

Electronic Supporting Information

A two-enzyme cascade reaction consisting of two reaction pathways. Studies in bulk solution for understanding the performance of a flow-through device with immobilised enzymes.

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1. Acid-base equilibria of DCFH₂, DCFH[•], and DCF

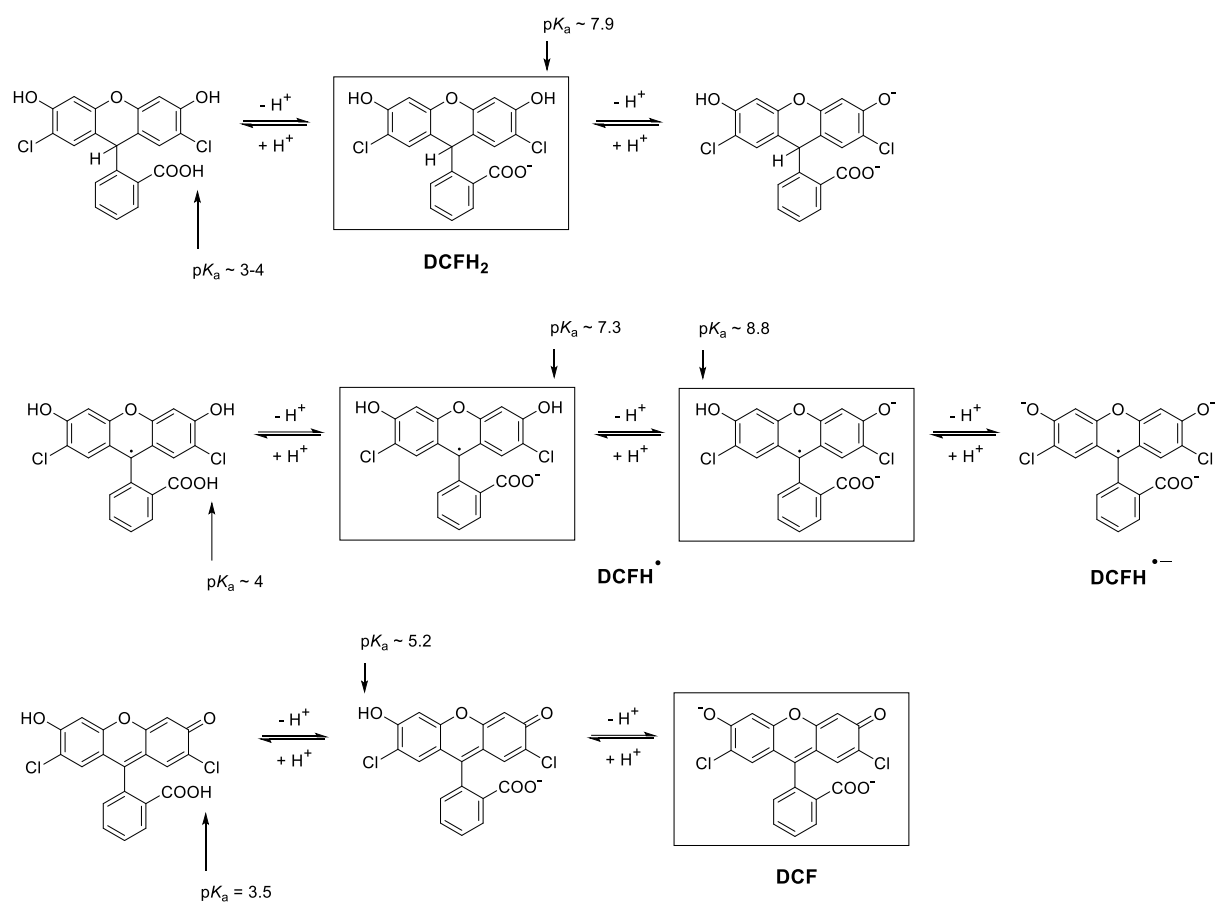
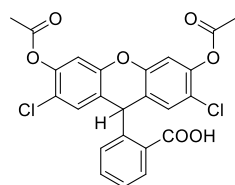
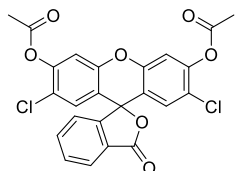


Figure S-1. Some of the relevant acid-base equilibria for DCFH₂, DCFH[•], and DCF. For pH = 7.2 conditions, the most abundant species are marked with a frame. Relevant pK_a values are indicated, as determined by (Wrona and Warman, 2006).^{S1}

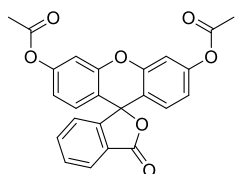
2. Chemical structures of different fluorescein derivatives



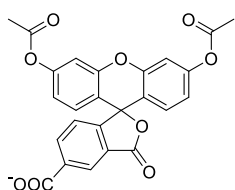
DCFH₂-DA
2',7'-dichlorodihydrofluorescein diacetate



DCF-DA
2'-7'-dichlorofluorescein diacetate



fluorescein diacetate



6-carboxyfluorescein diacetate

Figure S-2. Comparison of the chemical structures of DCFH₂-DA (2',7'-dichlorodihydrofluorescein diacetate), 2'-7'-dichlorofluorescein diacetate, fluorescein diacetate, and 6-carboxyfluorescein diacetate.

