



07.00.003	( / ) (17 Y- )	14870
07.00.004	( / ) (50-99,9%) ( 28 )	27870
07.00.005	( / ) (17 Y- )	14870
07.00.006	( / ) (17 Y- )	14870
<b>/ 08.</b>		
08.00.001		8000
08.00.002	ATP7B (1 .)	36500
<b>/ 09.</b>		
09.00.001	(4-5 ( 2 ), ( 5 ), ( 2 ), ( 2 ),	5000
<b>/ 10.</b>		
10.00.001	10	4000
10.00.002	1	450
10.00.003	" " 1	660
10.00.004	" " 10	6600
10.00.005	" " 20	11000
10.00.006	" " 10	6600
10.00.007	" " 20	11000
10.00.008	" " 10	6600
10.00.009	" " 20	11000
10.00.010	"HUR" 10 c	2000
10.00.011	( , ) SP 1000	200
10.00.012	"HUR" (60 )	200
10.00.014	3 ( 8 )-(60 -1 )	1600
10.00.015	( " 12 " )-(60 -1 )	4800
10.00.016	( " 8 " )-(60 -1 )	3200
10.00.017	5 ( 8 )-( 55 )-	900
10.00.018	-Trainer (30 )	300
10.00.019	-Trainer (60 )	550
10.00.020	( )	400
10.00.021		300
<b>/ 11.</b>		
11.00.025	- , 1,0 -	250
11.00.026	- , -	1000
11.00.027		2200
11.00.028	"VIBRAMATIC" ( )	450
11.00.029	"VIBRAMATIC" ( )	450
11.00.030	"VIBRAMATIC"	250
11.00.031	"VIBRAMATIC" ( , , )	800
11.00.032	( 1-3 )-	500
11.00.033	( 3-7 )	700
11.00.034	( 7-12 )	900
11.00.035	1	250
11.00.036	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
<b>/ 12.</b>		
12.00.032	( 500 )	430

12.00.033	( 1000 )	570
12.00.001	, , (10 )	600
12.00.002	, , c (10 )	970
12.00.003	(10 )	950
12.00.004	(10 )	1300
12.00.005	(1 )-(10 )	770
12.00.006	(2 )-(10 )	970
12.00.007	-(10 )	700
12.00.008	(10 )	150
12.00.009	(10 )	220
12.00.010	(10 )	165
12.00.011	( 400 )-(10 )	415
12.00.012	( 2,0 )-(10 )	230
12.00.013	(40 )	350
12.00.014	(2 )-(40 )	420
12.00.015	( 100 )-(40 )	480
12.00.016	( 200 )-(40 )	590
12.00.017	( 400 )(60 )	630
12.00.018	2,0 (10 )	677
12.00.019	(10 )	380
12.00.020	( 2,0 )-(10 )	415
12.00.021	( 2,0 )-(10 )	220
12.00.022	( 1,0 )-(10 )	309
12.00.023	( 5,0 )-(10 )	215
12.00.024	( 2,0 )-(10 )	200
12.00.025	( 50 )-(40 )	570
12.00.026	( 10,0 )-(40 )	700
12.00.027	( 10 )-(40 )	500
12.00.028	( 1,5 )-(10 )	405
12.00.029	( 8 )-(10 )	315
12.00.030	( 2,0 )-(10 )	190
12.00.031	( 400 )	695
<b>/ 13.</b>		
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	( ) 2- 1	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
<b>/ 14.</b>				
14.00.001			+	1000
14.00.002	-	-		700
14.00.003		-		500
14.00.004				500
14.00.005				500
<b>/ 15.</b>				
15.00.001	-	( )	" " 1	300
15.00.002	-	( )	" " 2	350
15.00.003			" -1"	350
15.00.004			" -1 "	350
15.00.005			" -1	330
15.00.006			" "	500
15.00.007	-		" " 1	450
15.00.008	-		" "2	500
15.00.009		1	" "	370
15.00.010		2	" "	400
15.00.011		3	" "	550
15.00.012			" "	400
15.00.013			" "	165
15.00.014			" " 1	300
15.00.015			" " 2	350
15.00.016			" "	330
15.00.017			" "	220
15.00.018			" "	330
15.00.019			-1,01	550
15.00.020		-1,01	1	400
15.00.021		-1,01	2	450
15.00.022			" "	330
15.00.023			" "	400
15.00.024			" -1"	350
15.00.025			" -1 "	220
15.00.026			" -1 "	400
15.00.027			" -1" " "	440
15.00.028			" -1"	440
15.00.029			" "	250
15.00.030			" -1"	468
15.00.031		1	" "	370
15.00.032		2	" "	400
15.00.033		3	" "	550
15.00.045		-		300
15.00.046			-	400
15.00.047			- +	500
15.00.048		( )	(15 )	350
15.00.049		( )	(20 )	400
15.00.050		( )	(30 )	450
15.00.051		1	(15 )	300
15.00.052		1	(30 )	350
15.00.053			- (20 )	780
15.00.034		-1,01	( )	230
15.00.035		-1,01		270
15.00.036	-	( )	" " -( )	380

15.00.037	-1,01	-( )	330
15.00.038	-1,01	(1 - 3000 )	730
15.00.039	" -1"	(1 -3000 )	780
15.00.040	" "	(1 -3000 )	870
15.00.041	-1,01		260
15.00.042	-1,01		205
15.00.043	-1,01		205
15.00.044	-1,01		300
<b>16.</b>	<b>/ 16.01.</b>		
16.01.001			1300
16.01.002			800
16.01.003			1000
16.01.004			650
16.01.005			500
<b>16.</b>	<b>/ 16.02.</b>		
16.02.001			1300
16.02.002			2000
16.02.003			800
<b>16.</b>	<b>/ 16.03.</b>		
16.03.001			1300
16.03.002			800
<b>16.</b>	<b>/ 16.04.</b>		
16.04.001			1300
16.04.002			800
<b>16.</b>	<b>/ 16.05.</b>		
16.05.001			1300
16.05.002			900
<b>16.</b>	<b>/ 16.06.</b>		
16.06.001			1300
16.06.002			800
<b>16.</b>	<b>/ 16.07.</b>		
16.07.001			1300
16.07.002			800
<b>16.</b>	<b>/ 16.08.</b>		
16.08.001		+	1100
16.08.002			500
16.08.003			600
16.08.004			450
<b>16.</b>	<b>/ 16.09.</b>		
16.09.001			1300
16.09.002			650
16.09.003			400
16.09.004			800
16.09.005			500
16.09.006			300
<b>/ 17.</b>			
17.00.003		(2 )	2400
17.00.004		(30 )	900
17.00.005		(60 )	1100
17.00.006		(2 )	2400
17.00.001			1300
17.00.002			800
<b>21.</b>	<b>(General Clinical Blood Tests) / 21.01.</b>		
<b>(General Clinical Blood Tests)</b>			

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
<b>22. (Immunohematology) / 22.01. (Immunohematology)</b>		
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
<b>23. (Assessment of Coagulation System) / 23.01. (Assessment of Coagulation System)</b>		
23.01.001	( ) ( ), - (Activated Partial Thromboplastin Time, APTT)	110
23.01.002	( ), ( ) (Prothrombin, rothrombin Time, PT, International Normalized Ratio, INR)	150
23.01.003	(Fibrinogen, FG)	120
23.01.004	III, % ( III, Antithrombin III, % Activity)	250
23.01.005	( ) (Thrombin Time, TT)	100
23.01.006	D- (D-Dimer)	520
23.01.007	( ) (Lupus Anticoagulant, LA )	500
23.01.008	C, % (Protein C, % Activity)	1300
23.01.009	S (Protein S, Free)	1300
23.01.010	(Plasminogen)	300
23.01.011	(Induced Platelet Aggregation)	100
23.01.012	( ) (Prothrombin Index, PTI)	100
23.01.013	( , , D- , , III, ) (Hemostatic System: PTI, INR, PT, )	2500
23.01.014	( ) (Coagulation (without Description))	2000
<b>24. (Blood Chemistry Tests) / 24.01. (Carbohydrates)</b>		
24.01.001	(Glucose)	70
24.01.002	(Fructosamine)	200
24.01.003	HbA1 (HbA1 , Glycated Hemoglobin, GHB)	350
24.01.004	(Lactate)	350
24.01.005	- 2 (2-Hour Oral Glucose Tolerance Test, OGTT, Glucose Concentration (Fasting and 2 Hours after Load), Venous Blood)	200
24.01.006	- 2 (2-Hour Oral Glucose Tolerance Test, OGTT, Glucose and C-Protein Concentration (Fasting and 2 Hours after Load), Venous Blood)	700
<b>24. (Blood Chemistry Tests) / 24.02. (Lipids)</b>		
24.02.001	( ) (Triglycerides)	70
24.02.002	( ) (Cholesterol Total)	70
24.02.003	( , , ?- ) (High-Density Lipoprotein Cholesterol, HDL Cholesterol)	110
24.02.004	( , , ?- ) (Low-Density Lipoprotein Cholesterol, LDL Cholesterol)*	110

24.02.005	( , ) (Very Low-Density Lipoprotein Cholesterol, VLDL cholesterol)	300
24.02.006	(a, ) (Lipoprotein (a), Lp (a))	450
24.02.007	1 ( 1, 1) (Apolipoprotein A1, Apo A1)	350
24.02.008	B ( B, ) (Apolipoprotein B, Apo B)	350
24.02.009	(Bile Acids)	2100
24.02.010	( ) (Cholesterol/HDL Ratio)*	0
<b>24. (Blood Chemistry Tests) / 24.03. (Proteins and Amino Acids)</b>		
24.03.001	(Albumin)	70
24.03.002	(Protein Total)	70
24.03.003	(Serum Protein Electrophoresis, SPE, SPE )*	200
24.03.004	- , . ( , ) (M-Gradient, Screening, Serum Protein Electrophoresis (SPEP), Immunofixation with Polyvalent Antiserum, Quan	2000
24.03.005	- , ( IgG, IgA, IgM, , ), (M-Gradient, Typing, Serum Protein Electrophoresis (SPEP), Immunofixation with Antisera	3200
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
<b>24. (Blood Chemistry Tests) / 24.04. (Assessment of renal function)</b>		
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 ystatin C Equation)	750
24.04.005	(Urea)	70
24.04.006	(Uric cid)	70
<b>24. (Blood Chemistry Tests) / 24.05. (Pigments)</b>		
24.05.001	(Bilirubin Total)	70
24.05.002	( Conjugated Bilirubin ) (Direct Bilirubin, DBIL,	70
24.05.003	( Indirect Bilirubin, IB, Free Blood Bilirubin)*	0
<b>24. (Blood Chemistry Tests) / 24.06. (Enzymes)</b>		
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	- (?- , ) ( lpha- milase, ?-Amylase)	80
24.06.004	- (P- ) (Pancreatic ?-Amylase)	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	( , L- , + ) (Lactate Dehydrogenase, LDH)	120
24.06.010	-1 ( -1, 1- , - ) (Lactate Dehydrogenase, Isoforms 1, Lactic Acid Dehydrogenase, LDH1, Alpha-Hydroxybutyrate Dehydrogenase, Alpha-HBDH)	200
24.06.011	(S- , II, S- , ) (Cholinesterase, Pseudocholinesterase, PCHE)	200

