

CDLT FACA Panel meeting results; July 23,
2025

1. 0533U: Drug metabolism (adverse drug reactions and drug response), genotyping of 16 genes, reported as metabolizer status and transporter function

Activity type	Multiple choice	
Total responses		10
Unique participants		10
Response options	Count	
Abstain		0
Gapfill		0
Crosswalk to: 0516U		8
Crosswalk to: 0434U		0
Crosswalk to: 0476U		2

2. 0423U: Psychiatry (e.g., depression, anxiety), genomic analysis panel, including variant analysis of 26 genes, buccal swab, report including metabolizer status and risk of drug toxicity by condition

Activity type	Multiple choice	
Total responses		10
Unique participants		10
Response options	Count	
Abstain		0
Gapfill		3
Crosswalk to: 0476U		6
Crosswalk to: 0516U		1
Crosswalk to: 0345U		0
Crosswalk to: 0411U		0

3. 0552U: Testing for genetic disorders from outer layer of embryo cells, reported as low-risk or high-risk for familial genetic disorder in preimplantation genetic assessment

Activity type	Multiple choice	
Total responses		10
Unique participants		10
Response options	Count	
Abstain		0
Gapfill		4
Crosswalk to: 81425		0
Crosswalk to: 0254U		6

4. 0553U: Analysis of 24 chromosomes using DNA genomic sequence analysis of outer layer of embryo cells, results reported as normal/balanced or balanced per embryo tested in preimplantation genetic assessment

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

0

Gapfill

0

Crosswalk to: 0254U

10

5. 0554U: Analysis of 24 chromosomes using DNA genomic sequence analysis of outer layer of embryo cells, results reported as normal, monosomy, trisomy, segmental aneuploidy, triploid, haploid, or mosaic, per embryo testing in preimplantation assessment

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

0

Gapfill

0

Crosswalk to: 0254U

10

6. 0555U: Analysis of 24 chromosomes using DNA genomic sequence analysis of outer layer of embryo cells, results reported as reported as normal/balanced (euploidy/balanced), unbalanced structural rearrangement...

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

0

Gapfill

0

Crosswalk to: 0254U

10

7. 81195: Cytogenomic genome-wide analysis, hematologic malignancy, structural variations and copy number variations, optical genome mapping (OGM)	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 0331U		10

8. 8XXX4: Cytogenomic (genome-wide) analysis for constitutional chromosomal abnormalities; interrogation of structural and copy number variants, optical genome mapping (OGM)	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 81229 + 88262 + 81404 + 81243 + 88275		0
	Crosswalk to: 0260U		10

9. X159U: Transplantation medicine (liver allograft rejection), miRNA gene expression profiling by RT-PCR of 4 genes (miR-122, miR-885, miR-23a housekeeping, spike-in control), serum, algorithm reported as risk of liver allograft rejection	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		1
	Crosswalk to: 0018U x 0.85		3
	Crosswalk to: 0069U		6

10. 0534U: Oncology (prostate), microRNA, single-nucleotide polymorphisms (SNPs) analysis by RT-PCR of 32 variants, using buccal swab, algorithm reported as a risk score	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		1
	Crosswalk to: 0018U		9
11.0531U: Infectious disease (acid-fast bacteria and invasive fungi), DNA (673 organisms), next-generation sequencing, plasma	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		10
	Crosswalk to: 0326U		0
	Crosswalk to: 81520		0
	Crosswalk to: 0480U		0
12. X177U: Infectious disease (bacterial and fungal), DNA of 44 organisms (34 bacteria, 10 fungi), urine, next-generation sequencing, reported as positive or negative for each organism	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		10
	Crosswalk to: 0504U		0
13. 871XX: Susceptibility studies, antimicrobial agent; carbapenemase enzyme detection (eg, Klebsiella pneumoniae carbapenemase [KPC], New Delhi metallo-beta-lactamase [NDM], Verona integron-encoded...	Activity type	Multiple choice	

