (Staphylococcus aureus Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	760
30.54.007		(Staphylococcus aureus), (Staphylococcus aureus Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)	800
30.54.008		(Staphylococcus aureus), (Staphylococcus aureus Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)	900
30. (Serology Diagnosis of Infectious Disease) / 30.55. (Tetanus)			
30.55.001		IgG (anti-Tetanus toxoid IgG)	1100
30. (Serology Diagnosis of Infectious Disease) / 30.56. (Streptococcal Infection) A B (Streptococcus group A, B, S. pyogenes, S. agalactiae)			
30.56.001		(Streptococcus spp., DNA, Sputum)*	800
30.56.002		(Streptococcus spp., DNA, Plasma)*	670
30.56.003		(Streptococcus spp., DNA, Scrape of Faucial Epithelial Cells)*	400
30.56.004		(Streptococcus spp., DNA, Saliva)*	400
30.56.005		(Streptococcus group A, Streptococcus pyogenes) (Streptococcus pyogenes Culture. Bacteria Identification)	900
30.56.006		(Streptococcus group A, Streptococcus pyogenes), (Streptococcus pyogenes Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	1100

30.56.007		(Streptococcus group B, Streptococcus agalactiae) (Streptococcus agalactiae Culture. Bacteria Identification)	500
30.56.008		(Streptococcus group B, Streptococcus agalactiae), (Streptococcus agalactiae Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	900
30.56.009		(Streptococcus Group B. One Step Rapid Immunochromatographic Assay)	760
30. (Strongyloides stercoralis) (Serology Diagnosis of Infectious Disease) / 30.57.			
30.57.001		IgG (Anti-Strongyloides stercoralis IgG)	1300
30. (Deltaretrovirus, Human T-Lymphotropic Virus, HTLV) (Serology Diagnosis of Infectious Disease) / 30.58.			
30.58.001		IgG - 1 2 (Anti-HTLV-1, 2 IgG)	400
30. (Toxocara canis) (Serology Diagnosis of Infectious Disease) / 30.59.			
30.59.001		IgG (Anti-Toxocara IgG)	220
30. (Toxoplasmosis, Toxoplasma gondii) (Serology Diagnosis of Infectious Disease) / 30.60.			
30.60.001		IgG Toxoplasma gondii (Anti-Toxoplasma gondii IgG)	300
30.60.002		Ig Toxoplasma gondii (Anti-Toxoplasma gondii Ig)	300
30.60.003		IgM IgG Toxoplasma gondii, (Anti-Toxoplasma gondii IgM, IgG, Avidity)	760
30.60.004		(Toxoplasma gondii, DNA, Exudate)*	170
30.60.005		(Toxoplasma gondii, DNA, Serum)*	400
30.60.006		(Toxoplasma gondii, DNA, Cerebrospinal Fluid)*	170
30. (Trichinella spiralis) (Serology Diagnosis of Infectious Disease) / 30.61.			
30.61.001		IgG (Anti-Trichinella IgG)	500
30.61.002		IgM IgG (Anti-Trichinella IgM, IgG)	900
30. (Trichomonas vaginalis, Trichomoniasis) (Serology Diagnosis of Infectious Disease) / 30.62.			
30.62.001		IgG Trichomonas vaginalis (Anti-Trichomonas vaginalis IgG)	260
30.62.002		IgG IgA Trichomonas vaginalis (Anti-Trichomonas vaginalis IgG, IgA)	260
30.62.003		(Trichomonas vaginalis, DNA, Urine)*	250
30.62.004		(Trichomonas vaginalis, DNA, Prostatic Fluid, Semen)*	250
30.62.005		(Trichomonas vaginalis, DNA, Scrape of Urogenital Epithelial Cells)*	250
30. (Mycobacterium Tuberculosis) (Serology Diagnosis of Infectious Disease) / 30.63.			
30.63.001		IgM, IgA, IgG Mycobacterium tuberculosis, (Anti-Mycobacterium tuberculosis IgM, IgA, IgG, Total)	1500
30.63.002		(Mycobacterium tuberculosis, DNA, Exudate)*	300
30.63.003		(Mycobacterium tuberculosis, DNA, Sputum)*	500
30.63.004		(Mycobacterium tuberculosis, DNA, Menstrual Blood)*	300
30.63.005		(Mycobacterium tuberculosis, DNA, Urine)*	250
30.63.006		(Mycobacterium tuberculosis, DNA, Serum)*	400

30.63.007		(Mycobacterium tuberculosis, DNA, Synovial Fluid)*	400
30.63.008		(Mycobacterium tuberculosis, DNA, Cerebrospinal Fluid)*	300
30.63.009		(Mycobacterium tuberculosis, DNA, Prostatic Fluid, Semen)*	250
30. () (Serology Diagnosis of Infectious Disease) / 30.64. (Ureaplasmosis, Ureaplasma spp., Ureaplasma urealyticum, Ureaplasma parvum)			
30.64.001		IgG Ureaplasma urealyticum (anti-Ureaplasma urealyticum IgG)	280
30.64.002		IgA Ureaplasma urealyticum (anti-Ureaplasma urealyticum IgA)	280
30.64.003		IgG IgA Ureaplasma urealyticum (anti-Ureaplasma urealyticum IgG, IgA)	280
30.64.004		(Ureaplasma urealyticum) (-960), (Ureaplasma urealyticum (T-960), DNA, Urine)*	250
30.64.005		(Ureaplasma urealyticum) (-960), (Ureaplasma urealyticum (T-960), DNA, Prostatic Fluid, Semen)*	250
30.64.006		(Ureaplasma urealyticum) (-960), (Ureaplasma urealyticum (T-960), DNA, Scrape of Urogenital Epithelial Cells)*	250
30.64.007		(Ureaplasma parvum), (Ureaplasma parvum, DNA, Urine)*	250
30.64.008		(Ureaplasma arvum), (Ureaplasma parvum, DNA, Prostatic Fluid, Semen)*	250
30.64.009		(Ureaplasma arvum), (Ureaplasma parvum, DNA, Scrape of Urogenital Epithelial Cells)*	250
30.64.010		(Ureaplasma urealyticum + Ureaplasma arvum), (Ureaplasma urealyticum + Ureaplasma arvum, DNA, Urine)*	250
30.64.011		(Ureaplasma urealyticum + Ureaplasma arvum), (Ureaplasma urealyticum + Ureaplasma arvum, DNA, Prostatic Fluid, Semen)*	250
30.64.012		(Ureaplasma urealyticum + Ureaplasma arvum), (Ureaplasma urealyticum + Ureaplasma arvum, DNA, Scrape of Urogenital Epithelial Cells)*	250
30. () (Serology Diagnosis of Infectious Disease) / 30.65. (Helicobacter pylori)			
30.65.001	A	IgG Helicobacter ylori (Anti-Helicobacter pylori IgG)	300
30.65.002	A	IgM Helicobacter ylori (Anti-Helicobacter pylori IgM)	700
30.65.003	A	IgA Helicobacter ylori (Anti-Helicobacter pylori IgA)	700
30.65.004	A	IgG Helicobacter pylori, (Anti-Helicobacter pylori IgG, Immunoblot)	2200
30.65.005	A	IgA Helicobacter pylori, (Anti-Helicobacter pylori IgA, Immunoblot)	2200
30. () (Serology Diagnosis of Infectious Disease) / 30.66. Chlamydophila pneumoniae (Chlamydiosis, Chlamydia Infection, Chlamydia trachomatis,			
30.66.001	A	IgA Chlamydia trachomatis (Anti-Chlamydia trachomatis IgA)	200
30.66.002	A	IgG Chlamydia trachomatis (Anti-Chlamydia trachomatis IgG)	200
30.66.003	A	IgA IgG Chlamydia trachomatis, (Anti-Chlamydia trachomatis IgA, IgG)	250
30.66.004	A	IgM Chlamydia trachomatis (Anti-Chlamydia trachomatis IgM)	500
30.66.005		IgM IgG Chlamydophila pneumoniae (anti-Chlamydophila pneumoniae IgM, IgG)	350
30.66.006		IgG () Chlamydia trachomatis (Anti-cHSP60 IgG)	900
30.66.007		/ (Helicobacter pylori, DNA, Biopates of Gastric Mucosa and/or Duodenum, PCR)	700

30.66.008		(Chlamydia trachomatis), trachomatis, DNA, Exudate)*	(Chlamydia	250
30.66.009		(Chlamydia trachomatis), Conjunctiva Epithelial Cells)*	(Chlamydia trachomatis, DNA, Scrape of	250
30.66.010		(Chlamydia trachomatis), trachomatis, DNA, Urine)*	(Chlamydia	250
30.66.011		(Chlamydia trachomatis), DNA, Scrape of Rectal Epithelial Cells)*	(Chlamydia trachomatis,	250
30.66.012		(Chlamydia trachomatis), Faucial Epithelial Cells)*	(Chlamydia trachomatis, DNA, Scrape of	250
30.66.013		(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Synovial Fluid)*		400
30.66.014		(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Cerebrospinal Fluid)*		250
30.66.015		(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Prostatic Fluid, Semen)*		250
30.66.016		(Chlamydia trachomatis), Scrape of Urogenital Epithelial Cells)*	(Chlamydia trachomatis, DNA,	250
30.66.017		A IgA Chlamydophila pneumoniae (Anti-Chlamydophila pneumoniae IgA)		550
30.66.018		A IgM Chlamydophila pneumoniae (Anti-Chlamydophila pneumoniae IgM)		200
30.66.019		A IgG Chlamydophila pneumoniae (Anti-Chlamydophila pneumoniae IgG)		200
30.66.020		(Chlamydia pneumoniae), (Chlamydophila pneumoniae, DNA, Sputum)*		900
30.66.021		(Chlamydia pneumoniae), (Chlamydophila pneumoniae, DNA, Plasma)*		550
30.66.022		(Chlamydia pneumoniae), of Faucial Epithelial Cells)*	(Chlamydophila pneumoniae, DNA, Scrape	500
30.66.023		(Chlamydia pneumoniae), (Chlamydophila pneumoniae, DNA, Saliva)*		500
30. (Serology Diagnosis of Infectious Disease) / 30.67. (Cytomegalovirus, CMV, Cytomegalovirus Infection)				
30.67.001		A IgG (Anti-CMV IgG)		250
30.67.002		A IgM (Anti-CMV IgM)		280
30.67.003		IgM IgG (Anti-CMV IgM, IgG, Avidity)		500
30.67.004		, (CMV DNA, Exudate)*		250
30.67.005		, (CMV DNA, Scrape of Conjunctiva Epithelial Cells)*		250
30.67.006		, (CMV DNA, Scrape of Skin Epithelial Cells)*		250
30.67.007		, (CMV DNA, Blood)*		400
30.67.008		, (CMV DNA, Urine)*		250
30.67.009		, (CMV DNA, Scrape of Nasal Epithelial Cells)*		250
30.67.010		, (CMV DNA, Scrape of Faucial Epithelial Cells)*		250
30.67.011		, Serum)*	(CMV DNA,	3350
30.67.012		, (CMV DNA, Saliva)*		250
30.67.013		, DNA, Cerebrospinal Fluid)*	(CMV	250
30.67.014		, DNA, Prostatic Fluid, Semen)*	(CMV	250
30.67.015		, (CMV DNA, Scrape of Urogenital Epithelial Cells)*		250

30. (Serology Diagnosis of Infectious Disease) / 30.68.			
(Enterobius spp.)			
30.68.001		(), (nterobiasis, Swab)	380
30. (Serology Diagnosis of Infectious Disease) / 30.69.			
(Epstein-Barr Virus, EBV, EBV Infection)			
30.69.001		(EBV DNA, Exudate)*	250
30.69.002		(EBV DNA, Blood)*	250
30.69.003		(EBV DNA, Urine)*	250
30.69.004		(EBV DNA, Scrape of Nasal Epithelial Cells)*	250
30.69.005		(EBV DNA, Scrape of Faucial Epithelial Cells)*	250
30.69.006		(EBV DNA, Serum)*	400
30.69.007		(EBV DNA, Saliva)*	250
30.69.008		(EBV DNA, Cerebrospinal Fluid)*	250
30.69.009		(EBV DNA, Prostatic Fluid, Semen)*	250
30.69.010		(EBV DNA, Scrape of Urogenital Epithelial Cells)*	250
30.69.011		Ig (nti-EBV Viral Capsid Antigens (VCA) Ig)	280
30.69.012		IgG (nti-EBV Nuclear Antigen (EBNA) IgG)	250
30.69.013		IgG (nti-EBV Early Antigen (EA) IgG)	250
30.69.014		IgG (nti-EBV Viral Capsid Antigens (VCA) IgG)	280
30.69.015		IgG (Epstein-Barr Virus Nuclear Antigen IgG, EBNA IgG)	350
30.69.016		(), IgG (EBV IgG, Immunoblot)	1300
30.69.017		(), Ig (EBV IgM, Immunoblot)	1300
30. (Serology Diagnosis of Infectious Disease) / 30.70.			
(Echinococcus spp.)			
30.70.001		IgG (Anti-Echinococcus IgG)	280
30. (Serology Diagnosis of Infectious Disease) / 30.71.			
(Escherichia coli)			
30.71.001		(Escherichia coli O157:H7,), (Escherichia coli O157:H7. One Step Rapid Immun hromotographic Assay)	700
30. (Serology Diagnosis of Infectious Disease) / 30.10.			
(Helminthiasis)			
30.10.003		PARASEP	676
31. (Microbiological investigations) / 31.01.			
(Microbiological investigations)			
31.01.047		(Eye Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*	0
31.01.001		(Stool Culture)	1300
31.01.002		(Stool Culture. Intestinal Bacterial Overgrowth, Bacteriophage Efficiency Testing)	1500
31.01.003		(Vaginal Biocenosis: Bacteriophage Susceptibility Testing (Gram Stain, Bacterioscopic Examination of Smear))*	1100
31.01.004		(Gram Stain. Bacterioscopic Examination of Smear)	150

31.01.005		(Bordetella pertussis/parapertussis, /) (Bordetella pertussis/parapertussis, Nasopharyngeal Culture. Bacteria Identification)	900
31.01.006		(Candida, Cryptococcus) (Yeast Culture. Identification and Antimycotic Susceptibility testing)	500
31.01.007		(Clostridium difficile,) (Clostridium difficile Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	600
31.01.008		(Escherichia coli O157:H7,), (Escherichia coli O157:H7. One Step Rapid Immun hromotographic Assay)	700
31.01.009		(Mycoplasma hominis Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	400
31.01.010		(Neisseria meningitidis Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	700
31.01.011		(Trichomonas vaginalis,) (Trichomonas vaginalis Culture)	700
31.01.012		(Ureaplasma spp.), (Ureaplasma spp. Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	700
31.01.013		(Mycoplasma hominis) (Ureaplasma spp.), (Mycoplasma hominis Culture, Ureaplasma spp. Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	950
31.01.014		(Neisseria gonorrhoeae,), (GC, Neisseria gonorrhoeae Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	500
31.01.015		(Stool Culture (Salmonella spp., Shigella spp.). Bacteria Identification)	550
31.01.016		(Stool Culture (Salmonella spp., Shigella spp.). Bacteria Identification and Antibiotic Susceptibility Testing)	700
31.01.017		(Stool Culture, Salmonella s p., Shigella s p. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)	700
31.01.018		(Corynebacterium diphtheriae Culture)	650
31.01.019		(-) (Staphylococcus aureus) (Staphylococcus aureus (Methicillin-Resistant Staphylococcus aureus – MRSA) Culture. Bacteria Identification)	500
31.01.020		(-) (Staphylococcus aureus), (Staphylococcus aureus (Methicillin-Resistant Staphylococcus aureus – MRSA) Culture. Ba	500
31.01.021		(-) (Staphylococcus aureus), (Staphylococcus aureus (Methicillin-Resistant Staphylococcus aureu	700
31.01.022		(-) (Staphylococcus aureus), (Staphylococcus aureus (Methicillin-Resistant Staphylococcus aureus – M	700
31.01.023		(Staphylococcus aureus) (Staphylococcus aureus Culture. Bacteria Identification)	700
31.01.024		(Staphylococcus aureus), (Staphylococcus aureus Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	700
31.01.025		(Staphylococcus aureus), (Staphylococcus aureus Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)	750

31.01.026		(Staphylococcus aureus, (Staphylococcus aureus Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)	900
31.01.027		(Yersinia enterocolitica, (Yersinia enterocolitica, Stool Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	750
31.01.028		(Campylobacter s p.) (Campylobacter spp., Stool Culture. Bacterial Identification)	500
31.01.029		(Listeria monocytogenes,) (Listeria monocytogenes Culture. Bacteria Identification)	750
31.01.030		(Wound/Pus/Aspirate/Tissue Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	650
31.01.031		(Wound/Pus/Aspirate/Tissue Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	700
31.01.032		(Wound/Pus/Aspirate/Tissue Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*	750
31.01.033		(Breast Milk Culture. Bacteria Identification)	600
31.01.034		(Breast Milk Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	750
31.01.035		(Breast Milk Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	900
31.01.036		(Breast Milk Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*	900
31.01.037		(Bile Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	900
31.01.038		(Bile Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	1000
31.01.039		(Upper Respiratory Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	600
31.01.040		(Upper Respiratory Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	750
31.01.041		(Upper Respiratory Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*	750
31.01.042		(Urine Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	600
31.01.043		(Urine Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	700
31.01.044		(Urine Culture. Bacteria Identification, Antibiotic susceptibility and Bacteriophage Efficiency Testing)*	700
31.01.045		(Eye Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	500

