## Single molecule image processing

## **Track Features:**

- 1. linfit slope: slope of the linear fit of the log intensity profile over normalized time
- 2. linfit\_amplitude: background corrected exp of the constant of the linear fit of the log intensity profile over normalized time
- 3. amplitude\_mean: mean amplitude of each track (normalized by the median amplitude of all tracks on the image stack)
- 4. amplitude\_mean2: mean amplitude of the first half of the track over the sum of the mean of amplitude for the second half and the first half;
- 5. charact\_exp: characteristic parameter in the exponential fit of the (background corrected) track amplitude over time
- 6. (iniamp\_exp: initial intensity (at t=0) from the exponential fit of the (background corrected) track amplitude over time)
- 7. amplitude\_std: std of amplitude for the first half of the track over the sum of the std of amplitude for the second half and the first half.
- 8. position std: std of the x-, y-position over time
- 9. psf\_std: linear slope of the std of the position over time (i.e. spot diameter increases for out of focus particles)
- 10. pos\_std: std of the position;
- 11. pos\_std2: std of the position for the first half of the track over the sum of the std of position for the second half and the first half;