3. Absorption spectra of DCF

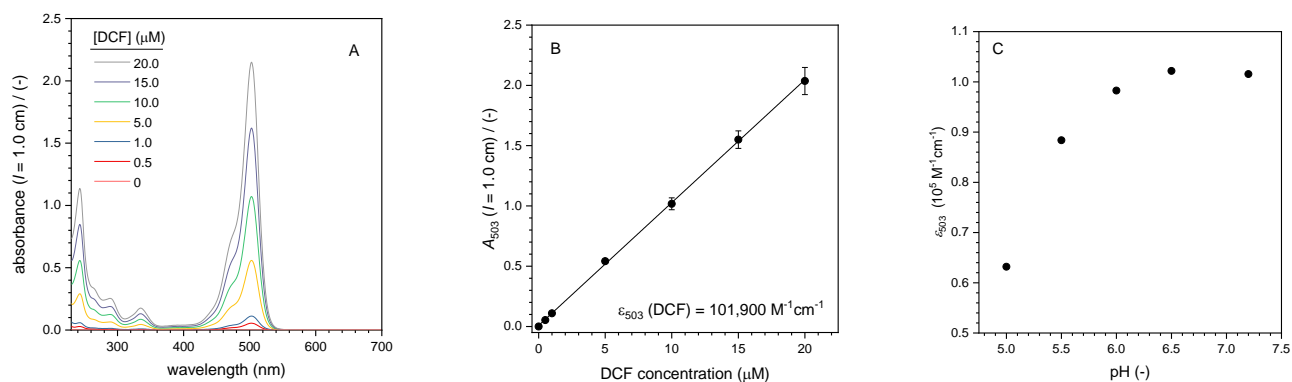


Figure S-3. Concentration dependence of the absorption spectrum of DCF dissolved in PB (10 mM sodium phosphate buffer solution, pH = 7.2). **(A)** Spectra for [DCF] = 0, 0.5, 1.0, 5.0, 10.0, 15.0, and 20.0 μM , path length $l = 1.0 \text{ cm}$, $T = 25 \text{ }^\circ\text{C}$. **(B)** A_{503} vs. [DCF], plotted as mean values \pm standard deviations ($n = 3$, separately prepared solutions from three different DCF stock solutions). The molar absorption coefficient for $\lambda = 503 \text{ nm}$ was determined, $\epsilon_{503}(\text{DCF}) = 1.02 \cdot 10^5 \text{ M}^{-1}\text{cm}^{-1}$. **(C)** Dependence of the molar absorption coefficient at $\lambda = 503 \text{ nm}$, $\epsilon_{503}(\text{DCF})$, on pH.

4. Details about the preparation of *de*-PG2₁₀₀₀-BAH-enzyme conjugates

4.1. Conjugates *de*-PG2₁₀₀₀-BAH₁₅₂-BCA₁₀₁ and *de*-PG2₁₀₀₀-BAH₂₀₇-BCA₁₅₂

The denpol-enzyme conjugates *de*-PG2₁₀₀₀-BAH₁₅₂-BCA₁₀₁ and *de*-PG2₁₀₀₀-BAH₂₀₇-BCA₁₅₂ were obtained in a similar way as described before for the preparation of *de*-PG2₁₀₀₀-BAH₁₇₅-BCA₁₁₅, see (Yoshimoto et al., 2018).^{S2} Besides, there were small experimental variations (slightly higher amounts of BAH formed during the conjugation reaction as compared to the amounts of added 4FB; [4FB] being determined by spectral fitting). All materials used were the same as before,^{S2} except *de*-PG₁₀₀₀; it was the same as the one used in the work of (Hou et al., 2019).^{S3} For the spectral changes during conjugate formation, see **Figure S-4**. For comparison, some of the characteristic data for the conjugate formation are provided:

de-PG2₁₀₀₀-BAH₁₇₅-BCA₁₁₅ (Yoshimoto et al., 2018):^{S2}

- MSR (4FB/BCA) = 1.02
- *de*-PG2₁₀₀₀-HyNic₄₄₀
- used for conjugation reaction: [4FB] = 67 μM, [HyNic] = 120 μM, pH = 7.2
- observed in conjugation reaction: [BAH] = 48 μM
- [BCA]/[BAH] in conjugate = 0.66 (66 %)

de-PG2₁₀₀₀-BAH₁₅₂-BCA₁₀₁ (this work, used for measurement with *p*-NA in a BCA loaded glass fiber filter only):

- MSR (4FB/BCA) = 0.99
- *de*-PG2₁₀₀₀-HyNic₄₃₆
- used for conjugation reaction: [4FB] = 37 μM, [HyNic] = 120 μM, pH = 7.2
- observed in conjugation reaction: [BAH] = 42 μM
- [BCA]/[BAH] in conjugate = 0.66 (66 %)

de-PG2₁₀₀₀-BAH₂₀₇-BCA₁₅₂ (this work, used for the cascade reaction in the glass fiber filter flow-through device):

