

Objectives

- ATP process for USEPA approval of Colilert-18 for fecal coliforms
- What is a number- MPN and MF & 95% confidence limits
- Review of calculations for microbiology tests



Colilert-18 Fecal Coliform ATP Study

- The study was based on USEPA ATP protocol (40CFR136)
- Colilert-18/Quanti-Tray compared against m-FC to determine fecals in waste water
- Based on the ATP Study, the EPA determined that Colilert-18 is a suitable method for the detection of fecal coliforms in wastewater.

Protocol Summary

- Primary wastewater effluent samples from 10 different locations within the New England area were approved for use as sources of fecal coliform bacteria for this study.
- Effluent samples were collected, processed and tested at a certified laboratory within six hours.



Colilert-18 -Fecal Coliform ATP Study

- Effluent samples were diluted to approximately 20-50 organisms per 100 mL of sample.
- Each sample was analyzed by Colilert-18 and SM9222D in replicates of 20 after incubation at 44.5 ±0.2°C.
- Sensitivity of fecal coliform detection was determined by comparing the number of presumptive positive responses for the Colilert-18 against the number of presumptive positive responses from m-FC
- Specificity was determined by confirming the identity of at least 200 presumptive positive and 200 presumptive negative determinations from each test method.



Separation of Snap Packs



Quanti-Tray Demonstration



Add Colilert/Colilert-18 to sample and shake to dissolve



Pour mixture into a Quanti-Tray



Quanti-Tray Demonstration



Seal and incubate at 44.5°C for fecal coliforms for 18 hours



Incubate Samples at $44.5 \pm 0.2^\circ\text{C}$
for Colilert-18 for Fecal coliforms



DEX

Or A Water Bath for Fecal
Coliforms at $44.5 \pm 0.2^\circ\text{C}$



DEX

Quanti-Tray Demonstration

Count positive wells and
refer to MPN table



DEX

Quality Control for Colilert-18

- Use of positive and negative controls
 - For Colilert-18, test upon receipt of each lot of product
- Quanti-Cult
 - 3 sets of a positive Coliform
 - 3 sets of positive coliform/E.coli
 - 3 sets of a negative control
- ATCC Strains



IDEXX

Internal Requirements for Products

- IDEXX is an ISO 9001-2009 certified facility
- IDEXX manufactures under 6 Sigma
 - = or < 3.4 defects/million opportunities
 - Goal is to improve processes to that level of quality or better
- All manufacturing follows approved manufacturing documentation
- All products must meet strict QC requirements
- QA responsible to review all batch records prior to releasing the product for sale.

IDEXX

Certificates of Quality

- **Obtain a C of Q for all our products from our website**
 - <http://www.idexx.com/water/certificates/>
 - Indicates all the testing performed for the specific product and lot
 - Signed by QC for completion date of testing along with QA approval
 - All products are under quarantined until the batch records are reviewed and approved by QA
 - Once approved by QA, the specific product and lot is released for distribution

IDEXX

Requirements to use Colilert-18 & Quanti-Tray from USEPA region 4 and FL

- Region 4 Science and Division approves the use of IDEXX Colilert-18 for the detection and enumeration of fecal coliforms in wastewater samples as an Alternate Test Procedure (ATP) until it has been formally promulgated nationally as part of 40 CFR Part 136 under the Clean Water Act.
- Labs are to send their requests to use Colilert-18 for fecal coliform detection to Region 4 (next slides)
- FL is aware of this approval by Region 4; Contact accreditation body to confirm what is required such as DOC.
- **Private labs will need to list all facilities for which they will use this method**



Requirements to use Colilert-18 & Quanti-Tray from USEPA Region 4 and FL.

- This letter should be submitted in writing via fax, e-mail, or standard mail to Ray Terhune, the Regional DMR-QA Coordinator, for Regional approval of this ATP process.
- EPA Region 4
- Science and Ecosystem Support Division
- Quality Assurance Section
- 980 College Station Road
- Athens, Georgia 30605-2720
- Telephone: (706) 355-8557
- Email address: Terhune.Ray@epa.gov
- The facility should keep a copy of their request and approval letter on file for State Inspection purposes.