24.06.012	( ) (Acid Phosphatase, ACP)	180
24.06.013	( ) (Alkaline Phosphatase, ALP)	70
<b>24. (Blood Chemistry Tests) / 24.07. (Assessment of endocrine and secretory functions of gastrointestinal tract (GI Tract))</b>		
24.07.001	(Gastrin)	600
24.07.002	I (Pepsinogen I)	400
24.07.003	II (Pepsinogen II)	400
24.07.004	I II ( 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
<b>24. (Blood Chemistry Tests) / 24.08. (Bone Metabolism)</b>		
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
<b>24. (Blood Chemistry Tests) / 24.10. (Vitamins)</b>		
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
<b>24. (Blood Chemistry Tests) / 24.11. (Inorganic Matters)</b>		
24.11.001	/ / ( +/Potassium, Na+/Sodium, l-/Chloride, Serum)	220
24.11.002	(Ca) (Calcium Total)	70
24.11.003	(Ca <sup>2+</sup> , 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
<b>24. (Blood Chemistry Tests) / 24.12. (Specific proteins)</b>		
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
<b>24. (Blood Chemistry Tests) / 24.13. (Tumor Markers)</b>		



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
<b>25. (Assessment of endocrine system function) / 25.01. (Assessment of Pituitary Function)</b>		
25.01.001	( , ) (Adrenocorticotrophic 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
<b>25. (Assessment of endocrine system function) / 25.02. (Assessment of Thyroid Function)</b>		
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 T4Index, 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
<b>25. (Assessment of endocrine system function) / 25.03. (Assessment of adrenocortical function)</b>		
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
<b>25. (Assessment of endocrine system function) / 25.04. (Assessment of Androgen Status)</b>		
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 $\beta$ -Androstanediol Glucuronid, 3 $\beta$ -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
<b>25. (Assessment of endocrine system function) / 25.05. (Estrogens and Progestins)</b>		
25.05.001	( 2) (Estradiol, E2)	280
25.05.002	(Progesterone)	280
<b>25. (Assessment of endocrine system function) / 25.06. (Nonsteroidal Gonadal Factors)</b>		
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
<b>25. (Assessment of endocrine system function) / 25.07. (Monitoring of Pregnancy, Biochemical Markers of Fetal Status)</b>		
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
25.07.009	1- (First-Trimester Biochemical Markers (Astraia))	3200

<b>25. (Assessment of endocrine system function) / 25.08. (Assessment of Pancreatic Endocrine Function)</b>		
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
<b>25. (Assessment of endocrine system function) / 25.09. (Biogenic Amines, BA)</b>		
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	(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
<b>25. (Assessment of endocrine system function) / 25.10. (Calcium-regulating hormones)</b>		
25.10.001	(Calcitonin)	650
25.10.002	(Parathyroid Hormone, PTH)	300
<b>25. (Assessment of endocrine system function) / 25.11. (Adipose Tissue Hormones)</b>		
25.11.001	(Leptin)	700
<b>25. (Assessment of endocrine system function) / 25.12. (Regulation of Erythropoiesis)</b>		
25.12.001	(Erythropoetin)	900
<b>26. (Microelements) / 26.01. (Microelements, Serum, Venous Blood)</b>		
26.01.001	(Essential Vital Elements, Essential Trace Elements, Serum)	1500
26.01.002	(Toxic Trace Elements, Toxic Heavy Metals, Venous Blood)	1500
26.01.003	(Trace Elements, Serum, Venous Blood: Screening)	3300
26.01.004	(Cadmium (Cd), Serum)	850
26.01.005	(Cobalt (Co), Serum)	850
26.01.006	(Copper (Cu), Serum)	850
26.01.007	(Manganese (Mn), Serum)	850
26.01.008	(Selenium (Se), Serum)	850
26.01.009	(Zinc (Zn), Serum)	850
26.01.010	(Nickel (Ni), Serum)	850
26.01.011	(Gold (Au), Serum)	850
26.01.012	(Molybdenum (Mo), Serum)	850
26.01.013	(Thallium (Tl), Serum)	850
26.01.014	(Arsenic (As), Serum)	850
26.01.015	(Iodine, serum)	850
26.01.016	(Lithium (Li), serum)	900
26.01.017	(Cadmium (Cd), load)	850
26.01.018	(Cobalt (Co), load)	850
26.01.019	(Copper (Cu), load)	850

26.01.020	(Mn) (Manganese (Mn), food)	850
26.01.021	(Ni) (Nickel (Ni), food)	850
26.01.022	(Se) (Selenium (Se), food)	850
26.01.023	(Zn) (Zinc (Zn), food)	850
26.01.024	(Pb) (Lead (Pb), food)	850
26.01.025	(Hg) (Mercury (Hg), food)	850
<b>26.</b>	<b>(Microelements) / 26.02. (Microelements, Urine)</b>	
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	(Cu) (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
<b>26.</b>	<b>(Microelements) / 26.03. (Microelements, Hair)</b>	
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 (Rb), 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
<b>26.</b>	<b>(Microelements) / 26.04. (Microelements, Nails)</b>		
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
<b>27.</b>	<b>(Drug Monitoring) / 27.01.</b>	<b>(Drug Monitoring)</b>
27.01.001	( ) (Phenobarbitalum)	2100
27.01.002	( , ) (Phenytoin)	950
27.01.003	( , ) (Acidum Valproicum, Depakin, Convulex)	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
<b>28.</b>	<b>(Examinations of Urine) / 28.01.</b>	<b>(Clinical Tests)</b>
28.01.001	( ) (Complete Urinalysis, Microscopic Examination)	180
28.01.002	(Nechiporenko's Urine Test)	180
28.01.003	( ) (Sulkowitch Urine Calcium Test)	200
<b>28.</b>	<b>(Examinations of Urine) / 28.02.</b>	<b>( ) (Urine Biochemistry (Daily Excretion))</b>
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 acid, 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	( oxalates, Urine)	1400
28.02.014	- (Calcium-Creatinine Ratio, Random Urine)	400
<b>28.</b>	<b>(Examinations of Urine) / 28.03.</b>	<b>(Psychoactive Substances, Urine)</b>
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
<b>28.</b>	<b>(Examinations of Urine) / 28.04.</b>	<b>(Study of Kidney Stone)</b>
28.04.001	(Kidney Stone Analysis)	3000
<b>29.</b>	<b>(Examinations of Feces, Stool Examinations) / 29.01.</b>	<b>(Examinations of Feces, Stool Examinations)</b>
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 (Elastase 1, E1)	1800
29.01.010	-1- (Alpha-1-Antitrypsin, Feces)	2000
29.01.011	(Fecal Calprotectin)	2000
<b>30. (Serology Diagnosis of Infectious Disease) / 30.01.</b>		
<b>(Adenovirus Infection)</b>		
30.01.001	IgA (nti-Adenovirus IgA)	590
30.01.002	IgG (nti-Adenovirus IgG)	590
<b>30. (Serology Diagnosis of Infectious Disease) / 30.02.</b>		
<b>(Ascariasis)</b>		
30.02.001	IgG (Anti-Ascaris lumbricoides IgG)	600
<b>30. (Serology Diagnosis of Infectious Disease) / 30.03.</b>		
<b>(Aspergillosis)</b>		
30.03.001	Aspergillus fumigatus ( 3), - IgG (Aspergillus fumigatus, IgG, M3)	400
<b>30. (Serology Diagnosis of Infectious Disease) / 30.04.</b>		
<b>(Bacterial flora (Bacteroides spp.))</b>		
30.04.001	, (Bacteroides spp., DNA, Scrape of Urogenital Epithelial Cells)*	300
<b>30. (Serology Diagnosis of Infectious Disease) / 30.05.</b>		
<b>( ) (Borreliosis, Lyme Borreliosis, Lyme Disease, LD)</b>		
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	, ( orrelia burgdorferi, DNA, Cerebrospinal Fluid)*	500
<b>30. (Serology Diagnosis of Infectious Disease) / 30.06.</b>		
<b>(Brucellosis)</b>		
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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.07.</b>		
<b>- ( ) (Human Immunodeficiency Virus, HIV)</b>		
30.07.001	1 2 1 2 (HIV Ag/Ab Combo)	300
30.07.002	-1, (HIV RNA, Plasma)*	11000
<b>30. (Serology Diagnosis of Infectious Disease) / 30.08.</b>		
<b>- , ( ) (Human papillomavirus, HPV, HPV Infection)</b>		
30.08.001	39, 45, 51, 52, 56, 58, 59, 66, 68 + , o , 14 : 16, 18, 31, 33, 35, (HPV DNA, Scrape of Rectal Epithelial Cells, 14 Ty	800
30.08.002	39, 45, 51, 52, 56, 58, 59, 66, 68 + , o , 14 : 16, 18, 31, 33, 35, (HPV DNA, Scrape of Faucial Epithelial Cells, 14 Types (16, 18,	800
30.08.003	39, 45, 51, 52, 56, 58, 59, 66, 68 + , o , 14 : 16, 18, 31, 33, 35, (HPV DNA, Scrape of Urogenital Epithelial Cells, 14	800