- MSR (4FB/BCA) = 0.84
- *de*-PG2₁₀₀₀-HyNic₃₆₂
- used for conjugation reaction: [4FB] = 60 μM, [HyNic] = 120 μM, pH = 7.2
- observed in conjugation reaction: [BAH] = 68 μM
- [BCA]/[BAH] in conjugate = 0.73 (73 %)

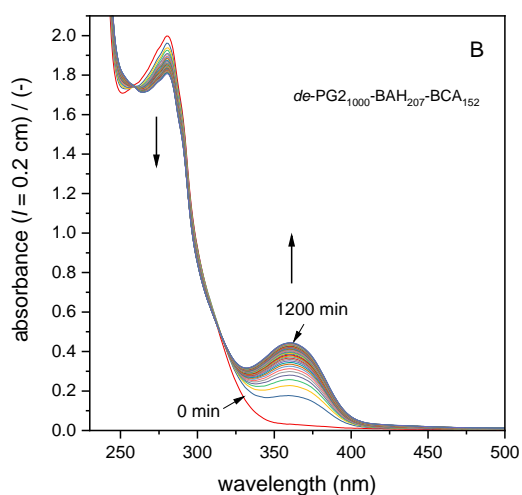
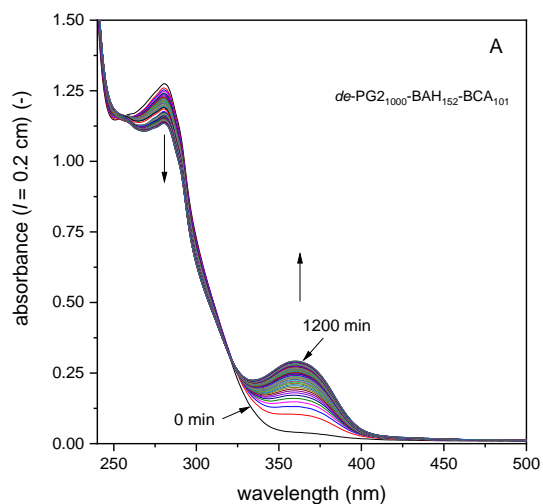


Figure S-4. Conjugation reaction between **(A)** *de*-PG2₁₀₀₀-HyNic₄₃₆ and BCA-4FB (MSR(4FB/BCA) = 0.99), and **(B)** *de*-PG2₁₀₀₀-HyNic₃₆₂ and BCA-4FB (MSR(4FB/BCA) = 0.84). The absorption spectrum was recorded every 30 min during a total reaction time of 20 h (reaction between HyNic and 4FB to form stable BAH (bis-aryl hydrazone) bonds between the denpol and BCA). The absorption band of the BAH bond formation (centered around $\lambda = 354$ nm) as well as a decrease in band intensity around $\lambda = 280$ nm (HyNic and 4FB) are visible and indicate the ongoing conjugation reaction.

From $A_{354}(20 \text{ h}) - A_{354}(0 \text{ h})$, the concentration of BAH formed could be calculated. With the known amounts of added denpol r.u. and BCA (as well the free, unbound BCA recovered from the reaction mixture during purification), the ratio of BAH to BCA in the conjugate could be estimated to *de*-PG2₁₀₀₀-BAH₁₅₂-BCA₁₀₁ **(A)** and *de*-PG2₁₀₀₀-BAH₂₀₇-BCA₁₅₂ **(B)**.

4.2. Conjugate *de*-PG2₁₀₀₀-BAH₈₆-HRP₉₀

The denpol-enzyme conjugate *de*-PG2₁₀₀₀-BAH₈₆-HRP₉₀ was obtained in a similar way as described before for the preparation of *de*-PG2₁₀₀₀-BAH-HRP₇₁, see (Hou et al., 2019).^{S3} This time, the pH value of the conjugation reaction buffer used was a bit lower: pH = 4.4 instead of 4.7, noted as “*MesB1*” in (Hou et al., 2019).^{S3} This deviation did not lead to significant changes in the conjugation reaction. The filtrates obtained during purification of the conjugate solution by ultrafiltration were quantified for HRP content by measuring the heme group absorption at A_{403} . Assuming that non-conjugated HRP could pass the ultrafiltration membrane quantitatively (100 kDa MWCO), the total amount of HRP in the ultrafiltrate was determined and subtracted from the total amount of HRP used for the conjugation reaction. In this way, the amount of conjugated HRP was determined and compared with the amount of BAH bonds formed (subscripts of HRP and BAH, respectively). With this, the previously made assumption that for each BAH bond formed about one HRP molecule is conjugated, see (Hou et al., 2019),^{S3} could be confirmed (86 BAH and 90 HRP per 1000 r.u.). The higher amount of conjugated HRP (and BAH bonds) per 1000 r.u. – around 90 per average denpol chain in this work, around 70 previously^{S3} – can be explained by the higher content of HyNic in the denpol used in this work (*de*-PG2₁₀₀₀-HyNic₃₁₀) as compared to the one in our previous work (*de*-PG2₁₀₀₀-HyNic₂₄₀).^{S3} In both denpol-HRP conjugate reactions, the same concentrations of HyNic and 4FB were used, leading to similar extent of BAH bond formation, while the concentration of r.u. during the conjugation reaction was lower in the work presented, as compared to the previous preparation. All materials used were the same as in (Hou et al., 2019).^{S3} For the spectral changes during conjugate formation, see **Figure S-5**. For comparison, some of the characteristic data for the conjugate formation are provided:

de-PG2₁₀₀₀-BAH-HRP₇₁ (Hou et al., 2019).^{S3}

- MSR (4FB/HRP) = 0.73
- *de*-PG2₁₀₀₀-HyNic₂₄₀
- used for conjugation reaction: [4FB] = 50 μ M, [HyNic] = 100 μ M, pH = 4.7
- observed in conjugation reaction: [BAH] = 29.5 μ M
- [HRP]/[BAH] in conjugate assumed to be 1:1 as MSR(4FB/HRP) < 1.

de-PG2₁₀₀₀-BAH₈₆-HRP₉₀ (this work):

- MSR (4FB/HRP) = 0.62
- *de*-PG2₁₀₀₀-HyNic₃₁₀
- used for conjugation reaction: [4FB] = 50 μ M, [HyNic] = 100 μ M, pH = 4.4
- observed in conjugation reaction: [BAH] = 27.9 μ M
- [HRP]/[BAH] in conjugate = 1.05 (105 %) (slight excess, experimentally determined)

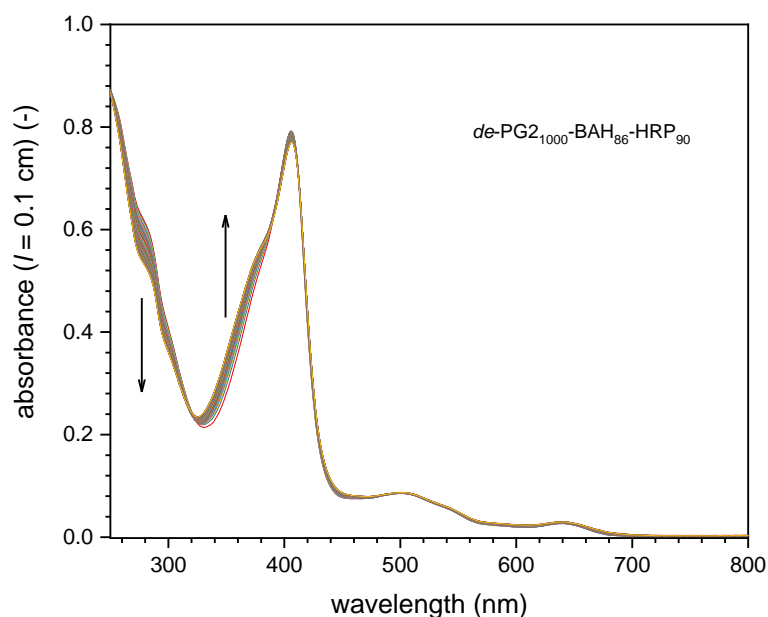


Figure S-5. Conjugation reaction between *de*-PG2₁₀₀₀-HyNic₃₁₀ and HRP-4FB (MSR(4FB/HRP) = 0.62). The absorption spectrum was recorded every 30 min during a total reaction time of 20 h (reaction between HyNic and 4FB to form stable BAH (bis-aryl hydrazone) bonds between the denpol and HRP). The absorption band of the BAH bond formation (centered around $\lambda = 354$ nm) as well as a decrease in band intensity around $\lambda = 280$ nm (HyNic and 4FB) are visible and indicate the ongoing conjugation reaction. The strong band at $\lambda = 403$ nm is due to the heme group of HRP.

From $A_{354}(20 \text{ h}) - A_{354}(0 \text{ h})$, the concentration of BAH formed could be calculated. With the known amounts of added denpol r.u. and HRP (as well the free, unbound HRP recovered from the reaction mixture during purification), the ratio of BAH to HRP in the conjugate could be estimated to *de*-PG2₁₀₀₀-BAH₈₆-HRP₉₀.

5. Details about deviations from the general procedure applied for the immobilisation of denpol-enzyme conjugates in glass fiber filters

Compared to the general procedure described in section 2.5. for the immobilisation of denpol-enzyme conjugates in glass fiber filters, there were a few variations, as indicated below.

Deviations in experiment ① (see **Figure 11**): One of the two independent experiments using sequentially- and co-immobilized denpol-BCA and denpol-HRP loaded filters for activity measurements with DCFH₂-DA and H₂O₂ (see **Figure 12**, triangles, “exp. 1”). Instead of keeping the incubation buffer fixed for sequential and co-immobilization (as used for “exp. 2”, 80% PB, 20% of 10mM phosphate, pH 7.0, 0.15 M NaCl, see 2.5.2 and 2.5.3), for the sequential immobilization each denpol-enzyme was incubated with the same buffer as for the individual incubations of one filter per holder (no NaCl in the BCA-denpol incubation buffer, see 2.5.1). In case of the co-immobilisation, a buffer ratio was used as close to PB as the denpol-HRP stock solution (in other buffer) allowed (finally 88.5% PB, 11.5% of 10mM phosphate, pH 7.0, 0.15 M NaCl). By that slight variation within “exp. 1” and “exp. 2” it could be checked if the NaCl contained in the buffer used for immobilization of BCA-conjugate in this work (in contrast to the NaCl-less incubation buffer, as introduced in (Yoshimoto et al. 2018))^{S2} had any influence on the cascade experiment. From the results obtained (**Figure 12** and **Figure S-21**, comparison of **Figure 13** and **Figure S-22**), no significant change upon buffer modification was visible.