Estimation of Bacterial Densities by MPN

- Dates back to 1915 when the concept was introduced by M.H. McCrady (J of Infectious Disease Vol 17, 1915*)
- Prior to this novel concept; no means of direct counting
 - Only had presence-absence of fermentation tubes
- The method is a means for estimating without any direct count, the density of organisms in a liquid.
- Multiple samples of the liquid are taken and incubated in suitable media
 - Record presence or absence of growth in each sample tube
 - Ingenious application of probability theory
 - Estimate the number of organisms from the number of negative tubes

*The Numerical Interpretation of Fermentation Tube Results

MPN Theory – a short math lesson

Basic assumptions:

- The organisms are distributed randomly throughout the liquid
 - Sample is well shaken (important and often neglected)
- A sample will exhibit growth (in the culture media) whenever one or more of the target organisms is present
 - Media should be selective and sensitive
- Requires that at least 1 tube shows no growth (sterile)

If n samples, each of volume v mL are taken from a liquid, and s of these are sterile, then an estimate of the organism density d per mL in the original sample is:

$$d = -2.303/v \log(s/n) \text{ [Poisson distribution]}$$

d can be shown statistically to have the highest probability of estimating the actual organism density – the "Most Probable Number" - MPN

Single dilution – Maximum counting range

Number of Tubes	Maximum MPN/100mL	Number of Tubes	Maximum MPN/100mL
5	8	35	125
10	23	40	148
15	40	45	171
20	60	50	196
25	80	75	324
30	102	100	461

$n = 50, v = 2 \text{ mL}, s = 1$

$$d = -2.303/v \log(s/n)$$

$$d = -2.303/2 \log(1/50) = 1.956 \text{ MPN/mL} = 195.6 \text{ MPN/100 mL}$$

Uncertainty of MPN

•Skewed distribution & Uncertainty of Result

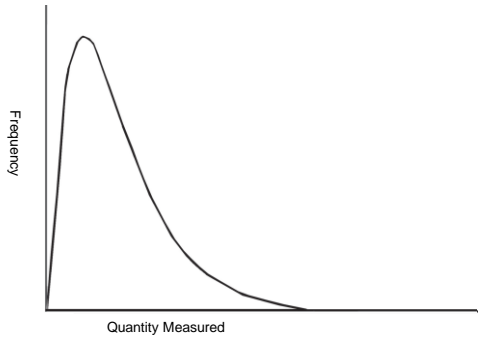
For microbiological tests, because of the complications of the skewed distribution, the estimate of uncertainty from a well-designed precision experiment will be regarded as a reasonable estimate of the uncertainty of results

➤ When MPN tables are used to obtain results, the 95% CL in the table will be regarded as a reasonable estimate of the uncertainty of the results

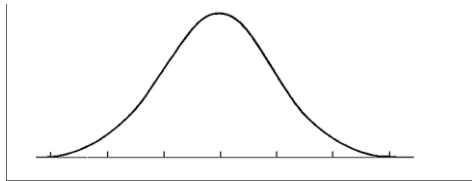
➤ Labs are encourage to identify and question unusual combinations of positive tubes in the tables.

- 15 tube MPN combination of + tubes
 - » 1-2-3
 - » 3-5-4

Frequency curve (positively skewed distribution)



Normal Distribution (Bell Shape)



Standard Methods MPN Table 15 Tube

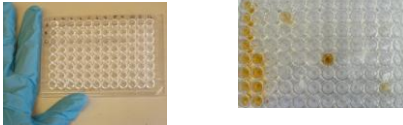
9-54 MICROBIOLOGICAL EXAMINATION 8000

Table 15-11 (V) MPN Index and 95% Confidence Limits for Various Combinations of Positive Results in Three Tubes per Dilution

