


Recording beads to obtain an experimental PSF

Background

Beads as PSFs

To measure the **Point Spread Function** (PSF) of a microscope, one or more **3D images** of **Sub Resolution** beads are needed. For a typical confocal system that means that the bead *should* have a diameter below 50 nm. So far we have not seen images from such small beads which had a sufficient **Signal To Noise Ratio** to extract a PSF directly, nor a sufficient signal to accurately determine their position so as to average them.

Quantum dots  have been considered for this purpose, but good preparations of these are hard to obtain as they easily form aggregates, thus not being "single points" anymore.

If your beads are not point-like they **can not be directly used as PSFs**. In current **Fluorescence Microscopes** this happens with almost any bead, unless you are using quantum dots. They are **too large**, and if you use them as PSFs in **DeConvolution** you will wash out from your data any feature that is of a size comparable to the bead image, or smaller. Moreover, the remaining data will suffer from **Over Restoration**.

A single bead image will also have a lot of **Photon Noise**, thus an averaging procedure using many beads is in any case very much recommended.

Method

*One solution is to use large beads with a diameter from 100 nm to 200 nm, and to correct for the bead size. The correction is done with a inverse deconvolution run. Given a model of the bead shape, the PSF is computed 'distilled' which its convolution with the bead model matches the measured bead image. (See **Huygens Deconvolution** for more details).*

In case single bead images have insufficient signal for the PSF extraction, the individual bead images can be accurately aligned and averaged to enhance the signal and increase the **Signal To Noise Ratio**. All this is automatically done with the **Huygens Essential Psf Distiller**.

Very important

- Beads should be recorded with **the same Microscopic Parameters** that you will use later to image your specimens, and at least at the **Ideal Sampling** (see **PSF zoom**).
- Record **3D stacks**, not only a single 2D plane. Do not to restrict the number of z-sections to the exact size of the beads, but make a good recording also a little away from them (about 10 extra planes above and below).
- Do not use **too large beads**, they lack sufficient high spatial frequency components. See **Sub Resolution**.
- Do not use **too dense** preparations: beads should be far away from each other to properly probe the PSF. (See images below).

(OK, but those points are very difficult to achieve!!! Inside what ranges can I move to still trust my microscope calibration? See → [Parameter Variation](#)).

Procedure outline

Briefly, the procedure has the following sequence:

1. Prepare a specimen containing fluorescent latex beads.
2. Record a number of images at the microscope settings you also use when recording the images you want to restore.
3. Align and average the individual bead images from single or multiple files.
4. Reconstruct the PSF from the averaged bead image.
5. Store the resulting PSF in ICSfileFormat for later usage.

This post describes steps 1 and 2 of the procedure. To see steps 3-5 using the [Huygens Software](#), go to [Psf From Beads](#).

Practical beads

- [Beads For Widefield Psf](#)
- [Bright Beads For Confocal Psf](#)

Multicolor beads are very useful to calibrate also the → [Color Shift](#).

The beads are appropriately diluted, absorbed to a glass slide, embedded in your usual embedding medium and covered with a cover slip.

When prepared properly, you should hardly see any [Bleaching Effects](#) during the recording of an image. This allows you to use a high averaging count during image acquisition.

Sample preparation

(From an Invitrogen [Product Information Sheet](#) )

Experimental protocols depend somewhat on the instrument and software used; please refer to the materials applicable to your particular instrument. The following serves as a guideline for mounting Tetraspeck microspheres on microscope slides.

1. Use clean glass microscope slides, i.e. oil and dust-free. Special cleaning is usually not required.
2. If desired, the beads in suspension can be diluted with distilled water before use. Before sampling, be sure that the beads are uniformly suspended by mixing on a vortex mixer or by sonicating.
3. Apply 5 μ l of the Tetraspeck bead suspension to the surface of a slide and spread with the pipette tip. Wait for the droplet to dry and then apply \sim 5 μ l of glycerol or other mounting medium, such as water or immersion oil over the dry sample of beads. Some immersion oils may gradually extract dye from the microspheres, resulting in diminished bead fluorescence and

increased background fluorescence. Consequently, the durability of slides prepared using oil may be limited.