	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	0
	Gapfill	1
	Crosswalk to: 87185 x 3	9
14. 8XXXX: Infectious agent antigen detection by immunoassay with direct optical (ie, visual) observation; severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) (coronavirus disease [COVID-19]) and influenza virus types A and B	Activity type	Multiple choice
	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	0
	Gapfill	1
	Crosswalk to: 87811 + 87804	1
	Crosswalk to: 87811 + 87804 x 2	8
15. X166U: Borrelia burgdorferi, antibody detection of 24 recombinant protein groups, by immunoassay, IgG	Activity type	Multiple choice
	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	0
	Gapfill	10
16. X182U: Infectious disease (sepsis), semiquantitative measurement of pancreatic stone protein concentration, whole blood, reported as risk of sepsis	Activity type	Multiple choice
	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	0
	Gapfill	10
17. 0574U: Culture filtrate protein-10-kDa testing in serum or plasma for tuberculosis	Activity type	Multiple choice
	Total responses	10

Unique participants	10
Response options	Count
Abstain	0
Gapfill	10

18. 0556U: Testing of pathogen-specific DNA and RNA by real-time PCR for 12 targets from nasal or throat swab, reported as detected or not detected in bacterial or viral respiratory tract infection

Activity type	Multiple choice
Total responses	10
Unique participants	10
Response options	Count
Abstain	0
Gapfill	0
Crosswalk to: 87633	10

19. 0557U: Real-time amplification testing for bacterial DNA markers in vaginal fluid reported as detected or not detected for each organism in bacterial vaginosis and vaginitis

Activity type	Multiple choice
Total responses	10
Unique participants	10
Response options	Count
Abstain	0
Gapfill	0
Crosswalk to: 81514	10

20. 0563U: Pathogen-specific nucleic acid DNA or RNA testing for 11 viral targets and 4 bacterial targets in upper respiratory specimen for each pathogen, reported as positive or negative...

Activity type	Multiple choice
Total responses	10
Unique participants	10
Response options	Count
Abstain	0
Gapfill	0
Crosswalk to: 87633	10
Crosswalk to: 0115U	0

21. 0564U: Pathogen-specific nucleic acid DNA or RNA testing for 10 viral targets and 4 bacterial targets in upper respiratory specimen, each pathogen...

Activity type	Multiple choice	
Total responses		10
Unique participants		10
Response options	Count	
Abstain		0
Gapfill		0
Crosswalk to: 87633		10
Crosswalk to: 0115U		0

23. X171U: Infectious disease (bacterial or viral), 32 genes (29 informative and 3 housekeeping), immune response mRNA, gene expression profiling by split-well multiplex reverse transcription loop-mediated isothermal amplification (RT-LAMP)...

Activity type	Multiple choice	
Total responses		10
Unique participants		10
Response options	Count	
Abstain		1
Gapfill		0
Crosswalk to: 0203U		9

24. X183U: Infectious disease (tropical fever pathogens), vector-borne and zoonotic pathogens, including 2 viruses (Chikungunya virus and Dengue virus serotypes 1, 2, 3, and 4), 1 bacterium...

Activity type	Multiple choice	
Total responses		10
Unique participants		10
Response options	Count	
Abstain		0
Gapfill		0
Crosswalk to: 87506		10

25. 0536U: Red blood cell antigen (fetal RhD), PCR analysis of exon 4 of RHD gene and housekeeping control gene GAPDH from whole blood in pregnant individuals at 10+ weeks gestation known to be RhD negative, reported as fetal RhD status	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 0453U		10
	Crosswalk to: 0488U		0
	Crosswalk to: 0494U		0

26. 0523U: Oncology (solid tumor), DNA, qualitative, next-generation sequencing (NGS) of single- nucleotide variants (SNV) and insertion/deletions in 22 genes utilizing formalin-fixed paraffin- embedded tissue...	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 81445		10
	Crosswalk to: 0022U MINUS 81449		0

27. 0538U: Oncology (solid tumor), next-generation targeted sequencing analysis, formalin-fixed paraffin- embedded (FFPE) tumor tissue, DNA analysis of 600 genes, interrogation for single-nucleotide variants...	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 0334U		1
	Crosswalk to: 81459		9

28. 0543U: Oncology (solid tumor), next-generation sequencing of DNA from formalin-fixed paraffin-embedded (FFPE) tissue of 517 genes, interrogation for single-nucleotide variants, multi- nucleotide variants...	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 81459		9
	Crosswalk to: 0379U		1
29. X164U: Oncology (cutaneous melanoma), RNA, gene expression profiling by real-time qPCR of 10 genes (8 content and 2 housekeeping), utilizing formalin-fixed paraffin-embedded (FFPE) tissue...	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 81525		10
	Crosswalk to: 81529		0
30. X172U: Oncology, mRNA, gene expression profiling of 216 genes (204 targeted and 12 housekeeping genes), RNA expression analysis, formalin-fixed paraffin-embedded (FFPE) tissue, quantitative, reported as log2 ratio per gene	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 81456		10