Deviations in experiment ② (see **Figure 11**): single denpol-BCA loaded filter for activity measurements with *p*-NA (see **Figure 9**). The BCA-conjugate stock solution used was *de*-PG2₁₀₀₀-BAH₁₅₂-BCA₁₀₁ (instead of *de*-PG2₁₀₀₀-BAH₂₀₇-BCA₁₅₂) and was diluted to [BAH] = 3.3 μM and [BCA] = 2.2 μM (instead of [BAH] = 12 μM and [BCA] = 8.8 μM) for the incubation. For the washing step, the filter containing the conjugate was immersed three times in 1.5 mL fresh PB for 15 min (instead of washing by pumping PB through the filter placed in the filter holder).

Deviations in experiment ③ (see **Figure 11**): single denpol-HRP loaded filters for activity measurements with ABTS²⁻/H₂O₂ (see **Figure 10**). The filters were loaded with 50 or 250 μL of the conjugate solution. In the case of loading with 50 μL, after incubation for 1 h, the filter was washed for 5 h (instead of 3 h). For the loading with 250 μL, after incubation for 1 h, the filter was first washed by 8 times immersion in 1.5 mL fresh phosphate buffer solution (100 mM sodium phosphate, 1.15 M NaCl, pH = 7.2) and storage for 15 min. Afterwards, the filter was placed in the filter holder and washed by pumping the phosphate buffer solution through the filter for 3 h.

6. Details about the visualisation of the denpol-BAH-enzyme conjugates prepared

In an attempt to visualise the space occupied by enzymes bound onto the surface of the polymer (*de*-PG2), we represent the situation using a solid cylinder for a fraction carrying 100 r.u. of *de*-PG2, and identical, non-overlapping spheres for the proteins a variable distance (Δ) between 0 and 1 nm away from the polymer surface (see **Figure 1**, right). The conformation of 100 r.u. of PG2 can be considered cylindrical, as the contour length of its backbone does not exceed the persistence length of PG2, $\lambda^* \approx 25$ nm,⁵⁴ and because the radial density profile of the polymer is known to exhibit a plateau at small distances before it decays beyond approximately 1.5 nm, see Figure 9 in (Bertran et al., 2013).⁵⁵ The corresponding effective PG2 radius is in rough agreement with an estimate of 2.5 nm.⁵⁴ We use this radius for the visualisation, assuming the deprotection of PG2 (thus *de*-PG2, without the Boc protection group from PG2) did not significantly change it. The volume of the spheres we choose to match the volume of the protein (BCA or HRP) using the empirical expression from Fischer et al., 2004)⁵⁶ for the effective mass density ρ_p of proteins, $\rho_p \approx [1.41 + 0.145 \exp(-M_p/13)]$ g/cm³, where M_p is the molecular mass of a protein in units of kDa. Using $M_p = 29$ for BCA (RCSB PDB: 1V9E) and $M_p = 44$ for HRP (RCSB PDB: 1HCH) the resulting sphere radii are $R_p = 2.0$ nm (BCA) and $R_p = 2.3$ nm (HRP). The distance Δ takes into account the linkers located between polymer and enzyme. Each BAH linker is modelled as a flexible trimer with two bonds of fixed length 0.5 nm, where the first bond is taken normal to the *de*-PG2 surface, and the second bond chosen randomly. The mean amount of bound enzymes per 100 r.u. is provided by our experimental values for 1000 r.u. *de*-PG2. The centers of the spheres are determined using a random number generator, that specifies a location on the surface of a cylinder with radius 2.5 nm + Δ + R_p and length λ^* . Attempted locations that lead to overlap between existing spheres are rejected, while non-overlapping spheres are accepted. Every visualised configuration is thus different (if repeated with same input values), but each of them is a representant of the ensemble.

7. UV/vis absorption spectra of BCA, HRP, DCFH₂-DA, DCFH₂-MA, DCFH₂, DCF-MA, DCF, and DCF_{ox}

Table S-1. Molar absorption coefficients ϵ_{λ} ($M^{-1}cm^{-1}$) for BCA (measured),^a HRP (measured),^b H₂O₂ (measured), DCFH₂-DA (measured), DCFH₂-MA (fitted), DCFH₂ (measured), DCF-MA (fitted), DCF (measured), and DCF_{ox} (measured). PB (10 mM sodium phosphate, pH = 7.2), $T = 25$ °C. Wavelength range: $\lambda = 240 - 600$ nm.

^a: ϵ_{280} (BCA) set to $5.6 \cdot 10^4$ ($M^{-1}cm^{-1}$);^{S7} and ^b: ϵ_{403} (HRP) set to $1.02 \cdot 10^5$ ($M^{-1}cm^{-1}$).^{S8}

Please note that negative values for absorption coefficients given in the table have no physical meaning. They are mainly the result from measurement background/noise or spectral fitting where they reflect the background/noise of the actual measurements from which the spectra were fitted. The data in the table are the ones used for the actual quantitative analysis of the reaction mixtures (including the negligible, small negative values).