Combinations of Positives	MPN Index/100 ml	Confidence Limits		Combinations of Positives	MPN Index/100 ml	Confidence Limits	
		Low	High			Low	High
0-0-0	< 0.1	0	0.6	4-0-0	75	50	100
0-0-1	1.8	0.000	6.8	4-1-0	17	6.0	40
0-0-2	3.6	0.000	10	4-1-1	24	8.5	50
0-0-3	5.7	0.70	10	4-1-2	33	10	50
0-1-0	3.6	1.0	13	4-1-3	45	10	50
0-1-1	6.0	0.60	15	4-2-0	32	10	50
0-1-2	10.0	0.70	15	4-2-1	45	14	100
0-1-3	16.0	1.0	20	4-2-2	60	14	100
1-0-0	3.6	0.75	13	4-2-3	80	14	100
1-0-1	6.0	1.0	15	4-3-0	110	14	100
1-0-2	9.0	1.0	15	4-3-1	150	14	100
1-0-3	14.0	1.0	20	4-3-2	200	14	100
1-1-0	6.0	1.0	15	4-3-3	280	14	100
1-1-1	10.0	1.0	20	4-4-0	400	14	100
1-1-2	16.0	1.0	25	4-4-1	550	14	100
1-1-3	25.0	1.0	35	4-4-2	750	14	100
2-0-0	9.0	1.0	20	4-4-3	1000	14	100
2-0-1	15.0	1.0	25	4-5-0	1400	14	100
2-0-2	22.0	1.0	30	4-5-1	1900	14	100
2-0-3	34.0	1.0	45	4-5-2	2600	14	100
2-1-0	15.0	1.0	35	4-5-3	3600	14	100
2-1-1	25.0	1.0	50	4-6-0	5000	14	100
2-1-2	38.0	1.0	70	4-6-1	7000	14	100
2-1-3	57.0	1.0	100	4-6-2	10000	14	100
3-0-0	18.0	1.0	45	4-6-3	14000	14	100
3-0-1	30.0	1.0	70	4-7-0	20000	14	100
3-0-2	45.0	1.0	100	4-7-1	28000	14	100
3-0-3	68.0	1.0	150	4-7-2	40000	14	100
3-1-0	30.0	1.0	75	4-7-3	55000	14	100
3-1-1	50.0	1.0	120	4-8-0	75000	14	100
3-1-2	75.0	1.0	180	4-8-1	105000	14	100
3-1-3	110.0	1.0	250	4-8-2	145000	14	100
4-0-0	36.0	1.0	90	4-8-3	200000	14	100
4-0-1	60.0	1.0	150	4-9-0	280000	14	100
4-0-2	90.0	1.0	220	4-9-1	400000	14	100
4-0-3	135.0	1.0	330	4-9-2	550000	14	100
4-1-0	60.0	1.0	150	4-9-3	750000	14	100
4-1-1	100.0	1.0	250	4-10-0	1050000	14	100
4-1-2	150.0	1.0	370	4-10-1	1450000	14	100
4-1-3	225.0	1.0	550	4-10-2	2000000	14	100
5-0-0	72.0	1.0	180	4-10-3	2800000	14	100
5-0-1	120.0	1.0	300	4-11-0	4000000	14	100
5-0-2	180.0	1.0	450	4-11-1	5500000	14	100
5-0-3	270.0	1.0	670	4-11-2	7500000	14	100
5-1-0	120.0	1.0	300	4-11-3	10500000	14	100
5-1-1	200.0	1.0	500	4-12-0	14500000	14	100
5-1-2	300.0	1.0	750	4-12-1	20000000	14	100
5-1-3	450.0	1.0	1100	4-12-2	28000000	14	100
6-0-0	84.0	1.0	210	4-12-3	40000000	14	100
6-0-1	140.0	1.0	350	4-13-0	55000000	14	100
6-0-2	210.0	1.0	520	4-13-1	75000000	14	100
6-0-3	315.0	1.0	770	4-13-2	105000000	14	100
6-1-0	140.0	1.0	350	4-13-3	145000000	14	100
6-1-1	230.0	1.0	570	4-14-0	200000000	14	100
6-1-2	340.0	1.0	850	4-14-1	280000000	14	100
6-1-3	510.0	1.0	1250	4-14-2	400000000	14	100
7-0-0	108.0	1.0	270	4-14-3	550000000	14	100
7-0-1	180.0	1.0	450	4-15-0	750000000	14	100
7-0-2	270.0	1.0	670	4-15-1	1050000000	14	100
7-0-3	405.0	1.0	1000	4-15-2	1450000000	14	100
7-1-0	180.0	1.0	450	4-15-3	2000000000	14	100
7-1-1	300.0	1.0	750	4-16-0	2800000000	14	100
7-1-2	450.0	1.0	1100	4-16-1	4000000000	14	100
7-1-3	675.0	1.0	1600	4-16-2	5500000000	14	100
8-0-0	144.0	1.0	360	4-16-3	7500000000	14	100
8-0-1	240.0	1.0	600	4-17-0	10500000000	14	100
8-0-2	360.0	1.0	900	4-17-1	14500000000	14	100
8-0-3	540.0	1.0	1350	4-17-2	20000000000	14	100
8-1-0	240.0	1.0	600	4-17-3	28000000000	14	100
8-1-1	400.0	1.0	1000	4-18-0	40000000000	14	100
8-1-2	600.0	1.0	1500	4-18-1	55000000000	14	100
8-1-3	900.0	1.0	2250	4-18-2	75000000000	14	100
9-0-0	180.0	1.0	450	4-18-3	105000000000	14	100
9-0-1	300.0	1.0	750	4-19-0	145000000000	14	100
9-0-2	450.0	1.0	1100	4-19-1	200000000000	14	100
9-0-3	675.0	1.0	1600	4-19-2	280000000000	14	100
9-1-0	300.0	1.0	750	4-19-3	400000000000	14	100
9-1-1	500.0	1.0	1250	4-20-0	550000000000	14	100
9-1-2	750.0	1.0	1875	4-20-1	750000000000	14	100