31.01.046		(Eye Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	650
31.01.048		(Genitourinary Tract Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	900
31.01.049		(Genitourinary Tract Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	1000
31.01.050		(Genitourinary Tract Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*	1000
31.01.051		(Streptococcus Group B. One Step Rapid Immunohromotographic Assay)	650
31.01.052		(Eye Culture. Bacteria Identification and Antibiotic Susceptibility Testing)*	550
31.01.053		(Eye Culture. Bacteria Identification, Antibiotic Susceptibility, Enlarged Testing)*	650
31.01.054		(Eye Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*	700
31.01.055		(Sputum and Tracheobronchial washings Culture. Bacteria Identification and Antibiotic Susceptibility Testing, Microscopy)	600
31.01.056		(Sputum and Tracheobronchial washings Culture. Bacteria Identification, Antibiotic Susceptibility, E	1000
31.01.057		(Rotavirus), (Rotavirus Direct Detection by Latex Agglutination)	500
31.01.058		(Adenovirus), (Adenovirus. One Step Rapid Immunohromotographic Assay)	650
31.01.059		(Giardia liamblia), (Giardia liamblia. One Step Rapid Immunohromotographic Assay)	650
31.01.060		(Streptococcus group B, Streptococcus agalactiae) (Streptococcus agalactiae Culture. Bacteria Identification)	500
31.01.061		(Streptococcus group B, Streptococcus agalactiae), (Streptococcus agalactiae Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	900
31.01.062		(Streptococcus group A, Streptococcus pyogenes) (Streptococcus pyogenes Culture. Bacteria Identification)	900
31.01.063		(Streptococcus group A, Streptococcus pyogenes), (Streptococcus pyogenes Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	1000
31.01.064		(Cryptosporidium parvum), (Cryptosporidium parvum. One Step Rapid Immunohromotographic Assay)	650
31.01.065		(Helicobacter pylori), (Helicobacter pylori. One Step Rapid Immunohromotographic Assay)	650
31.01.066		Toxin A Clostridium difficile, (Toxin A Clostridium difficile. One Step Rapid Immunohromotographic Assay)	650
31.01.067		(Urogenital Microorganisms: Detection, Count and Antimicrobial Agent Susceptibility, Colorimetric Method)	2000
31.01.068		Gardnerella vaginalis (Gardnerella vaginalis Culture)	600

31.01.069		Candida albicans (Candida albicans Culture)	400
31.01.070		Mycoplasma genitalium (Mycoplasma genitalium Culture, Qualitative, Quantitative)	400
32. (Immunoglobulins, Total) (Immunological Studies) / 32.02.			
32.02.001		(Immunoglobulin A, IgA)	300
32.02.002		(Immunoglobulin , Ig)	300
32.02.003		G (Immunoglobulin G, IgG)	300
32.02.004		E (IgE, (Immunoglobulin Total, IgE Total)	300
32. (Immunological Studies) / 32.03. (Specific Proteins)			
32.03.001		(Eosinophil Cationic Protein, ECP)	900
32. (Immunological Studies) / 32.04. (Complement Components)			
32.04.001		3, 4 (Complement components C3, C4)	650
32. (Immunological Studies) / 32.05. (Cytokines)			
32.05.001		-1? (-1?) (Interleukin 1 Beta, IL-1)	1700
32.05.002		-6 (-6) (Interleukin 6, IL-6)	1700
32.05.003		-8 (-8) (Interleukin 8, IL-8)	1700
32.05.004		-10 (-10) (Interleukin 10, IL-10)	1700
32.05.005		-? (-?) (Tumor Necrosis Factor Alpha, TNF-?, Cachectin)	1700
34. (Diagnosis of Autoimmune Diseases) / 34.01. (Connective-Tissue Diseases, CTDs)			
34.01.001		(Anti-Nuclear Antibodies, ANA, Screening)	400
34.01.002		IgG (- IgG, -) (Double-Stranded (Native) DNA IgG Antibodies, Anti-dsDNA IgG)	400
34.01.003		, HEp-2 (, , HEp-2-) (Antinuclear Antibodies, ANA, Hep-2 Substrate, ANA-Hep2, Fluorescent Anti-Nuclear Antibodies detection, FANA, iter	1300
34.01.004		IgG () (Extractable Nuclear Antigen, ENA, Anti-Ribonucleoprotein Antibodies, Anti-RNP)	1350
34.01.005		(Scl-70, ENP-A, CENP-B, RP11, RP155, NOR90, Th/To, PM-Sc100, PM-Sc175, Ku, PDGFR, Ro-52), (Scleroderma (Systemic Sclerosis) Antibody Panel: Ant	3650
34.01.006		(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-Sc	2900
34.01.007		IgG (a), (Anti-Nuclear Antibodies, ANA, IgG, Screening)	1300
34.01.008		()	2000
34.01.009		, IgG; , Ig ; ()	3550
34.01.010		()	2000
34.01.011		« (- - IgG, 3, 4)» (Systemic lupus erythematosus (SLE) profile, activity monitoring (anti-double-stranded DNA IgG, C3 and C4 complement components)	1300
34. (Diagnosis of Autoimmune Diseases) / 34.02. (Antiphospholipid Syndrome, APS)			
34.02.001		IgM IgG (Anti-Phospholipid Antibodies, APA, IgM, IgG)	600
34.02.002		() (Lupus Anticoagulant, LA)	550

34.02.003		IgG, IgM (Anti- phosphatidylserine,	1500
34.02.004		IgA, IgM, IgG (Anti-cardiolipin Antibodies IgA, IgM, IgG, aCL, Screening)	500
34.02.005		IgA (Anticardiolipin IgA, aCL IgA)	700
34.02.006		IgG (Anticardiolipin IgG, aCL IgG)	700
34.02.007		IgM (Anticardiolipin IgM, aCL IgM)	1250
34.02.008		IgG, IgA, IgM -2- 1, (Anti-?-Glycoprotein 1 Antibodies, anti-?-G 1, IgG, IgA, IgM, Total)	1400
34.02.009		IgG, IgM (Anti- phosphatidylserine/ prothrombin antibodies, Anti-PS/PT, IgG, IgM, Total)	1400
34.02.010		IgG V (Annexin V antibodies, aAnV, IgG)	1300
34.02.011		Ig V (Annexin V antibodies, aAnV, Ig)	1400
34.02.012		, IgG, IgM	1500
34.02.013		(, , IgG, IgM)	2650
34.02.014		(; , IgG, IgM; -2- 1)	3600
34. (Diagnosis of Autoimmune Diseases) / 34.03. (Immunological Infertility factors)			
34.03.001		(Anti-Spermatozoa Antibodies, ASA, Serum)	500
34.03.002		(Anti-Spermatozoa Antibodies, ASA, Semen)	1450
34. (Diagnosis of Autoimmune Diseases) / 34.04. (Antiphospholipid Syndrome, APS)			
34.04.001		() (Rheumatoid Factor, RF)	150
34.04.002		(Anti- cyclic Citrullinated Peptide, anti-CCP)	1200
34.04.003		IgG (Anti- eratin antibodies, AKA, Anti-Filaggrin antibodies, AFA, IgG)	1700
34.04.004		IgG (-M) (Anti-Mutated Citrullinated Vimentin Antibodies, Anti-MCV, Anti-Modified Citrullinated Vimentin Antibodies, Anti-Sa Antibodies, IgG)	1450
34.04.005		(), (Rheumatoid Factor, RF, urbidimetry)	1350
34.04.006		(Synovial Fluid Smear, Crystals)	1450
34.04.007		- HLA-B27 (Molecular Genetic Testing HLA-B27)	800
34. (Diagnosis of Autoimmune Diseases) / 34.05. (Vasculitis and Renal Lesions)			
34.05.001		IgG (-) (Glomerular Basement Membrane antibodies, Anti-GBM, IgG)	1400
34.05.002		IgG () (Anti-Neutrophil cytoplasmic antibodies, ANCA, IgG)	1700
34.05.003		IgG, IgA, IgM (HUVEC), (Anti-Endothelial Cell Antibodies, AECA, IgG, IgA, IgM, Total)	1500
34.05.004		IgG, IgA, IgM 2 (PLA2R), (Anti-Phospholipase A2 Receptor Antibodies, Anti-PLA2R, IgG, IgA, IgM, Total)	2800
34.05.005		IgG (- PO) (Myeloperoxidase Antibody, MPO)	1200
34.05.006		IgG C1q (Anti-Complement 1q Antibodies, Anti-C1q, IgG)	1200
34.05.007		IgG -3 (-PR-3) (Anti- roteinase-3 antibodies, PR-3- antibodies, PR-3 ANCA, IgG)	1200
34.05.008		(, IgG (Anti-Neutrophil Cytoplasmic Antibodies, ANCA, IgG, Panel)	2900

34.05.009		(/pANCA, cANCA), IgG	2100
34.05.010		(/pANCA, cANCA), IgG	1500
34.05.011		(/pANCA, cANCA), IgG	3270
34. (Diagnosis of Autoimmune Diseases) / 34.06. (Autoimmune Endocrinopathies: Diabetes mellitus type 1)			
34.06.001		IgG (Anti-Islet Cell Antibodies, Islet Cell Autoantibodies, ICA)	1400
34.06.002		IgG (-GAD) (Anti-GAD Antibodies, Glutamate Decarboxylase Antibodies, AT-GAD, IgG)	1600
34.06.003		IgG (Insulin Autoantibodies, IAA, IgG)	400
34.06.004		(IA-2) (Islet Antigen 2 Antibodies, Anti-IA2 antibodies, IA-2 Ab, Tyrosine Phosphatase Antibodies)	1600
34.06.005		GAD/IA-2, (Anti-GAD/IA2 Antibodies Pool, Glutamic Acid Decarboxylase-65, GAD and Insulinoma Antigen 2 (Tyrosine Phosphatase, IA2, ICA-512) Autoantibodies, Total)	1500
34. (Diagnosis of Autoimmune Diseases) / 34.07. (Autoimmune Thyroid diseases)			
34.07.001		(Anti-Thyroid Peroxidase Autoantibodies, Antimicrosomal Antibodies, TPO Antibodies, TPOAb, Anti-TPO)	280
34.07.002		(Anti-Thyroglobulin Autoantibodies, Thyroglobulin Antibodies, Tg Autoantibodies, TgAb, Anti-Tg Ab, ATG)	280
34.07.003		(Anti-Thyroid Microsomal Antibodies)	650
34.07.004		(Thyroid-Stimulating Hormone Receptor Antibodies, TSH Receptor Antibodies, TSHRABs, TSH binding inhibitor immunoglobulin, TBII)	1100
34. (Diagnosis of Autoimmune Diseases) / 34.08. (Autoimmune adrenal insufficiency)			
34.08.001		IgA, IgM, IgG (Anti-Ovarian Antibodies, AOA, IgA, IgM, IgG, Total)	1100
34.08.002		IgA, IgM, IgG (Anti-Steroidal Cell Antibodies, StCAB, Steroidal Cell Autoantibodies, SCA, IgA, IgM, IgG, Total)	1150
34.08.003		IgA, IgM, IgG (Anti-Testicular Steroid-Cell Antibodies, Testicular Anti-Steroidal Cell Antibodies, Testicular StCAB, Steroidal Cell Autoantibodies, SCA against Testis, IgA, IgM, IgG, Total)	1500
34.08.004			2500
34. (Diagnosis of Autoimmune Diseases) / 34.09. (Autoimmune Skin Diseases)			
34.09.001		IgG (Basement Membrane Zone Antibodies, BMZ, IgG)	1700
34.09.002		IgG (Desmoglein Antibodies, Desmoglein 1, DSG1 and Desmoglein 3, DSG3 Antibodies, IgG)	2300
34.09.003		IgG -1 (Desmoglein 1, DSG1 Antibodies, IgG)	2300
34.09.004		IgG -3 (Desmoglein 3, DSG3 Antibodies, IgG)	2300
34.09.005		IgG BP180 (Anti-Bp180 antibodies, Bullous Pemphigoid (180 kDa) Antibodies, Antibodies to BP Antigen 2, IgG)	2300
34.09.006		IgG BP230 (Anti-Bp230 antibodies, Bullous Pemphigoid (230 kDa) Antibodies, Antibodies to BP Antigen 1, IgG)	2300
34.09.007		« (Bullous Dermatitis Diagnostics profile (antibodies to epidermis desmosomes, antibodies to skin basal membrane)) »	3700

34. (Diagnosis of Autoimmune Diseases) / 34.10. (Autoimmune Disorders of Gastrointestinal Tract (GI Tract, GIT). Coeliac Disease)			
34.10.001		IgA IgG (Anti-ndomysial ntibodies, Anti-EMA, IgA, IgG, Total)	1200
34.10.002		Ig (Anti-ndomysial ntibodies, Anti-EMA, Ig)	1300
34.10.003		IgA, IgG, IgM (Gastric Parietal Cell Antibodies, GPA, Anti- arietal cell antibodies, APCA, IgA, IgG, IgM, Total)	1300
34.10.004		IgA IgG (Anti-Reticulin Antibodies, ARA, IgA, IgG, Total)	1200
34.10.005		IgG (Anti-Deaminated Gliadin Peptide, Anti-DGP, IgG)	470
34.10.006		Ig (Anti-Deaminated Gliadin Peptide, Anti-DGP, IgA)	470
34.10.007		Ig (Anti-Tissue Transglutaminase Antibodies, Anti-tTG, tTGA, IgA)	900
34.10.008		IgG (Anti-Tissue Transglutaminase Antibodies, Anti-tTG, tTGA, IgG)	900
34.10.009		IgG (Anti-Intrinsic Factor, IFAb, Intrinsic Factor Antibodies, IgG)	1400
34.10.010		IgG (Anti-Sacch romyces Cerevisiae Antibodies, ASCA, IgG)	1150
34.10.011		Ig (Anti-Sacch romyces Cerevisiae Antibodies, ASCA, IgA)	1200
34.10.012		Ig (Anti-Neutrophil Cytoplasmic Antibodies, ANCA, IgA)	1200
34.10.013		IgG4- (Diagnosis of Autoimmune Pancreatitis and other IgG4-Related Diseases)	1700
34.10.014		() ; ()	1600
34.10.015		() (/pANCA, cANCA, IgG; /ANCA, IgA; ASCA, IgG, IgA)	3600
34.10.016		, (, IgA; , IgG; IgA)	1500
34.10.017		, (, IgA; , IgA, IgG; IgA)	2600
34.10.018		(Fecal Calprotectin)	2000
34.10.019		IgG IgA (Anti-Tissue Transglutaminase, tTG, Anti-tTG, IgG, Ig)	1200
34.10.020		« » (IgG, IgA, IgG, IgA) (Cereal Intolerance profile (gliadin IgG antibodies, gliadin IgA antibodies, tissue transglutamase antibodies))	1500
34.10.021		IgA IgG (Anti-Intestinal Goblet Cells Antibodies, GAB, IgA, IgG, Total)	1400
34.10.022		IgG IgA GP2 (Anti-GP2, IgG, IgA)	1800
34.10.023		, IgG IgA (, Autoantibodies against Exocrine Pancreas, Pancreatic Antibodies, PAB)	1400
34. (Diagnosis of Autoimmune Diseases) / 34.11. (Autoimmune Liver Diseases)			
34.11.001		IgA, IgG, IgM (Anti-Mitochondrial Antibodies, AMA, IgA, IgG, IgM, Total)	1250
34.11.002		IgA, IgG, IgM (Smooth Muscle Antibodies, SMA, Anti-Smooth Muscle Antibodies, ASMA, IgA, IgG, IgM, Total)	1250
34.11.003		IgA, IgG, IgM (-LKM), (Antibodies to Liver/Kidney Microsome, aLKM, LKM, IgA, IgG, IgM, Total)	1250

34.11.004		IgG (- 2, 2-3, Sp100, PML, gp210, LKM-1, LC-1, SLA/LP, SSA/Ro-52), (Autoimmune Disease Liver Panel: AMA-M2, M2-3E (BPO), Sp100, PML, gp210, LKM-1, LC-1, SLA/LP, SSA/	3300
34.11.005		IgG (-ASGPR) (Autoantibodies Against Asialoglycoprotein Receptor, Anti-ASGPR, IgG)	1740
34.11.006		(Markers of Autoimmune Hepatitis)	1250
34. (Diagnosis of Autoimmune Diseases) / 34.12. (Autoimmune Neurological Diseases)			
34.12.001		IgA, IgG, IgM 4, (, NMO) (Aquaporin-4Receptor Antibodies, anti-AQP4, Neuromyelitis Optica, NMO, IgA, IgG, IgM, Total)	2400
34.12.002		IgG () (Anti-Skeletal Muscle Antibodies, ASMA, IgG)	1300
34.12.003		IgG IgM (- : GM1; GM2-GM3-GM4; GD1a, GD1b, GD2-GD3, GT1a, GT1b, GQ1b,), (Anti-GM1 Antibodies, Anti-GQ1b Antibodies, Anti-Gangliosideantibodies, Ganglioside Antibodies Panel, Total)	4900
34.12.004		- IgG (- : Mi-2, Ku, PM-Scl 100/75; Jo1 PL-7 PL-12 EJ OJ; SRP, SSA (Ro52)) (Myositis-Specific Panel)	3200
34.12.005		(,), o (Acetylcholine Receptor Antibodies, Anti-AChR, Total)	4650
34.12.006		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))	4650
34.12.007		IgA, IgG, IgM NMDA (N- -D-) (N-Methyl-D-Aspartate Receptor Antibodies, IgA, IgG, IgM, Total)	3700
34.12.008		IgG () (Oligoclonal IgG, Cerebrospinal Fluid (CSF), Serum)	3700
34.12.009		N- 1 (Procollagen Type 1 N-terminal Propeptide, P1NP, Total)	1000
34. (Diagnosis of Autoimmune Diseases) / 34.13. (Autoimmune Lung Diseases, Autoimmune Heart Diseases)			
34.13.001		IgG (), (Anti-Heart Antibodies, IgG)	1300
34.13.002		- () (Angiotensin Converting Enzyme, ACE, Serum)	2400
34.13.003		() (Neopterin, Serum)	1700
34.13.004		() (),)	2100
34. (Diagnosis of Autoimmune Diseases) / 34.14. (Immune Thrombocytopenia, ITP)			
34.14.001		IgG , (Platelet ntibodies IgG, Indirect)	1400
35. (Allergy Research) / 35.01. (IgE, Total)			
35.01.001		E (IgE,) (Immunoglobulin Total, IgE Total)	300
35. (Allergy Research) / 35.02. IgE: (Determination of Specific IgE: Comprehensive Allergy Panels)			
35.02.001		, IgE (Panel Different Allergens, IgE)	2300
35.02.002		, IgE (Pediatric Panel, IgE)	2300
35.02.003		, IgE (Respiratory Panel, IgE)	2300
35. (Allergy Research) / 35.03. IgE: (Determination of Specific IgE: Food Allergens)			
35.03.001		, IgE (Food Allergy Panel, IgE)	2300
35.03.002		1: , , , , IgE (FP15 (F33, F49, F92, F95), Food Panel: Orange, Banana, Apple, Peach, IgE)*	500
35.03.003		2: , , , , IgE (FP50 (F84, F91, F92, F210), Food Panel: Kiwi Fruit, Mango, Banana, Pineapple, IgE)*	500

