

Supporting Materials and Methods

Analysis of time profiles based on the multistep hopping mechanism was performed with numerical analysis by using MATLAB software. A kinetic model of the multistep hole-transfer process is shown in Scheme 1.

Charge recombination process (k_{cr}) can be ignored because the charge-separated state persisted over several hundred microseconds when naphthalimide (NI) and the nearest G are separated by six A bases. According to Scheme 1, simultaneous differential equations are shown as follows:

$$\begin{aligned}\frac{d[G_1]}{dt} &= -k[G_1] + k[G_2] \\ \frac{d[G_2]}{dt} &= k[G_1] - 2k[G_2] + k[G_3] \\ &\vdots \\ \frac{d[G_n]}{dt} &= k[G_{n-1}] - (k + k_1)[G_n] \\ \frac{d[PTZ]}{dt} &= k_1[G_n]\end{aligned}\quad [2]$$

where $[G_i (i = 1 \dots n)]$ corresponds to the hole population at each G-site, k is the hole-transfer rate constant between Gs, and k_1 is hole transfer from G_n^{*+} to PTZ. Fitting results for 5'-(GA) $_n$ and (GT) $_n$ ($n = 12$) according to Eq. 2 are presented in Fig. 8, providing the rate constants of $k = 4 \times 10^7 \text{ s}^{-1}$ and $k_1 = 6 \times 10^7 \text{ s}^{-1}$ for (GA) $_{12}$ and $k = 6.5 \times 10^5 \text{ s}^{-1}$ and $k_1 = 5 \times 10^6 \text{ s}^{-1}$ for (GT) $_{12}$, respectively. The obtained values of k for (GA) $_{12}$ and (GT) $_{12}$ were similar with the values obtained for (GT) $_2$ and (GA) $_2$ with single-exponential analysis according to Eq. 1 in the main text, showing the validity of approximate analysis performed with single-exponential fitting. Difference in the k_1 values between (GA) $_{12}$ and (GT) $_{12}$ is attributed to the difference in the nucleobase that is stacked to PTZ.