Wavelength (nm)	BCA (stock solution)	HRP (stock solution)	H ₂ O ₂ (stock solution)	DCFH ₂ -DA (stock solution)	DCFH ₂ -MA (fitted)	DCFH ₂ (prepared <i>in situ</i>)	DCF-MA (fitted)	DCF (stock solution)	DCF-fully oxidized (prepared <i>in situ</i>)
600	476	2340	0	-16	-17	-19	-96	-176	-81
599	463	2350	0	-17	-18	-17	-88	-173	-84
598	451	2330	0	-19	-16	-17	-110	-159	-85
597	460	2340	0	-17	-13	-13	-76	-145	-65
596	474	2390	0	-18	-36	-18	-56	-137	-75
595	472	2410	0	-16	-21	-14	-61	-142	-65
594	473	2410	0	-16	-23	-14	-74	-154	-53
593	465	2440	0	-19	-17	-15	-115	-155	-57
592	477	2460	0	-21	-19	-19	-115	-155	-48
591	490	2490	0	-18	-23	-15	-133	-157	-52
590	477	2530	0	-15	-22	-13	-140	-165	-47
589	462	2500	0	-15	-22	-13	-107	-163	-31
588	462	2530	0	-14	-24	-16	-107	-166	-37
587	458	2530	0	-14	-10	-14	-85	-174	-26
586	476	2560	0	-16	-8	-12	-54	-150	-16
585	473	2600	0	-15	-14	-13	-63	-148	-28
584	461	2640	0	-15	-13	-13	-80	-166	-16
583	474	2670	0	-16	-9	-12	-73	-150	-10
582	478	2690	0	-19	-23	-13	-98	-153	11
581	471	2720	0	-17	-14	-14	-103	-158	9
580	453	2740	0	-15	-23	-17	-110	-152	12
579	469	2790	0	-16	-24	-13	-110	-160	34
578	467	2830	0	-13	-21	-8	-87	-163	55
577	478	2880	0	-14	-23	-10	-41	-150	63
576	467	2870	0	-16	-18	-11	-27	-147	67
575	484	2960	0	-16	-18	-11	-48	-134	77
574	496	3020	0	-16	-22	-9	-52	-139	96
573	477	3070	0	-15	-19	-11	-81	-147	100
572	477	3120	0	-13	-11	-13	-75	-135	117
571	472	3190	0	-14	-16	-12	-75	-144	140
570	483	3270	0	-16	-22	-12	-83	-133	166
569	490	3320	0	-15	-14	-10	-101	-125	199
568	496	3430	0	-12	-13	-11	-87	-119	206
567	481	3500	0	-13	-21	-11	-60	-114	219
566	498	3660	0	-15	-10	-9	-59	-123	242
565	490	3710	0	-15	-3	-8	-72	-137	278
564	455	3810	0	-13	-11	-11	-93	-139	293
563	475	3920	0	-14	-15	-12	-92	-140	323
562	488	4090	0	-16	-29	-12	-73	-136	348