35.03.004		3: IgE (FP73 (F26, F27, F83, F88), Food Panel: Pork, Beef, Chicken Meat, Lamb, IgE)*	500
35.03.005		, IgE (Pineapple, IgE, F210)	370
35.03.006		, IgE (Orange, IgE, F33)	370
35.03.007		, IgE (Peanut, IgE, F13)	370
35.03.008		, IgE (Banana, IgE, F92)	370
35.03.009		, IgE (Lamb, IgE, F88)	370
35.03.010		- , IgE (Beta Lactoglobulin, IgE, F77)	370
35.03.011		, IgE (Beef, IgE, F27)	370
35.03.012		, IgE (Grapefruit, IgE, F209)	370
35.03.013		, IgE (Buckwheat, IgE, F11)	370
35.03.014		, IgE (Casein, IgE, F78)	370
35.03.015		, IgE (Cabbage, IgE, F216)	370
35.03.016		, IgE (Potato, IgE, F35)	370
35.03.017		, IgE (Kiwi Fruit, IgE, F84)	370
35.03.018		, IgE (Strawberry, IgE, F44)	370
35.03.019		, IgE (Milk, IgE, F2)	370
35.03.020		, IgE (Crab, IgE, F23)	370
35.03.021		, IgE (Shrimp, IgE, F24)	370
35.03.022		, IgE (Chicken Meat, IgE, F83)	370
35.03.023		, IgE (Lemon, IgE, F208)	370
35.03.024		, IgE (Mango, IgE, F91)	370
35.03.025		, IgE (Carrot, IgE, F31)	370
35.03.026		, IgE (Oat, IgE, F7)	370
35.03.027		, IgE (Baker's Yeast, IgE, F45)	370
35.03.028		, IgE (Peach, IgE, F95)	370
35.03.029		, IgE (Brewer's Yeast, IgE, F403)	370
35.03.030		, IgE (Common Millet, IgE, F55)	370
35.03.031		, IgE (Wheat, IgE, F4)	370
35.03.032		, IgE (Rice, IgE, F9)	370
35.03.033		, IgE (Pork, IgE, F26)	370
35.03.034		, IgE (Celery, IgE, F85)	370
35.03.035		, IgE (Soybean, IgE, F14)	370
35.03.036		, IgE (Tomato, IgE, F25)	370
35.03.037		, IgE (Codfish, IgE, F3)	370
35.03.038		, IgE (Pumpkin, IgE, F225)	370
35.03.039		, IgE (Hazelnut, IgE, F17)	370
35.03.040		, IgE (Chocolate, IgE, F105)	370
35.03.041		, IgE (Apple, IgE, F49)	370
35.03.042		, IgE (Egg White, IgE, F1)	370
35.03.043		, IgE (Egg Yolk, IgE, F75)	370
35.		(Allergy Research) / 35.04.	IgE:
		(Determination of Specific IgE: Household Allergens)	
35.04.001		: 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 albic	500
35.04.002		Penicillium notatum, IgE (Penicillium notatum, IgE, M1)	370
35.04.003		Cladosporium herbarum, IgE (Cladosporium herbarum, IgE, M2)	370
35.04.004		Aspergillus fumigatus, IgE (Aspergillus fumigatus, IgE, M3)	370
35.04.005		Candida albicans, IgE (Candida albicans, IgE, M5)	370
35.04.006		Alternaria tenuis, IgE (Alternaria tenuis, IgE, M6)	370
35.04.007		/H1-Greer, IgE (House Dust – Greer, IgE, H1)	370
35.04.008		/H2-Stier, IgE (House Dust – Hollister-Stier, IgE, H2)	370

35.04.009		Dermatophagoides pteronyssinus (D1), IgE (Dermatophagoides pteronyssinus, IgE, D1)	370
35.04.010		Dermatophagoides farinae (D2), IgE (Dermatophagoides farinae, IgE, D2)	370
35.04.011		Dermatophagoides microceras (D3), IgE (Dermatophagoides microceras, IgE, D3)	370
35. (Allergy Research) / 35.05. (Determination of Specific IgE: Allergens of Animal Origin)			IgE:
35.05.001		: , IgE (EP70 (E6, E82, E84, E87, E88), Animal Panel: Guinea Pig Epithelium, Rabbit Epithelium, Hamster Epithelium, Rat, Mouse, IgE)*	900
35.05.002		, IgE (Cat Dander-Epithelium, IgE, E1)	300
35.05.003		, IgE (Dog Epithelium, IgE, E2)	370
35.05.004		, IgE (Cockroach, IgE, I6)	370
35.05.005		, IgE (Guinea Pig Epithelium, IgE, 6)	370
35.05.006		, IgE (Budgerigar Feathers, IgE, 78)	370
35.05.007		, IgE (Sheep Epithelium, IgE, 81)	370
35.05.008		, IgE (Chicken Feathers, IgE, 85)	370
35. (Allergy Research) / 35.06. (Determination of Specific IgE: Allergens of Plant Pollen)			IgE:
35.06.001		: , IgE (GP1 (G3, G4, G5, G6, G8), Grass Panel 1: Orchard Grass, Meadow Fescue, Perennial Rye Grass, Timothy Grass, June Grass (Kentucky Bluegrass), IgE)*	500
35.06.002		: , IgE (GP3 (G1, G5, G6, G12, G13), Grass Panel: Sweet Vernal Grass, Perennial Rye Grass, Timothy Grass, Cultivated Rye Grass, Velvet Grass, I	500
35.06.003		: , IgE (TP9 (T2, T4, T12, T3, T7), Tree Panel: Alder, Hazelnut, Willow, Birch, Oak, IgE)*	500
35.06.004		: , IgE (WP1 (W1, W6, W9, W10, W11), Weed Panel: Common Ragweed, Mugwort, English Plantain, Lamb's Quarters, Russian Thistle, IgE)*	500
35.06.005		, IgE (Birch, IgE, 3)	370
35.06.006		, IgE (Cottonwood, IgE, T14)	370
35.06.007		, IgE (Timothy Grass, IgE, G6)	370
35.06.008		, IgE (Mugwort, IgE, W6)	370
35.06.009		, IgE (Wormwood, IgE, W5)	370
35. (Allergy Research) / 35.07. (Determination of specific IgE: professional allergens)			IgE:
35.07.001		, IgE (Latex, IgG, K82)	370
35. (Allergy Research) / 35.08. (Determination of Specific IgG: Food Allergens)			IgG:
35.08.001		1: , IgG (FP15 (F33, F49, F92, F95), Food Panel: Orange, Banana, Apple, Peach, IgG)*	500
35.08.002		2: , IgG (FP50 (F84, F91, F92, F210), Food Panel: Kiwi Fruit, Mango, Banana, Pineapple, IgG)*	500
35.08.003		3: , IgG (FP73 (F26, F27, F83, F88), Food Panel: Pork, Beef, Chicken Meat, Lamb, IgG)*	500
35.08.004		G (IgG) (Basic Food Profile, IgG)	9000
35.08.005		, IgG (Pineapple, IgG, F210)	370
35.08.006		, IgG (Orange, IgG, F33)	370
35.08.007		, IgG (Peanut, IgG, F13)	370
35.08.008		, IgG (Banana, IgG, F92)	370
35.08.009		, IgG (Lamb, IgG, F88)	370
35.08.010		- , IgG (Beta Lactoglobulin, IgG, F77)	370
35.08.011		(F27), - IgG (Beef, IgG, F27)	370

35.08.012		(F209), -	IgG (Grapefruit, IgG, F209)	370
35.08.013		(F11), -	IgG (Buckwheat, IgG, F11)	370
35.08.014		(F78), -	IgG (Casein, IgG, F78))	370
35.08.015		(F216), -	IgG (Cabbage, IgG, F216)	370
35.08.016		(F35), -	IgG (Potato, IgG, F35)	370
35.08.017		(F84), -	IgG (Kiwi Fruit, IgG, F84)	370
35.08.018		(F44), -	IgG (Strawberry, IgG, F44)	370
35.08.019		(F2), -	IgG (Milk, IgG, F2)	370
35.08.020		(F24), -	IgG (Shrimp, IgG, F24)	370
35.08.021		(F83), -	IgG (Chicken Meat, IgG, F83)	370
35.08.022		(F208), -	IgG (Lemon, IgG, F208)	370
35.08.023		(F91), -	IgG (Mango, IgG, F91)	370
35.08.024		(F31), -	IgG (Carrot, IgG, F31)	370
35.08.025		(F7), -	IgG (Oat, IgG, F7)	370
35.08.026		(F45), - F45)	IgG (Baker's Yeast, IgG,	370
35.08.027		(F95), -	IgG (Peach, IgG, F95)	370
35.08.028		(F403), - F403)	IgG (Brewer's Yeast, IgG,	370
35.08.029		(F55), -	IgG (Common Millet, IgG, F55)	370
35.08.030		(F4), -	IgG (Wheat, IgG, F4)	370
35.08.031		(F9), -	IgG (Rice, IgG, F9)	370
35.08.032		(F26), -	IgG (Pork, IgG, F26)	370
35.08.033		(F14), -	IgG (Soybean, IgG, F14)	370
35.08.034		(F25), -	IgG (Tomato, IgG, F25)	370
35.08.035		(F3), -	IgG (Codfish, IgG, F3)	370
35.08.036		(F225), -	IgG (Pumpkin, IgG, F225)	370
35.08.037		(F17), -	IgG (Hazelnut, IgG, F17)	370
35.08.038		(F105), -	IgG (Chocolate, IgG, F105)	370
35.08.039		(F49), -	IgG (Apple, IgG, F49)	370
35.08.040		(F1), -	IgG (Egg White, IgG, F1)	370
35.08.041		(F75), -	IgG (Egg Yolk, IgG, F75)	370
35. (Allergy Research) / 35.09. IgG:				
(Determination of Specific IgG: Household Allergens)				
35.09.001			: 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 albic	500
35.09.002		Penicillium notatum (1), - notatum, IgG, M1)	IgG (Penicillium	370
35.09.003		Cladosporium herbarum (2), - (Cladosporium herbarum, IgG, M2)	IgG	370
35.09.004		Aspergillus fumigatus (3), - fumigatus, IgG, M3)	IgG (Aspergillus	370
35.09.005		Alternaria tenuis (6), - tenuis, IgG, M6)	IgG (Alternaria	370
35.09.006		Dermatophagoides pteronyssinus (D1), - (Dermatophagoides pteronyssinus, IgG, D1)	IgG	370
35.09.007		Dermatophagoides farinae (D2), - (Dermatophagoides farinae, IgG, D2)	IgG	370
35.09.008		Dermatophagoides microceras (D3), - (Dermatophagoides microceras, IgG, D3)	IgG	370
35.09.009		/Greer (1), - Greer, IgG, H1)	IgG (House Dust –	370
35.09.010		/Stier (2), - Hollister-Stier, IgG, H2)	IgG (House Dust –	370
35. (Allergy Research) / 35.10. IgG:				
(Determination of Specific IgG: allergens of Animal Origin)				

35.10.001		(1), - IgG (Cat Dander-Epithelium, IgG, E1)	370
35.10.002		(2), - IgG (Dog Epithelium, IgG, E2)	370
36. (Study of Sperm) / 36.03. (Antispermatozoal Antibodies)			
36.03.001		(Anti-Spermatozoa Antibodies, ASA, Serum)	500
36.03.002		(Anti-Spermatozoa Antibodies, ASA, Semen)	1450
37. (Cytological Research) / 37.01. (Liquid-Based Cytology, LBC, Pap Stain)			
37.01.001		(NovaPrep ®) (Liquid Based Cytology, LBC. Cervical Screening (Papanicolaou Stain (Pap Stain), Technology NovaPrep ®))	1450
37. (Cytological Research) / 37.02. (Papanicolaou Stain) ? -			
37.02.001		() (Cytological Examination: Cervix, Pap-test)	900
37.02.002		() (Cytological Examination: Different Biomaterial exopt Cervix, Pap-test)	900
37. (Cytological Research) / 37.03. (Romanowsky-Giemsa Stain) -			
37.03.001		(Cytological Examination of Material Obtained during Surgical Procedures and Other Urgent Research)	370
37.03.002		(Examination of Scrapings and Prints of Skin and Mucous Membranes)	370
37.03.003		(Examination of Scrapings and Prints Tumor and Tumor Like Formations)	500
37.03.004		(Examination of Endoscopic Material)	500
37.03.005		() (Examination of Bronchial Washouts)	500
37.03.006		(Examination of Scrapings: Cervix and Cervical Canal)	300
37.03.007		(The Bethesda System ? TBS) (Cytological Examination of Cervical Epithelium with Description on The Bethesda System, TBS)	550
37.03.008		() (Examination of Uterine Aspiration)	370
37.03.009		() (Examination of Imprint Intrauterine Device, IUD)	370
37.03.010		(Examination of Urine)	370
37.03.011		(Examination of Breast Discharge)	370
37.03.012		(Examination of Transudates, Exudates, Secrets)	370
37.03.013		(Examination of Sputum)	370
37.03.014		(Examination of Punctates: Breast)	370
37.03.015		(Examination of Punctates: Skin)	370
37.03.016		(Examination of Punctates: Other Organs and Tissues)	1400
37.03.017		(The Bethesda System for Reporting Thyroid Cytopathology (TBSRTC), Fine-Needle Aspiration (FNA))	600
37.03.018		() (Cytological Examination: Scrapings (Smear) of Nasal Mucous Membrane (1 Localization))	250

37.03.019		Helicobacter pylori (Examination of Endoscopic Material: Presence of Helicobacter pylori)	600
37. (Cytological Research) / 37.04. (Cytological Examination of Early Stained Slides with Cytological Smear)			
37.04.001		() (Consultation of Finished Cytological Preparations (1 Glass))	500
38. (Histological Examination) / 38.01. (Histology, hematoxylin and eosin stain (H&E stain, HE stain))			
38.01.001		, ()	1500
38.01.002		Helicobacter pylori () (Helicobacter pylori Associated Gastritis)	3300
38.01.003		Helicobacter pylori () (Helicobacter pylori, Mucus, Histochemical Study)	1500
38.01.005		(,) (Estrogen and Progesterone Receptors, Immunohistochemical Study (Fixed Biomaterial in Paraffin Block))	5500
38.01.006		(1 + 1) (Consultation of Finished Histological Preparations (1 Glass + 1 Block))	1300
38.01.007		Ki-67 (MIB-1) , Ki-67 () (Ki-67 (MIB-1) Expression, Assessment of Proliferative Activity by Expression Ki-67, Imm	4400
38.01.008		Ki-67 (MIB-1) , Ki-67 () (Ki-67 (MIB-1) Expression, Assessment of Proliferative Activity by Expression Ki-67, Immun	4400
38.01.009		HER2/neu , HER2- , () (HER2/neu Expression, HER2 Status, Immunohistochemical Study (Fixed Biomaterial in Formalin Buffer))	4400
38.01.010		HER2 in situ (FISH) (Determination of HER2 Status of Tumor, Fluorescence In Situ Hybridization)	18000
38.01.011		HER2/neu , HER2- , () (HER2/neu Expression, HER2 Status, Immunohistochemical Study (Fixed Biomaterial in Paraffin Block))	4400
38.01.012		(CD138) (, -) (Chronic Endometritis, Identification of Plasma Cells CD138, Immunohistochemical Diagnosis (Fixed Bioma	4400
38.01.013		(CD138) (, -) (Chronic Endometritis, Identification of Plasma Cells CD138, Immunohistochemical Diagnosis (Fixed Biomate	4400
38.01.014		: p16INK4a (,) (Early Diagnosis Marker of Dysplasia with High Risk Malignancy: p16INK4a, Immuno	3900
38.01.016		- : p16INK4a + Ki-67 (,) (Cervical Cancer – Study of Two Ma	6500
38.01.017		- : p16INK4a + Ki-67 (,) (Cervical Cancer – Study of Two Mark	6500
38.01.018		- (,) (Breast Cancer, Immunohistochemistry, IHC (Formalin-Fixed Biomaterial))	13000