30.08.004	, o 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,	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,	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	, 3 : 6, 11, 44 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 3 Types (6, 11, 44))	500
30.08.009	, 3 : 6, 11, 44 + (HPV DNA, Scrape of Rectal Epithelial Cells, 3 Types (6, 11, 44))	500
30.08.010	, 3 : 6, 11, 44 (HPV DNA, Scrape of Faucial Epithelial Cells, 3 Types (6, 11, 44))	500
30.08.011	, o 4 : 6, 11, 16, 18 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 4 Types (6, 11, 16, 18) Screening )	900
30.08.012	, o 15 : 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,	1000
30.08.013	, o 21 : 6, 11, 16, 18, 26, 31, 33, 35, 39, 44, 45, 51, 52, 53, 56, 58, 59, 66, 68, 73, 82 + (HPV DNA, Scrape of Urogenital Epithelial Cells, 21 Types (6,	2400
<b>30. (Serology Diagnosis of Infectious Disease) / 30.09.</b>		
<b>(Gardnerella vaginalis)</b>		
30.09.001	, (Gardnerella vaginalis, 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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.10.</b>		
<b>(Helminthiasis)</b>		
30.10.001	( ) (PRO Stool, Helminth Eggs)	300
30.10.002	« » ( , , ) (Parasites: Opisthorchis, Echinococcosis, Toxocara, Trichinella)	400
<b>30. (Serology Diagnosis of Infectious Disease) / 30.11. A</b>		
<b>(Hepatitis A Virus, HAV)</b>		
30.11.001	IgG (Anti-HAV IgG)	500
30.11.002	IgM (Anti-HAV IgM)	600
30.11.003	, (HAV RNA, Serum)*	500
<b>30. (Serology Diagnosis of Infectious Disease) / 30.12. B</b>		
<b>(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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.13. (Hepatitis C Virus, HCV)</b>		
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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.14. (Hepatitis D Virus, HDV) D</b>		
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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.15. (Hepatitis E Virus, HEV)</b>		
30.15.001	IgM E (Anti-HEV IgM)	800
30.15.002	IgG E (Anti-HEV IgG)	800
<b>30. (Serology Diagnosis of Infectious Disease) / 30.16. (G) (HGV, Hepatitis G Virus) G</b>		
30.16.001	G, (HGV RNA, Serum)*	600
<b>30. (Serology Diagnosis of Infectious Disease) / 30.17. ( 1 2 , Herpes simplex virus, HSV-1, HSV-2)</b>		
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 IgG, 1 2 (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, yping)*	450
30.17.019	1 2 , (HSV-1, 2 DNA, Scrape of Conjunctiva Epithelial Cells, yping)*	450
30.17.020	1 2 , (HSV-1, 2 DNA, Scrape of Skin Epithelial Cells, yping)*	450
30.17.021	1 2 , (HSV-1, 2 DNA, Blood, yping)*	600
30.17.022	1 2 , (HSV-1, 2 DNA, Urine, yping)*	500
30.17.023	1 2 , (HSV-1, 2 DNA, Scrape of Nasal Epithelial Cells, yping)*	450
30.17.024	1 2 , (HSV-1, 2 DNA, Scrape of Faucial Epithelial Cells, yping)*	450
30.17.025	1 2 , (HSV-1, 2 DNA, Serum, yping)*	600
30.17.026	1 2 , (HSV-1, 2 DNA, Saliva, yping)*	450
30.17.027	1 2 , (HSV-1, 2 DNA, Cerebrospinal Fluid, yping)*	450
30.17.028	1 2 , (HSV-1, 2 DNA, Prostatic Fluid, Semen, yping)*	450
30.17.029	1 2 , (HSV-1, 2 DNA, Scrape of Urogenital Epithelial Cells, yping)*	450
<b>30. (Serology Diagnosis of Infectious Disease) / 30.18.</b>		
<b>- 6 (Human herpesvirus 6, HHV-6)</b>		
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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.19.</b>		
<b>- 8 (Human herpesvirus 8, HHV-8)</b>		
30.19.001	IgG 8 (Anti-HHV-8 IgG)	600
<b>30. (Serology Diagnosis of Infectious Disease) / 30.20.</b>		
<b>( ) (Neisseria gonorrhoeae)</b>		
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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.23. (Entamoeba histolytica)</b>		
30.23.001	IgG (Anti-Entamoeba histolytica IgG)	600
<b>30. (Serology Diagnosis of Infectious Disease) / 30.24. (Shigella flexneri, Shigella sonnei)</b>		
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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.25. (Corynebacterium diphtheriae)</b>		
30.25.001	IgG (Anti-Diphtheria Toxoid IgG)	1100
30.25.002	(Corynebacterium diphtheriae Culture)	700
<b>30. (Serology Diagnosis of Infectious Disease) / 30.26. (Yersinia enterocolitica)</b>		
30.26.001	IgA Yersinia nterocolitica ( nti-Yersinia enterocolitica IgA)	550
30.26.002	IgG Yersinia nterocolitica ( nti-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 nterocolitica :3 (Yersinia enterocolitica O:3, IHA)	550
30.26.006	Yersinia nterocolitica :9 (Yersinia enterocolitica O:9, IHA)	550
<b>30. (Serology Diagnosis of Infectious Disease) / 30.27. (Campylobacter spp.)</b>		
30.27.001	(Campylobacter s p.) (Campylobacter spp., Stool Culture. Bacterial Identification)	550
<b>30. (Serology Diagnosis of Infectious Disease) / 30.28. (Candidiasis, Candida)</b>		
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	Semen)*	(Candida albicans, DNA, Prostatic Fluid,	250
30.28.010	albicans, DNA, Scrape of Urogenital Epithelial Cells)*	(Candida	250
30.28.011	(Candida albicans, Candida crusei, Candida glabrata), Candida crusei, Candida glabrata, DNA)*	(Candida albicans,	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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.29. (Flavivirus, Tick-borne encephalitis virus)</b>			
30.29.001	IgG	(Anti-Tick-borne Encephalitis Virus (TBEV)	300
30.29.002	IgM	(Anti-Tick-borne Encephalitis Virus (TBEV)	300
<b>30. (Serology Diagnosis of Infectious Disease) / 30.30. (Clostridium difficile)</b>			
30.30.001		(Clostridium difficile, Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	800
30.30.002	Toxin A Clostridium difficile, difficile. One Step Rapid Immun	chromotographic Assay)	700
<b>30. (Serology Diagnosis of Infectious Disease) / 30.31. (Bordetella pertussis, Whooping Cough)</b>			
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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.32. (Morbillivirus, Measles virus)</b>			
30.32.001		IgG ( nti-Measles IgG)	1100
30.32.002		Ig ( nti-Measles IgM)	800
<b>30. (Serology Diagnosis of Infectious Disease) / 30.33. (Rubella virus)</b>			
30.33.001		IgG ( nti-Rubella IgG)	380
30.33.002		Ig ( nti-Rubella Ig )	280
30.33.003		IgG ( nti-Rubella IgG, Immunoblot )	2200
30.33.004		IgM IgG Avidity)	900
30.33.005		(Rubella Virus, DNA, Serum)*	450
<b>30. (Serology Diagnosis of Infectious Disease) / 30.34. (Cryptosporidium parvum)</b>			
30.34.001		(Cryptosporidium parvum), (Cryptosporidium parvum. One Step Rapid Immun	600
<b>30. (Serology Diagnosis of Infectious Disease) / 30.35. (Lactobacillus spp.)</b>			
30.35.001		(Lactobacillus spp., DNA, Scrape of Urogenital Epithelial Cells)*	260
<b>30. (Serology Diagnosis of Infectious Disease) / 30.36. (Listeria monocytogenes, Listeriosis)</b>			
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	Fluid)*	(Listeria monocytogenes, DNA, Synovial	450
30.36.005	Cerebrospinal Fluid)*	(Listeria monocytogenes, DNA,	300
30.36.006	monocytogenes Culture. Bacteria Identification)	(Listeria monocytogenes, ) (Listeria	800
<b>30. (Serology Diagnosis of Infectious Disease) / 30.37.</b>			
<b>(Giardia lamblia intestinalis)</b>			
30.37.001	IgA, Total)	IgM, IgG, IgA ( nti-Giardia lamblia IgM, IgG,	600
30.37.002	Rapid Immun hromotographic Assay)	(Giardia liamblia), (Giardia liamblia. One Step	800
30.37.003	IgM	( nti-Giardia lamblia IgM)	300
<b>30. (Serology Diagnosis of Infectious Disease) / 30.38.</b>			
<b>(Neisseria meningitides)</b>			
30.38.001	meningitidis Culture. Bacteria Identification and Antibiotic Susceptibility Testing)	(Neisseria	800
<b>30. (Serology Diagnosis of Infectious Disease) / 30.39.</b>			
<b>(Skin and Nail Mycoses)</b>			
30.39.001		(Fungal Infections of Nails)	1000
30.39.002		(Fungal Infections of Skin)	1000
30.39.003	(skin))	( ) (Parasitic Fungi, Microscopy and Culture	1700
30.39.004	(nails))	( ) (Parasitic Fungi, Microscopy and Culture	1700
<b>30. (Serology Diagnosis of Infectious Disease) / 30.40.</b>			
<b>(Mycoplasmosis, Mycoplasma Infection)</b>			
<b>(Mycoplasma genitalium, Mycoplasma pneumoniae, Mycoplasma hominis)</b>			
30.40.001	Urine)*	(Mycoplasma hominis), (Mycoplasma hominis, DNA,	250
30.40.002	hominis, DNA, Prostatic Fluid, Semen)*	(Mycoplasma	250
30.40.003		(Mycoplasma hominis), (Mycoplasma hominis, DNA, Scrape of Urogenital Epithelial Cells)*	250
30.40.004	Urine)*	(Mycoplasma genitalium), (Mycoplasma genitalium, DNA,	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	DNA, Sputum)*	(Mycoplasma pneumoniae), (Mycoplasma pneumoniae,	430
30.40.008	pneumoniae, DNA, Plasma)*	(Mycoplasma	430
30.40.009		(Mycoplasma pneumoniae), (Mycoplasma pneumoniae, DNA, Scrape of Faucial Epithelial Cells)*	380
30.40.010	DNA, Saliva)*	(Mycoplasma pneumoniae), (Mycoplasma pneumoniae,	380
30.40.011		Ig IgG Mycoplasma hominis ( nti-Mycoplasma hominis Ig , IgG)	1100
30.40.012		Ig Mycoplasma hominis ( nti-Mycoplasma hominis Ig )	650
30.40.013		IgG Mycoplasma hominis ( nti-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 ( nti-Mycoplasma hominis IgA)	400
30.40.018		IgG IgA Mycoplasma hominis ( nti-Mycoplasma hominis IgG, IgA)	900
<b>30. (Serology Diagnosis of Infectious Disease) / 30.41.</b>			
<b>(Urogenital Tract Microbiocenosis)</b>			
30.41.001	(Lactob cillus 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 + , (FEMOFLOR?9, Scrape of Urogenital Epithelial Cells)*	1100
30.41.007	-17 + , (FEMOFLOR?17, Scrape of Urogenital Epithelial Cells)*	1760
30.41.008	? ? Comprehensive Study of Microflora Composition of Urogenital Tract (UGT)) (INBIOFLOR	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	? - ( ) (INBIOFLOR ? 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 ur alyticum) (Ureaplasma ur alyticum, 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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.42.</b>		
<b>(Mobiluncus curtisii, M. mulieris)</b>		
30.42.001	(Mobiluncus curtisii, DNA, Scrape of Urogenital Epithelial Cells)	250
<b>30. (Serology Diagnosis of Infectious Disease) / 30.43.</b>		
<b>( 2- , Norovirus, Group 2)</b>		
30.43.001	2- , (Norovirus, Group 2, RNA, Fecal)	500
<b>30. (Serology Diagnosis of Infectious Disease) / 30.44.</b>		
<b>( , Opisthorchis felineus)</b>		
30.44.001	IgG (Anti-Opisthorchis felineus IgG)	300
30.44.002	IgM, IgG ( ) (Anti-Opisthorchis felineus IgM, IgG, Circulating Immune Complexes (CIC))	300
<b>30. (Serology Diagnosis of Infectious Disease) / 30.45.</b>		
<b>(Acute Intestinal Infections)</b>		
30.45.001	, (Enterovirus, RNA, Fecal)	500
30.45.002	2- , (Norovirus, Group 2, RNA, Fecal)	500
30.45.003	, - (Acute Intestinal Infections, PCR, Fecal)	1300
30.45.004	, - , (Acute Intestinal Infections, PCR, Fecal)	1400
<b>30. (Serology Diagnosis of Infectious Disease) / 30.46.</b>		
<b>(Bordetella parapertussis)</b>		
30.46.001	(Bordetella pertussis/parapertussis, / ) (Bordetella pertussis/parapertussis, Nasopharyngeal Culture. Bacteria Identification)	900
<b>30. (Serology Diagnosis of Infectious Disease) / 30.47.</b>		
<b>(Parotitis epidemica, Mumps, Epidemic Parotitis)</b>		
30.47.001	A IgG (Anti-Mumps IgG)	700
30.47.002	A IgM (Anti-Mumps IgM)	700
<b>30. (Serology Diagnosis of Infectious Disease) / 30.48.</b>		
<b>(Entamoeba histolytica, Giardia intestinalis, Balantidium coli)</b>		
30.48.001	: , , (PRO Stool)	300