4. Cover the sample with a coverslip. Seal the coverslip with nail polish, quick-drying glue or melted paraffin.

N.B.: as the beads should be measured in the same conditions you will use later for your samples, the same mounting medium should be used as well!!!

Record images

To record bead images, follow these steps:

1. Set the recording parameters to the **same values** you will use for recording the object you want to restore. **Take care that you do not undersample the image.** Recommended **Sampling Distances** are about 50 nm lateral sample interval and 100 nm axial sampling interval for confocals, the double for widefields. (But see *Imaged volume* below for details!!!).
 2. Average the image as much as possible, for instance 16 or 32 times. **Avoid conditions where the intensities of the image before averaging are clipped.** **Clipped Images** are obtained when the **Dynamic Range** of the input signal exceeds that of the analog to digital converter (ADC). In practice this means that the lowest values in the input image are actually negative, and/or that the highest values should in fact be represented by a number greater than 255. Negative values are usually converted to 0; values above 255 to 255. (Some microscopes have ADCs with a larger range than 0-255).
- Confocal data: Record a number of images of lateral size in the order of 512x512 and 40-50 sections. There should be 1-10 beads in each image. If the beads are too close to each other or to an edge they will be rejected at a later stage. In addition you can cut away empty areas in an image later on, so it is not necessary to search for images with say 5 equally spaced beads. In total you need between 5 to 10 usable beads.
 - Widefield data: You should be able to obtain a strong signal from a single bead with not too much bleaching. Make sure that it is indeed a single bead and not a cluster. Record the bead in a 512x512 image. With an objective with **Numerical Aperture** NA = 1.3 50-100 sections at 200 nm distance should be sufficient, to cover at least 12 μm along Z. Although you have just a single bead, you still need to apply the Average bead tool to centre the bead and to remove the **BackGround**.

Imaged volume

If the number of planes recorded along Z is too low, you will get an error when distilling the PSF: "the axial image size is too small". It can also happen that this error is not displayed but no usable bead is found in your image.

What matters for a correct PSF distillation is not (only) the number of slices, but the total physical volume that is actually imaged. A good bead image includes information of the cone of blur around it, that specially in **Wide Field Microscopes** and in low **Numerical Aperture** (NA) cases can be very large. For example, when using a 0.95 NA objective in a widefield microscope the PSF is very large along Z, and a larger volume must be imaged in this direction than in the NA = 1.3 example above to register all the relevant information. You could indeed acquire more planes, but you can also combine that with a reduction of the **Sampling Density** along Z. The ideal sampling rate (that depends on the NA) for normal imaging can be found using the **Nyquist Calculator**, but when recording beads some **OverSampling** is recommended.

If you ask the calculator to show also a PSF, you can find out how large it is expected to be, both in real volume and in number of samples when imaged at the **Nyquist Rate**. (Experimentally you can record some less planes along Z than that is shown there, because it is not recommended to acquire

very low intensity regions much affected by noise. But that theoretical PSF is a good guide to see how much volume is necessary to distill a PSF).