9-1-3	1125.0	1.0	2800	4-20-2	1050000000000	14	100
10-0-0	360.0	1.0	900	4-20-3	1450000000000	14	100
10-0-1	600.0	1.0	1500	4-21-0	2000000000000	14	100
10-0-2	900.0	1.0	2250	4-21-1	2800000000000	14	100
10-0-3	1350.0	1.0	3375	4-21-2	4000000000000	14	100
10-1-0	360.0	1.0	900	4-21-3	5500000000000	14	100
10-1-1	600.0	1.0	1500	4-22-0	7500000000000	14	100
10-1-2	900.0	1.0	2250	4-22-1	10500000000000	14	100
10-1-3	1350.0	1.0	3375	4-22-2	14500000000000	14	100
11-0-0	420.0	1.0	1050	4-22-3	20000000000000	14	100
11-0-1	700.0	1.0	1750	4-23-0	28000000000000	14	100
11-0-2	1050.0	1.0	2625	4-23-1	40000000000000	14	100
11-0-3	1575.0	1.0	3938	4-23-2	55000000000000	14	100
11-1-0	420.0	1.0	1050	4-23-3	75000000000000	14	100
11-1-1	700.0	1.0	1750	4-24-0	105000000000000	14	100
11-1-2	1050.0	1.0	2625	4-24-1	145000000000000	14	100
11-1-3	1575.0	1.0	3938	4-24-2	200000000000000	14	100
12-0-0	480.0	1.0	1200	4-24-3	280000000000000	14	100
12-0-1	800.0	1.0	2000	4-25-0	400000000000000	14	100
12-0-2	1200.0	1.0	3000	4-25-1	550000000000000	14	100
12-0-3	1800.0	1.0	4500	4-25-2	750000000000000	14	100
12-1-0	480.0	1.0	1200	4-25-3	1050000000000000	14	100
12-1-1	800.0	1.0	2000	4-26-0	1450000000000000	14	100
12-1-2	1200.0	1.0	3000	4-26-1	2000000000000000	14	100
12-1-3	1800.0	1.0	4500	4-26-2	2800000000000000	14	100
13-0-0	540.0	1.0	1350	4-26-3	4000000000000000	14	100
13-0-1	900.0	1.0	2250	4-27-0	5500000000000000	14	100
13-0-2	1350.0	1.0	3375	4-27-1	7500000000000000	14	100
13-0-3	2025.0	1.0	5063	4-27-2	10500000000000000	14	100
13-1-0	540.0	1.0	1350	4-27-3	14500000000000000	14	100
13-1-1	900.0	1.0	2250	4-28-0	20000000000000000	14	100
13-1-2	1350.0	1.0	3375	4-28-1	28000000000000000	14	100
13-1-3	2025.0	1.0	5063	4-28-2	40000000000000000	14	100
14-0-0	600.0	1.0	1500	4-28-3	55000000000000000	14	100
14-0-1	1000.0	1.0	2500	4-29-0	75000000000000000	14	100
14-0-2	1500.0	1.0	3750	4-29-1	105000000000000000	14	100
14-0-3	2250.0	1.0	5625	4-29-2	145000000000000000	14	100
14-1-0	600.0	1.0	1500	4-29-3	200000000000000000	14	100
14-1-1	1000.0	1.0	2500	4-30-0	280000000000000000	14	100
14-1-2	1500.0	1.0	3750	4-30-1	400000000000000000	14	100
14-1-3	2250.0	1.0	5625	4-30-2	550000000000000000	14	100
15-0-0	660.0	1.0	1650	4-30-3	750000000000000000	14	100
15-0-1	1100.0	1.0	2750	4-31-0	1050000000000000000	14	100
15-0-2	1650.0	1.0	4125	4-31-1	1450000000000000000	14	100
15-0-3	2475.0	1.0	6188	4-31-2	2000000000000000000	14	100
15-1-0	660.0	1.0	1650	4-31-3	2800000000000000000	14	100
15-1-1	1100.0	1.0	2750	4-32-0	4000000000000000000	14	100
15-1-2	1650.0	1.0	4125	4-32-1	5500000000000000000	14	100
15-1-3	2475.0	1.0	6188	4-32-2	7500000000000000000	14	100
16-0-0	720.0	1.0	1800	4-32-3	10500000000000000000	14	100
16-0-1	1200.0	1.0	3000	4-33-0	14500000000000000000	14	100
16-0-2	1800.0	1.0	4500	4-33-1	20000000000000000000	14	100
16-0-3	2700.0	1.0	6750	4-33-2	28000000000000000000	14	100
16-1-0	720.0	1.0	1800	4-33-3	40000000000000000000	14	100
16-1-1	1200.0	1.0	3000	4-34-0	55000000000000000000	14	100
16-1-2	1800.0	1.0	4500	4-34-1	75000000000000000000	14	100
16-1-3	2700.0	1.0	6750	4-34-2	105000000000000000000	14	100
17-0-0	780.0	1.0	1950	4-34-3	145000000000000000000	14	100
17-0-1	1300.0	1.0	3250	4-35-0	200000000000000000000	14	100
17-0-2	1950.0	1.0	4875	4-35-1	280000000000000000000	14	100
17-0-3	2925.0	1.0	7313	4-35-2	400000000000000000000		