561	477	4210	0	-18	-19	-11	-49	-109	398
560	458	4310	0	-19	-17	-11	-59	-117	434
559	472	4510	0	-13	-21	-9	-55	-108	480
558	510	4740	0	-11	-7	-8	-60	-96	527
557	503	4860	0	-9	-7	-4	-58	-102	589
556	496	5070	0	-12	-14	-6	-55	-94	637
555	470	5210	0	-10	-18	-5	-66	-90	694
554	471	5390	0	-9	-18	-3	-91	-90	759
553	523	5650	0	-18	-17	-11	-82	-76	823
552	505	5730	0	-15	-28	-13	-67	-47	919
551	493	5910	0	-12	-12	-10	-62	-35	1010
550	476	6090	0	-15	-24	-8	-65	-22	1110
549	508	6300	0	-14	-13	-6	-75	2	1200
548	516	6480	0	-12	-4	-5	-42	28	1310
547	509	6620	0	-11	-13	-5	-37	70	1460
546	512	6810	0	-15	-9	-7	-27	117	1570
545	512	6980	0	-16	-6	-4	17	175	1720
544	488	7110	0	-13	-10	-1	-26	229	1900
543	492	7250	0	-10	-9	0	-55	290	2060
542	513	7420	0	-12	-5	3	-15	401	2250
541	511	7550	0	-13	-1	-1	-53	491	2450
540	514	7650	0	-14	-4	-1	-51	613	2650
539	515	7810	0	-9	-2	3	18	787	2900
538	494	7890	0	-15	-3	-1	-4	972	3130
537	509	8030	0	-13	-2	3	9	1210	3380
536	504	8150	0	-15	-7	2	15	1500	3660
535	504	8230	0	-8	4	12	-13	1840	3980
534	502	8350	0	-5	-7	8	-23	2230	4290
533	524	8510	0	-12	-8	7	-49	2710	4580
532	533	8650	0	-11	5	10	-57	3310	4920
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527	534	9260	0	-12	10	20	-201	8540	6710
526	547	9390	0	-12	24	24	-256	10180	7090
525	544	9560	0	-9	15	26	-320	12090	7480
524	521	9620	0	-9	18	31	-356	14330	7860
523	528	9770	0	-10	19	36	-406	16900	8250
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521	538	10060	0	-10	28	39	-432	23190	8980
520	532	10030	0	-9	42	47	-5	27040	9350
519	538	10190	0	-7	47	54	29	31240	9710
518	535	10330	0	-7	45	59	16	35830	10050
517	529	10510	0	-11	49	63	10	40780	10370
516	553	10640	0	-7	65	71	101	46080	10710
515	569	10770	0	-11	59	75	111	51680	11010
514	557	10900	0	-7	69	84	100	57500	11280
513	540	10950	0	-9	83	93	47	63450	11530
512	565	11070	0	-8	98	100	-73	69410	11770
511	554	11140	0	-8	102	106	-72	75270	12020
510	551	11230	0	-3	110	116	-189	80820	12200
509	564	11380	0	-6	111	116	-183	85970	12340
508	568	11440	0	-7	121	126	-126	90580	12440
507	572	11460	0	-1	123	135	-14	94540	12560
506	579	11510	0	-6	130	135	259	97710	12590
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500	551	11670	0	-6	142	133	3340	98220	12070
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486	575	11260	0	2	81	73	12020	49340	9390
485	603	11200	0	-4	64	65	12350	47050	9230
484	608	11150	0	-2	67	64	12630	45040	9100
483	583	11080	0	-3	65	62	12900	43290	8990
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476	590	10650	0	-2	56	55	14470	36360	8440
475	600	10580	0	-1	61	52	14740	35710	8370
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470	618	10440	0	-5	37	45	16520	32030	8000
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467	626	10410	0	-7	23	41	17860	28860	7720
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465	630	10360	0	-3	26	39	18680	26350	7490
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458	642	10350	0	0	16	28	20560	17320	6600
457	649	10330	0	4	16	30	20630	16200	6480
456	658	10300	0	-1	17	25	20610	15130	6320
455	627	10280	0	-4	10	19	20520	14150	6200
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447	652	10740	0	-2	4	12	18810	8850	5380
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437	705	14750	0	-1	0	13	16990	5380	4750
436	687	15470	0	-3	-2	8	16820	5070	4680
435	702	16280	0	-1	0	12	16680	4790	4630
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433	658	18250	0	-1	-1	10	16410	4260	4480
432	710	19490	0	-1	5	8	16300	4000	4410
431	677	20750	0	-3	6	5	16170	3750	4340
430	693	22230	0	1	14	8	15940	3530	4260
429	713	23920	0	2	16	10	15740	3340	4180
428	727	25730	0	3	-2	12	15440	3150	4110
427	741	27670	0	0	-2	11	15120	2960	4030
426	754	29880	0	0	-3	6	14810	2790	3950
425	742	32190	0	-1	-18	4	14460	2600	3860
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422	752	40560	0	1	-9	9	13570	2280	3660
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420	829	47430	0	5	9	4	12980	2110	3490
419	837	51020	0	7	9	14	12720	2040	3460
418	809	54840	0	6	-13	8	12480	1970	3400

417	766	58890	0	0	-30	2	12260	1920	3340
416	799	63180	0	2	-17	4	12100	1880	3280
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377	941	70400	0	-3	-5	7	14530	1330	3430
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336	1020	21750	0	19	30	40	6400	7900	5310
335	980	21230	0	26	25	51	6290	7920	5380
334	965	20630	0	22	31	53	6150	7880	5390
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330	954	18920	0	15	83	99	5790	7240	5420
329	1010	18450	0	26	106	118	5630	6890	5440
328	1010	18280	0	37	120	151	5460	6500	5510
327	1010	18100	0	41	147	187	5330	6230	5550
326	1050	17800	0	39	179	207	5240	5950	5490
325	1020	17490	0	26	188	232	5240	5750	5470
324	1030	17180	0	19	217	290	5230	5590	5590
323	1030	16880	0	20	279	354	5140	5400	5610
322	1020	16840	0	31	336	420	5030	5220	5640
321	1030	16570	0	39	358	487	4960	5060	5680
320	1030	16490	0	36	410	562	4870	4910	5770
319	1060	16390	0	40	438	626	4830	4760	5790
318	1130	16300	0	35	510	715	4870	4630	5800
317	1080	16180	0	35	572	795	4800	4470	5910
316	1150	16110	0	39	594	859	4740	4330	5950
315	1180	16060	0	56	622	933	4680	4210	5980
314	1120	16080	0	56	670	1010	4650	4130	6010
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311	1350	16320	0	132	893	1240	4740	4090	6230
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307	2200	17060	1	616	1600	1920	4860	4390	6620
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305	3420	17970	1	1210	2180	2510	4980	4930	6810
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302	6520	18890	1	2260	2940	3570	5050	6140	7240
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299	11750	20420	1	2720	3460	4510	5090	7740	7680
298	14020	20910	1	2800	3640	4790	5080	8350	7880
297	16700	21400	1	2860	3800	5040	5120	8960	8020
296	20050	22180	1	2930	3930	5300	5260	9550	8210
295	24010	22840	2	2980	4080	5540	5340	10070	8320
294	28660	23680	2	3050	4240	5760	5490	10550	8480
293	33750	24800	2	3120	4430	6020	5720	10930	8670
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273	50150	35440	7	6900	5790	4680	19030	11950	11300
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263	34230	35780	12	11130	7410	5380	18830	15680	13060
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259	28490	36200	16	11430	7780	5910	17630	16660	14180
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257	25460	35790	18	11190	7940	6260	17180	17970	14890
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251	21310	37390	24	9600	8810	8100	17810	28520	18180
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249	22170	39470	27	9020	9340	9180	19720	35880	19650
248	23250	40940	29	8760	9720	9810	20830	40120	20440
247	24660	42610	30	8550	10120	10490	22360	44320	21220
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245	29610	47700	34	8350	11200	12100	26370	50890	22680
244	33460	51290	35	8370	11840	13010	29250	52510	23290
243	38550	55920	37	8490	12540	14030	32110	52920	23860
242	45110	61450	39	8740	13390	15170	34850	52240	24360
241	53450	67980	41	9120	14440	16400	37060	50840	24770
240	63850	75670	44	9630	15560	17690	38830	48960	25080