38.01.019		() (Breast Cancer, Immunohistochemistry, IHC (Paraffin-Embedded Tissue Block))	13000
38.01.020		() (Immunohistochemical diagnosis of lymphoproliferative diseases (Tissue Embedded in Paraffin Block))	15500
38.01.021		() (Immunohistochemical diagnosis of lymphoproliferative diseases (Fixed Biomaterial in Formalin Buffer))	15500
38.01.022		() (Immunohistochemical diagnosis in cancer)	15500
38.01.023		() (Immunohistochemical diagnosis in c	15500
38.01.004		(Estrogen and Progesterone Receptors, Immunohistochemical Study)	0
38.01.015		: p16INK4a () (Early Diagnosis Marker of Dysplasia with High Risk Malignancy: p16INK4a, Immunohi	0
39. (Cytogenetic Research) / 39.01.			
39.01.001		() (Karyotype)	5000
40. (Genetic Predispositions) / 40.01. VIP-			
(Genetic VIP-Profiles)			
40.01.001		(ACE, AGT, F2, F5, ApoE, MTHFR, MTRR, MTR, NOS3, FGB, ITGB3, ITGA2, GP1BA, CYP2C9, GSTT1, GSTM1, GSTP1, LCT, CALCR, COL1A1, CFTR, GJB2, PAH, SMN, RHD, HLA II, VKORC1, NAT2, BRCA1, BRCA2, DLG5, NOD2, POMC, U	84000
40.01.002		(ACE, AGT, F2, F5, ApoE, MTHFR, MTRR, MTR, NOS3, FGB, ITGB3, ITGA2, GP1BA, CYP2C9, GSTT1, GSTM1, GSTP1, LCT, CALCR, COL1A1, CFTR, GJB2, PAH, SMN, RHD, HLA II, VKORC1, NAT2, BRCA1, BRCA2, DLG5, NOD2, POMC, U	61000
40.01.004		() (ACE, AGT, F2, F5, ApoE, MTHFR, MTRR, MTR, NOS3, FGB, ITGB3, ITGA2, GP1BA, CYP2C9, GSTT1, GSTM1, GSTP1, LCT, CALCR, COL1A1, CFTR, GJB2, PAH, SMN, RHD, HLA II, VKORC1, NAT2, BRCA1, BRCA2, DLG	74500
40.01.005		() (ACE, AGT, F2, F5, ApoE, MTHFR, MTRR, MTR, NOS3, FGB, ITGB3, ITGA2, GP1BA, CYP2C9, GSTT1, GSTM1, GSTP1, LCT, CALCR, COL1A1, CFTR, GJB2, PAH, SMN, RHD, HLA II, VKORC1, NAT2, BRCA1, BRCA2, DLG5, NOD2,	65000
40. (Genetic Predispositions) / 40.02.			
(Lifestyle and Genetic Factors)			
40.02.001		(ACE, AGT, F2, F5, ApoE, MTHFR, MTRR, MTR, NOS3, GSTT1, GSTM1, GSTP1, NAT2, ITGB3, ITGA2, GP1BA, FGB, UGT1A1, DAT, OPRM1, ANKK1, ALDH2, ADH2) (Healthy Lifestyle (Genes ACE, AGT, F2, F5, ApoE, MTHFR, MTRR, MTR, NOS3, GSTT1, GSTM1,	39000
40.02.002		(F2, F5, ApoE, MTHFR, MTRR, MTR, ACE, AGT, GSTT1, GSTM1, GSTP1, NAT2, UGT1A1, AR, CFTR; AZF-) (I'm Healthy (for Men) (Genes F2, F5, ApoE, MTHFR, MTRR, MTR, ACE, AGT, GSTT1, GSTM1, GSTP1, NAT2, UGT1A1, AR, CFTR; AZF-Region))	31500
40.02.003		(F2, F5, ApoE, MTHFR, MTRR, MTR, ACE, AGT, GSTT1, GSTM1, GSTP1, NAT2, BRCA1, BRCA2, UGT1A1) (I'm Healthy (for Women) (Genes F2, F5, ApoE, MTHFR, MTRR, MTR, ACE, AGT, GSTT1, GSTM1, GSTP1, NAT2, BRCA1, BRCA2, UGT1A1))	27500

40.02.004		(ACE, AGT, F2, F5, ApoE, MTHFR, MTRR, MTR, NOS3, ITGB3, ITGA2, GP1BA, FGB) (Higher Physical Activity: Risk Assessment (Genes ACE, AGT, F2, F5, ApoE, MTHFR, MTRR, MTR, NOS3, ITGB3, ITGA2, GP1	17500
40.02.005		(ACE, AGT, F2, F5, ApoE, MTHFR, MTRR, MTR, NOS3, ITGB3, ITGA2, GP1BA, FGB) ((Higher Physical Activity: Risk Assessment (Genes ACE, AGT, F2, F5, Apo	15000
40.02.006		(MTHFR, MTRR, MTR, F2, F5) (Preparation for Surgery (Genes MTHFR, MTRR, MTR, F2, F5))	3700
40.02.007		(MTHFR, MTRR, MTR, F2, F5) ((Preparation for Surgery (Genes MTHFR, MTRR, MTR, F2, F5) (without Description))	2000
40.02.008		(F2, F5) (Risk of Oral Contraceptives, Ocs (Genes F2, F5))	2100
40.02.009		(F2, F5) ((Risk of Oral Contraceptives, OCs (Genes F2, F5) (without Description))	1900
40.02.010		(GSTT1, GSTM1, GSTP1, NAT2) (Risk of Cancer in Smoking (Genes GSTT1, GSTM1, GSTP1, NAT2))	9900
40.02.011		(GSTT1, GSTM1, GSTP1, NAT2) ((Risk of Cancer in Smoking (Genes GSTT1, GSTM1, GSTP1, NAT2) (without Description))	8100
40.02.012		(GSTT1, GSTM1, GSTP1, NAT2) (Tanning Risks: Protection of Skin (Genes GSTT1, GSTM1, GSTP1, NAT2))	9900
40.02.013		(GSTT1, GSTM1, GSTP1, NAT2) ((Tanning Risks: Protection of Skin (Genes GSTT1, GSTM1, GSTP1, NAT2) (without Description))	8100
40.02.014		(GSTT1, GSTM1, GSTP1, NAT2) (Intake of Grilled and Smoked Foods and Cancer Risk (Genes GSTT1, GSTM1, GSTP1, NAT2))	9900
40.02.015		(GSTT1, GSTM1, GSTP1, NAT2) ((Intake of Grilled and Smoked Foods and Cancer Risk (Genes GSTT1, GSTM1, GSTP1, NAT2) (without Description))	8100
40.02.016		(LCT) (Lactose Intolerance in Adulthood (Gene LCT))	1500
40. (Genetic Predispositions) / 40.03.			
(Reproductive Health)			
40.03.001		(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))	20000
40.03.002		(F2, F5, MTHFR, MTRR, MTR, ACE, AGT, RHD) (Want to Become a Mother: Pregnancy Complications (Genes F2, F5, MTHFR, MTRR, MTR, ACE, AGT, RHD))	15000
40.03.003		(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))	13000
40.03.004		(F2, F5) (Thrombotic Tendency in Pregnancy: Minimum (Genes F2, F5))	2300
40.03.005		(F2, F5) ((Thrombotic Tendency in Pregnancy: Minimum (Genes F2, F5) (without Description))	1900
40.03.006		(ACE, AGT, MTHFR, MTRR, MTR, F2, F5) (Gestosis and Placental Insufficiency (Genes ACE, AGT, MTHFR, MTRR, MTR, F2, F5))	9000
40.03.007		(ACE, AGT, MTHFR, MTRR, MTR, F2, F5) ((Gestosis and Placental Insufficiency (Genes ACE, AGT, MTHFR, MTRR, MTR, F2, F5) (without Description))	8000

40.03.008		(MTHFR, MTRR, MTR, F2, F5) (Habitual Miscarriage, Thrombotic Tendency in Pregnancy: Extended Panel (Genes MTHFR, MTRR, MTR, F2, F5))	4000
40.03.009		(MTHFR, MTRR, MTR, F2, F5) (Habitual Miscarriage, Thrombotic Tendency in Pregnancy: Extended Panel (3000
40.03.010		(MTHFR, MTRR, MTR) (Isolated Malformations in Fetus (Genes MTHFR, MTRR, MTR))	2800
40.03.011		(MTHFR, MTRR, MTR) (-) (Isolated Malformations in Fetus (Genes MTHFR, MTRR, MTR) (without Description))	2200
40.03.012		(F2, F5) (Thrombotic Complications of Ovulation Induction (Genes F2, F5))	2200
40.03.013		(F2, F5) (-) (Thrombotic Complications of Ovulation Induction (Genes F2, F5) (without Description))	1800
40.03.014		(). CYP21OHB, 9 . . (Congenital Adrenal Hyperplasia (CAH), Gene CYP21OHB, 9 Freq. Mut.)	15500
40.03.015		e (F2, F5) (Risk of Oral Contraceptives, Ocs (Genes F2, F5))	2200
40.03.016		e (F2, F5) (-) (Risk of Oral Contraceptives, OCs (Genes F2, F5) (without Description))	1800
40.03.017		(MTHFR, MTRR, MTR) (Folic Acid Metabolism (Genes MTHFR, MTRR, MTR))	2800
40.03.018		(MTHFR, MTRR, MTR) (-) (Folic Acid Metabolism (Genes MTHFR, MTRR, MTR) (without Description))	2200
40.03.019		/ (BRCA1, BRCA2) (Hereditary Breast and/or Ovarian Cancer, HBOC (Genes BRCA1, BRCA2))	4300
40.03.020		/ (BRCA1, BRCA2) (-) (Hereditary Breast and/or Ovarian Cancer, HBOC (Genes BRCA1, BRCA2) (without Description))	3700
40.03.021		/ BRCA1, BRCA2, CHEK2, NBN (Hereditary Breast and/or Ovarian Cancer)	8000
40.03.022		(AR, CFTR; AZF- ;) (Male Sterility (Genes AR, CFTR; AZF-Region; Karyotype))	16300
40.03.023		(AR, CFTR; AZF-) (Genetic Factors of Male Infertility (Genes AR, CFTR; AZF-Region))	13500
40.03.024		: (AZF-) (Impairment of Spermatogenesis: Full Panel (AZF-Region))	8900
40.03.025		: (AZF-) (-) (Impairment of Spermatogenesis: Full Panel (AZF-Region) (without Description))	9000
40. (Genetic Predispositions) / 40.04. HLA-			
- (HLA (Human Leukocyte Antigens) Typing and Rh factor)			
40.04.001		- (Rh factor Definition)	4000
40.04.002		- (RH factor Genotype)	8000
40.04.003		- (-) (Genotype of RH factor Definition (without Description))	8000
40.04.004		- HLA-B27 (Molecular Genetic Testing HLA-B27)	900
40.04.005		HLA II (DRB1, DQA1, DQB1) (System Human Leukocyte Antigen (HLA) Class II, Typing (Genes DRB1, DQA1, DQB1))	6000
40.04.006		HLA II - DRB1 (System Human Leukocyte Antigen (HLA) Class II, Typing)	1600
40. (Genetic Predispositions) / 40.05.20.5			
(Blood-Clotting Sequence)			

40.05.001		: (F2, F5, MTHFR, MTRR, MTR) (Thrombosis: Advanced Panel (Genes F2, F5, MTHFR, MTRR, MTR))	3200
40.05.002		: (F2, F5, MTHFR, MTRR, MTR) (-) (Thrombosis: Advanced Panel (Genes F2, F5, MTHFR, MTRR, MTR) (without Description))	3000
40.05.003		(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))	5000
40.05.004		(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, F	4000
40.05.005		: (F2, F5) (Thrombosis: Minimum (Genes F2, F5))	2200
40.05.006		: (F2, F5) (-) (Thrombosis: Minimum (Genes F2, F5) (without Description))	2000
40.05.007		(FGB) (Fibrinogen (Gene FGB))	2700
40.05.008		(FGB) (-) (Fibrinogen (Gene FGB) (without Description))	2700
40.05.009		(MTHFR, MTRR, MTR) (Hyperhomocysteinemia (Genes MTHFR, MTRR, MTR))	2700
40.05.010		(MTHFR, MTRR, MTR) (-) (Hyperhomocysteinemia (Genes MTHFR, MTRR, MTR) (without Description))	2200
40.05.011		(ITGA2, GP1BA) (Platelet Hyperaggregation (Genes ITGA2, GP1BA))	2800
40.05.012		(ITGA2, GP1BA) (-) (Platelet Hyperaggregation (Genes ITGA2, GP1BA) (without Description))	2500
40.05.013		(ITGB3) (Platelet Fibrinogen Receptor (Gene ITGB3))	1800
40.05.014		(ITGB3) (-) (Platelet Fibrinogen Receptor (Gene ITGB3) (without Description))	1500
40. (Genetic Predispositions) / 40.06. (Heart and Blood Vessel (Cardiovascular) Diseases)			
40.06.001		- (ACE, AGT, F2, F5, ApoE, MTHFR, MTRR, MTR, NOS3, ITGB3, ITGA2, GP1BA, FGB) (Cardio-Vascular Diseases (Genes ACE, AGT, F2, F5, ApoE, MTHFR, MTRR, MTR, NOS3, ITGB3, ITGA2, GP1BA, FGB))	15700
40.06.002		, (ACE, AGT, NOS3) (Arterial Hypertension: Full Panel (Genes ACE, AGT, NOS3))	4150
40.06.003		, (ACE, AGT, NOS3) (-) (Arterial Hypertension: Full Panel (Genes ACE, AGT, NOS3) (without Description))	3700
40.06.004		, (ACE, AGT) (Arterial Hypertension, Renin-Angiotensin System Disorder (Genes ACE, AGT))	2700
40.06.005		, (ACE, AGT) (-) (Arterial Hypertension, Renin-Angiotensin System Disorder (Genes ACE, AGT) (without Description))	2700
40.06.006		NO-, (NOS3) (Arterial Hypertension, Endothelial NO-Synthase Disturbance (Gene NOS3))	1800
40.06.007		NO-, (NOS3) (-) (Arterial Hypertension, Endothelial NO-Synthase Disturbance (Gene NOS3) (without Description))	1500
40.06.008		, (ITGA2, GP1BA, ACE, AGT, NOS3, ApoE) (Ischemic Heart Disease, Myocardial Infarction (Genes ITGA2, GP1BA, ACE, AGT, NOS3, ApoE))	7800

40.06.009		(ITGA2, GP1BA, ACE, AGT, NOS3, ApoE) (-) (Ischemic Heart Disease, Myocardial Infarction (Genes ITGA2, GP1BA, ACE, AGT, NOS3, ApoE) (without Description))	7500
40.06.010		(ITGB3, ITGA2, GP1BA, FGB) (Ischemic Stroke (Genes ITGB3, ITGA2, GP1BA, FGB))	6000
40.06.011		(ITGB3, ITGA2, GP1BA, FGB) (-) (Ischemic Stroke (Genes ITGB3, ITGA2, GP1BA, FGB) (without Description))	5500
40.06.012		(ApoE) (Atherosclerosis (Gene ApoE))	3000
40. (Genetic Predispositions) / 40.07. (Gastrointestinal Diseases)			
40.07.001		(DLG5, NOD2, OCTN1, OCTN2) (Crohn's Disease (Genes DLG5, NOD2, OCTN1, OCTN2))	5300
40. (Genetic Predispositions) / 40.08. (Central Nervous System (CNS) Diseases)			
40.08.001		(ApoE) (Alzheimer's Disease (Gene ApoE))	3000
40. (Genetic Predispositions) / 40.09. (Cancer Diseases)			
40.09.001		(MTHFR, MTRR, MTR, GSTT1, GSTM1, GSTP1, NAT2, BRCA1, BRCA2, CHEK2, NBS1) (Cancer in Women (Genes MTHFR, MTRR, MTR, GSTT1, GSTM1, GSTP1, NAT2, BRCA1, BRCA2, CHEK2, NBS1))	19000
40.09.002		(MTHFR, MTRR, MTR, GSTT1, GSTM1, GSTP1, NAT2, AR) (Cancer in Men (Genes MTHFR, MTRR, MTR, GSTT1, GSTM1, GSTP1, NAT2, AR))	14000
40.09.003		(GSTT1, GSTM1, GSTP1, NAT2) (Cancer Associated with Environmental Toxins (Genes GSTT1, GSTM1, GSTP1, NAT2))	8400
40.09.004		BRCA1, BRCA2) (Hereditary Breast and/or Ovarian Cancer, HBOC (Genes BRCA1, BRCA2))	4500
40.09.005		BRCA1, BRCA2) (-) (Hereditary Breast and/or Ovarian Cancer, HBOC (Genes BRCA1, BRCA2) (without Description))	3800
40.09.006		BRCA- : BRCA1, BRCA2) (Hereditary Breast Cancer In Men: Cancer of Breast, Pancreatic, Prostate, Testicular Cancer (Genes BRCA1, BRCA2))	4400
40.09.007		BRCA- : BRCA1, BRCA2) (-) (Hereditary Breast Cancer In Men: Cancer of Breast, Pancreatic, Prostate	4000
40.09.008		BRCA2, CHEK2, NBN (Hereditary Breast and/or Ovarian Cancer)	7900
40.09.009		(10, 11, 13, 14, 15 RET) (Familial Medullary Thyroid Cancer (Exons 10, 11, 13, 14, 15 Gene RET))	13000
40.09.010		2B (RET) (Multiple Endocrine Neoplasia Type 2B (Gene RET))	4100
40.09.011		2A (10, 11 RET) (Multiple Endocrine Neoplasia Type 2A (Exons 10, 11 Gene RET))	8600
40.09.012		Ph'- (): 617V/617F JAK2 (Marker of Ph-negative Chronic Myeloproliferative Disorders (cMPD): quantifica	4500
40.09.013		Ph'- (): 617F JAK2 (Marker of Ph-negative Chronic Myeloproliferative Disorders (cMPD): qualitative assessment of presence of	3600
40. (Genetic Predispositions) / 40.11. (Metabolic Disease, Metabolic Disturbance)			

