## The two-sample Kolmogorov-Smirnov Test

We used a two-sample Kolmogorov-Smirnov test to quantitatively compare the virtual test data set  $X_i$  with simulated probe data-sets  $Y_j$ . The principle of the test can be found in text books on statistics (e.g. (1, 2)), and shall be briefly sketched here.

Let us assume two samples  $X_i$  and  $Y_j$  with size n and m drawn from continuous distributions. The two samples can be characterized by their empirical cumulative density functions

$$cdf_X(x) = \frac{\#i: X_i \le x}{n}$$
 and  $cdf_Y(x) = \frac{\#i: Y_i \le x}{m}$ 

which are defined as the proportion of observed values that are less than or equal to x. The test is based on finding the maximum distance  $D = \max(|cdf_x(x) - cdf_y(x)|)$ . Suppose that both X<sub>i</sub> and Y<sub>j</sub> are drawn from the same distribution, and their values are such that D=d. Since a large value of D would appear to be inconsistent with the null hypothesis that both samples are drawn from the same distribution, it follows that the P-value for this data-set is given by

$$P-value = prob(D \ge d)$$

Note that prob specifies the probability for  $D \ge d$  under the assumption that  $H_0$  is correct. The key argument for the versatility of the Kolmogorov-Smirnov test is the proposition that  $prob(D \ge d)$  is the same for any continuous distribution. The distribution of D as function of n and m can be found in tables (2) and is implemented for calculation of the P-value in the Matlab function kstest2 (The MathWorks, Natrick, MA).



1. Ross, S. M. Introduction to probability and statistics for engineers and scientists, 3<sup>rd</sup> edition, Elsevier, San Diego

2. Conover, W. J.. Practical nonparametric statistics, 3<sup>rd</sup> edition, Wiley, New York

## **Supplemental Figure 1**



**Supplemental Figure 1.** Escape-probability  $\eta$  as a function of  $\hat{\tau}$ . Data were obtained from Monte Carlo simulations of hop diffusion, by determining  $\hat{\tau}$  for various  $\eta$ .



![](_page_2_Figure_1.jpeg)

**Supplemental Figure 2:** Correction of the P-value distribution. We simulated free diffusion according to Fig. 3 and calculated the minimum  $P_{min}$  of the P-values obtained for n=1, n=2 and n=3. The distribution is clearly non-uniform (**A**). Using the transformation  $P_{corr} = 1 - (1 - P_{min})^{n_{max}}$ , a uniform distribution of  $P_{corr}$  is generated, which allows for interpretation of  $P_{corr}$  as P-value (**B**).

## **Supplemental Figure 3**

![](_page_3_Figure_1.jpeg)

**Supplemental Figure 3:** P-values as function of K and  $\tau_{off}$  for various ratios  $D_A/D_{AB}$ . Tests were performed on the data-sets described in Fig. 12C. The panels show results for  $D_A/D_{AB}=2$  (A), 3 (B), 4 (C) and 10 (D). Interestingly, already a ratio of three is sufficient to significantly restrict the parameter range for  $\tau_{off}$ . Data obtained for a ratio of 2, however, contain hardly any information on the interaction lifetime.