<b>30. (Serology Diagnosis of Infectious Disease) / 30.49.</b>			
<b>(Respiratory Syncytial Virus, RSV)</b>			
30.49.001	A (RSV) IgG	IgG	(Anti-Respiratory Syncytial Virus) 700
30.49.002	A (RSV) IgM	IgM	(Anti-Respiratory Syncytial Virus) 700
<b>30. (Serology Diagnosis of Infectious Disease) / 30.50.</b>			
<b>(Typhus, Rickettsia prowazekii)</b>			
30.50.001			(Rickettsia prowazekii, IHA) 550
<b>30. (Serology Diagnosis of Infectious Disease) / 30.51.</b>			
<b>(Rotavirus)</b>			
30.51.001	(Rotavirus), Agglutination)		(Rotavirus Direct Detection by Latex) 500
<b>30. (Serology Diagnosis of Infectious Disease) / 30.52.</b>			
<b>(Salmonella spp.)</b>			
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	Identification)		(Stool Culture (Salmonella spp., Shigella spp.). Bacteria) 600
30.52.009	Susceptibility Testing)		(Stool Culture (Salmonella spp., Shigella spp.). Bacteria Identification and Antibiotic) 800
30.52.010	Antibiotic Susceptibility and Bacteriophage Efficiency Testing)		(Stool Culture, Salmonella s p., Shigella s p. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing) 800
<b>30. (Serology Diagnosis of Infectious Disease) / 30.53.</b>			
<b>(Treponema pallidum, Syphilis)</b>			
30.53.001	RPR – est)		(Syphilis RPR (Rapid Plasma Reagins), nticardiolipin) 100
30.53.002	A IgG, Total)	IgM IgG Treponema pallidum,	(Anti-Treponema pallidum IgM, 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)		(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	Cerebrospinal Fluid)*		(Treponema pallidum, DNA, 250
30.53.014	Prostatic Fluid, Semen)*		(Treponema pallidum, DNA, 250
30.53.015			(Treponema pallidum, DNA, Scrape of Urogenital Epithelial Cells)* 250
<b>30. (Serology Diagnosis of Infectious Disease) / 30.54.</b>			
<b>(Staphylococcus aureus)</b>			

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 aureu	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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.55.</b>		
<b>(Tetanus)</b>		
30.55.001	IgG (Anti-Tetanus toxoid IgG)	1100
<b>30. (Serology Diagnosis of Infectious Disease) / 30.56.</b>		
<b>(Streptococcal Infection) (Streptococcus group A, B, S. pyogenes, S. agalactiae)</b>		
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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.57.</b>		
<b>(Strongyloides stercoralis)</b>		
30.57.001	IgG (Anti-Strongyloides stercoralis IgG)	1300
<b>30. (Serology Diagnosis of Infectious Disease) / 30.58.</b>		
<b>(Deltaretrovirus, Human T-Lymphotropic Virus, HTLV)</b>		
30.58.001	IgG (Anti-HTLV-1, 2 IgG)	400
<b>30. (Serology Diagnosis of Infectious Disease) / 30.59.</b>		
<b>(Toxocara canis)</b>		
30.59.001	IgG (Anti-Toxocara IgG)	220
<b>30. (Serology Diagnosis of Infectious Disease) / 30.60.</b>		
<b>(Toxoplasmosis, Toxoplasma gondii)</b>		
30.60.001	IgG (Anti-Toxoplasma gondii IgG)	300
30.60.002	IgM (Anti-Toxoplasma gondii IgM)	300
30.60.003	IgM, IgG (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	, Cerebrospinal Fluid)*	(Toxoplasma gondii, DNA,	170
<b>30. ( ) (Serology Diagnosis of Infectious Disease) / 30.61.</b>			
<b>( , Trichinella spiralis)</b>			
30.61.001	IgG	( nti-Trichinella IgG)	500
30.61.002	IgM IgG	(Anti-Trichinella IgM, IgG)	900
<b>30. ( ) (Serology Diagnosis of Infectious Disease) / 30.62.</b>			
<b>( Trichomonas vaginalis, Trichomoniasis)</b>			
30.62.001	IgG Trichomonas vaginalis ( nti-Trichomonas vaginalis IgG)		260
30.62.002	IgG IgA Trichomonas vaginalis ( nti-Trichomonas vaginalis IgG, IgA)		260
30.62.003	,	(Trichomonas vaginalis, DNA, Urine)*	250
30.62.004	, Prostatic Fluid, Semen)*	, (Trichomonas vaginalis, DNA,	250
30.62.005	, (Trichomonas vaginalis, DNA, Scrape of Urogenital Epithelial Cells)*		250
<b>30. ( ) (Serology Diagnosis of Infectious Disease) / 30.63.</b>			
<b>( Mycobacterium Tuberculosis)</b>			
30.63.001	IgM, IgA, IgG Mycobacterium tuberculosis, ( nti-Mycobacterium tuberculosis IgM, IgA, IgG, otal)		1500
30.63.002	, Exudate)*	(Mycobacterium tuberculosis, DNA,	300
30.63.003	, Sputum)*	(Mycobacterium tuberculosis, DNA,	500
30.63.004	, tuberculosis, DNA, Menstrual Blood)*	(Mycobacterium	300
30.63.005	, (Mycobacterium tuberculosis, DNA, Urine)*		250
30.63.006	, DNA, Serum)*	(Mycobacterium tuberculosis,	400
30.63.007	, tuberculosis, DNA, Synovial Fluid)*	(Mycobacterium	400
30.63.008	, tuberculosis, DNA, Cerebrospinal Fluid)*	(Mycobacterium	300
30.63.009	, tuberculosis, DNA, Prostatic Fluid, Semen)*	, (Mycobacterium	250
<b>30. ( ) (Serology Diagnosis of Infectious Disease) / 30.64.</b>			
<b>( Ureaplasmosis, Ureaplasma spp., Ureaplasma urealyticum, Ureaplasma parvum)</b>			
30.64.001	IgG Ureaplasma urealyticum ( nti-Ureaplasma urealyticum IgG)		280
30.64.002	IgA Ureaplasma urealyticum ( nti-Ureaplasma urealyticum IgA)		280
30.64.003	IgG IgA Ureaplasma urealyticum ( nti-Ureaplasma urealyticum IgG, IgA)		280
30.64.004	(Ureaplasma urealyticum) ( -960), urealyticum (T-960), DNA, Urine)*	(Ureaplasma	250
30.64.005	(Ureaplasma urealyticum) ( -960), (Ureaplasma urealyticum (T-960), DNA, Prostatic Fluid, Semen)*	,	250
30.64.006	(Ureaplasma urealyticum) ( -960), Urogenital Epithelial Cells)*	(Ureaplasma urealyticum (T-960), DNA, Scrape of	250
30.64.007	(Ureaplasma parvum), (Ureaplasma parvum, DNA, Urine)*		250
30.64.008	(Ureaplasma arvum), parvum, DNA, Prostatic Fluid, Semen)*	, (Ureaplasma	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), DNA, Scrape of Urogenital Epithelial Cells)*	(Ureaplasma urealyticum + Ureaplasma arvum,	250
<b>30. ( ) (Serology Diagnosis of Infectious Disease) / 30.65.</b>			
<b>( Helicobacter pylori)</b>			
30.65.001	A IgG Helicobacter ylori (Anti-Helicobacter pylori IgG)		300

30.65.002	A	IgM Helicobacter pylori (Anti-Helicobacter pylori IgM)	700
30.65.003	A	IgA Helicobacter pylori (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
<b>30. (Serology Diagnosis of Infectious Disease) / 30.66. Chlamydia pneumoniae, (Chlamydia pneumoniae) (Chlamydia pneumoniae, Chlamydia pneumoniae, Chlamydia pneumoniae)</b>			
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 Chlamydia pneumoniae (anti-Chlamydia pneumoniae IgM, IgG)	350
30.66.006		IgG ( ) Chlamydia trachomatis (Anti-cHSP60 IgG)	900
30.66.007		(Helicobacter pylori, DNA, Biopsies of Gastric Mucosa and/or Duodenum, PCR)	700
30.66.008		(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Exudate)*	250
30.66.009		(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Scrape of Conjunctiva Epithelial Cells)*	250
30.66.010		(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Urine)*	250
30.66.011		(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Scrape of Rectal Epithelial Cells)*	250
30.66.012		(Chlamydia trachomatis), (Chlamydia trachomatis, DNA, Scrape of Faucial Epithelial Cells)*	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), (Chlamydia trachomatis, DNA, Scrape of Urogenital Epithelial Cells)*	250
30.66.017	A	IgA Chlamydia pneumoniae (Anti-Chlamydia pneumoniae IgA)	550
30.66.018	A	IgM Chlamydia pneumoniae (Anti-Chlamydia pneumoniae IgM)	200
30.66.019	A	IgG Chlamydia pneumoniae (Anti-Chlamydia pneumoniae IgG )	200
30.66.020		(Chlamydia pneumoniae), (Chlamydia pneumoniae, DNA, Sputum)*	900
30.66.021		(Chlamydia pneumoniae), (Chlamydia pneumoniae, DNA, Plasma)*	550
30.66.022		(Chlamydia pneumoniae), (Chlamydia pneumoniae, DNA, Scrape of Faucial Epithelial Cells)*	500
30.66.023		(Chlamydia pneumoniae), (Chlamydia pneumoniae, DNA, Saliva)*	500
<b>30. (Serology Diagnosis of Infectious Disease) / 30.67. Cytomegalovirus, CMV, Cytomegalovirus Infection</b>			
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	(CMV DNA, Serum)*	3350
30.67.012	(CMV DNA, Saliva)*	250
30.67.013	(CMV DNA, Cerebrospinal Fluid)*	250
30.67.014	(CMV DNA, Prostatic Fluid, Semen)*	250
30.67.015	(CMV DNA, Scrape of Urogenital Epithelial Cells)*	250
<b>30. ( , Enterobius spp.) (Serology Diagnosis of Infectious Disease) / 30.68.</b>		
30.68.001	( ), ( nterobiasis, Swab)	380
<b>30. ( ) (Serology Diagnosis of Infectious Disease) / 30.69. (Epstein-Barr Virus, EBV, EBV Infection)</b>		
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 (Anti-EBV Viral Capsid Antigens (VCA) Ig )	280
30.69.012	IgG (Anti-EBV Nuclear Antigen (EBNA) IgG )	250
30.69.013	IgG (Anti-EBV Early Antigen (EA) IgG )	250
30.69.014	IgG (Anti-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
<b>30. ( , Echinococcus spp.) (Serology Diagnosis of Infectious Disease) / 30.70.</b>		
30.70.001	IgG (Anti-Echinococcus IgG)	280
<b>30. ( , Escherichia coli) (Serology Diagnosis of Infectious Disease) / 30.71.</b>		
30.71.001	(Escherichia coli O157:H7, ), (Escherichia coli O157:H7. One Step Rapid Immun chromatographic Assay)	700
<b>31. : (Microbiological investigations) / 31.01. (Microbiological investigations)</b>		
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 chromatographic 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 Immun chromatographic 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 Immunochromatographic Assay)	650
31.01.059	(Giardia lamblia), (Giardia lamblia. One Step Rapid Immunochromatographic 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 Immunochromatographic Assay)	650
31.01.065	(Helicobacter pylori), (Helicobacter pylori. One Step Rapid Immunochromatographic Assay)	650
31.01.066	Toxin A Clostridium difficile, (Toxin A Clostridium difficile. One Step Rapid Immunochromatographic 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
31.01.047	(Eye Culture. Bacteria Identification, Antibiotic Susceptibility and Bacteriophage Efficiency Testing)*	0
<b>32. (Immunoglobulins, Total) (Immunological Studies) / 32.02.</b>		
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
<b>32. (Immunological Studies) / 32.03. (Specific Proteins)</b>		
32.03.001	(Eosinophil Cationic Protein, ECP)	900
<b>32. (Immunological Studies) / 32.04. (Complement Components)</b>		
32.04.001	3, 4 (Complement components C3, C4)	650
<b>32. (Immunological Studies) / 32.05. (Cytokines)</b>		
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
<b>34. (Diagnosis of Autoimmune Diseases) / 34.01. (Connective-Tissue Diseases, CTDs)</b>		