31. X185U: Oncology (breast), RNA expression profiling of 329 genes by targeted next-generation sequencing and 20 proteins by multiplex immunofluorescence, formalin-fixed paraffin-embedded (FFPE)...	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		7
	Crosswalk to: 81521		3
32. 0527U: Herpes simplex virus (HSV) types 1 and 2 and Varicella zoster virus (VZV), amplified probe technique, each pathogen reported as detected or not detected	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 87631		10
	Crosswalk to: 0455U		0
33. 0528U: Lower respiratory tract infectious agent detection, 18 bacteria, 8 viruses, and 7 antimicrobial- resistance genes, amplified probe technique, including reverse transcription for RNA targets...	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 0321U		10
	Crosswalk to: 87633 + 87632		0
34. 871X1: Susceptibility studies, antimicrobial agent; carbapenem resistance genes, amplified probe technique, per isolate	Activity type	Multiple choice	
	Total responses		10

	Unique participants	Count	10
	Response options	Count	
	Abstain		0
	Gapfill		1
	Crosswalk to: 87150 x 5		8
	Crosswalk to: 87505		1
35. 87513: Infectious agent detection by nucleic acid (DNA or RNA); Helicobacter pylori (H. pylori), clarithromycin resistance, amplified probe technique	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		1
	Gapfill		0
	Crosswalk to: 87564		9
36. 8XXX9: Infectious agent detection by nucleic acid (DNA or RNA); joint space pathogens and drug resistance genes, multiplex amplified probe technique, 26 or more targets	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 0505U		9
	Crosswalk to: 87633		1
37. 8XXX2: Infectious agent detection by nucleic acid (DNA or RNA); Chlamydia trachomatis and Neisseria gonorrhoeae, multiplex amplified probe technique	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 87491 + 87591		9
	Crosswalk to: 0402U x 0.5		0
	Crosswalk to: 87801		1

38. X180U: Infectious disease (genitourinary pathogens), DNA, 46 targets (28 pathogens, 18 resistance genes), RT-PCR amplified probe technique, urine, each analyte reported as detected or not detected

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

0

Gapfill

0

Crosswalk to: 0321U

10

**Crosswalk to: 0374U-deleted code

0

39. G0567: Infectious agent detection by nucleic acid (DNA or RNA); hepatitis C, screening, amplified probe technique

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

0

Gapfill

0

Crosswalk to: 87521

10

Crosswalk to: 87521 x 1.5

0

40. 0540U: Transplantation medicine, quantification of donor- derived cell-free DNA using next-generation sequencing analysis of plasma, reported as percentage of donor- derived cell free DNA to determine probability of rejection

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

0

Gapfill

0

Crosswalk to: 0493U

10

41. 0530U: Oncology (pan-solid tumor), ctDNA, utilizing plasma, next- generation sequencing (NGS) of 77 genes, 8 fusions, microsatellite instability, and tumor mutation burden...

Activity type

Multiple choice

Total responses	10
Unique participants	10
Response options	Count
Abstain	1
Gapfill	0
Crosswalk to: 81464	9
Crosswalk to: 81455	0

42. 0539U: Oncology (solid tumor), cell- free circulating tumor DNA (ctDNA), 152 genes, next- generation sequencing, interrogation for single- nucleotide variants, insertions/deletions, gene rearrangements...

Activity type	Multiple choice
Total responses	10
Unique participants	10
Response options	Count
Abstain	0
Gapfill	0
Crosswalk to: 81464	10
Crosswalk to: 0326U	0

43. 0560U: Genomic sequence analysis in cell-free DNA in whole blood and tumor tissue, baseline assessment for design and construction of a personalized variant panel to evaluate current minimal...