40.11.001		HLA II (DRB1, DQA1, DQB1) (Hereditary Predisposition to Diabetes Type 1 (Insulin-Dependent Diabetes), HLA Class II (Genes DRB1, DQA1, DQB1))	6400
40.11.002		HLA II (DQA1, DQB1)	6000
40.11.003		ADAMTS9, KCNJ11, KCNQ1, PPARG (Non-Insulin-Dependent Diabetes Type 2)	10300
40.11.004		ADAMTS9, KCNJ11, KCNQ1, PPARG () (Non-Insulin-Dependent Diabetes Type 2 (without Description))	7600
40.11.005		(UGT1A1) (Gilbert's Syndrome (Gene UGT1A1))	4000
40.11.006		: (CALCR, COL1A1, VDR) (Osteoporosis: Full Panel (Genes CALCR, COL1A1, VDR))	5400
40.11.007		: (CALCR, COL1A1, VDR) () (Osteoporosis: Full Panel (Genes CALCR, COL1A1, VDR) (without Description))	4400
40.11.008		: (CALCR, COL1A1) (Osteoporosis: Abridged Panel (Genes CALCR, COL1A1))	3700
40.11.009		: (CALCR, COL1A1) () (Osteoporosis: Abridged Panel (Genes CALCR, COL1A1) (without Description))	3500
40.11.010		: D (VDR) (Osteoporosis, Vitamin D Receptor (VDR) (Gene VDR))	1500
40.11.011		: D (VDR) () (Osteoporosis, Vitamin D Receptor (VDR) (Gene VDR) (without Description))	1400
40.11.012		(MTHFR, MTRR, MTR) (Folic Acid Metabolism (Genes MTHFR, MTRR, MTR))	2600
40.11.013		(MTHFR, MTRR, MTR) () (Folic Acid Metabolism (Genes MTHFR, MTRR, MTR) (without Description))	2200
40.11.014		(LCT) (Adult Lactase Deficiency (Gene LCT))	1700
40.11.015		, I (HFE) (Hemochromatosis Type 1 (Gene HFE))	3000
40. (Genetic Predispositions) / 40.12. (Pharmacogenetics)			
40.12.001		(ITGB3) (Aspirin and Plavix. Genetic Marker of Resistance to Antiplatelet	1600
40.12.002		: (CYP2C9, VKORC1, CYP4F2, GGCX) (Warfarin Dose Panel (Genes CYP2C9, VKORC1, CYP4F2, GGCX))	12400
40.12.003		:	5500
40.12.004		N- 2 (Hydr	5500
40.12.005		, , ATII.	2800
40.12.006		(UGT1A1) (Metabolism of Irinotecan (Gene UGT1A1))	4000
40.12.007		/ II:	3000

40.12.008			2700
40.12.009			2800
40.12.010		-1- -S- (GSTM1) (Penicillamine. Genetic Factors of En	5100
40.12.011		(ApoE) (Statins. Genetic Factors of Decrease or Enhance Clinical Effectiveness in Use of	3000
40.12.012		() . _	4400
40.12.013		: _ , , N-	3000
40.12.014		-28 (-28), () (Interleukin 28 Beta IL28B, Genotyping (Study of Genetic Markers Determining Effectiveness	1000
40.12.015		- (?-). CYP2D6. (?-Adrenergic Blockers. Gene CYP2D6. Pharmacogenetics)	10000
40. (Genetic Predispositions) / 40.13. (Detoxification System of Xenobiotics and Carcinogens)			
40.13.001		YP2D6 (YP2D6) (Cytochrome YP2D6 (Gene YP2D6))	10000
40.13.002		CYP2C9 (CYP2C9) (Cytochrome CYP2C9 (Gene CYP2C9))	3000
40.13.003		(GSTT1, GSTM1, GSTP) (Glutathionetransferases (Genes GSTT1, GSTM1, GSTP))	5200
40.13.004		N- 2 (NAT2) (N-Acetyltransferase 2 (Gene NAT2))	4500
40.13.005		(UGT1A1) (Uridine Diphosphate Glucuronyltransferase (Gene UGT1A1))	4100
40. (Genetic Predispositions) / 40.14. (Services of Physician-Geneticist)			
40.14.001		1 (7201 , 7611 , 7014 , 7030 , 125 / , 7207) (Genetic Test Results: Description of the 1-st Category Complexity)	800
40.14.002		2 (118 / , 121 / , 123 / , 131 / , 141 / , 149 / , 150 / , 115 / , 152 / , 124 / , 154 /) (Genetic Test Results: Description of the 2-nd Category Complexity)	1500
40.14.003		3 (122 / , 129 / , 120 / , 137 / , 138 / , 153 / , 151 / , 110 / , 114 / , 140 / , 7661 , 7258 , 134 / , 135 / , 136 /) (Genetic Test Results: Description of the	2800
40.14.004		4 (144 / , 143 / , 139 / , 145 / , 108 / , 19 /) (Genetic Test Results: Description of the 4-th Category Complexity)	4400
40. (Genetic Predispositions) / 40.01. (Genetic VIP-Profiles) VIP-			
40.01.003		() (ACE, AGT, F2, F5, ApoE, MTHFR, MTRR, MTR, NOS3, FGB, ITGB3, ITGA2, GP1BA, CYP2C9, GSTT1, GSTM1, GSTP1, LCT, CALCR, COL1A1, CFTR, GJB2, PAH, SMN, RHD, HLA II, VKORC1, NAT2, BRCA1, BRCA2, DLG	0

40.01.006		() (ACE, AGT, F2, F5, ApoE, MTHFR, MTRR, MTR, NOS3, FGB, ITGB3, ITGA2, GP1BA, CYP2C9, GSTT1, GSTM1, GSTP1, LCT, CALCR, COL1A1, CFTR, GJB2, PAH, SMN, RHD, HLA II, VKORC1, NAT2, BRCA1, BRCA2, DLG5, NOD2,	0
41. (Hereditary Monogenic Diseases) / 41.01. (Hereditary Monogenic Diseases)			
41.01.104		« » TRPV4, « . » . (Distal Spinal Muscular Atrophy Congenital Non-Progressive, Gene TRPV4, Hot-Point Mut.)	0
41.01.239		MEFV, . . (Familial Mediterranean Fever, FMF, Gene MEFV, Freq. Mut.)	0
41.01.264		(). LPIN1, . (Myoglobinuria Acute Recurrent Autosomal Recessive, Gene LPIN1, Mut.)	0
41.01.001		(CFTR, GJB2, PAH, SMN) (Main Hereditary Diseases (Genes CFTR, GJB2, PAH, SMN))	13600
41.01.002		, (1-). ABCA4, . . (Stargardt Disease 1, STGD1, Fundus Flavimaculatus Included, Gene ABCA4, Freq. Mut.)	8800
41.01.004		(). CYP21OHB, 9 . . (Congenital Adrenal Hyperplasia (CAH), Gene CYP21OHB, 9 Freq. Mut.)	15400
41.01.005		SLC39A4, . (Acrodermatitis Enteropathica, Gene SLC39A4, Mut.)	29600
41.01.006		GPR143, . (Albinism Ocular Type I, Gene GPR143, Mut.)	8600
41.01.007		2 . OCA2, . (Albinism Oculocutaneous Type IIA, Gene OCA2, Mut.)	97300
41.01.008		1 . TYR, . (Albinism Oculocutaneous Type IA, Gene TYR, Mut.)	17900
41.01.009		CRB1, . (Leber Congenital Amaurosis 1, LCA, Gene CRB1, Mut.)	55300
41.01.013		RPE65, . (Leber Congenital Amaurosis 1, LCA, Gene RPE65, Mut.)	38000
41.01.014		RPS19, . (Diamond-Blackfan Anemia 1, DBA1, Gene RPS19, Mut.)	17900
41.01.015		FBN2, . (Contractural Congenital Arachnodactyly, CCA, Gene FBN2, Mut.)	41600
41.01.016		/ SCN5A, . (Arrhythmogenic Right Ventricular Dysplasia 1, Gene SCN5A, Mut.)	50600
41.01.017		(-). MYH3, . . (Arthrogryposis Distal Type 2A, Gene MYH3, Freq. Mut.)	9600
41.01.018		FXN, . (Friedrich Ataxia, Gene FXN, Mut.)	17600
41.01.019		FXN, . . (Friedrich Ataxia, Gene FXN, Freq. Mut.)	4400
41.01.020		(). SLC26A2, . (Atelosteogenesis II, De la Chapelle Dysplasia, Gene SLC26A2, Mut.)	23250
41.01.021		OPA1, . (Optic Atrophy 1, Gene OPA1, Mut.)	96250
41.01.022		, 12 . . (Leber Hereditary Optic Neuropathy, LHON, Mitochondrial DNA, 12 Freq. Mut.)	13350
41.01.023		, 3 . . (Leber Hereditary Optic Neuropathy, LHON, Mitochondrial DNA, 3 Freq. Mut.)	4400
41.01.024		« » OPA1, « . » . (Optic Atrophy With Or Without Deafness, Ophthalmoplegia, Myopathy, Ataxia And Neuropathy, Gene OPA1, Hot-Point Mut.)	8600
41.01.025		OPA3, . (Optic Atrophy 1, Gene OPA3, Mut.)	7300

41.01.026		TMEM126A, . (Optic Atrophy 1, Gene TMEM126A, Mut.)	19550
41.01.027		TNFRSF6, . (Autoimmune Lymphoproliferative Syndrome, ALPS, Gene TNFRSF6, Mut.)	28100
41.01.028		« » TNFRSF6, « .» . (Autoimmune Lymphoproliferative Syndrome, ALPS, Gene TNFRSF6, Hot-Point Mut.)	6100
41.01.029		GRN, . (Aphasia Primary Progressive, Gene GRN, Mut.)	22250
41.01.030		II. Col2A1, . (Achondrogenesis Type II, Gene Col2A1, Mut.)	119335
41.01.031		I. SLC26A2, . (Achondrogenesis, Fraccaro Type, Gene SLC26A2, Mut.)	23250
41.01.032		FGFR3, . . (Achondroplasia, Gene FGFR3, Freq. Mut.)	8830
41.01.033		CNGB3, . (Achromatopsia 3, ACHM, Gene CNGB3, Mut.)	67650
41.01.034		FIG4, . (Amyotrophic Lateral Sclerosis, Gene FIG4, Mut.)	97250
41.01.035		ALSIN, . (Amyotrophic Lateral Sclerosis, Gene ALSIN, Mut.)	119330
41.01.036		VAPB, . (Amyotrophic Lateral Sclerosis, Gene VAPB, Mut.)	22250
41.01.037		VAPB, . (Amyotrophic Lateral Sclerosis, Gene VAPB, Freq. Mut.)	6000
41.01.038		BEST1, . (Best Vitelliform Macular Dystrophy, All Known Mutations, Gene BEST1, Mut.)	34800
41.01.039		ATP7B, . (Wilson Disease, Gene ATP7B, Mut.)	81450
41.01.040		ATP7B, . . (Wilson Disease, Gene ATP7B, Freq. Mut.)	7800
41.01.041		PANK2, (Neurodegeneration With Brain Iron Accumulation 1, Gene PANK2, Mut.)	30650
41.01.042		PANK2, . . (Neurodegeneration With Brain Iron Accumulation 1, Gene PANK2, Freq. Mut.)	6600
41.01.043		PRNP, . (Gerstmann-Straussler Disease, Gene PRNP, Mut.)	10600
41.01.044		EDNRB, . (Hirschprung Disease, HSCR, Gene EDNRB, Mut.)	24730
41.01.045		NTRK1, . (Hirschprung Disease, HSCR, Gene NTRK1, Mut.)	41600
41.01.046		ZEB2, . (Hirschprung Disease, HSCR, Gene ZEB2, Mut.)	39100
41.01.047		10, 11, 13, 14, 15 RET, . (Hirschprung Disease, HSCR, Exons 10, 11, 13, 14, 15 Gene RET, Mut.)	17900
41.01.048		PTEN, . (Cowden Syndrome 1, Gene PTEN, Mut.)	33350
41.01.049		PRNP, . (Creutzfeldt-Jakob Disease, Gene PRNP, Mut.)	10600
41.01.050		PTEN, . (Lhermitte-Duclos Syndrome, Gene PTEN, Mut.)	33350
41.01.051		NDP, . (Norrie Disease, Gene NDP, Mut.)	8300
41.01.052		CAV3, . (Rippling Muscle Disease, Gene CAV3, Mut.)	8300
41.01.053		GAA, . (Pompe Disease, Gene GAA, Mut.)	4400
41.01.054		CSTB, . (Progressive Myoclonic Epilepsy 1A Unverricht and Lundborg, Gene CSTB, Mut.)	12340
41.01.055		CSTB, . . (Progressive Myoclonic Epilepsy 1A Unverricht and Lundborg, Gene CSTB, Freq. Mut.)	4400

41.01.056		ABCA4, . . (Stargardt Disease 1, STGD1, Fundus Flavimaculatus Included, Gene ABCA4, Freq. Mut.)	8800
41.01.057		CNGB3, . (Stargardt Disease 1, STGD1, Fundus Flavimaculatus Included, Gene CNGB3, Mut.)	67650
41.01.058		« » ELOVL4, « » . . (Stargardt Disease 1, STGD1, Fundus Flavimaculatus Included, Gene ELOVL4, Hot-Point Mut.)	6100
41.01.059		B1. ROR2, . (Brachydactyly Type B1, Gene ROR2, Mut.)	12340
41.01.060		KRT5, . (Epidermolysis Bullosa, Gene KRT5, Mut.)	22250
41.01.061		« » LAMB3, « » . . (Epidermolysis Bullosa, Gene LAMB3, without Hot-Point Mut.)	45450
41.01.062		« » LAMB3, « » . . (Epidermolysis Bullosa, Gene LAMB3, Hot-Point Mut.)	28100
41.01.063		TBX1, . (Velocardiofacial Syndrome, Gene TBX1, Mut.)	34350
41.01.064		NTRK1, . (Congenital Insensitivity To Pain With Anhidrosis, CIPA, Gene NTRK1, Mut.)	41630
41.01.065		ADAMTSL2, . (Geleophysic Dysplasia 1, Gene ADAMTSL2, Mut.)	12340
41.01.066		FBN1, . (Geleophysic Dysplasia 1, Gene FBN1, Mut.)	216000
41.01.067		IX B, . (Hemophilia B, Gene Factor IX, Mut.)	26000
41.01.068		-IgD « » MVK, « » . . (Hyper-IgD Syndrome, Gene MVK, Hot-Point Mut.)	9000
41.01.069		-IgD CD40LG, . (Hyper-IgD Syndrome, Gene CD40LG, Mut.)	37000
41.01.070		-IgM CD40LG, . (Hyper-IgM Syndrome, Gene CD40LG, Mut.)	17900
41.01.071		13 24 SCN4A, . (Hyperkalemic Periodic Paralysis Type 2, Exons 13, 24 Gene SCN4A, Mut.)	9650
41.01.072		SCN4A, . (Hyperkalemic Periodic Paralysis Type 2, Gene SCN4A, Mut.)	81450
41.01.073		KRT1, . (Epidermolytic Hyperkeratosis, Gene KRT1, Mut.)	22250
41.01.074		KRT9, . (Epidermolytic Hyperkeratosis, Gene KRT9, Mut.)	27000
41.01.075		CAV3, . (Familial Hypertrophic Cardiomyopathy, Gene CAV3, Mut.)	8320
41.01.076		TNNT2, . (Familial Hypertrophic Cardiomyopathy, Gene TNNT2, Mut.)	41640
41.01.080		12, 18 19 SCN4A, . (Hypokalemic Periodic Paralysis Type 1, Exons 12, 18, 19 Gene SCN4A, Mut.)	12350
41.01.081		D- PHEX, . (Hypophosphatemic Vitamin D-Resistant Rickets, Gene PHEX, Mut.)	82450
41.01.082		FGFR3, . . (Hypochondroplasia, Gene FGFR3, Freq. Mut.)	8800
41.01.083		CYP1B1, . (Primary Congenital Glaucoma 3A, PCG 3A, Gene CYP1B1, Mut.)	12380
41.01.084		(). CYP1B1, . (Primary Open Angle Glaucoma 1A, POAG 1A, Gene CYP1B1, Mut.)	12380
41.01.086		HNF1B, . (Renal Cysts And Diabetes Syndrome, Gene HNF1B, Mut.)	33250
41.01.087		S H, . (Holoprosencephaly 3, Gene S H, Mut.)	14850