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
<b>34. (Diagnosis of Autoimmune Diseases) / 34.02. (Antiphospholipid Syndrome, APS)</b>		
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- hosphatidylserine, IgG, IgM)	1500
34.02.004	IgA, IgM, IgG , ( ardiolipin 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- hosphatidylserine/ rothrombin ntibodies, Anti-PS/PT, IgG, IgM, Total)	1400
34.02.010	IgG V (Annexin V ntibodies, aAnV, IgG)	1300
34.02.011	Ig V (Annexin V ntibodies, aAnV, Ig )	1400
34.02.012	, IgG, IgM	1500
34.02.013	( , , IgG, IgM)	2650
34.02.014	, IgG, IgM; -2- 1) ( ;	3600
<b>34. (Diagnosis of Autoimmune Diseases) / 34.03. (Immunological Infertility factors)</b>		
34.03.001	(Anti-Spermatozoa Antibodies, ASA, Serum)	500
34.03.002	(Anti-Spermatozoa Antibodies, ASA, Semen)	1450
<b>34. (Diagnosis of Autoimmune Diseases) / 34.04. (Antiphospholipid Syndrome, APS)</b>		
34.04.001	( ) (Rheumatoid Factor, RF)	150
34.04.002	( ) (Anti- yclic Citrullinated Peptide, anti-CCP)	1200
34.04.003	IgG ( , ) (Anti- eratin ntibodies, AKA, Anti-Filaggrin ntibodies, 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
<b>34. (Diagnosis of Autoimmune Diseases) / 34.05. (Vasculitis and Renal Lesions)</b>		
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- proteinase-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
<b>34. (Diagnosis of Autoimmune Diseases) / 34.06. (Autoimmune Endocrinopathies: Diabetes mellitus type 1)</b>		
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
<b>34. (Diagnosis of Autoimmune Diseases) / 34.07. (Autoimmune Thyroid diseases)</b>		
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
<b>34. (Diagnosis of Autoimmune Diseases) / 34.08. (Autoimmune adrenal insufficiency)</b>		
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
<b>34. (Diagnosis of Autoimmune Diseases) / 34.09. (Autoimmune Skin Diseases)</b>		



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 Dermatoses Diagnostics profile (antibodies to epidermis desmosomes, antibodies to skin basal membrane))	3700
<b>34. (Diagnosis of Autoimmune Diseases) / 34.10. (Autoimmune Disorders of Gastrointestinal Tract (GI Tract, GIT). Coeliac Disease)</b>		
34.10.001	IgA IgG (Anti- endomysial antibodies, Anti-EMA, IgA, IgG, Total)	1200
34.10.002	Ig (Anti- endomysial antibodies, 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-Saccharomyces Cerevisiae Antibodies, ASCA, IgG)	1150
34.10.011	Ig (Anti-Saccharomyces 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	(Anti-pancreatic autoantibodies, pANCA)	1600
34.10.015	cANCA, IgG; /ANCA, IgA; ASCA, IgG, IgA	3600
34.10.016	(Anti-pancreatic autoantibodies, pANCA, IgG; IgA)	1500
34.10.017	(Anti-pancreatic autoantibodies, pANCA, IgG; IgA)	2600
34.10.018	(Fecal Calprotectin)	2000
34.10.019	IgG IgA (Anti-Tissue Transglutaminase, tTG, Anti-tTG, IgG, IgA)	1200
34.10.020	« » (Cereal Intolerance profile (gliadin IgG antibodies, gliadin IgA antibodies, tissue transglutaminase 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	(Anti-pancreatic autoantibodies, pANCA, IgG IgA, Autoantibodies against Exocrine Pancreas, Pancreatic Antibodies, PAB)	1400

<b>34. (Diagnosis of Autoimmune Diseases) / 34.11. (Autoimmune Liver Diseases)</b>		
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 (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
<b>34. (Diagnosis of Autoimmune Diseases) / 34.12. (Autoimmune Neurological Diseases)</b>		
34.12.001	IgA, IgG, IgM (Anti-Aquaporin-4 Receptor 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 (Anti-GM1 Antibodies, Anti-GQ1b Antibodies, Anti-Gangliosideantibodies, Ganglioside Antibodies Panel, Total)	4900
34.12.004	IgG (Myositis-Specific Panel: Mi-2, Ku, PM-Scl 100/75; Jo1 PL-7 PL-12 EJ OJ; SRP, SSA (Ro52))	3200
34.12.005	(Acetylcholine Receptor Antibodies, Anti-AChR, Total)	4650
34.12.006	IgG (Anti-Neuronal Antibodies, Blot-Line (Hu (ANNA1), Yo-1 (PCA1), CV2, 2, Ri (ANNA2), 2, Ri (ANNA2), Amphiphysin))	4650
34.12.007	IgA, IgG, IgM (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-P1NP, Total)	1000
<b>34. (Diagnosis of Autoimmune Diseases) / 34.13. (Autoimmune Lung Diseases, Autoimmune Heart Diseases)</b>		
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
<b>34. (Diagnosis of Autoimmune Diseases) / 34.14. (Immune Thrombocytopenia, ITP)</b>		
34.14.001	IgG (Platelet antibodies IgG, Indirect)	1400
<b>35. (Allergy Research) / 35.01. (IgE, Total)</b>		
35.01.001	E (IgE, (Immunoglobulin Total, IgE Total)	300
<b>35. (Allergy Research) / 35.02. (Determination of Specific IgE: Comprehensive Allergy Panels) IgE:</b>		
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
<b>35. (Allergy Research) / 35.03. (Determination of Specific IgE: Food Allergens) IgE:</b>		
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

<b>35.</b>	<b>(Allergy Research) / 35.04.</b>	<b>IgE:</b>
<b>(Determination of Specific IgE: Household Allergens)</b>		

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

<b>35. (Allergy Research) / 35.05. (Determination of Specific IgE: Allergens of Animal Origin) IgE:</b>		
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
<b>35. (Allergy Research) / 35.06. (Determination of Specific IgE: Allergens of Plant Pollen) IgE:</b>		
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
<b>35. (Allergy Research) / 35.07. (Determination of specific IgE: professional allergens) IgE:</b>		
35.07.001	, IgE (Latex, IgG, K82)	370
<b>35. (Allergy Research) / 35.08. (Determination of Specific IgG: Food Allergens) IgG:</b>		
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), -	IgG (Baker's Yeast, IgG, F45)	370
35.08.027	(F95), -	IgG (Peach, IgG, F95)	370
35.08.028	(F403), -	IgG (Brewer's Yeast, IgG, F403)	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
<b>35. (Allergy Research) / 35.09. (Determination of Specific IgG: Household Allergens)</b>			<b>IgG:</b>
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), -	IgG (Penicillium notatum, IgG, M1)	370
35.09.003	Cladosporium herbarum ( 2), - M2)	IgG (Cladosporium herbarum, IgG,	370
35.09.004	Aspergillus fumigatus ( 3), - M3)	IgG (Aspergillus fumigatus, IgG,	370
35.09.005	Alternaria tenuis ( 6), -	IgG (Alternaria tenuis, IgG, M6)	370
35.09.006	Dermatophagoides pteronyssinus (D1), - pteronyssinus, IgG, D1)	IgG (Dermatophagoides	370
35.09.007	Dermatophagoides farinae (D2), - IgG, D2)	IgG (Dermatophagoides farinae,	370
35.09.008	Dermatophagoides microceras (D3), - microceras, IgG, D3)	IgG (Dermatophagoides	370
35.09.009	/Greer ( 1), -	IgG (House Dust – Greer, IgG, H1)	370
35.09.010	/Stier ( 2), -	IgG (House Dust – Hollister-Stier, IgG, H2)	370
<b>35. (Allergy Research) / 35.10. (Determination of Specific IgG: allergens of Animal Origin)</b>			<b>IgG:</b>
35.10.001	, ( 1), -	IgG (Cat Dander-Epithelium, IgG, E1)	370
35.10.002	, ( 2), -	IgG (Dog Epithelium, IgG, E2)	370
<b>36. (Study of Sperm) / 36.03. (Antispermatozoal Antibodies)</b>			
36.03.001	(Anti-Spermatozoa Antibodies, ASA, Serum)		500
36.03.002	(Anti-Spermatozoa Antibodies, ASA, Semen)		1450
<b>37. (Cytological Research) / 37.01. (Liquid-Based Cytology, LBC, Pap Stain)</b>			
37.01.001	( NovaPrep ®) (Liquid Based Cytology, LBC. Cervical Screening (Papanicolaou Stain (Pap Stain), Technology NovaPrep ®))		1450
<b>37. (Cytological Research) / 37.02. (Papanicolaou Stain)</b>			<b>? -</b>
37.02.001	( ) (Cytological Examination: Cervix, Pap-test)		900
37.02.002	( , - ) (Cytological Examination: Different Biomaterial except Cervix, Pap-test)		900

<b>37. (Cytological Research) / 37.03.</b>		
<b>(Romanowsky-Giemsa Stain)</b>		
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
<b>37. (Cytological Research) / 37.04.</b>		
<b>(Cytological Examination of Early Stained Slides with Cytological Smear)</b>		
37.04.001	( ) (Consultation of Finished Cytological Preparations (1 Glass))	500
<b>38. (Histological Examination) / 38.01.</b>		
<b>(Histology, hematoxylin and eosin stain (H&amp;E stain, HE stain))</b>		
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
<b>39. (Cytogenetic Research) / 39.01. (Cytogenetic Research)</b>		
39.01.001	(Karyotype)	5000
<b>40. (Genetic Predispositions) / 40.01. (Genetic VIP-Profiles) VIP-</b>		
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	e ( F2, F5) (Risk of Oral Contraceptives, Ocs (Genes F2, F5))	2100
40.02.009	e ( 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
40.02.017	- : ( DAT, OPRM1, ANKK1, ALDH2, ADH2) (Alcoholism and Drug Addiction, Tendency to Develop: Full Panel (Genes DAT, OPRM1, ANKK1, ALDH2, ADH2))	6800
40.02.018	- : ( DAT, OPRM1, ANKK1, ALDH2, ADH2) ( - ) (Alcoholism and Drug Addiction, Tendency to Develop: Full Panel (Genes DAT, OPRM1, ANKK1, ALDH2, ADH2) (without	5800
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	(Thrombotic Tendency in Pregnancy: Minimum (Genes F2, F5) (without Description))	1900
40.03.006	(Gestosis and Placental Insufficiency (Genes ACE, AGT, MTHFR, MTRR, MTR, F2, F5))	9000
40.03.007	(Gestosis and Placental Insufficiency (Genes ACE, AGT, MTHFR, MTRR, MTR, F2, F5) (without Description))	8000
40.03.008	(Habitual Miscarriage, Thrombotic Tendency in Pregnancy: Extended Panel (Genes MTHFR, MTRR, MTR, F2, F5))	4000
40.03.009	(Habitual Miscarriage, Thrombotic Tendency in Pregnancy: Extended Panel (	3000
40.03.010	(Isolated Malformations in Fetus (Genes MTHFR, MTRR, MTR))	2800
40.03.011	(Isolated Malformations in Fetus (Genes MTHFR, MTRR, MTR) (without Description))	2200
40.03.012	(Thrombotic Complications of Ovulation Induction (Genes F2, F5))	2200
40.03.013	(Thrombotic Complications of Ovulation Induction (Genes F2, F5) (without Description))	1800
40.03.014	(Congenital Adrenal Hyperplasia (CAH), Gene CYP21OHB, 9 Freq. Mut.)	15500
40.03.015	(Risk of Oral Contraceptives, Ocs (Genes F2, F5))	2200
40.03.016	(Risk of Oral Contraceptives, OCs (Genes F2, F5) (without Description))	1800
40.03.017	(Folic Acid Metabolism (Genes MTHFR, MTRR, MTR))	2800
40.03.018	(Folic Acid Metabolism (Genes MTHFR, MTRR, MTR) (without Description))	2200
40.03.019	(Hereditary Breast and/or Ovarian Cancer, HBOC (Genes BRCA1, BRCA2))	4300
40.03.020	(Hereditary Breast and/or Ovarian Cancer, HBOC (Genes BRCA1, BRCA2) (without Description))	3700
40.03.021	(Hereditary Breast and/or Ovarian Cancer) / BRCA1, BRCA2, CHEK2, NBN	8000
40.03.022	(Male Sterility (Genes AR, CFTR; AZF-Region; Karyotype)) ;	16300
40.03.023	(Genetic Factors of Male Infertility (Genes AR, CFTR; AZF-Region))	13500
40.03.024	(Impairment of Spermatogenesis: Full Panel (AZF-Region))	8900
40.03.025	(Impairment of Spermatogenesis: Full Panel (AZF-Region) (without Description))	9000
<b>40. (Genetic Predispositions) / 40.04. HLA- (HLA (Human Leukocyte Antigens) Typing and Rh factor)</b>		
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
<b>40. (Genetic Predispositions) / 40.05.20.5 (Blood-Clotting Sequence)</b>		
40.05.001	(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, F7, PAI-1))	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
<b>40. (Genetic Predispositions) / 40.06. (Heart and Blood Vessel (Cardiovascular) Diseases)</b>		
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
<b>40. (Genetic Predispositions) / 40.07. (Gastrointestinal Diseases)</b>		
40.07.001	( DLG5, NOD2, OCTN1, OCTN2) (Crohn's Disease (Genes DLG5, NOD2, OCTN1, OCTN2))	5300
<b>40. (Genetic Predispositions) / 40.08. (Central Nervous System (CNS) Diseases)</b>		
40.08.001	( ApoE) (Alzheimer's Disease (Gene ApoE))	3000