Activity type	Multiple choice
Total responses	10
Unique participants	10
Response options	Count
Abstain	0
Gapfill	0
Crosswalk to: 0306U	10

44. 0561U: Genomic sequence analysis in cell-free DNA in whole blood, subsequent assessment with comparison to initial assessment to evaluate for minimal residual disease in cancer

Activity type	Multiple choice
Total responses	10
Unique participants	10
Response options	Count
Abstain	0

	Gapfill	0
	Crosswalk to: 0307U	10
45. 0562U: Targeted genomic sequence analysis of 33 genes, detection of single-nucleotide variants (SNVs) in plasma, reported as presence of actionable variants in solid tumor cancer	Activity type	Multiple choice
	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	1
	Gapfill	0
	Crosswalk to: 0388U	9
46. 0571U: Next-generation sequencing of DNA in 80 genes and RNA in 10 genes in plasma, reported as clinically actionable variants in solid tumor cancer	Activity type	Multiple choice
	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	1
	Gapfill	0
	Crosswalk to: 81455	9
47. X169U: Targeted genomic sequence analysis panel, solid organ neoplasm, circulating cell-free DNA (cfDNA) analysis from plasma of 521 genes, interrogation for sequence variants, gene copy number...	Activity type	Multiple choice
	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	1
	Gapfill	0
	Crosswalk to: 0239U	9
48. 0524U: Obstetrics (preeclampsia), sFlt-1/PlGF ratio, immunoassay, utilizing serum or plasma, reported as a value	Activity type	Multiple choice
	Total responses	10

	Unique participants	Count	10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 0482U		10
49. 0550U: Oncology (prostate), enzyme-linked immunosorbent assays (ELISA) for total prostate- specific antigen (PSA) and free PSA, serum, combined with age, previous negative prostate biopsy status...	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 81539 x 0.5		10
50. 0558U: Quantitative enzyme-linked immunosorbent assay (ELISA) for secreted colorectal cancer protein marker in serum, reported as response/no response to therapy or disease progression/regression	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		1
	Crosswalk to: 86316		9
51. 0559U: Quantitative enzyme-linked immunosorbent assay (ELISA) for secreted breast cancer protein marker in serum, result reported as indicative of response/no response to therapy or disease progression/regression	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		1
	Crosswalk to: 86316		9

52. 0573U: Testing for 3 biomarkers in pancreatic cyst lesion fluid, algorithm reported as categorical mucinous or non-mucinous in pancreatic cancer	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		10
	Crosswalk to: 82948 + 82378+ 0080U x .66		0
53. X178U: Oncology (prostate cancer), biochemical analysis of 3 proteins (total PSA, free PSA, and HE4), plasma, serum, prognostic algorithm incorporating 3 proteins and digital rectal examination...	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 81539		10
54. X187U: Oncology (pancreatic cancer), multiplex immunoassay of ICAM1, TIMP1, CTSD, THBS1, and CA 19-9, serum, diagnostic algorithm reported as positive or negative	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 81539		10
	Crosswalk to: 0342U		0
55. X162U: Oncology (ovarian), serum, analysis of 39 glycoproteins by liquid chromatography with tandem mass spectrometry (LC-MS/MS) in multiple reaction monitoring mode, reported as likelihood of malignancy	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	

Abstain	0
Gapfill	1
Crosswalk to: 81503	9

56. 0549U: Oncology (urothelial), DNA, quantitative methylated real- time PCR of TRNA-Cys, SIM2, and NKX1-1, using urine, diagnostic algorithm reported as a probability index for bladder cancer and/or upper tract urothelial carcinoma (UTUC)

Activity type	Multiple choice
Total responses	10
Unique participants	10
Response options	Count
Abstain	0
Gapfill	0
Crosswalk to: 0012M	10
Crosswalk to: 0229U	0

57. 0565U: Next-generation sequencing methylation pattern assay to detect 6626 epigenetic alterations in cell-free DNA plasma, algorithm reported as cancer signal detected or not detected

Activity type	Multiple choice
Total responses	10
Unique participants	10
Response options	Count
Abstain	0
Gapfill	1
Crosswalk to: 0410U	9
Crosswalk to: 0507U	0

58. 0566U: qPCR-based analysis of 13 differentially methylated regions in pleural fluid, algorithm reported as a qualitative result in lung cancer

Activity type	Multiple choice
Total responses	10
Unique participants	10
Response options	Count
Abstain	0
Gapfill	7
Crosswalk to: 0398U	3