Overcoming the tube numbers problem

- To increase counting range need more tubes
- For 100mL sample could use 50 tubes x 2mL
- Would give counting range of ~200MPN/100mL
 - Big headache for lab – lots of tubes – lots of pipettes -lots of washing! Not really practical.
 - Microtitre plate (96 wells) – MPN range 438/100mL



Quanti-Tray®

- Use multiple well concept but automate the sealing



51 well Quanti-Tray
Total sample volume 100mL

$$d = -2.303/v \log(s/n)$$

$$n = 51, v = 1.96 \text{ mL}, s = 1$$

$$d = -2.303/1.96 \log(1/51) = 2.006 \text{ MPN/mL} = 200.6 \text{ MPN/100 mL}$$



Multiple Dilutions

- Not practical to extend the counting range by increasing the number of wells beyond a certain point.
- Sample serially diluted and each dilution is inoculated into a similar number of tubes. Commonly used configurations:
 - 3 tubes x 3 decimal dilutions – food industry
 - 5 tubes x 3 decimal dilutions – food, water wastewater (Max MPN = 1600/100mL)
- With multiple dilutions the statistical calculation of the MPN value becomes more complex. **Thomas** provided an **approximation** that could be used for any combination of tubes

$$\text{MPN/100mL} = \frac{\# \text{ of positive tubes} \times 100}{\sqrt{(\# \text{ of mL in negative tubes}) \times (\# \text{ of mL in all tubes})}}$$



MPN vs. CFU

- Is there a difference different between the two?
 - A simple answer is that the use of either term is based on the method the lab uses for the detection of bacteria
 - [Additional information](#)



Distribution of Bacterial Populations

- For chemical analysis the distribution of results usually follows a bell shape curve-symmetrical distribution about the mean
- Microbial distribution are usually not symmetrical and have a skewed distribution . This can be a result of many low values and some high values
 - This can lead to the arithmetic mean to be much larger than the median
- Statistical techniques requires the assumption of symmetrical distributions such as a normal curve and the skewed data is converted to log values resemble a normal distribution.



Calculate the coliform density of all the samples and convert to Log₁₀ to obtain a geometric mean. Average the 7 samples and take the anti-log to obtain the final value for the 7 samples.