<b>40. (Genetic Predispositions) / 40.09.</b>		
<b>(Cancer Diseases)</b>		
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) ( 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	/ BRCA1, 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 <sup>-</sup> ( ): 617V/617F JAK2 (Marker of Ph-negative Chronic Myeloproliferative Disorders (cMPD): quantifica	4500
40.09.013	Ph <sup>-</sup> ( ): 617F JAK2 (Marker of Ph-negative Chronic Myeloproliferative Disorders (cMPD): qualitative assessment of presence of	3600
<b>40. (Genetic Predispositions) / 40.10.</b>		
<b>(Alcohol and Drug Addiction)</b>		
40.10.001	- : ( DAT, OPRM1, ANKK1, ALDH2, ADH2) (Alcoholism and Drug Addiction, Tendency to Develop: Full Panel (Genes DAT, OPRM1, ANKK1, ALDH2, ADH2))	6600
40.10.002	- : ( DAT, OPRM1, ANKK1, ALDH2, ADH2) ( Alcoholism and Drug Addiction, Tendency to Develop: Full Panel (Genes DAT, OPRM1, ANKK1, ALDH2, ADH2) (withou	5700
40.10.003	: ( ) ( DAT, OPRM1, ANKK1) (Alcoholism and Drug Addiction: Abridged Panel, without Genetics of Alcohol Metabolism (Genes DAT, OPRM1, ANKK1))	4300
40.10.004	: ( ) ( DAT, OPRM1, ANKK1) ( Alcoholism and Drug Addiction: Abridged Panel, without Genetics of Alcohol Metabolism (Genes DAT, OPRM1,	3600
40.10.005	: ( OPRM1, ANKK1) (Alcoholism and Drug Addiction: Physical Dependence (Genes OPRM1, ANKK1))	3000
40.10.006	: ( OPRM1, ANKK1) ( Alcoholism and Drug Addiction: Physical Dependence (Genes OPRM1, ANKK1) (without Description))	2500
40.10.007	: ( DAT) (Alcoholism and Drug Addiction: Psychological Dependence (Gene DAT))	2000
40.10.008	: ( DAT) ( - ) (Alcoholism and Drug Addiction: Psychological Dependence (Genes DAT) (without Description))	1400
40.10.009	( ALDH2, ADH2) (Genetic Factors on Metabolism of Alcohol (Genes ALDH2, ADH2))	2700

40.10.010	( ALDH2, ADH2) ( - ) (Genetic Factors on Metabolism of Alcohol (Genes ALDH2, ADH2) (without Description))	2500
<b>40. (Genetic Predispositions) / 40.11. (Metabolic Disease, Metabolic Disturbance)</b>		
40.11.001	1- 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
<b>40. (Genetic Predispositions) / 40.12. (Pharmacogenetics)</b>		
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	N- 2 (Hydr	5500
40.12.004	N- 2 ( NAT2) (Isoniazidum. Genetic Markers of Increased Risk	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 ), ( )	1000
40.12.015	(?- ) CYP2D6. (?-Adrenergic Blockers. Gene CYP2D6. Pharmacogenetics)	10000
<b>40. (Genetic Predispositions) / 40.13. (Detoxification System of Xenobiotics and Carcinogens)</b>		
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
<b>40. (Genetic Predispositions) / 40.14. (Services of Physician-Geneticist)</b>		
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
<b>40. (Genetic Predispositions) / 40.01. VIP- (Genetic VIP-Profiles)</b>		
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
<b>41. (Hereditary Monogenic Diseases) / 41.01. (Hereditary Monogenic Diseases)</b>		
41.01.104	« » TRPV4, « . » . (Distal Spinal Muscular Atrophy Congenital Non-Progressive, Gene TRPV4, Hot-Point Mut.)	0
41.01.113	CRYAA, . (Cataract, Gene CRYAA, 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
41.01.003	(1-). « » ELOVL4, « » . . (Stargardt Disease 1, STGD1, Fundus Flavimaculatus Included, Gene ELOVL4, Hot-Point Mut.)	6000
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
41.01.010	. CRX, . (Leber Congenital Amaurosis 1, LCA, Gene CRX, Mut.)	10700
41.01.011	. LCA5, . (Leber Congenital Amaurosis 1, LCA, Gene LCA5, Mut.)	30600
41.01.012	. LRAT, . (Leber Congenital Amaurosis 1, LCA, Gene LRAT, Mut.)	10700
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, Mut.)	FBN1, . (Geleophysic Dysplasia 1, Gene	216000
41.01.067	IX, Mut.)	IX B, . (Hemophilia B, Gene Factor	26000
41.01.068	-IgD Syndrome, Gene MVK, Hot-Point Mut.)	« » MVK, « .» . . (Hyper-IgD	9000
41.01.069	-IgD Mut.)	CD40LG, . (Hyper-IgD Syndrome, Gene CD40LG,	37000
41.01.070	-IgM Mut.)	CD40LG, . (Hyper-IgM Syndrome, Gene CD40LG,	17900
41.01.071	(Hyperkalemic Periodic Paralysis Type 2, Exons 13, 24 Gene SCN4A, Mut.)	13 24 SCN4A, .	9650
41.01.072	Periodic Paralysis Type 2, Gene SCN4A, Mut.)	SCN4A, . (Hyperkalemic	81450
41.01.073	KRT1, . (Epidermolytic Hyperkeratosis, Gene KRT1, Mut.)		22250
41.01.074	KRT9, . (Epidermolytic Hyperkeratosis, Gene KRT9, Mut.)		27000
41.01.075	Cardiomyopathy, Gene CAV3, Mut.)	CAV3, . (Familial Hypertrophic	8320
41.01.076	Cardiomyopathy, Gene TNNT2, Mut.)	TNNT2, . (Familial Hypertrophic	41640
41.01.077	(Hyperphenylalaninemia BH4-Deficient, Gene GCH1, Mut.)	GCH1, .	23250
41.01.078	(Hyperphenylalaninemia BH4-Deficient, Gene PTS, Mut.)	PTS, .	19550
41.01.079	(Hyperphenylalaninemia BH4-Deficient, Gene QDPR, Mut.)	QDPR, .	26950
41.01.080	(Hypokalemic Periodic Paralysis Type 1, Exons 12, 18, 19 Gene SCN4A, Mut.)	12, 18 19 SCN4A,	12350
41.01.081	D- (PHEX, . (Hypophosphatemic Vitamin D-Resistant Rickets, Gene PHEX, Mut. )		82450
41.01.082	Freq. Mut.)	FGFR3, . . (Hypochondroplasia, Gene FGFR3,	8800
41.01.083	3A, Gene CYP1B1, Mut.)	CYP1B1, . (Primary Congenital Glaucoma 3A, PCG	12380
41.01.084	(Primary Open Angle Glaucoma 1A, POAG 1A, Gene CYP1B1, Mut.)	CYP1B1, .	12380
41.01.085	(Primary Open Angle Glaucoma 1A, POAG 1A, Gene MYOC, Mut.)	MYOC, .	12380
41.01.086	And Diabetes Syndrome, Gene HNF1B, Mut.)	HNF1B, . (Renal Cysts	33250
41.01.087	S H, . (Holoprosencephaly 3, Gene S H, Mut.)		14850
41.01.088	(Medium-Chain Acyl-CoA Dehydrogenase Deficiency, MCAD-Deficiency, Gene ACADM, Freq. Mut.)	ACADM, . .	4460
41.01.089	Primary Carnitine Deficiency, SPCD, Carnitine Deficiency Systemic Primary, CDSP, Gene SLC22A5, Mut.)	SLC22A5, . (Systemic	3850
41.01.090	SLC26A2, Mut.)	SLC26A2, . (Diastrophic Dysplasia, Gene	23250
41.01.091	Dilated, Gene DES, Mut.)	( ). DES, . (Cardiomyopathy	24730
41.01.092	Dilated, Gene EYA4, Mut.)	( ). EYA4, . (Cardiomyopathy	41630
41.01.093	Dilated, Gene FKTN, Mut.)	( ). FKTN, . (Cardiomyopathy	41630
41.01.094	Dilated, Gene LMNA, Mut.)	( ). LMNA, . (Cardiomyopathy	34800
41.01.095	Dilated, Gene SCN5A, Mut.)	( ). SCN5A, . (Cardiomyopathy	50660
41.01.096	Dilated, Gene SGCD, Mut.)	( ). SGCD, . (Cardiomyopathy	34350
41.01.097	Dilated, Gene TAZ, Mut.)	( ). TAZ, . (Cardiomyopathy	21370