59. 0569U: Next-generation sequencing analysis of tumor methylation markers present in cell-free circulating tumor DNA (ctDNA) in whole blood, algorithm reported as...	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		10
	Crosswalk to: 0493U		0
	Crosswalk to: 0537U		0
60. 0572U: High-throughput telomere length quantification by FISH testing in whole blood, diagnostic algorithm reported as risk of prostate cancer	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		10
	Crosswalk to: 0537U		0
62. 0420U: Oncology (urothelial), mRNA expression profiling by real-time quantitative PCR of MDK, HOXA13, CDC2, IGFBP5, and CXCR2 in combination with droplet digital PCR ddPCR) analysis of 6 single-nucleotide polymorphisms (SNPs) genes...	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		1
	Crosswalk to: 0012M + 0356U x 0.35		9
63. 8XXX0: Oncology (central nervous system tumor), DNA methylation analysis of at least 10,000 methylation sites, utilizing DNA extracted from formalin-fixed tumor tissue, algorithm(s) reported...	Activity type	Multiple choice	
	Total responses		10

	Unique participants	10
	Response options	Count
	Abstain	0
	Gapfill	8
	Crosswalk to: 0537U	2
64. X179U: Oncology (hematolymphoid neoplasms), DNA, targeted genomic sequence of 417 genes, interrogation for gene fusions, translocations, rearrangements...	Activity type	Multiple choice
	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	0
	Gapfill	0
	Crosswalk to: 81455	10
65. 0535U: Perfluoroalkyl substances (PFAS) (eg, perfluorooctanoic acid, perfluorooctane sulfonic acid), by liquid chromatography with tandem mass spectrometry (LC-MS/MS), plasma or serum, quantitative	Activity type	Multiple choice
	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	0
	Gapfill	0
	Crosswalk to: 0457U	10
66. 0541U: Cardiovascular disease (HDL reverse cholesterol transport), cholesterol efflux capacity, LC-MS/MS, quantitative measurement of 5 distinct HDL-bound apolipoproteins...	Activity type	Multiple choice
	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	0
	Gapfill	10
	Crosswalk to: 82172 x 5	0

67. 0542U: Nephrology (renal transplant), urine, nuclear magnetic resonance (NMR) spectroscopy measurement of 84 urinary metabolites, combined with patient data, quantification of BK virus...

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

0

Gapfill

9

Crosswalk to: 81558

1

68. X161U: Perfluoroalkyl substances (PFAS) (eg, perfluorooctanoic acid, perfluorooctane sulfonic acid), 24 PFAS compounds by high-performance liquid chromatography with tandem mass spectrometry (LC-MS/MS), plasma or serum, quantitative

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

0

Gapfill

0

Crosswalk to: 0457U

10

Crosswalk to: 0457U x 1.15

0

69. X165U: Nephrology (diabetic chronic kidney disease), enzyme-linked immunosorbent assay (ELISA) of apolipoprotein A4 (APOA4), CD5 antigen-like (CD5L) combined with estimated glomerular filtration rate (GFR)...

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

0

Gapfill

0

Crosswalk to: 0385U

10

70. X170U: Therapeutic drug monitoring, 60-150 drugs and metabolites, urine, saliva, quantitative liquid chromatography with tandem mass spectrometry (LC-MS/MS), specimen validity...

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

0

Gapfill

0

Crosswalk to: 0328U

10

71. X181U: Transplantation medicine, antibody to non-human leukocyte antigens (non-HLA), blood specimen, flow cytometry, single-antigen bead technology, 39 targets, individual positive antibodies reported

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

1

Gapfill

1

Crosswalk to: 86833

8

72. X186U: Gastroenterology (irritable bowel syndrome), IgG antibodies to 18 food items by microarray-based immunoassay, whole blood or serum, report as elevated (positive) or normal (negative) antibody levels

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

0

Gapfill

2

Crosswalk to: 0360U

8

Crosswalk to: 0399U x 2

0

Crosswalk to: 0399U

0

73. 0529U: Hematology (venous thromboembolism [VTE]), genome-wide single-nucleotide polymorphism variants, including F2 and F5 gene analysis, and Leiden variant, by microarray analysis, saliva, report as risk score for VTE

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

0

Gapfill

9

Crosswalk to: 81229

1

74. 0532U: Rare diseases (constitutional disease/hereditary disorders), rapid whole genome and mitochondrial DNA sequencing for single- nucleotide variants, insertions/deletions, copy number variations...

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

0

Gapfill

1

Crosswalk to: 0094U

9

75. X168U: Rare diseases (constitutional disease/hereditary disorders), rapid whole genome comparator DNA sequencing for single-nucleotide variants, insertions/deletions, copy number variations...