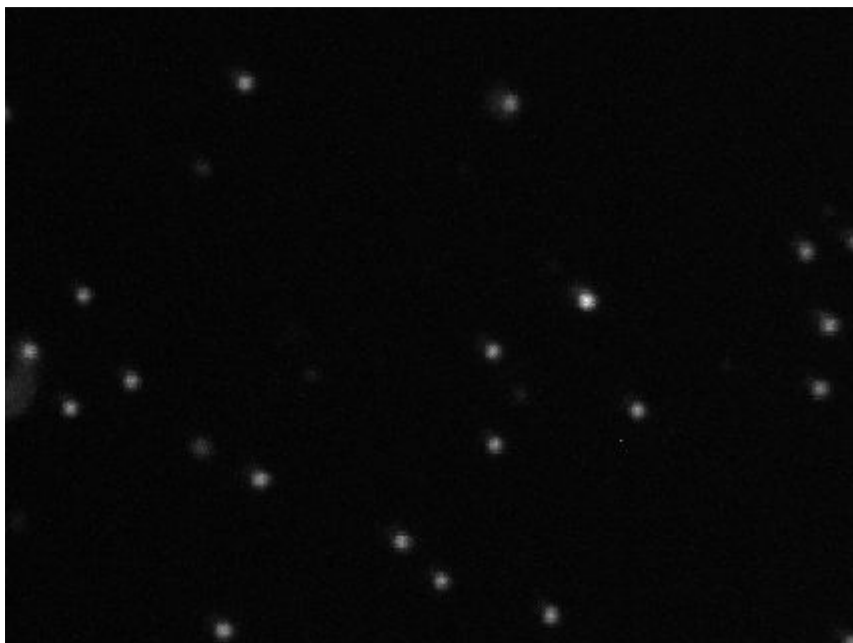
For a 0.95 NA typical widefield microscope the beads should be recorded in a 3D image that covers at least $\sim 21 \mu\text{m}$ along the optical axis. The **Nyquist Rate** along Z for these conditions is one plane every $\sim 700 \text{ nm}$. If we record the beads with some **OverSampling** (recommended), at one plane every 500 nm, we need at least 42 planes to cover that volume. At that rate, despite we recorded less planes than in the $\text{NA} = 1.3$ example above, we have covered much more 3D volume!!!

Bead density

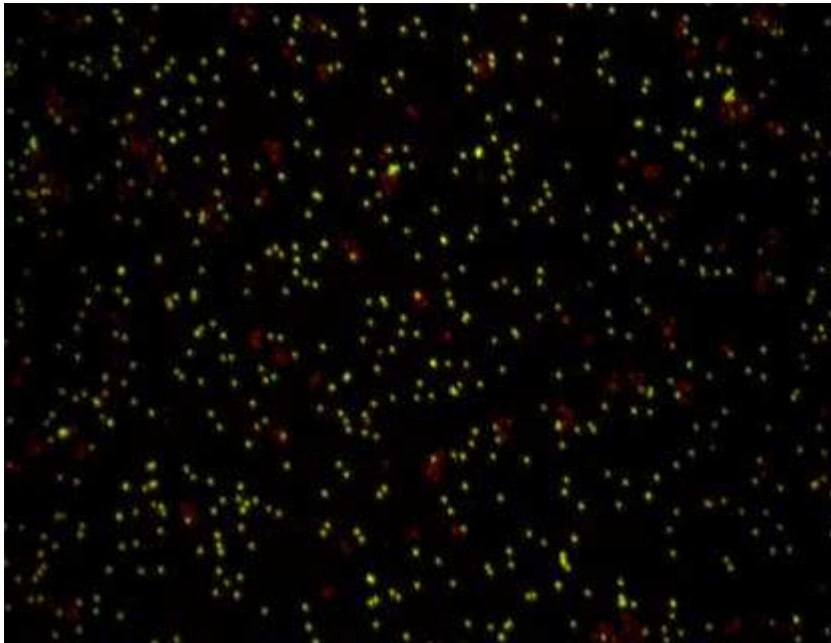
If the number of beads in the image is of an order much larger than 10, chances are that they are too close to each other. (Therefore their blurred images overlap, and it is no longer possible to disentangle their PSF's). The **Psf Distiller** will reject beads that are too close to each other. You can make its criterion less strict by using the "reduce PSF size" parameter, but there is a limit for this to work properly.

Examples

This could be an example of a bead image, is a top view **Fast Mip**. It is quite OK, but some less recorded beads would be even better.



On the other hand, the following image has such a high bead density that is useless for a PSF distillation:



The procedure to obtain a PSF from the recorded beads using the [Huygens Software](#) continues in [Psf From Beads](#).

More information:

- [Deconvolving Beads](#)
- [Theo Vs Exp Psf](#)

Search the FAQ for "bead".

The original document is available at <http://www.svi.nl/RecordingBeads>