41.01.098	( ). Dilated, Gene TNNT2, Mut.)	TNNT2, . (Cardiomyopathy	41630
41.01.099	( ). , . (Cardiomyopathy Dilated, X-Linked Gene Emerine, Mut.)	-	10700
41.01.100	.	Col2A1, . (Kniest Dysplasia, Gene Col2A1, Mut.)	119350
41.01.101	V. Motor Neuropathy, DHMN, Gene GARS, Mut.)	GARS, . (Distal Hereditary	61900
41.01.102	V. Motor Neuropathy, DHMN, Gene BSCL2, Mut.)	BSCL2, . (Distal Hereditary	6670
41.01.103	.	IGHMBP2, . (Distal Spinal Muscular Atrophy 1, DSMA1, Gene IGHMBP2, Mut.)	56550
41.01.105	.	COL8A2, . (Macular Corneal Dystrophy, Gene COL8A2, Mut.)	18550
41.01.106	.	SLC4A11, . (Macular Corneal Dystrophy, Gene SLC4A11, Mut.)	44450
41.01.107	.	CHST6, . (Macular Corneal Dystrophy, Gene CHST6, Mut.)	12340
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 congenital Ichthyosis, ARCI 1, All Known Mutations, Gene TGM1, Mut.)	34800
41.01.114	.	CRYAB, . (Cataract, Gene CRYAB, Mut.)	10700
41.01.115	.	CRYBA1, . (Cataract, Gene CRYBA1, Mut.)	23250
41.01.116	.	CRYBB1, . (Cataract, Gene CRYBB1, Mut.)	23250
41.01.117	.	CRYGC, . (Cataract, Gene CRYGC, Mut.)	9650
41.01.118	.	CRYGD, . (Cataract, Gene CRYGD, Mut.)	9650
41.01.119	.	GJA3, . (Cataract, Gene GJA3, Mut.)	12340
41.01.120	.	GJA8, . (Cataract, Gene GJA8, Mut.)	6670
41.01.121	.	MIP, . (Cataract, Gene MIP, Mut.)	15850
41.01.122	.	GNAS, . (Progressive Osseous Heteroplasia, POH, Gene GNAS, Mut.)	34350
41.01.123	.	« » ANKH, « .» . (Cranio metaphyseal Dysplasia, Gene ANKH, Hot-Point Mut.)	9650
41.01.124	.	ANKH, . (Cranio metaphyseal Dysplasia, Gene ANKH, Mut.)	45450
41.01.125	.	TWIST1, . (Craniosynostosis Type 2, Gene TWIST1, Mut.)	13900
41.01.126	.	MSX2, . (Craniosynostosis Type 2, Gene MSX2, Mut.)	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
41.01.134	.	PRPH2, . (Vitelliform Macular Dystrophy, Gene PRPH2, Mut.)	12380

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	CACNAIA, (Familial Hemiplegic Migraine, FHM, Gene CACNAIA, 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
41.01.163	« » CAPN3, « » (Muscular Dystrophy Limb-Girdle Type 2A, Gene CAPN3, Hot-Point Mut.)	17900
41.01.164	CAPN3, (Muscular Dystrophy Limb-Girdle Type 2A, Gene CAPN3, Freq. Mut.)	5400
41.01.165	CAPN3, « » (Muscular Dystrophy Limb-Girdle Type 2A, Gene CAPN3, except Hot-Point Mut.)	50700
41.01.166	CAV3, (Muscular Dystrophy Limb-Girdle Type 2A, Gene CAV3, Mut.)	8320

