

Q1 2018 Urine Culture



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Specimen 1 - 76 year old Male, Post TURP, Foley catheter

Organisms	Extent 1	2	3	4	5	Total
943 - Aerobe found; but referred for ID	9	0	1	0	1	11
993 - Growth of gram-negative organisms	4	0	0	0	0	4
814 - <i>Klebsiella pneumoniae</i>	2	0	2	0	0	4
991 - <i>Klebsiella</i> or <i>Enterobacter</i>	0	0	3	0	0	3
811 - <i>Klebsiella</i> sp.; NOS	0	0	2	0	0	2
989 - <i>Klebsiella</i> ; <i>Staphylococcus</i> or <i>Streptococcus</i>	0	0	1	0	0	1
TOTAL PARTICIPANTS						25

Flagging appears for failure to report 811, 814, 942, 919, 942, 943, 983, 989, 991 or 993.

In addition to the required organism, participants in all extents may report (No additional codes)

This sample contained *Klebsiella pneumoniae*.

Specimen 2 - 81 year old Female, Incontinency

Organisms	Extent 1	2	3	4	5	Total
943 - Aerobe found; but referred for ID	9	0	1	0	1	11
987 - <i>E.coli</i> ; <i>Citrobacter</i> or <i>Enterobacter</i>	1	0	3	0	1	5
993 - Growth of gram-negative organisms	4	0	0	0	0	4
799 - <i>Escherichia coli</i>	1	0	3	0	0	4
798 - <i>Escherichia</i> sp.; NOS	0	0	1	0	0	1
718 - Normal flora found, not normally reported	0	0	1	0	0	1
994 - Growth of gram-positive organisms	0	0	0	0	1	1
TOTAL PARTICIPANTS						27

Flagging appears for failure to report 798, 799, 919, 942, 943, 987 or 993.

In addition to the required organism, participants in all extents may report 718, 777, and 994

This sample contained *Escherichia coli* and *Corynebacterium* sp.

Specimen 3 - 20 year old Male, Fever pain

Organisms	Extent 1	2	3	4	5	Total
943 - Aerobe found; but referred for ID	10	0	1	0	1	12
994 - Growth of gram-positive organisms	4	0	1	0	0	5
989 - <i>Klebsiella</i> ; <i>Staphylococcus</i> or <i>Streptococcus</i>	0	0	2	0	0	2
873 - <i>Staphylococcus</i> sp.; NOS	0	0	2	0	0	2
879 - <i>Staphylococcus saprophyticus</i>	0	0	2	0	0	2
942 - Primary culture only, refer for ID	1	0	0	0	0	1
874 - <i>Staphylococcus</i> sp.; coagulase-negative; NOS	0	0	1	0	0	1
TOTAL PARTICIPANTS						25

Flagging appears for failure to report 873, 874, 879, 942, 943, 989 or 994.

In addition to the required organism, participants in all extents may report (No additional codes)

This sample contained *Staphylococcus saprophyticus*.

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Specimen 4 - 35 year old Female, Hematuria

Organisms	Extent	1	2	3	4	5	Total
943 - Aerobe found; but referred for ID		10	0	1	0	1	12
994 - Growth of gram-positive organisms		5	0	1	0	0	6
873 - Staphylococcus sp.; NOS		0	0	3	0	0	3
989 - Klebsiella; Staphylococcus or Streptococcus		0	0	2	0	0	2
878 - Staphylococcus epidermidis		0	0	1	0	0	1
874 - Staphylococcus sp.; coagulase-negative; NOS		0	0	1	0	0	1
		TOTAL PARTICIPANTS					25

Flagging appears for failure to report 873, 874, 878, 943, 989, or 994

In addition to the required organism, participants in all extents may report (No additional codes)

This sample contained *Staphylococcus epidermidis*.

Specimen 5 - 45 year old Female, Painful urination

Organisms	Extent	1	2	3	4	5	Total
943 - Aerobe found; but referred for ID		9	0	2	0	1	12
987 - E.coli; Citrobacter or Enterobacter		1	0	3	0	1	5
993 - Growth of gram-negative organisms		4	0	0	0	0	4
799 - Escherichia coli		1	0	3	0	0	4
994 - Growth of gram-positive organisms		2	0	0	0	1	3
798 - Escherichia sp.; NOS		0	0	1	0	0	1
718 - Normal flora found, not normally reported		0	0	1	0	0	1
		TOTAL PARTICIPANTS					30

Flagging appears for failure to report 798, 799, 943, 987 or 993.

In addition to the required organism, participants in all extents may report 718, and 994.

This sample contained *Escherichia coli* and *Corynebacterium* sp.