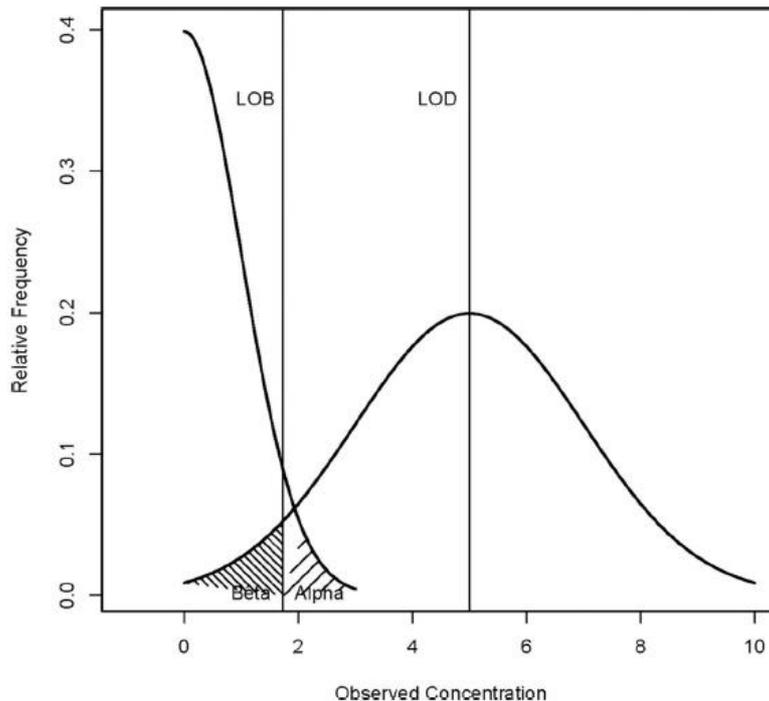


## MEMO - How to calculate the limit of detection (LOD)

For automated computation, see the statistical tools available on [www.gene-pi.com](http://www.gene-pi.com)

- a) **Definition:** the limit of Detection  $LOD$  with a confidence level  $(1 - \beta)$  is defined (in concentration unit) as the minimum concentration for which detecting the target sequence in a well is possible with a probability of  $1 - \beta$ .

In other words, this is the minimum concentration that can be said to be non-zero and statistically higher than the limit of blank  $LOB$  with a  $1 - \beta$  probability (typically 95% for  $\beta = 5\%$ ).



### b) Calculation method:

- 1) Determine the limit of blank  $LOB(95\%)$  of your experiment by following the method "How to calculate the limit of blank".

- 2) Calculate 
$$p_0 = \frac{2b + z^2 + z \sqrt{z^2 + 4b(1 - b/N)}}{2N(1 + z^2/N)}$$

where:

- $b = LOB(95\%)$  is the 95% limit of blank
- $z = 1.645$  is the "one-tail" quantile at 95%
- $N$  is the total number of partitions that are generated on average in a well (typically  $N = 28000$ )

- $p_0$  is the higher-value solution of the following equation (which can be simplified as a second degree equation in  $p$ ):

$$p = b/N + z \sqrt{p(1-p)/N}$$

- 3) The LOD with a 95% confidence level is determined as follows (calculation based on the Normal Law approximation and the Poisson Law):

- In concentration in the well (cp / uL):

$$LOD(95\%) = -N \ln(1 - p_0)/V$$

$$LOD(95\%) = -\ln(1 - p_0)/v$$

where  $v$  is the average partition volume in  $\mu L$  (for example  $v = 0,00058592 \mu L$  in a Sapphire chip), so that  $V = N v$  is the analyzed volume (i.e. the cumulative total volume of all the partitions generated in a well).

- In number of copies included in the volume analyzed in the well:

$$LOD_{cp}(95\%) = [-N \ln(1 - p_0)]$$

knowing that this unit in number of copies can be approximated as “number of positive partitions” in the well.

- For example: if  $LOB(95\%) = 2$  and  $N = 28000$ , then  $LOD(95\%) = 0.37$  cp / uL and  $LOD_{cp}(95\%) = 7$  copies (or "positive partitions")
- Special case: if  $LOB(95\%) = 0$  (i.e. false positives are never present) then  $LOD(95\%) = 3/V = 0.18$  cp / uL and  $LOD_{cp}(95\%) = 3$  copies.  
In fact, if there are never false positives, then the 95% detection limit is equal to the 95% sampling limit which is:  $LOS(95\%) = 3/V$  cp / uL.
- **Definition of the sampling limit**  $LOS$  at a confidence level  $(1 - \alpha) = 95\%$ : the lowest concentration in the well for which the probability of having no target sequence in the analyzed sub-sample is less than  $\alpha = 5\%$ .