

Estimation of clones numbers needed for Library

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Estimation of number of clones needed for Library

Clark & Carbon equation 1976

$$N = \frac{\ln(1-P)}{\ln(I-n)}$$

حيث إن :

N = عدد الكلونات التي يجب اختيارها لتمثيل معظم مورثات الجين في المكتبة .

n = نسبة حجم قطع الحامض النووي الهندسة وراثياً إلى جين الكائن .

p = احتمالية وجود أي مورث في المكتبة .

Example: ✕

Library size needed for mouse genomic Library: ✕

--- size of DNA fragments 20 Kb ✕

--- Mice genome 30000000000 pb ✕

--- Probability (P) 95% ✕

$$\frac{\ln(1-P)}{\ln(1-n)} \quad 20 \quad ✕$$

$$N = \frac{\ln(1-0.95)}{\ln(1-1.4 \times 10^{10})} = 1.4 \times 10^{10} \text{ Cells} \quad ✕$$

$$\frac{\ln(1-0.95)}{\ln(1-1.4 \times 10^{10})} = 4.2 \times 10^{10} \text{ Cells} \quad ✕$$

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How can you calculate your library size? ✘

After transformation assay: ✘

1. Make 10 -1; 10 -2,.....10 -10 dilutions by adding 100 ul from the library to 900 ul from LB or agar broth to have the dilution 10 -1 then make the rest dilutions from this stock.
2. Choose the dilutions from 10 -7 to 10 -10. ✘
3. Take 100 ul from each dilution and plate it on LB or Agar plate – 3 plates each-. ✘
4. Incubate plate at 37.5C for 24 hrs. ✘
5. Copy all plates colonies and re-plate them on AMP LB or Agar plates. ✘
6. Incubate the plates at 37.5C for 24 hrs. ✘
7. Calculate the AMP resist colonies, compare these colonies with those in steps 4-5. ✘
8. The number of AMP sensitive colonies will refer to positive clones = Library size ✘

<i>Haemophilus influenzae</i>	1,830,135	✗
<i>Helicobacter pylori</i>	1,667,867	✗
<i>Bacillus subtilis</i>	4,214,814	✗
<i>Mycoplasma genitalium</i>	580,073	✗
<i>Archaeoglobus fulgidus</i>	2,178,400	✗
Eukaryotes	✗	
<i>Saccharomyces cerevisiae</i>	12,069,313	✗
<i>Caenorhabditis elegans</i>	97,000,000	✗
<i>Drosophila melanogaster</i>	180,000,000	✗
<i>Arabidopsis thaliana</i>	115,500,000	✗
<i>Homo sapiens</i>	3,200,000,000	✗
<i>Mus domesticus</i>	3,000,000,000	✗

Thank you a lot for listening