Activity type

Multiple choice

Total responses

10

Unique participants

10

Response options

Count

Abstain

1

Gapfill

1

Crosswalk to: 81426

8

Crosswalk to: 0094U x 0.5

0

76. 0567U: Whole-genome sequence analysis in blood, saliva, amniocentesis, chorionic villus sample or tissue, identification and categorization of genetic variants in constitutional/heritable disorder	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		1
	Crosswalk to: 0212U		7
	Crosswalk to: 0212U + 0318U		1
	Crosswalk to: 0212U (or 81425) + 0212U + 0318U		1

77. X167U: Rare diseases (constitutional disease/hereditary disorders), rapid whole genome DNA sequencing for single-nucleotide variants, insertions/deletions, copy number variations, blood, saliva, tissue sample, variants reported	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		1
	Gapfill		0
	Crosswalk to: 0094U		9

78. 0544U: Nephrology (transplant monitoring), 48 variants by digital PCR, using cell-free DNA from plasma, donor-derived cell-free DNA, percentage reported as risk for rejection	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		1
	Crosswalk to: 0493U		9

79. X160U: Transplantation medicine (liver allograft rejection), quantitative donor-derived cell-free DNA (cfDNA) by whole genome next-generation sequencing, plasma and mRNA gene expression profiling...	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 81558		9
	Crosswalk to: 81558 + 0118U		1
80. X163U: Neurology (prion disease), cerebrospinal fluid, detection of prion protein by quaking-induced conformational conversion, qualitative	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		1
	Gapfill		0
	Crosswalk to: 0035U		9
81. 0570U: Testing for proteins in whole blood or plasma, individual components reported with overall result of elevated or non-elevated based on threshold comparison in traumatic brain injury	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 0358U		10
82. X184U: Neurology (Alzheimer disease), plasma, 3 distinct isoform-specific peptides (APOE2, APOE3, and APOE4) by liquid chromatography with tandem mass spectrometry (LC-MS/MS), reported as an APOE prototype	Activity type	Multiple choice	

	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	1
	Gapfill	2
	Crosswalk to: 0412U	2
	Crosswalk to: 0412U x .33	5
83. 0521U: Rheumatoid factor IgA and IgM, cyclic citrullinated peptide (CCP) antibodies, and scavenger receptor A (SR-A) by immunoassay, blood	Activity type	Multiple choice
	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	0
	Gapfill	0
	Crosswalk to: 86200 x 4	9
	Crosswalk to: 0312U x 0.5	1
84. 0522U: Carbonic anhydrase VI, parotid specific/secretory protein and salivary protein 1 (SP1), IgG, IgM, and IgA antibodies, chemiluminescence, semiquantitative, blood	Activity type	Multiple choice
	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	0
	Gapfill	9
	Crosswalk to: 81490	1
85. 0525U: Oncology, spheroid cell culture, 11-drug panel (carboplatin, docetaxel, doxorubicin, etoposide, gemcitabine, niraparib, olaparib, paclitaxel, rucaparib, topotecan, veliparib) ovarian...	Activity type	Multiple choice
	Total responses	10
	Unique participants	10
	Response options	Count
	Abstain	0
	Gapfill	0
	Crosswalk to: 0248U	10

86. 0526U: Nephrology (renal transplant), quantification of CXCL10 chemokines, flow cytometry, urine, reported as pg/mL creatinine baseline and monitoring over time	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		1
	Gapfill		9
87. 0545U: Acetylcholine receptor (AChR), antibody identification by immunofluorescence, using live cells, reported as positive or negative	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		10
88. 0546U: Low-density lipoprotein receptor-related protein 4 (LRP4), antibody identification by immunofluorescence, using live cells, reported as positive or negative	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		10
89. 0547U: Neurofilament light chain (NfL), chemiluminescent enzyme immunoassay, plasma, quantitative	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 0361U		10
90. 0548U: Glial fibrillary acidic protein (GFAP), chemiluminescent enzyme immunoassay, using plasma	Activity type	Multiple choice	
	Total responses		10

	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 0361U		10
91. 0551U: Tau, phosphorylated, pTau217, by single-molecule array (ultrasensitive digital protein detection), using plasma	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		10
92. 0568U: Testing for proteins by ultra high-sensitivity molecule array detection in plasma, algorithm reported as positive, intermediate, or negative for Alzheimer pathology	Activity type	Multiple choice	
	Total responses		10
	Unique participants		10
	Response options	Count	
	Abstain		0
	Gapfill		0
	Crosswalk to: 81503		10