41.01.088		ACADM, . . (Medium-Chain Acyl-CoA Dehydrogenase Deficiency, MCAD-Deficiency, Gene ACADM, Freq. Mut.)	4460
41.01.089		SLC22A5, . (Systemic Primary Carnitine Deficiency, SPCD, Carnitine Deficiency Systemic Primary, CDSP, Gene SLC22A5, Mut.)	3850
41.01.090		SLC26A2, . (Diastrophic Dysplasia, Gene SLC26A2, Mut.)	23250
41.01.091		(). DES, . (Cardiomyopathy Dilated, Gene DES, Mut.)	24730
41.01.092		(). EYA4, . (Cardiomyopathy Dilated, Gene EYA4, Mut.)	41630
41.01.093		(). FKTN, . (Cardiomyopathy Dilated, Gene FKTN, Mut.)	41630
41.01.094		(). LMNA, . (Cardiomyopathy Dilated, Gene LMNA, Mut.)	34800
41.01.095		(). SCN5A, . (Cardiomyopathy Dilated, Gene SCN5A, Mut.)	50660
41.01.096		(). SGCD, . (Cardiomyopathy Dilated, Gene SGCD, Mut.)	34350
41.01.097		(). TAZ, . (Cardiomyopathy Dilated, Gene TAZ, Mut.)	21370
41.01.098		(). TNNT2, . (Cardiomyopathy Dilated, Gene TNNT2, Mut.)	41630
41.01.099		(). Emerine, Mut.) , . (Cardiomyopathy Dilated, X-Linked Gene	10700
41.01.100		Col2A1, . (Kniest Dysplasia, Gene Col2A1, Mut.)	119350
41.01.101		V. GARS, . (Distal Hereditary Motor Neuropathy, DHMN, Gene GARS, Mut.)	61900
41.01.102		V. BSCL2, . (Distal Hereditary Motor Neuropathy, DHMN, Gene BSCL2, Mut.)	6670
41.01.103		IGHMBP2, . (Distal Spinal Muscular Atrophy 1, DSMA1, Gene IGHMBP2, Mut.)	56550
41.01.108		SCN5A, . (Ventricular Fibrillation, Familial 1, Gene SCN5A, Mut.)	50660
41.01.109		BTK, . (Hypogammaglobulinemia and Isolated Growth Hormone Deficiency, Fleisher Syndrome, Gene BTK, Mut.)	61900
41.01.110		KRT2, . (Ichthyosis Bullosa Of Siemens, Gene KRT2, Mut.)	25950
41.01.111		FLG, . (Ichthyosis Vulgaris, Gene FLG, Freq. Mut.)	9650
41.01.112		TGM1, . (Autosomal Recessive ongenital Ichthyosis, ARCI 1, All Known Mutations, Gene TGM1, Mut.)	34800
41.01.122		GNAS, . (Progressive Osseous Heteroplasia, POH, Gene GNAS, Mut.)	34350
41.01.123		ANKH, « » . (Cranioetaphyseal Dysplasia, Gene ANKH, Hot-Point Mut.)	9650
41.01.124		ANKH, . (Cranioetaphyseal Dysplasia, Gene ANKH, Mut.)	45450
41.01.125		Gene TWIST1, Mut.) TWIST1, . (Craniosynostosis Type 2,	13900
41.01.126		Gene MSX2, Mut.) MSX2, . (Craniosynostosis Type 2,	8320
41.01.127		39 LAMA3, . (Laryngoonychocutaneous Syndrome, Laryngo-Onycho-Cutaneous Syndrome, Exon 39 Gene LAMA3, Mut.)	6670
41.01.128		GJC2, . (Leukodystrophy Hypomyelinating 2, Gene GJC2, Mut.)	25950

41.01.129		FLT4, . (Lymphedema, Gene FLT4, Mut.)	88880
41.01.130		GJC2, . (Hereditary Lymphedema Type 1C, Gene GJC2, Mut.)	25950
41.01.131		« » LMNA, « » . . (Familial Partial Lipodystrophy 2, FPLD 2, Gene LMNA, Hot-Point Mut.)	6670
41.01.132		BSCL2, . (Congenital Generalized Lipodystrophy, CGL, Type 1, Gene BSCL2, Mut.)	26950
41.01.133		LMNA, . (Familial Partial Lipodystrophy 2, Gene LMNA, Mut.)	34810
41.01.135		8, 9 LMNA, . (Mandibuloacral Dysplasia, Exons 8, 9 Gene LMNA, Mut.)	6670
41.01.136		MVK, . (Mevalonic Aciduria, Gene MVK, Mut.)	37050
41.01.137		, CYB5R3 . (Methemoglobinemia, Gene CYB5R3, Mut.)	28100
41.01.138		, CYB5R3 . . (Methemoglobinemia, Gene CYB5R3, Freq. Mut.)	4400
41.01.139		OPA3, . (3-Methylglutaconic Aciduria Type III, Gene OPA3, Mut.)	8320
41.01.140		CACNA1A, . (Familial Hemiplegic Migraine, FHM, Gene CACNA1A, Mut.)	162000
41.01.141		GDF6, . (Microphthalmia Isolated 4, Gene GDF6, Mut.)	10690
41.01.142		CRYBA4, . (Microphthalmia with Cataract, Gene CRYBA4, Mut.)	21370
41.01.143		SGCE, . (Myoclonic Dystonia, Gene SGCE, Mut.)	45450
41.01.144		ITGA7, . (Congenital Muscular Dystrophy, CMD, Gene ITGA7, Mut.)	55550
41.01.145		SEPN1, . (Congenital Fiber Type Disproportion, CFTD, Gene SEPN1, Mut.)	34800
41.01.146		DMPK, . . (Myotonic Dystrophy 1, Gene DMPK, Freq. Mut.)	4200
41.01.147		ZNF9, . . (Myotonic Dystrophy 2, Gene ZNF9, Freq. Mut.)	4200
41.01.148		CLCN1, . . (Myotonia Congenita, Gene CLCN1, Freq. Mut.)	9900
41.01.149		CLCN1, . (Myotonia Congenita, Gene CLCN1, Mut.)	61900
41.01.150		DES, . (Myofibrillar Myopathy, Gene DES, Mut.)	24730
41.01.151		CRYAB, . (Myofibrillar Myopathy, Gene CRYAB, Mut.)	10690
41.01.152		, MYOT . (Myofibrillar Myopathy, Gene MYOT, Mut.)	34350
41.01.153		CFTR, . (Cystic Fibrosis, Gene CFTR, Mut.)	104650
41.01.154		CFTR, . . (Cystic Fibrosis, Gene CFTR, Freq. Mut.)	12040
41.01.155		7 ITGA7, . (Congenital Muscular Dystrophy with Integrin Alpha-7 Deficiency, Congenital Muscular Dystrophy with ITGA7-Deficiency, Gene ITGA7, Mut.)	55550
41.01.156		« » LAMA2, « » . . (Merosin-Deficient Congenital Muscular Dystrophy, Gene LAMA2, Hot-Point Mut.)	38050
41.01.157		LAMA2, « » . . (Merosin-Deficient Congenital Muscular Dystrophy, Gene LAMA2, without Hot-Point Mut.)	190000

41.01.158		LAMA2, . (Merosin-Deficient Congenital Muscular Dystrophy, Gene LAMA2)	208000
41.01.159		FKRP, . (Muscular Dystrophy-Dystroglycanopathy, Gene FKRP, Mut.)	10690
41.01.160		FKRP, . (Muscular Dystrophy-Dystroglycanopathy, Gene FKRP, Freq. Mut.)	6000
41.01.161		(Duchenne Muscular Dystrophy, Dystrophin Gene Deletion, Boys)	7800
41.01.162		(Duchenne Muscular Dystrophy, X-Lyonization, Girls)	6670
41.01.164		CAPN3, . . (Muscular Dystrophy Limb-Girdle Type 2A, Gene CAPN3, Freq. Mut.)	5400
41.01.167		FKRP, . (Muscular Dystrophy Limb-Girdle Type 2A, Gene FKRP, Mut.)	10700
41.01.168		FKRP, . . (Muscular Dystrophy Limb-Girdle Type 2A, Gene FKRP, Freq. Mut.)	6000
41.01.171		POMT1, . (Muscular Dystrophy Limb-Girdle Type 2A, Gene POMT1, Mut.)	61900
41.01.172		SGCA, . (Muscular Dystrophy Limb-Girdle Type 2A, Gene SGCA, Mut.)	23250
41.01.173		SGCA, . . (Muscular Dystrophy Limb-Girdle Type 2A, Gene SGCA, Freq. Mut.)	9900
41.01.174		SGCB, . (Muscular Dystrophy Limb-Girdle Type 2A, Gene SGCB, Mut.)	23250
41.01.178		FKTN, . (Muscular Dystrophy Fukuyama-Type, Gene FKTN, Mut.)	41630
41.01.179		FHL1, . (Emery-Dreifuss Muscular Dystrophy, Gene FHL1, Mut.)	30650
41.01.180		LMNA, . (Emery-Dreifuss Muscular Dystrophy, Gene LMNA, Mut.)	34800
41.01.181		, . (Emery-Dreifuss Muscular Dystrophy, X-Linked Gene Emerine, Mut.)	10700
41.01.182		MULIBRAY. TRIM37, . (Muscle-Liver-Brain-Eye, Gene TRIM37, Mut.)	9000
41.01.183		SRY, . (Disorders Sex Determination, Gene SRY, Mut.)	6200
41.01.184		SRY, . (Disorders Sex Determination, Analysis Gene SRY, Mut.)	4400
41.01.185		I. EGR2, . (Charcot-Marie-Tooth Disease Type 1B, Gene EGR2, Mut.)	14450
41.01.186		I. GJB1, . (Charcot-Marie-Tooth Disease Type 1B, Gene GJB1, Mut.)	9000
41.01.187		I. LITAF, . (Charcot-Marie-Tooth Disease Type 1B, Gene LITAF, Mut.)	14450
41.01.188		I. PRPS1, . (Charcot-Marie-Tooth Disease Type 1B, Gene PRPS1, Mut.)	24730
41.01.189		I. SH3TC2, FIG4, FGD4 GDAP1, . . (Charcot-Marie-Tooth Disease Type 1B, Gene GDAP1, Freq. Mut.)	8900
41.01.190		I. YARS, . (Charcot-Marie-Tooth Disease Type 1B, Gene YARS, Mut.)	41630
41.01.191		I. 17 22, . (Charcot-Marie-Tooth Disease Type 1B, Duplication on Chromosome 17 Gene 22, Mut.)	4400

41.01.192		I. 0, . (Charcot-Marie-Tooth Disease Type 1B, Gene 0, Mut.)	14550
41.01.193		I. 22, . (Charcot-Marie-Tooth Disease Type 1B, Gene 22, Mut.)	12380
41.01.194		I. NDRG1 SH3TC2, . . (Charcot-Marie-Tooth Disease Type 1B, Genes NDRG1, SH3TC2, Mut.)	4400
41.01.195		II. DNMT2, . (Charcot-Marie-Tooth Disease Type 2A1, Gene DNMT2, Mut.)	75450
41.01.196		II. FIG4, . (Charcot-Marie-Tooth Disease Type 2A1, Gene FIG4, Mut.)	97250
41.01.197		II. GARS, . (Charcot-Marie-Tooth Disease Type 2A1, Gene GARS, Mut.)	62000
41.01.198		II. GDAP, . (Charcot-Marie-Tooth Disease Type 2A1, Gene GDAP, Mut.)	21370
41.01.199		II. HSPB1, . (Charcot-Marie-Tooth Disease Type 2A1, Gene HSPB1, Freq. Mut.)	9000
41.01.200		II. LMNA, . (Charcot-Marie-Tooth Disease Type 2A1, Gene LMNA, Mut.)	34800
41.01.201		II. MFN2, . (Charcot-Marie-Tooth Disease Type 2A1, Gene MFN2, Mut.)	51720
41.01.202		II. MFN2, . . (Charcot-Marie-Tooth Disease Type 2A1, Gene MFN2, Freq. Mut.)	4400
41.01.203		II. NEFL, . (Charcot-Marie-Tooth Disease Type 2A1, Gene NEFL, Mut.)	21370
41.01.204		22, . (Hereditary Neuropathy with Liability to Pressure Palsies, HNPP, Gene 22, Mut.)	12380
41.01.205		22 (Hereditary Neuropathy with Liability to Pressure Palsies, HNPP, Gene 22, Copy Number Variation)	13500
41.01.206		TTR, . (Transthyretin-Related Hereditary Amyloidosis, Gene TTR, Mut.)	6670
41.01.207		C1NH, . (Hereditary Angioedema Type I, Gene C1NH, Mut.)	24730
41.01.208		, I (HFE) (Hemochromatosis Type 1 (Gene HFE))	3000
41.01.209		ALX4, . (Parietal Foramina, PFM, Gene ALX4, Mut.)	14550
41.01.210		MSX2, . (Parietal Foramina, PFM, Gene MSX2, Mut.)	8320
41.01.212		EYA4, . (Deafness Autosomal Recessive 1, DFNB1A, Gene EYA4, Mut.)	41630
41.01.213		GJB3, . (Deafness Autosomal Recessive 1, DFNB1A, Gene GJB3, Mut.)	8320
41.01.214		GJB6, . (Deafness Autosomal Recessive 1, DFNB1A, Gene GJB6, Mut.)	8320
41.01.215		, SLC26A4 . (Deafness Autosomal Recessive 1, DFNB1A, Gene SLC26A4, Mut.)	46820
41.01.216		DFNB1, . (Deafness Autosomal Recessive 1, Frequent Deletions Locus Gene DFNB1, Mut.)	4400
41.01.217		ELA2, . (Neutropenia Severe Congenital 1 Autosomal Dominant, SCN1, Gene ELA2, Mut.)	13700

41.01.218		WAS, . (Neutropenia Severe Congenital 1 Autosomal Dominant, SCN1, Gene WAS, Mut.)	24730
41.01.219		NPHP1, . (Nephronophthisis 1, NPHP1, Gene NPHP1, Mut.)	12040
41.01.220		NPHS1, . (Nephrotic Syndrome Type 1, NPHS1, Gene NPHS1, Mut.)	67650
41.01.221		NPHS2, . (Nephrotic Syndrome Type 1, NPHS1, Gene NPHS2, Mut.)	30650
41.01.222		13 SCN4A, . (Normokalemic Periodic Paralysis, Exon 13 Gene SCN4A, Mut.)	6760
41.01.227		LEP, MC4R, POMC, . (Monogenic Obesity, Genes LEP, MC4R, POMC, Mut.)	16450
41.01.228		RABPN1, . . (Oculopharyngeal Muscular Dystrophy, OPMD, Gene RABPN1, Freq. Mut.)	4400
41.01.229		(TCIRG1, . . (Osteopetrosis Autosomal Recessive 1, OPTB1, Gene TCIRG1, Freq. Mut.)	4400
41.01.230		(TCIRG1, . (Osteopetrosis Autosomal Recessive 1, OPTB1, Gene TCIRG1, Mut.)	45450
41.01.231		- Gene CRX, Mut.)	10700
41.01.232		- Gene ADAM9, Mut.)	81450
41.01.233		- Gene RPGR, Mut.)	82450
41.01.234		SCN4A, . (Paramyotonia Congenita of Von Eulenburg, Gene SCN4A, Mut.)	82450
41.01.235		KTR6B, . (Pachyonychia Congenita 2, PC2, Gene KTR6B, Mut.)	22250
41.01.236		(HPGD, . (Hypertrophic Osteoarthropathy, Primary, Autosomal Recessive, 1, Gene HPGD, Mut.)	24730
41.01.237		BMPR2, . (Primary Pulmonary Hypertension 1, PPH1, Gene BMPR2, Mut.)	51720
41.01.238		MEFV, . (Familial Mediterranean Fever, FMF, Gene MEFV, Mut.)	34800
41.01.240		PRPH2, . (Retinitis Pigmentosa, Gene PRPH2, Mut.)	12380
41.01.241		BEST1, . (Retinitis Pigmentosa, Gene BEST1, Mut.)	38050
41.01.242		CA4, . (Retinitis Pigmentosa, Gene CA4, Mut.)	21370
41.01.243		CRB1, . (Retinitis Pigmentosa, Gene CRB1, Mut.)	55300
41.01.244		LRAT, . (Retinitis Pigmentosa, Gene LRAT, Mut.)	10700
41.01.245		NR2E3, . (Retinitis Pigmentosa, Gene NR2E3, Mut.)	21370
41.01.246		NRL, . (Retinitis Pigmentosa, Gene NRL, Mut.)	12340
41.01.247		RP2, . (Retinitis Pigmentosa, Gene RP2, Mut.)	17900
41.01.249		RPGR, . (Retinitis Pigmentosa, Gene RPGR, Mut.)	82450
41.01.250		CTSK, . (Pyknodysostosis, PKND, Gene CTSK, Mut.)	23250
41.01.251		FLCN, . (Primary Spontaneous Pneumothorax, PSP, Gene FLCN, Mut.)	41630
41.01.252		CAV3, . (Creatine Phosphokinase Elevated, Serum, Gene CAV3, Mut.)	8320
41.01.253		SHH, . (Polydactyly, Gene SHH, Mut.)	8650

