

Effect of Membrane Filter Pore Size on Microbial Recovery and Colony Morphology

Summary

Membrane filters with a $0.45 \, \mu m$ pore size have long been recognized as the standard for growth of microorganisms. However, there is little published literature comparing the effects of different pore sizes on colony size and recovery.

The 0.45 μ m pore size is used to recover bacteria and other microorganisms from many samples and environments—almost to the exclusion of other pore sizes. Only rarely are other sizes used for growth and recovery and there is little information available on the effects of different pore sizes on microorganisms. However, other pore sizes are commercially available for microbial enumeration and users will occasionally substitute a filter with a pore size larger or smaller than 0.45 μ m in an attempt to improve their results.

This study takes a broad look at the influence of different pore sizes on some common microorganisms. It provides data on the effect of pore size on growth and recovery. The study compares a variety of microorganism/media combinations on a range of pore sizes: 0.22, 0.45, 0.7, 0.8, and 1.2 μ m. The filtration method used in the study was a standard glass funnel and base with vacuum. Test filters were plated on solid (agar) media and compared against spread plates.

There was no universal pattern of results. Some microorganisms, such as *Micrococcus luteus* and *Candida albicans* showed no significant difference in recovery or colony size with membrane pore size. Other organisms such as *Pantoea agglomerans* showed no difference in colony size but had low recoveries on 1.2 and 0.22 µm membranes.

Acceptable recovery for membrane filters was defined as being $\geq 90\%$ versus the controls (spread plates).* The 0.45 μm membranes met this definition with all test systems. Some test systems showed equivalent recoveries with other pore sizes but in no case were the results significantly better. The lowest recoveries were seen with extremes of the pore size range (1.2 and 0.22 μm).

Materials and Methods

The pore sizes used in this study were selected from the various sizes of mixed esters of cellulose membranes manufactured by Millipore (Table 1).

The microorganisms and media combinations used in this study were chosen as a broad representation of common membrane filter applications: pharmaceutical, food and beverage, USP testing, water testing, and general microbiology (Table 2).

As an adjunct to the recovery and colony size experiments, the test filters were tested for their retentive capabilities under the conditions of average use. Each filter was challenged with a low level of *Brevundimonas (Pseudomonas) diminuta* (the standard organism for retention testing) and the filtrate was retained for enumeration.

Table 1. Test Filters

Pore Size (µm)	Millipore filter code	Flow Rate (sec/500 mL)	Bubble Point (psi)	Typical Applications	
0.22	GSWP	40 to 60	50 to 60	Sterile filtration	
0.45	HAWG	30 to 50	30 to 36	Microbial testing of water, beverages, and general micro- biology	
0.7	HCWG	15 to 23	19 to 23	Fecal coliform test- ing in surface and wastewater	
0.8	AAWG	10 to 16	14 to 18	Yeast and mold testing in beverages	
1.2	RAWG	7 to 11	11 to 13	Yeast and mold testing in "hard-to- filter" beverages	

Table 2. Test Systems

Microorganism	Source	Media	Temp. (°C)	Time (hours)
Primary effluent	Wastewater	m-Endo LES	35	24
Primary effluent	Wastewater	m-FC	44.5	24
Primary effluent	Wastewater	m-TEC	35 – 44.5	24
Bacillus subtilis	ATCC 13933	Tryptic soy agar	35	24
Brevundimonas* diminuta	ATCC 19146	Tryptic soy agar	30	48
Candida albicans	ATCC 10231	Tryptic soy agar	35	24
Clostridium sporogenes * *	ATCC 11437	Tryptic soy agar	35	48
Enterobacter aerogenes	ATCC 49701	Tryptic soy agar	35	24
Escherichia coli	ATCC 25922	m-FC	44.5	24
Micrococcus luteus	ATCC 9341	Tryptic soy agar	35	24
Pantoea agglomerans	Well water	m-Endo LES	35	48
Pantoea agglomerans	Well water	Tryptic soy agar	35	24

^{*}Previously grouped in the genera Pseudomonas

^{**}Grown anaerobically using a Gas Pak jar (BBL)

Results

The selection of test systems was not intended to be exhaustive but to give a broad overview of microbial recovery in relation to filter pore size. Six different filter pore sizes were tested with 12 microorganism/media combinations that are representative of the types of microorganisms encountered by those using the membrane filter (MF) technique.

