

Baylis, J. R., Jr. and A. G. DeBusk. Estimation of the frequency of multinucleate conidia in microconidiating strains.

Frequency of multinucleate microconidia.

Experiment Number	1	2	3
<u>ad-3 colonies</u> x 10 <sup>4</sup>			
per ml conidial suspension	7.42	6.40	76.00
<u>lys-3 colonies</u> x 10 <sup>4</sup>			
per ml conidial suspension	37.00	358.00	3.8
Ratio <u>ad-3 colonies</u> / <u>lys-3 colonies</u>	1/5	1/56	20/1
Total viable conidia x 10 <sup>4</sup>			
per ml conidial suspension	44.42	364.40	79.0
Wild type colonies			
per ml conidial suspension	83	53	19
Frequency @ colonies	0.836	0.982	0.048
Frequency <u>ad-3 colonies</u>	0.166	0.017	0.952
Per cent wild type colonies	0.0187	0.0014	0.0024
Estimated per cent multinucleate microconidia*	0.067	0.041	0.026

\* This value is obtained from the following equation:

$$\frac{\% \text{ wild type colonies}}{2(\text{lys-3 colony frequency}) (\text{ad-3 colony frequency})}$$

The frequency of multinucleate conidia in microconidiating strains of *N. crassa* was determined by means of a technique employing forced heterocaryons. The strains used were of the following genotypes: ad-3A; pe, fl (38701; Y8743m,L) and lys-3; pe, fl (4545; Y8743m, L).

Heterocaryons were formed by placing drops of a mixed microconidial suspension on plates of minimal medium. The heterocaryons formed on the plates were transferred to minimal agar slants and incubated. Microconidial suspensions from three independent heterocaryons were analyzed. Each was filtered through Nitex f53 mesh and glass wool to remove conidial clumps and mycelial fragments. Aliquots of the filtered suspension were plated on minimal, adenine-supplemented, and lysine-supplemented medium. From the plate counts and by application of the binomial theorem the frequency of multinucleate conidia was determined (Table). To simplify the calculations, all multinucleate microconidia were considered binucleate. The frequency of multinucleate microconidia varied little over a wide range of nuclear ratios. These percentages probably represent the upper limits of multinucleate microconidial frequencies since an undetermined fraction of the wild type colonies formed may have had their origin as multinucleate mycelial fragments or as newly-formed heterocaryons. The percentages of multinucleate microconidia obtained were less than 0.1% in all cases, and therefore somewhat lower than those reported by Barratt and Garnjobst (1949 Genetics 34: 351).