41.01.254		GLI3, . (Polydactyly, Gene GLI3, Mut.)	61900
41.01.255		PKHD1, « . » . . (Polycystic Kidney Disease Autosomal Recessive, Gene PKHD1, Hot-Point Mut.)	24730
41.01.256		UPK3A, . (Renal Hypodysplasia, Aplasia 1, Gene UPK3A, Mut.)	23250
41.01.257		10, 11, 13, 14, 15 RET, . (Renal Hypodysplasia, Aplasia 1, Exons 10, 11, 13, 14, 15 Gene RET, Mut.)	18550
41.01.258		LMNA, . (Hutchinson-Gilford Progeria Syndrome, Gene LMNA, Mut.)	38050
41.01.259		COMP, . . (Pseudoachondroplasia, Gene COMP, Freq. Mut.)	5400
41.01.260		GNAS, . (Pseudohypoparathyroidism, Type IA, Gene GNAS, Mut.)	34350
41.01.261		ABCC6, . (Pseudoxanthoma Elasticum, Gene ABCC6, Mut.)	105000
41.01.262		ABCC6, . . (Pseudoxanthoma Elasticum, Gene ABCC6, Freq. Mut.)	7200
41.01.263		GNAS, . (Pseudopseudohypoparathyroidism, Gene GNAS, Mut.)	34350
41.01.265		RS1, . (Retinoschisis 1 X-Linked Juvenile, RS1, Gene RS1, Mut.)	23250
41.01.266		SCN5A, . (Paroxysmal Familial Ventricular Fibrillation (VF), Gene SCN5A, Mut.)	50660
41.01.267		TNFRSF1A, . (TNF-Receptor-Associated Periodic Syndrome, TRAPS, Gene TNFRSF1A, Mut.)	22250
41.01.268		PRF1, . (Familial Hemophagocytic Lymphohistiocytosis, Gene PRF1, Mut.)	17900
41.01.269		STX11, . (Familial Hemophagocytic Lymphohistiocytosis, Gene STX11, Mut.)	12340
41.01.270		STXBP2, . (Familial Hemophagocytic Lymphohistiocytosis, Gene STXBP2, Mut.)	41630
41.01.271		UNC13D, . (Familial Hemophagocytic Lymphohistiocytosis, Gene UNC13D, Mut.)	61900
41.01.272		UNC13D, . . (Familial Hemophagocytic Lymphohistiocytosis, Gene UNC13D, Freq. Mut.)	4400
41.01.273		(10, 11, 13, 14, 15 RET) (Familial Medullary Thyroid Cancer (Exons 10, 11, 13, 14, 15 Gene RET))	13900
41.01.274		5, 8 RET, . (Familial Medullary Thyroid Cancer, Exons 5, 8 Gene RET, Mut.)	9000
41.01.275		NLRP3 . (Familial Cold Autoinflammatory Syndrome, FCAS, Gene NLRP3, Mut.)	44450
41.01.276		, WNK1 . (Hereditary Sensory and Autonomic olynuropathy, Gene WNK1, Mut.)	12350
41.01.277		, NGF . (Hereditary Sensory and Autonomic olynuropathy, Gene NGF, Mut.)	14850
41.01.278		CINCA, NLRP3 . (Chronic Infantile Neurologic Cutaneous Articular, Gene NLRP3, Mut.)	32970
41.01.279		ESC (-). NR2E3, . (Enhanced S- one Syndrome, Goldmann-Favre Syndrome, Gene NR2E3, Mut.)	22250
41.01.280		LEOPARD. 7, 12, 13 PTPN11, . (Lentigines, Electrocardiographic defects, Ocular hypertelorism, Pulmonary stenosis, Abnormalities of genitalia, Retardation of growth, sensorineural Deafness, Exons 7, 12, 13 Gene PTPN11,	10900
41.01.281		TAR. RBM8A, . (Thrombocytopenia-Absent Radius Syndrome, TAR-Syndrome, Gene RBM8A, Mut.)	23250

41.01.282		(). FGD1, . (Aarskog-Scott Syndrome, Faciodigitogenital Syndrome, Faciogenital Dysplasia, Gene FGD1, Mut.)	54300
41.01.283		« » ALMS1, « » . (Alstrom syndrome, Gene ALMS1, Hot-Point Mut.)	13350
41.01.284		KCNJ2, . (Andersen-Tawil Syndrome, Gene KCNJ2, Mut.)	12380
41.01.285		9 FGFR2, . (Antley-Bixler Syndrome, ABS, Exon 9 Gene FGFR2, Mut.)	7000
41.01.286		(). FGFR2, . (Apert Syndrome, AS, Gene FGFR2, Freq. Mut.)	9400
41.01.287		PRPS1, . (Art's Syndrome, Gene PRPS1, Mut.)	25950
41.01.288		PTEN, . (Bannayan-Ruvalcaba-Riley Syndrome, Gene PTEN, Mut.)	33350
41.01.289		TAZ, . (Barth Syndrome, Gene TAZ, Mut.)	22250
41.01.290		(). FLCN, . (Birt-Hogg-Dube Syndrome, BHD, Gene FLCN, Mut.)	44450
41.01.291		(). IKBK, . (Bloch-Sulzberger Syndrome, Familial Incontinentia Pigmenti, Gene IKBK, Freq. Mut.)	5400
41.01.292		(). EMG1, . (Bowen Conradi Syndrome, BCS, Gene EMG1, Mut.)	15850
41.01.293		SCN5A, . (Brugada Syndrome, BrS, Gene SCN5A, Mut.)	104000
41.01.294		(). BCS1L, . (Bjornstad Syndrome, Gene BCS1L, Mut.)	12380
41.01.295		PAX3, . (Waardenburg Syndrome, WS, Gene PAX3, Mut.)	29650
41.01.296		EDNRB, . (Waardenburg-Shah Syndrome, Gene EDNRB, Mut.)	25950
41.01.297		IRF6, . (Van der Woude Syndrome, Gene IRF6, Mut.)	34350
41.01.298		(). RECQL2, . (Werner Syndrome, Gene RECQL2, Mut.)	128000
41.01.299		(). WAS, . (Wiskott-Aldrich Syndrome, WAS, Gene WAS, Mut.)	25950
41.01.300		(). PHOX2B, . (Congenital Central Hypoventilation Syndrome, CCHS, Gene PHOX2B, Freq. Mut.)	3500
41.01.301		(). HPS1, . (Albinism culocutaneous, Hermansky-Pudlak type Gene HPS1, Freq. Mut.)	9650
41.01.302		(). GLI3, . (Greig Syndrome, Gene GLI3, Mut.)	61900
41.01.303		RAB27A, . (Griscelli Syndrome, Gene RAB27A, Mut.)	17900
41.01.304		9 FGFR2 7A FGFR1, . (Jackson-Weiss Syndrome, JWS, Exon 9 Gene FGFR2, Exon 7A Gene FGFR1, Mut.)	9000
41.01.305		TBX1, . (Di George Syndrome, DGS, Gene TBX1, Mut.)	34350
41.01.306		(UGT1A1) (Gilbert's Syndrome (Gene UGT1A1))	4000
41.01.307		(). NPHP1 (Joubert Syndrome, Cerebelloparenchymal Disorder IV, CPD IV, Classic Joubert Syndrome, Joubert Syndrome type A, Joubert-Boltshauser Syndrome, Pure Joubert Syndrome, Gene NPHP1, Mut.)	12040
41.01.308		(). RAB23, . (Carpenter Syndrome, Gene RAB23, Mut.)	26950
41.01.309		(). « » DNAI1 DNAH5, « » . (Kartagener Syndrome, Genes DNAI1, DNAH5, Hot-Point Mut.)	34350

41.01.310		(DNAI1, . (Kartagener Syndrome, Gene DNAI1, Mut.))	56550
41.01.311		GJB2, . (Keratitis-Ichthyosis-Deafness Syndrome, KID Syndrome, Gene GJB2, Mut.)	9000
41.01.312		GDF6, . (Klippel-Feil Syndrome, Gene GDF6, Mut.)	10690
41.01.313		ERCC6, . (Cockayne Syndrome, Gene ERCC6, Mut.)	75600
41.01.316		QT, KCNQ1 KCNE1, . (Short QT Syndrome, Short QTS, Genes KCNQ1, KCNE1, Mut.)	25880
41.01.317		HRAS, . (Costello Syndrome, Gene HRAS, Mut.)	7200
41.01.318		RPS6KA3, . (Coffin-Lowry Syndrome, Gene RPS6KA3, Mut.)	82450
41.01.319		PAX3, . (Craniofacial-Deafness-Hand Syndrome, CDHS, Gene PAX3, Mut.)	28090
41.01.320		UGT1, . (Crigler-Najjer Syndrome, Gene UGT1, Mut.)	17550
41.01.321		10 FGFR3, . (Crouzon Syndrome with Acanthosis Nigrificans, CAN, Exon 10 Gene FGFR3, Mut.)	6670
41.01.322		7 9 FGFR2, . (Crouzon Syndrome, Exons 7, 9 Gene FGFR2, Mut.)	9000
41.01.323		BCS1L, . (Leigh Syndrome, Gene BCS1L, Mut.) III.	12380
41.01.324		MED12, . . (Lujan-Fryns Syndrome, LFS, Gene MED12, Freq. Mut.)	6100
41.01.325		NLRP3, . (Muckle-Wells Syndrome, MWS, Gene NLRP3, Mut.)	44460
41.01.326		XK, . (McLeod Syndrome, Gene XK, Mut.)	15850
41.01.327		« » FBN1, « » . (Marfan Syndrome, Gene FBN1, Hot-Point Mut.)	27000
41.01.328		« » FBN1, « » . (Marfan Syndrome, Gene FBN1, without Hot-Point Mut.)	190000
41.01.329		FBN1, . (Marfan Syndrome, Gene FBN1, Mut.)	216000
41.01.331		2A (10, 11 RET) (Multiple Endocrine Neoplasia Type 2A (Exons 10, 11 Gene RET))	8600
41.01.332		2B (RET) (Multiple Endocrine Neoplasia Type 2B (Gene RET))	4200
41.01.333		ZEB2, . (Mowat-Wilson Syndrome, Gene ZEB2, Mut.)	51700
41.01.334		(TAZ, . (Left Ventricular Non-Compaction, LVNC, Gene TAZ, Mut.)	21370
41.01.335		, NBN . . (Nijmegen Breakage Syndrome, NBS, Gene NBN, Freq. Mut.)	4400
41.01.336		(LMX1B, . (Nail-Patella Syndrome, NPS, Onychoosteodysplasia, Gene LMX1B, Mut.)	26950
41.01.337		3, 7, 13 PTPN11, . (Syndrome Noonan, Exons 3, 7, 13 Gene PTPN11, Mut.)	9900
41.01.338		MED12, . . (Opiz-Kaveggia Syndrome, OKS, Gene MED12, Freq. Mut.)	6110
41.01.339		- a - a (ENG, . (Rendu-Osler-Weber Disease, Gene ENG, Mut.)	34350
41.01.340		TBX3, . (Pallister W Syndrome, Gene TBX3, Mut.)	24730
41.01.341		GLI3, . (Pallister-Hall Syndrome, Gene GLI3, Mut.)	61900

41.01.342		SLC26A4 . (Pendred Syndrome, Gene SLC26A4, Mut.)	46820
41.01.343		IRF6, . (Popliteal Pterygium Syndrome, PPS, Gene IRF6, Mut.)	34350
41.01.344		7, 9 FGFR2 7A FGFR1, . (Pfeiffer Syndrome, Exons 7, 9 Gene FGFR2, Exon 7A Gene FGFR1, Mut.)	12340
41.01.345		MECP2, . (Retts Syndrome, Gene MECP2, Mut.)	19550
41.01.346		(SEPNI, . (Stiff-M n Syndrome, SMS, Moersch-Woltmann Syndrome, Stiff-Trunk Syndrome, Gene SEPNI, Mut.)	34810
41.01.348		BSCL2, . (Silver Syndrome, Gene BSCL2, Mut.)	26950
41.01.349		GPC3, . (Simpson-Golabi-Behmel Syndrome, Type 1, SGBS1, Gene GPC3, Mut.)	30650
41.01.350		(SCN5A, . (Sick Sinus Syndrome, SSS, Gene SCN5A, Mut.)	50660
41.01.351		(DHCR7, . (Smith-Lemli-Opitz Syndrome, Gene DHCR7, Mut.)	31450
41.01.352		I. Col2A1, . (Stickler Syndrome, Type 1, Gene Col2A1, Mut.)	119500
41.01.353		(AR, . (Testicular Feminization Syndrome, Gene AR, Mut.)	34810
41.01.354		(TCOF1, . (Treacher-Collins Syndrome, Franceschetti-Klein Syndrome, Mandibulofacial Dysostosis without Limb Anomalies, Gene TCOF1, Mut.)	75550
41.01.355		QT. CAV3, . (Long QT Syndrome, LQTS, LQT, Gene CAV3, Mut.)	8320
41.01.356		QT. KCNJ2, . (Long QT Syndrome, LQTS, LQT, Gene KCNJ2, Mut.)	12380
41.01.357		QT. SCN4B, . (Long QT Syndrome, LQTS, LQT, Gene SCN4B, Mut.)	19000
41.01.358		QT. SCN5A, . (Long QT Syndrome, LQTS, LQT, Gene SCN5A, Mut.)	50660
41.01.359		QT. KCNH2 KCNE2, . (Long QT Syndrome, LQTS, LQT, Genes KCNH2, KCNE2, Mut.)	26000
41.01.360		QT. KCNQ1 KCNE1, . (Long QT Syndrome, LQTS, LQT, Genes KCNQ1, KCNE1, Mut.)	26000
41.01.361		(FKRP, . (Walker-Warburg Syndrome, WWS, Gene FKRP, Mut.)	10960
41.01.362		(POMT1, . (Walker-Warburg Syndrome, WWS, Gene POMT1, Mut.)	67000
41.01.363		(VHL, . (Von Hippel-Lindau Syndrome, VHL, Von Hippel-Lindau Hereditary Cancer Syndrome, Gene VHL, Mut.)	10960
41.01.364		(VHL, . (Von Hippel-Lindau Syndrome, VHL, Von Hippel-Lindau Hereditary Cancer Syndrome, Gene VHL, Copy Number Variation Gene VHL, Mut.)	13500
41.01.365		(TBX5, . (Holt-Oram Syndrome, Gene TBX5, Mut.)	34800
41.01.366		SBDS, . (Shwachman-Diamond Syndrome, Gene SBDS, Mut.)	19850
41.01.367		SBDS1, . . (Shwachman-Diamond Syndrome, Gene SBDS1, Freq. Mut.)	7200
41.01.368		SLC26A4 . (Large Vestibular Aqueduct Syndrome, LVAS, Gene SLC26A4, Mut.)	46820
41.01.369		VI. PLOD, . . (Ehlers-Danlos Syndrome, Type VI, Gene PLOD, Freq. Mut.)	7830
41.01.370		CHRNA1, . (Escobar Syndrome, Gene CHRNA1, Mut.)	28100

41.01.371		HOXD13, . (Synpolydactyly, Type 1, SPD1, Gene HOXD13, Mut.)	14650
41.01.372		FHL1, . (Scapuloperoneal Myopathy, SPM, Gene FHL1, Mut.)	30650
41.01.373		ROBO3, . (Horizontal Gaze Palsy with Progressive Scoliosis, HGPPS, Gene ROBO3, Mut.)	62000
41.01.374		(c GJC2, . (Westphal-Strumpell Disease, Gene GJC2, Mut.)	20470
41.01.375		, ATL1 . (Westphal-Strumpell Disease, Gene ATL1, Mut.)	51700
41.01.376		, SPAST . (Westphal-Strumpell Disease, Gene SPAST, Mut.)	51700
41.01.378		ALSIN, . (Infantile Ascending Hereditary Spastic Paralysis, IAHSP, Gene ALSIN, Mut.)	119500
41.01.379		IGHMBP2, . (Spinal Muscular Atrophy (SMA) with Diaphragmatic Paralysis, Gene IGHMBP2, Mut.)	51700
41.01.380		I, II, III, IV. SMN1, . (Spinal Muscular Atrophy, SMA, Type I, II, III, IV, Gene SMN1)	7830
41.01.381		I, II, III, IV. (Spinal Muscular Atrophy, SMA, Type I, II, III, IV, Copy Number Variation)	13500
41.01.382		I, II, III, IV. SMN1, . (() (Spinal Muscular Atrophy, SMA, Type I, II, III, IV, Gene SMN1, Mut. (Only Presence One Gene Copy))	30650
41.01.383		VAPB, (Spinal Muscular Atrophy, SMA, Late-Onset, Finkel Type, Gene VAPB, Mut.)	23000
41.01.384		VAPB, . . (Spinal Muscular Atrophy, SMA, Late-Onset, Finkel Type, Gene VAPB, Freq. Mut.)	6000
41.01.385		, X- « » UBA1, « » . . (Spinal Muscular Atrophy, SMA, X-Linked, Gene UBA1, Hot-Point Mut.)	7200
41.01.386		AR, . . (Kennedy Spinal and Bulbar Muscular Atrophy, Gene AR, Freq. Mut.)	4400
41.01.387		ATXN1, ATXN2, ATXN3, . . (Spinocerebellar Ataxia, Genes ATXN1, ATXN2, ATXN3, Freq. Mut.)	7830
41.01.388		ATXN7, . . (Spinocerebellar Ataxia, Gene ATXN7 Freq. Mut.)	5000
41.01.389		ATXN8, . . (Spinocerebellar Ataxia, Gene ATXN8, Freq. Mut.)	5000
41.01.390		PRNP, . (Spongiform Encephalopathy with Neuropsychiatric Features, Gene PRNP, Mut.)	10700
41.01.391		DLL3, . (Spondylocostal Dysostosis, Gene DLL3, Mut.)	27000
41.01.392		(). Col2A1, . (Spondyloepiphyseal Dysplasia Tarda, SEDT, Gene Col2A1, Mut.)	131000
41.01.393		(). TRAPPC2, . (Spondyloepiphyseal Dysplasia Tarda, SEDT, Gene TRAPPC2, Mut.)	14550
41.01.394		PRPS1, . (Phosphoribosylpyrophosphate Synthetase Superactivity, PRS Superactivity, Gene PRPS1, Mut.)	24730
41.01.395		, TOR1A . (Torsion Dystonia, Gene TOR1A, Mut.)	19550
41.01.396		GCH1, . (Torsion Dystonia, Gene GCH1, Mut.)	23250
41.01.397		PRRT2, . (Torsion Dystonia, Gene PRRT2, Mut.)	15850
41.01.398		SGCE, . (Torsion Dystonia, Gene SGCE, Mut.)	45450
41.01.399		SPR, . (Torsion Dystonia, Gene SPR, Mut.)	13350