41.01.167	Limb-Girdle Type 2A, Gene FKRP, Mut.)	FKRP, . (Muscular Dystrophy)	10700
41.01.168	Dystrophy Limb-Girdle Type 2A, Gene FKRP, Freq. Mut.)	FKRP, . . (Muscular)	6000
41.01.169	Dystrophy Limb-Girdle Type 2A, Gene FKTN, Mut.)	FKTN, . (Muscular)	41630
41.01.170	Dystrophy Limb-Girdle Type 2A, Gene LMNA, Mut.)	LMNA, . (Muscular)	38050
41.01.171	Dystrophy Limb-Girdle Type 2A, Gene POMT1, Mut.)	POMT1, . (Muscular)	61900
41.01.172	Dystrophy Limb-Girdle Type 2A, Gene SGCA, Mut.)	SGCA, . (Muscular)	23250
41.01.173	Dystrophy Limb-Girdle Type 2A, Gene SGCA, Freq. Mut.)	SGCA, . . (Muscular)	9900
41.01.174	Dystrophy Limb-Girdle Type 2A, Gene SGCB, Mut.)	SGCB, . (Muscular)	23250
41.01.175	Dystrophy Limb-Girdle Type 2A, Gene SGCD, Mut.)	SGCD, . (Muscular)	34350
41.01.176	Dystrophy Limb-Girdle Type 2A, Gene SGCG, Mut.)	SGCG, . (Muscular)	30650
41.01.177	Limb-Girdle Type 2A, Gene TTID, Mut.)	TTID, . (Muscular Dystrophy)	33350
41.01.178	Fukuyama-Type, Gene FKTN, Mut.)	FKTN, . (Muscular Dystrophy)	41630
41.01.179	Muscular Dystrophy, Gene FHL1, Mut.)	FHL1, . (Emery-Dreifuss)	30650
41.01.180	Muscular Dystrophy, Gene LMNA, Mut.)	LMNA, . (Emery-Dreifuss)	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, Mut.)	SRY, . (Disorders Sex Determination, Gene)	6200
41.01.184	Analysis Gene SRY, Mut.)	SRY, . (Disorders Sex Determination,	4400
41.01.185	EGR2, . (Charcot-Marie-Tooth Disease Type 1B, Gene EGR2, Mut.)	( - - ) I.	14450
41.01.186	GJB1, . (Charcot-Marie-Tooth Disease Type 1B, Gene GJB1, Mut.)	( - - ) I.	9000
41.01.187	LITAF, . (Charcot-Marie-Tooth Disease Type 1B, Gene LITAF, Mut.)	( - - ) I.	14450
41.01.188	PRPS1, . (Charcot-Marie-Tooth Disease Type 1B, Gene PRPS1, Mut.)	( - - ) I.	24730
41.01.189	SH3TC2, FIG4, FGD4 GDAP1, . . (Charcot-Marie-Tooth Disease Type 1B, Gene GDAP1, Freq. Mut.)	( - - ) I.	8900
41.01.190	YARS, . (Charcot-Marie-Tooth Disease Type 1B, Gene YARS, Mut.)	( - - ) I.	41630
41.01.191	17 Duplication on Chromosome 17 Gene 22, Mut.)	( - - ) I.	4400
41.01.192	0, . (Charcot-Marie-Tooth Disease Type 1B, Gene 0, Mut.)	( - - ) I.	14550
41.01.193	22, . (Charcot-Marie-Tooth Disease Type 1B, Gene 22, Mut.)	( - - ) I.	12380
41.01.194	Disease Type 1B, Genes NDRG1, SH3TC2, Mut.)	( - - ) I.	4400
41.01.195	DNM2, . (Charcot-Marie-Tooth Disease Type 2A1, Gene DNM2, Mut.)	( - - ) II.	75450
41.01.196	FIG4, . (Charcot-Marie-Tooth Disease Type 2A1, Gene FIG4, Mut.)	( - - ) II.	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
41.01.211	GJB2, . (Deafness Autosomal Recessive 1, DFNB1A, Gene GJB2, Mut.)	6200
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
41.01.223	LEP, . (Monogenic Obesity, Gene LEP, Mut.)	8650
41.01.224	POMC, . (Monogenic Obesity, Proopiomelanocortin, Gene POMC, Mut.)	12340
41.01.225	POMC (Monogenic Obesity, Proopiomelanocortin. Polymorphism of Gene POMC)	1600
41.01.226	MC4R, . (Monogenic Obesity, Melanocortin-4 Receptor, Gene MC4R, Mut.)	8650
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	- . CRX, . (CORD, Gene CRX, Mut.)	10700
41.01.232	- . ADAM9, . (CORD, Gene ADAM9, Mut.)	81450
41.01.233	- . RPGR, . (CORD, 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
41.01.248	. RPE65, . (Retinitis Pigmentosa, Gene RPE65, Mut.)	38050
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	- ( ). IKBKG, . . (Bloch-Sulzberger Syndrome, Familial Incontinentia Pigmenti, Gene IKBKG, 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	Gene EDNRB, Mut.)	EDNRB, . (Waardenburg-Shah Syndrome,	25950
41.01.297	Mut.)	IRF6, . (Van der Woude Syndrome, Gene IRF6,	34350
41.01.298	Gene RECQL2, Mut.)	RECQL2, . (Werner Syndrome,	128000
41.01.299	WAS, Gene WAS, Mut.)	WAS, . (Wiskott-Aldrich Syndrome,	25950
41.01.300	PHOX2B, . . (Congenital Central Hypoventilation Syndrome, CCHS, Gene PHOX2B, Freq. Mut.)		3500
41.01.301	(Albinism oculocutaneous, Hermansky-Pudlak type, Gene HPS1, Freq. Mut.)	HPS1, . .	9650
41.01.302	Gene GLI3, Mut.)	GLI3, . (Greig Syndrome,	61900
41.01.303	Mut.)	RAB27A, . (Griscelli Syndrome, Gene RAB27A,	17900
41.01.304	(Jackson-Weiss Syndrome, JWS, Exon 9 Gene FGFR2, Exon 7A Gene FGFR1, Mut.)	9 FGFR2 7A FGFR1, .	9000
41.01.305	Mut.)	TBX1, . (Di George Syndrome, DGS, Gene TBX1,	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	( Klippel-Feil Syndrome, Gene GDF6, Mut.)	GDF6, .	10690
41.01.313	ERCC6, . (Cockayne Syndrome, Gene ERCC6, Mut.)		75600
41.01.314	QT. KCNH2 KCNE2, . (Short QT Syndrome, Short QTS, Genes KCNH2, KCNE2, Mut.)		25880
41.01.315	QT. KCNJ2, . (Short QT Syndrome, Short QTS, Gene KCNJ2, Mut.)		12380
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	Gene RPS6KA3, Mut.)	RPS6KA3, . (Coffin-Lowry Syndrome,	82450
41.01.319	PAX3, . (Craniofacial-Deafness-Hand Syndrome, CDHS, Gene PAX3, Mut.)		28090
41.01.320	( Crigler-Najjar Syndrome, Gene UGT1, Mut.)	UGT1, .	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	LFS, Gene MED12, Freq. Mut.)	MED12, . . (Lujan-Fryns Syndrome,	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	(Marfan Syndrome, Gene FBN1, without Hot-Point Mut.)	190000
41.01.329	FBN1, . (Marfan Syndrome, Gene FBN1, Mut.)	216000
41.01.330	( 2). 15 RET, . (Multiple Endocrine Neoplasia Type 2B, Exon 15 Gene RET, Mut.)	7200
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
41.01.347	TWIST1, . (Saethre-Chotzen Syndrome, Gene TWIST1, Mut.)	10690
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	- ( ). WWS, Gene FKR, Mut.)	FKRP, . (Walker-Warburg Syndrome,	10960
41.01.362	- ( ). WWS, Gene POMT1, Mut.)	POMT1, . (Walker-Warburg Syndrome,	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	- ( - ). Syndrome, Gene TBX5, Mut.)	TBX5, . (Holt-Oram	34800
41.01.366	- . Syndrome, Gene SBDS, Mut.)	SBDS, . (Shwachman-Diamond	19850
41.01.367	- . (Shwachman-Diamond Syndrome, Gene SBDS1, Freq. Mut.)	SBDS1, . .	7200
41.01.368	- . Syndrome, LVA, Gene SLC26A4, Mut.)	SLC26A4 . (Large Vestibular Aqueduct	46820
41.01.369	- , VI. Syndrome, Type VI, Gene PLOD, Freq. Mut.)	PLOD, . . (Ehlers-Danlos	7830
41.01.370	- . CHRNG, . (Escobar Syndrome, Gene CHRNG, Mut.)		28100
41.01.371	- . HOXD13, Mut.)	HOXD13, . (Synpolydactyly, Type 1, SPD1, Gene	14650
41.01.372	- . SPM, Gene FHL1, Mut.)	FHL1, . (Scapuloperoneal Myopathy,	30650
41.01.373	- . Progressive Scoliosis, HGPPS, Gene ROBO3, Mut.)	ROBO3, . (Horizontal Gaze Palsy with	62000
41.01.374	- (c ). GJC2, . (Westphal-Strumpell Disease, Gene GJC2, Mut.)		20470
41.01.375	- , Mut.)	ATL1 . (Westphal-Strumpell Disease, Gene ATL1,	51700
41.01.376	- , Mut.)	SPAST . (Westphal-Strumpell Disease, Gene SPAST,	51700
41.01.377	- (c ). BSCL2, . (Westphal-Strumpell Disease, Gene BSCL2, Mut.)		27000
41.01.378	- . Paralysis, IAHS, Gene ALSIN, Mut.)	ALSIN, . (Infantile Ascending Hereditary Spastic	119500
41.01.379	- . Muscular Atrophy (SMA) with Diaphragmatic Paralysis, Gene IGHMBP2, Mut.)	IGHMBP2, . (Spinal	51700
41.01.380	- I, II, III, IV. Atrophy, SMA, Type I, II, III, IV, Gene SMN1)	SMN1, . (Spinal Muscular	7830
41.01.381	- I, II, III, IV. Atrophy, SMA, Type I, II, III, IV, Copy Number Variation)	(Spinal Muscular	13500
41.01.382	- I, II, III, IV. (Spinal Muscular Atrophy, SMA, Type I, II, III, IV, Gene SMN1, Mut. (Only Presence One Gene Copy))	SMN1, . (	30650
41.01.383	- . SMA, Late-Onset, Finkel Type, Gene VAPB, Mut.)	VAPB, (Spinal Muscular Atrophy,	23000
41.01.384	- . Atrophy, SMA, Late-Onset, Finkel Type, Gene VAPB, Freq. Mut.)	VAPB, . . (Spinal Muscular	6000
41.01.385	- , X- « . » . (Spinal Muscular Atrophy, SMA, X-Linked, Gene UBA1, Hot-Point Mut.)	UBA1,	7200
41.01.386	- . Spinal and Bulbar Muscular Atrophy, Gene AR, Freq. Mut.)	AR, . . (Kennedy	4400
41.01.387	- . (Spinocerebellar Ataxia, Genes ATXN1, ATXN2, ATXN3, Freq. Mut.)	ATXN1, ATXN2, ATXN3, . .	7830
41.01.388	- . Ataxia, Gene ATXN7 Freq. Mut.)	ATXN7, . . (Spinocerebellar	5000
41.01.389	- . Ataxia, Gene ATXN8, Freq. Mut.)	ATXN8, . . (Spinocerebellar	5000
41.01.390	- . PRNP, . (Spongiform Encephalopathy with Neuropsychiatric Features, Gene PRNP, Mut.)		10700
41.01.391	- . DLL3, Mut.)	DLL3, . (Spondylocostal Dysostosis, Gene	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
41.01.408	TSC1, . (Focal Cortical Dysplasia of Taylor, FCDT, Gene TSC1, Mut.)	82450
41.01.409	TSC1, . . (Focal Cortical Dysplasia of Taylor, FCDT, Gene TSC1, Freq. Mut.)	7200
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	CACNAIA, . (Episodic Ataxia, EA, Gene CACNAIA, 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	VHL, . (Autosomal Recessive Erythrocytosis, Gene VHL, Mut.)	10690
41.01.442	VHL, . . (Autosomal Recessive Erythrocytosis, Gene VHL, Freq. Mut.)	4500
<b>42. : (Definition of Biological Relationship: Paternity and Maternity) / 42.01. (Definition of Biological Relationship: Paternity and Maternity) :</b>		
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
<b>43. / « » (Hereditary Metabolic Disorders in Newborns. Newborn Screening "HEEL") / 43.01. « » (Hereditary Metabolic Disorders in Newborns. Newborn Screening "HEEL")</b>		
43.01.001	e « » (Newborn Screening "HEEL")	5200
43.01.002	( ) (Gas Chromatography of Urine (Organic Aciduria))	8100
43.01.003	( ) ( ) (Hereditary Diseases of Metabolism, Screening (Qualitative Urine Tests))	1800
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-Acetylaspartate)	(N- ) (HPLC-MS/MS Organic Acids	3730
43.01.008	(Acid))	( ) (HPLC-MS/MS Organic Acids (Orotic	3730
43.01.009	BD ( Freq. Mut.)	( ) (BD (Biotinidase Deficiency) Gene,	5520
43.01.010	GCDH ( Gene, Freq. Mut. (Glutaric Aciduria, Type 1))	1) (GCDH (Glutaryl-CoA Dehydrogenase)	4300
43.01.011	GCDH ( Gene (Glutaric Aciduria, Type 1))	1) (GCDH (Glutaryl-CoA Dehydrogenase)	38470
43.01.012	CBS ( Mut. (Homocystinuria))	( ) (CBS (Cystathionine-Beta-Synthase) Gene, Freq.	5600
43.01.013	( , « », BCKDHA, BCKDHB) (BCKDHA and BCKDHB Genes, Freq. Mut. (Maple Syrup Urine Disease, MSUD))		16000
43.01.014	MUT ( Aciduria, MMA))	( ) (MUT Gene, Freq. Mut. (Methylmalonic	16000
43.01.015	HADHA ( 3- - - ) (HADHA Gene, Freq. Mut. (Long-Chain 3-Hydroxyacyl-Coa Dehydrogenase (LCHAD) Deficiency))		4890
43.01.016	ACADM ( MCAD) (ACADM Gene, Freq. Mut. (Medium-Chain Acyl-CoA Dehydrogenase (MCAD) Deficiency))		4890
43.01.017	( Transcarbamylase (OTC) Deficiency))	( TC Gene (Ornithine	31200
43.01.018	FAH ( I) (FAH Gene, Freq. Mut. (Tyrosinemia, Type 1))		6880
43.01.019	FAH ( I) (FAH Gene (Tyrosinemia, Type 1))		47520
43.01.020	ASS ( ) (ASS Gene, Freq. Mut. (Citrullinemia))		6100
<b>44. (Investigation of Water and Soil Quality) / 44.01. (High Tech Investigations of Water)</b>			
44.01.001	: Assessment)	(Water: Complete and Comprehensive Quality	16500
44.01.002	:	(Water: Abridged Quality Assessment )	8700
44.01.003	: Contamination)	(Water: Suspicion Household Waste	7860
44.01.004	: Contamination )	(Water: Suspicion Industrial Waste	12300
44.01.005	: Suspicion Products of Combustion and Emissions from Motorways Contamination )	(Water:	9020
44.01.006	: Excessive Use of Chemicals for Water Treatment)	(Water: Suspicion	5150
<b>44. (Investigation of Water and Soil Quality) / 44.02. (Screening Investigations of Water)</b>			
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
<b>44. (Investigation of Water and Soil Quality) / 44.03. (High Tech Investigations of Soil)</b>			
44.03.001	:	(Soil: Comprehensive Toxicological Evaluation)	18200
44.03.002	:	(Soil: Agrochemical Evaluation)	9050
<b>45. ( ) (Profiles) / 45.01. (For children)</b>			
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
<b>45. ( ) (Profiles) / 45.02. (For women)</b>			
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
45.02.012	:	(Mastopathy: Extended Profile)	9500
45.02.013	:	(Mastopathy: Screening)	5300
<b>45. ( ) (Profiles) / 45.03. (For men)</b>			
45.03.001	VIP-	(VIP-Survey for Men)	6500
45.03.002	:	(Male oncologic risk: prostate)	800
<b>45. ( ) (Profiles) / 45.04. ( Annual preventive examinations)</b>			
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
<b>45. ( ) (Profiles) / 45.05. (Checkup of household staff)</b>			
45.05.001	«	» (My Healthy Nurse)	4740
<b>45. ( ) (Profiles) / 45.06. (Cardiovascular disease risk assessment)</b>			
45.06.001		(Preventing Heart and Blood Vessel Diseases )	1400
45.06.002	:	(Lipid Profile: Extended )	3100
45.06.003	:	(Lipid Profile: Screening)	350
<b>45. ( ) (Profiles) / 45.07. (Diagnosis of antiphospholipid syndrome (APS))</b>			
45.07.001		( ), (Antiphospholipid Syndrome, APS)	3500
<b>45. ( ) (Profiles) / 45.08. (Diagnosis of gastrointestinal tract (GI Tract))</b>			
45.08.001	:	( ) (Coeliac Disease: Gluten Intolerance )	4000
45.08.002		(GastroPanel)	3100
<b>45. ( ) (Profiles) / 45.09. (Assessment of liver function)</b>			
45.09.001	:	(Survey of Liver: Extended )	1550
45.09.002	:	(Survey of Liver: Screening)	350
45.09.003		(Autoimmune Liver Disease: Screening)	4700
<b>45. ( ) (Profiles) / 45.10. (Non-invasive Assessment of Liver Fibrosis)</b>			
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
<b>45. ( ) (Profiles) / 45.11. (Assessment of renal function)</b>			
45.11.001	:	(Survey of Kidneys: Extended )	1400

45.11.002	:	(Survey of Kidneys: Screening)	550
<b>45. ( ) (Profiles) / 45.12. (Hospital studies)</b>			
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
<b>45. ( ) (Profiles) / 45.13. (Diagnosis of connective tissue diseases (disorders of connective tissue))</b>			
45.13.001	:	(Joint Pain: Extended Survey)	3800
45.13.002	:	(Joint Pain: Screening)	950
<b>45. ( ) (Profiles) / 45.14. (Diagnosis of diabetes)</b>			
45.14.001	:	(Diabetes Control: Extended)	2000
45.14.002	:	(Diabetes Control: Screening)	550
45.14.003	:	(Diabetes: Autoimmune Markers)	3200
<b>45. ( ) (Profiles) / 45.15. (Diagnosis and monitoring therapy for osteoporosis)</b>			
45.15.001		(Diagnosis of Osteoporosis)	1800
<b>45. ( ) (Profiles) / 45.16. (Assessment of thyroid function)</b>			
45.16.001	:	(Thyroid Gland: Extended Survey)	1000
45.16.002	:	(Thyroid Gland: Screening)	700
<b>45. ( ) (Profiles) / 45.17. (Sexually transmitted infections (STI))</b>			
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
<b>45. ( ) (Profiles) / 45.18. (Weight Problems)</b>			
45.18.001	( )	(Weight Problems: Primary Survey)	2100
45.18.002	:	(Survey Before Diet: Minimum )	2000
45.18.003	:	(Survey Before Diet: Additional )	4700
<b>45. ( ) (Profiles) / 45.19. (Upper respiratory tract infection (URI))</b>			
45.19.001	,	( Acute Respiratory Infections, ARI: Runny Nose, Cough, Sore Throat)	3700
<b>45. ( ) (Profiles) / 45.20. (Allergy)</b>			
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
<b>45. ( ) (Profiles) / 45.21. (Hematological studies)</b>			
45.21.001		(Diagnosis of Anemia)	2100
<b>50. / 50.01.002.006.000</b>			
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
<b>50.</b>	<b>/ 50.02.002.024.000</b>	
50.02.018	- MTHFR: 677, MTR, MTRR	MTHFR: 1298, 1001
50.02.019	- (F2, F5, F7, F13, FGB, ITGA2, ITGB3, PAI-1)	1760