Sample No.	Coliform Density	Log ₁₀
1	6.0 x 10 ⁵	5.78
2	4.2 x 10 ⁶	6.63
3	1.6 x 10 ⁶	6.22
4	1.4 x 10 ⁵	6.14
5	4.0 x 10 ⁵	5.60
6	1.0 x 10 ⁶	6.02
7	5.1 x 10 ⁵	5.71

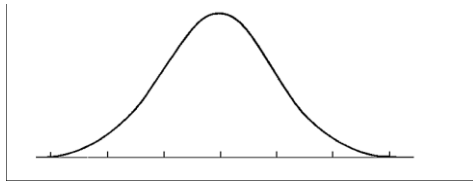
The average value Log₁₀ = 6.01 and the antilog = 1.03 x 10⁶



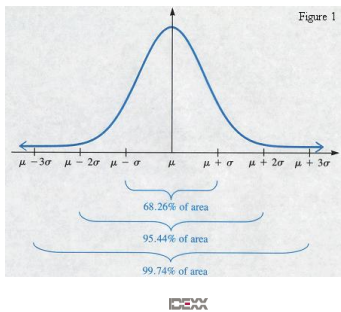
Frequency curve (positively skewed distribution)



Normal Distribution (Bell Shape)



•The graph of the distribution is a bell-shaped curve (a.k.a. a normal curve).



Precision of Quantitative Methods

- Perform duplicate analyses on first 15 positive samples of each matrix type, with each set of duplicates analyzed by a single analyst. If there is more than one analyst, include all analysts regularly running the tests, with each analyst performing an approximately equal number of tests
- Calculate precision of replicate analyses for each different type of sample examined
- Record duplicate analyses as *D1* and *D2* & calculate the logarithm of each result.
- If either of a set of duplicate results is <1 , add 1 to both values before calculating the log
- Calculate the range (*r*) for each pair of transformed duplicates as the overall mean (*R*) of these ranges. Transform the duplicates and calculate their range

Precision of Quantitative Methods

- If the range is greater than $>3.27R$, there is greater than 99% probability that the laboratory variability is excessive; in such a case, discard all analytical results since the last precision
- Identify and resolve the analytical problem before making further analyses.
- Example



Calculations

- Distribution of Bacterial Population-Geometric Mean
- Precision of Quantitative Methods
- Membrane Filtration
- MPN



Sample No.	Duplicate Analyses		Logarithms of Counts		Range of Logarithms (rlog) (L1-L2)
	D1	D2	L1	L2	
1	89	71	1.9494	1.8513	0.0981
2	38	34	1.5798	1.5315	0.0483
3	58	67	1.7634	1.8261	0.0627
14	7	6	0.8451	0.7782	0.0669
15	110	121	2.0414	2.0828	0.0414



Precision of Quantitative Methods

- Calculations from the table to determine the precision
 $\sum \text{r}_{\log} = 0.0981 + 0.0483 + 0.0627 + \dots + 0.0669 + 0.0414 = 0.7189$
 $R = \sum \text{r}_{\log} / n = 0.7189 / 15 = 0.479$
 Precision Criterion = $3.27(R) = 3.27(0.479) = 0.157$



Precision on Duplicates

- Thereafter, analyze 10% of routine samples in duplicate or one per test run
- Calculations:

Analyses	Duplicates		Log of		Range of Log	Results
	D1	D2	D1	D2		
1	71	65	1.85	1.81	0.04	A
2	74	50	1.88	1.70	0.18	U

Precision Criterion = $3.27(R) = 0.157$
 A = Acceptable, U = Unacceptable



Membrane Filtration

- Standard methods for Water & Waste Water (20 Edition)
 - MF for total coliforms requires counts between 20-80(coliforms) and 20-60 (fecal coliforms)with no more than 200 bacteria on a plate
 - If the total number exceeds 200 (coliforms & non-coliforms), or the colonies are not distinct enough to be counted &/or confluent growth, report results as TNTC or confluent
 - ill usually be absent of coliforms or minimal number
 - Clean/good quality water- If no colonies are observed, report the results as <1/100mL; if results are <20, calculate the results based on the number of colonies on the plate
 - Total colonies/100mL = Number of colonies x100/mL sample filtered
 - Example 8 colonies, 100 mL filtered:
Colonies/100mL = 8x100/100mL = 8/100mL



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- For duplicate samples; 2 X 50mL samples filtered and the 2 membranes had counts of 5 & 3 then calculate the colonies/100mL as:
 - total colonies/100mL = [(5+3 X100)]/(50 +50) = 8 colonies/100mL
- For waste, recreational or raw waters , 3 plates are usually necessary to achieve the 20-80 or 20-60 range to find the total colonies/100mL
 - Examples:



Number of Colonies on MF Plates; 3 dilutions/sample

Sample No.	0.1 mL Filtration	1 mL Filtration	10 mL Filtration
1	0	1	23
2	2	18	TNTC
3	0	8	65
4	0	5	58
5	0	1	17
6	0	1	39
7	0	1	20