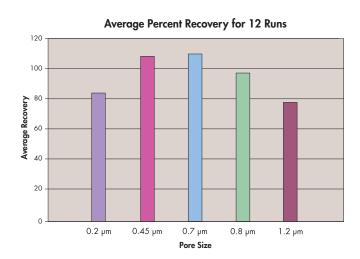
Colony Size

- Three test systems, Br. diminuta, Ent. aerogenes, and B. subtilis, showed differences in colony size with pore size
 - Colonies grown on 1.2 and 0.8 µm filters were larger than colonies grown on other filter pore sizes or spread plates.
 - Colonies grown on 0.7 µm filters were the same size as, or slightly larger than, colonies grown on spread plates.
 - Colonies on 0.45 and 0.2 µm filters were the same size as, or some what smaller than, colonies grown on spread plates.
- Other test systems showed virtually no difference in colony size with any of the other pore sizes as compared to colonies grown on spread plates.

Colony Size Versus Pore Size 0.2 µm 0.45 µm 0.7 µm 0.8 µm 1.2 µm Spread Plate 8 7 8 Br. diminuta Ent. aerogenes Microorganism

Microbial Recovery

- 0.45 µm and 0.7 µm filters demonstrated acceptable* recovery (≥ 90% versus spread plates) for all 12 test systems
- Although the average recovery for 0.8 µm filters was acceptable over the 12 test systems, the pore size had lower recoveries than 0.45 and 0.7 µm filters.
- Although 0.22 and 1.2 µm filters gave acceptable recoveries with some systems, their average recovery was significantly lower overall.



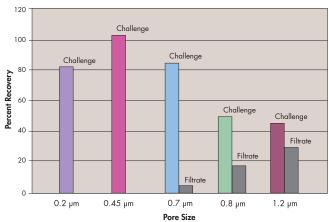
The average performance of each pore size was determined using all the test systems

^{*}Standard Methods defines acceptable recovery on membrane filters as 90% or greater compared to spread plates.

Retention

- The larger pore sizes (1.2 and 0.8 µm) allowed significant passage of a small organism at low challenge levels (starved *Br. diminuta*) but there was no passage with the 0.45 or 0.22 µm pore sizes.
- Although 0.22 µm filters retained the challenge, the average recovery across all test systems was lower than 0.45 µm filters.
- Passage might be one reason why larger pore sizes (> 0.7 µm) showed lower recoveries than smaller pore sizes.

Recovery of Challenge Versus Pore Size



Overall

Recovery is much more complex than the retention of microorganisms on the surface of a membrane filter and the influence of pore size. It is a combination of factors that may include:

- The microorganism species and its condition—each microorganism has the potential to react differently
- The sieving effects of the pore size as it relates to the retention of specific microorganisms
- Type of medium and selectivity
- Structure and chemistry of the membrane filter
- Environmental conditions (e.g., moisture, incubation, temperature)

The effect of filter pore size on any specific microorganism/medium combination is not always predictable. If pore sizes other than those indicated by industry standards are used, they should be validated on relevant samples and media and compared to $0.45~\mu m$.

This study confirmed that the standard 0.45 μm pore size is the most appropriate for general microbiological purposes. The 0.45 μm filters gave the most consistent recoveries across a variety of test systems and did not allow passage of the standard 0.2 μm sterilizing filter challenge microorganism, *B. diminuta*, under typical filtration conditions.

Conclusion

A membrane pore size larger than $0.45~\mu m$ can increase flow rate, throughput, and, occasionally, colony size (which makes the colonies easier to count). However, these larger pore sizes may not have sufficient retention for some microorganisms. Therefore, they are not well suited for total count applications.

Larger pore sizes can be used for enumerating specific organisms, such as fecal coliforms or yeast. They can also be used for difficult-to-filter samples where improved throughput or larger sample volumes are needed. In both cases, the filter's retention performance should be documented for the target microorganism(s).

Pore sizes smaller than $0.45~\mu m$ have the disadvantage of decreased flow rate, throughput, and, potentially, recovery. Therefore, the greater retentive properties of the $0.2~\mu m$ pore size have little benefit for the enumeration of bacteria, yeast, and molds in the variety of liquids considered in this study.

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