41.01.400		TRPS1, . (Trichorhinophalangeal Syndrome, TRPS, Gene TRPS1, Mut.)	37050
41.01.401		MPL, . (Congenital Amegakaryocytic Thrombocytopenia, CAMT, Gene MPL, Mut.)	30650
41.01.402		TSC1, . (Tuberous Sclerosis, Gene TSC1, Mut.)	82450
41.01.403		PRNP, . (Fatal Familial Insomnia, FFI, Gene PRNP, Mut.)	10700
41.01.404		PAH, . (Phenylketonuria, PKU, Gene PAH, Mut.)	41630
41.01.405		PAH, . . (Phenylketonuria, PKU, Gene PAH, Freq. Mut.)	12040
41.01.406		« » ACVR1, « » . . (Fibrodysplasia Ossificans Progressiva, FOP, Gene ACVR1, Hot-Point Mut.)	15850
41.01.407		« » ACVR1, « » . . (Fibrodysplasia Ossificans Progressiva, FOP, Gene ACVR1, without Hot-Point Mut.)	27000
41.01.410		RMRP, . (Metaphyseal Chondrodysplasia, McKusick Type, Gene RMRP, Mut.)	7000
41.01.411		EBP, . (Chondrodysplasia Punctata, CDP, Conradi-Hunermann Syndrome, Gene EBP, Mut.)	15850
41.01.412		ANKH, . (Chondrocalcinosis, Calcium Pyrophosphate Dihydrate, CPPD, Gene ANKH, Mut.)	45450
41.01.413		IT15, . . (Chorea Huntington, Gene IT15, Freq. Mut.)	5000
41.01.414		PRPH2, . (Choroidal Dystrophy, Gene PRPH2, Mut.)	12380
41.01.415		CHM, . (Choroideremia, CHM, Gene CHM, Mut.)	51700
41.01.416		CYBB, . (Chronic Granulomatous Disease, CGD, Gene CYBB, Mut.)	41630
41.01.417		BTK, . (X-Linked Agammaglobulinemia, XLA, Gene BTK, Mut.)	61900
41.01.418		RPS6KA3, . (X-Linked Mental Retardation, Gene RPS6KA3, Mut.)	82450
41.01.419		SLC9A6, . (X-Linked Mental Retardation, Gene SLC9A6, Mut.)	51700
41.01.420		ZDHHC9, . (X-Linked Mental Retardation, Gene ZDHHC9, Mut.)	9000
41.01.421		(, XIAP . (X-Linked Lymphoproliferative Syndrome, XLP, Gene XIAP, Mut.)	28090
41.01.422		(, SH2D1A, . (X-Linked Lymphoproliferative Syndrome, XLP, Gene SH2D1A, Mut.)	14550
41.01.423		FRMD7, . (X-Linked Nystagmus congenital 1, NYS1 X-Linked, Gene FRMD7, Mut.)	31630
41.01.424		IL2RG, . (X-Linked Severe Combined Immunodeficiency, Gene IL2RG, Mut.)	14550
41.01.425		DNM2, . (Centronuclear Myopathy, CNM, Gene DNM2, Mut.)	82450
41.01.426		MTM1, . (Centronuclear Myopathy, CNM, Gene MTM1, Mut.)	51700
41.01.427		ERCC6, . (Cerebrooculofacioskeletal Syndrome, COFS Syndrome, Gene ERCC6, Mut.)	75550
41.01.428		EXT1, . (Multiple Exostoses, Gene EXT1, Mut.)	41630
41.01.429		EXT2, . (Multiple Exostoses, Gene EXT2, Mut.)	51700
41.01.430		NDP, . (Familial Exudative Vitreoretinopathy, FEVR, Gene NDP, Mut.)	8320

41.01.431		EDA, . (Anhidrotic Ectodermal Dysplasia, Gene EDA, Mut.)	28100
41.01.432		GJB6, . (Hidrotic Ectodermal Dysplasia, Gene GJB6, Mut.)	8320
41.01.433		CACNA1A, . (Episodic Ataxia, EA, Gene CACNA1A, Mut.)	161850
41.01.434		COMP, . . (Multiple Epiphysial Dysplasia, MED, Gene COMP, Freq. Mut.)	5500
41.01.435		SLC26A2, . (Multiple Epiphysial Dysplasia, MED, Gene SLC26A2, Mut.)	23250
41.01.436		(). ALOXE3, . (Nonbullous Congenital Ichthyosiform Erythroderma, NBCIE, Gene ALOXE3, Mut.)	51720
41.01.437		(). LOX12B, . (Nonbullous Congenital Ichthyosiform Erythroderma, NBCIE, Gene LOX12B, Mut.)	34810
41.01.438		(). TGM1, . (Nonbullous Congenital Ichthyosiform Erythroderma, NBCIE, Gene TGM1, Mut.)	34810
41.01.439		GJB3, . (Erythrokeratoderma, Gene GJB3, Mut.)	8320
41.01.440		GJB4, . (Erythrokeratoderma, Gene GJB4, Mut.)	9000
41.01.441		(c). VHL, . (Autosomal Recessive Erythrocytosis, Gene VHL, Mut.)	10690
41.01.442		(c). VHL, . . (Autosomal Recessive Erythrocytosis, Gene VHL, Freq. Mut.)	4500
42. : (Definition of Biological Relationship: Paternity and Maternity) / 42.01. (Definition of Biological Relationship: Paternity and Maternity) :			
42.01.001		(3). (Urgent Establishment of Biological Relationship for One Parent at Indisputable Relationship of Another (3 Persons))	37210
42.01.002		(3). (Establishment of Biological Relationship for One Parent at Indisputable Relationship of Another (3 Persons))	18210
42.01.003		(2). (Urgent Establishment of Biological Relationship for One Parent in Absence of Another (2 Persons))	37210
42.01.004		(2). (Establishment of Biological Relationship for One Parent in Absence of Another (2 Persons))	16400
42.01.005		() (Additional research participant (child or mother or father))	6450
43. / « » (Hereditary Metabolic Disorders in Newborns. Newborn Screening "HEEL") / 43.01. « » (Hereditary Metabolic Disorders in Newborns. Newborn Screening "HEEL")			
43.01.001		e « » (Newborn Screening "HEEL")	5200
43.01.002		() (Gas Chromatography of Urine (Organic Aciduria))	8100
43.01.004		() (Biotin-Dependent Carboxylases Activity (Biotinidase Deficiency))	4700
43.01.005		- - () (HPLC-MS/MS Organic Acids (Succinylate))	4100
43.01.006		- - () (HPLC-MS/MS Organic Acids (Homogentisic Acid))	3730
43.01.007		- - (N-) (HPLC-MS/MS Organic Acids (N-Acetylaspartate))	3730
43.01.008		- - () (HPLC-MS/MS Organic Acids (Orotic Acid))	3730
43.01.009		BD () (BD (Biotinidase Deficiency) Gene, Freq. Mut.)	5520

43.01.010		GCDH () (GCDH (Glutaryl-CoA Dehydrogenase) Gene, Freq. Mut. (Glutaric Aciduria, Type 1))	4300
43.01.011		GCDH () (GCDH (Glutaryl-CoA Dehydrogenase) Gene (Glutaric Aciduria, Type 1))	38470
43.01.012		CBS () (CBS (Cystathionine-Beta-Synthase) Gene, Freq. Mut. (Homocystinuria))	5600
43.01.013		(, « », BCKDHA, BCKDHB) (BCKDHA and BCKDHB Genes, Freq. Mut. (Maple Syrup Urine Disease, MSUD))	16000
43.01.014		MUT () (MUT Gene, Freq. Mut. (Methylmalonic Aciduria, MMA))	16000
43.01.015		HADHA () (HADHA Gene, Freq. Mut. (Long-Chain 3-Hydroxyacyl-Coa Dehydrogenase (LCHAD) Deficiency))	4890
43.01.016		ACADM () (ACADM Gene, Freq. Mut. (Medium-Chain Acyl-CoA Dehydrogenase (MCAD) Deficiency))	4890
43.01.017		() (TC Gene (Ornithine Transcarbamylase (OTC) Deficiency))	31200
43.01.018		FAH () (FAH Gene, Freq. Mut. (Tyrosinemia, Type 1))	6880
43.01.019		FAH () (FAH Gene (Tyrosinemia, Type 1))	47520
43.01.020		ASS () (ASS Gene, Freq. Mut. (Citrullinemia))	6100
44. (Investigation of Water and Soil Quality) / 44.01. (High Tech Investigations of Water)			
44.01.001		: (Water: Complete and Comprehensive Quality Assessment)	16500
44.01.002		: (Water: Abridged Quality Assessment)	8700
44.01.003		: (Water: Suspicion Household Waste Contamination)	7860
44.01.004		: (Water: Suspicion Industrial Waste Contamination)	12300
44.01.005		: (Water: Suspicion Products of Combustion and Emissions from Motorways Contamination)	9020
44.01.006		: (Water: Suspicion Excessive Use of Chemicals for Water Treatment)	5150
44. (Investigation of Water and Soil Quality) / 44.02. (Screening Investigations of Water)			
44.02.001		: 30 (Water: Quality Assessment 30 Parameters)	6280
44.02.002		: 20 (Water: Quality Assessment 20 Parameters)	4750
44.02.003		- (Radiological Drinking Water Study – Basic Test ?- and ?-Activity)	6510
44.02.004		2 (Determination of Concentration 2 Radionuclides)	7500
44.02.005		4 (Determination of Concentration 4 Radionuclides)	14000
44.02.006		6 (Determination of Concentration 6 Radionuclides)	20400
44. (Investigation of Water and Soil Quality) / 44.03. (High Tech Investigations of Soil)			
44.03.001		: (Soil: Comprehensive Toxicological Evaluation)	18200
44.03.002		: (Soil: Agrochemical Evaluation)	9050
45. () (Profiles) / 45.01. (For children)			
45.01.001		(Testing for Kindergarten and School)	1200
45.01.002		(Pediatric Infections: Immune Response)	4540

45.01.003		: 0 14 (Healthy Child: for Children from 0 to 14 Years)	300
45. () (Profiles) / 45.02. (For women)			
45.02.001		VIP- (VIP-Survey for Women)	6700
45.02.002		: (Female Hormonal Profile: Ovarian Dysfunction, Menstrual Irregularities)	2100
45.02.003		: (Women's Oncorisk: Cervix)	1700
45.02.004		: (Miscarriage: Autoimmune Profile)	2700
45.02.005		(Assessment of Androgen Status)	1000
45.02.006		: (Pregnancy Planning: Diagnosis of Urogenital Tract Infection (UTI))	1500
45.02.007		: (Want to Become a Mother: Pregnancy Planning, Comprehensive Survey)	3500
45.02.008		TORCH- (ToRCH-Infections)	2000
45.02.009		: I (1-13) (Pregnancy: First Trimester (1-13 Weeks))	3500
45.02.010		: II (14-28) (Pregnancy: Second Trimester (14-28 Weeks))	400
45.02.011		: III (29-30) (Pregnancy: Third Trimester (29-30 Weeks))	2000
45. () (Profiles) / 45.03. (For men)			
45.03.001		VIP- (VIP-Survey for Men)	6500
45.03.002		: (Male oncologic risk: prostate)	800
45. () (Profiles) / 45.04. (Annual preventive examinations)			
45.04.001		(40) (Annual Check-Up after 40 Years of Age)	2000
45.04.002		: (40) (Healthy You ? Healthy Country: Annual Check-Up up to 40 Years of Age)	1500
45.04.003		: (Serum Biochemistry: Extended Profile)	1500
45.04.004		: (Serum Biochemistry: Minimum)	800
45.04.005		? (INBIOFLOR ? Comprehensive Study of Microflora Composition of Urogenital Tract (UGT))	3000
45. () (Profiles) / 45.05. (Checkup of household staff)			
45.05.001		« » (My Healthy Nurse)	4740
45. () (Profiles) / 45.06. (Cardiovascular disease risk assessment)			
45.06.001		(Preventing Heart and Blood Vessel Diseases)	1400
45.06.002		: (Lipid Profile: Extended)	3100
45.06.003		: (Lipid Profile: Screening)	350
45. () (Profiles) / 45.07. (Diagnosis of antiphospholipid syndrome (APS))			
45.07.001		(), (Antiphospholipid Syndrome, APS)	3500
45. () (Profiles) / 45.08. (Diagnosis of gastrointestinal tract (GI Tract))			
45.08.001		: () (Coeliac Disease: Gluten Intolerance)	4000
45.08.002		(GastroPanel)	3100
45. () (Profiles) / 45.09. (Assessment of liver function)			
45.09.001		: (Survey of Liver: Extended)	1550
45.09.002		: (Survey of Liver: Screening)	350

45.09.003		(Autoimmune Liver Disease: Screening)	4700
45. () (Profiles) / 45.10. (Non-invasive Assessment of Liver Fibrosis)			
45.10.001		(SteatoS reen)	5200
45.10.002		(FibroMax)	9900
45.10.003		(FibroTest)	8800
45.10.004		(FibroMax)	9900
45.10.005		(FibroTest)	7875
45. () (Profiles) / 45.11. (Assessment of renal function)			
45.11.001		: (Survey of Kidneys: Extended)	1400
45.11.002		: (Survey of Kidneys: Screening)	550
45. () (Profiles) / 45.12. (Hospital studies)			
45.12.001		(Hospitalization in Therapeutic Hospital)	1800
45.12.002		(Hospitalization in Surgical Hospital)	2400
45.12.003		: (Hospitalization in Surgical Hospital: Extended Survey)	4700
45. () (Profiles) / 45.13. (Diagnosis of connective tissue diseases (disorders of connective tissue))			
45.13.001		: (Joint Pain: Extended Survey)	3800
45.13.002		: (Joint Pain: Screening)	950
45. () (Profiles) / 45.14. (Diagnosis of diabetes)			
45.14.001		: (Diabetes Control: Extended)	2000
45.14.002		: (Diabetes Control: Screening)	550
45.14.003		: (Diabetes: Autoimmune Markers)	3200
45. () (Profiles) / 45.15. (Diagnosis and monitoring therapy for osteoporosis)			
45.15.001		(Diagnosis of Osteoporosis)	1800
45. () (Profiles) / 45.16. (Assessment of thyroid function)			
45.16.001		: (Thyroid Gland: Extended Survey)	1000
45.16.002		: (Thyroid Gland: Screening)	700
45. () (Profiles) / 45.17. (Sexually transmitted infections (STI))			
45.17.001		(HIV, Syphilis, Hepatitis B, C)	950
45.17.002		« : 6 () » (Comprehensive Study «Sex in City: 6 Infections (Blood Test)»)	1800
45.17.003		« : 14 + » (Comprehensive Study «Sex in City: 14 Infections + Smear on Flora»)	3200
45.17.004		« : 12 + () » (Comprehensive Study «Sex in City: 12 Infections (Urogenital Scraping)»)	2950
45.17.005		« : 8 + » (Comprehensive Study «Sex in City: 8 Infections + Smear on Flora»)	2400
45.17.006		(4 +): Chlamydia trachomatis, Neisseria gonorrhoeae, Trichomonas vaginalis, Mycoplasma genitalium, (Identification of Sexually Transmitted Infections (STI) Pathogens, Chlamydia trachomatis DNA,	1700
45.17.007		(Testing for Infection: Minimum)	550
45.17.008		(Annual Survey for Infections)	900
45. () (Profiles) / 45.18. (Weight Problems)			

45.18.001		() (Weight Problems: Primary Survey)	2100
45.18.002		: (Survey Before Diet: Minimum)	2000
45.18.003		: (Survey Before Diet: Additional)	4700
45. () (Profiles) / 45.19. (Upper respiratory tract infection (URI))			
45.19.001		(, ,) (Acute Respiratory Infections, ARI: Runny Nose, Cough, Sore Throat)	3700
45. () (Profiles) / 45.20. (Allergy)			
45.20.001		, , (Allergy to Animals, Dust, Mold)	3800
45.20.002		(Food Allergy)	4900
45.20.003		(Mold Allergy)	1700
45.20.004		(Plant Allergy)	2500
45. () (Profiles) / 45.21. (Hematological studies)			
45.21.001		(Diagnosis of Anemia)	2100
50. / 50.01.002.006.000			
50.01.002		() () ,	154
50.01.004			160
50.01.011			121
50.01.020		III	220
50.01.026		V C	550
50.01.028			253
50.01.035		-	1650
50.01.037			195
50.01.038			190
50.01.040			220
50.01.044		() -	132
50.01.045		C	528
50.01.047		(-)	660
50.01.050		-	440
50.01.051		(-)	341
50.01.052		()	880
50.01.055		()	176
50.01.056			605
50.01.059		VIII	1716
50.01.067		()	352
50. / 50.02.002.024.000			
50.02.018		- MTHFR: 1298, MTHFR: 677, MTR, MTRR	1001
50.02.019		- (F2, F5, F7, F13, FGB, ITGA2, ITGB3, PAI-1)	1760
/ 51.			
51.00.001			2000
51.00.002			2000
51.00.003		()	900
51.00.004		()	1500
51.00.005			1500
51.00.006			1500