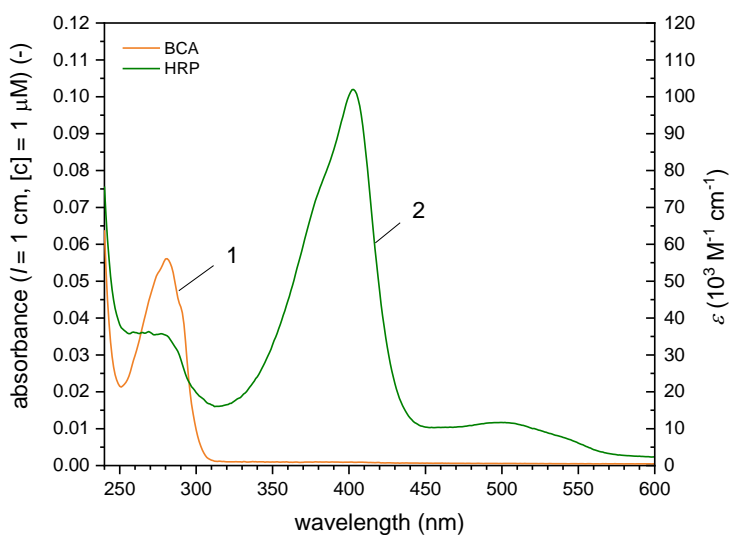


Figure S-6. UV/vis absorption spectra of BCA (1, orange) and HRP (2, green). The measured spectra were adjusted to ϵ_{280} (BCA) = $5.6 \cdot 10^4$ $M^{-1}cm^{-1}$ and ϵ_{403} (HRP) = $1.02 \cdot 10^5$ $M^{-1}cm^{-1}$, respectively (see section 2.1.).

8. Oxidation of DCF in the presence of HRP and H₂O₂

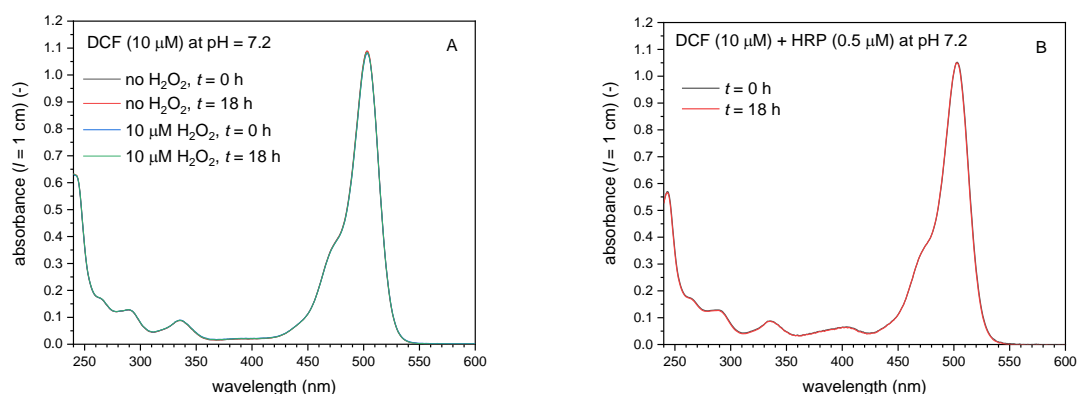


Figure S-7. Stability of DCF (10 μM) in PB (10 mM sodium phosphate buffer solution, pH = 7.2) at $T = 25\text{ }^{\circ}\text{C}$, **(A)** without added H₂O₂ or HRP (18 h), in the presence of H₂O₂ (9 μM, 72 h) without HRP; and **(B)** in the presence of HRP (0.5 μM, 18 h) without H₂O₂. For each stability test, the absorption spectrum was measured every 5 min against PB by keeping the solutions in the spectrophotometer (path length $l = 1\text{ cm}$). Under the condition used, DCF was stable as no significant change in the absorption spectrum could be detected.

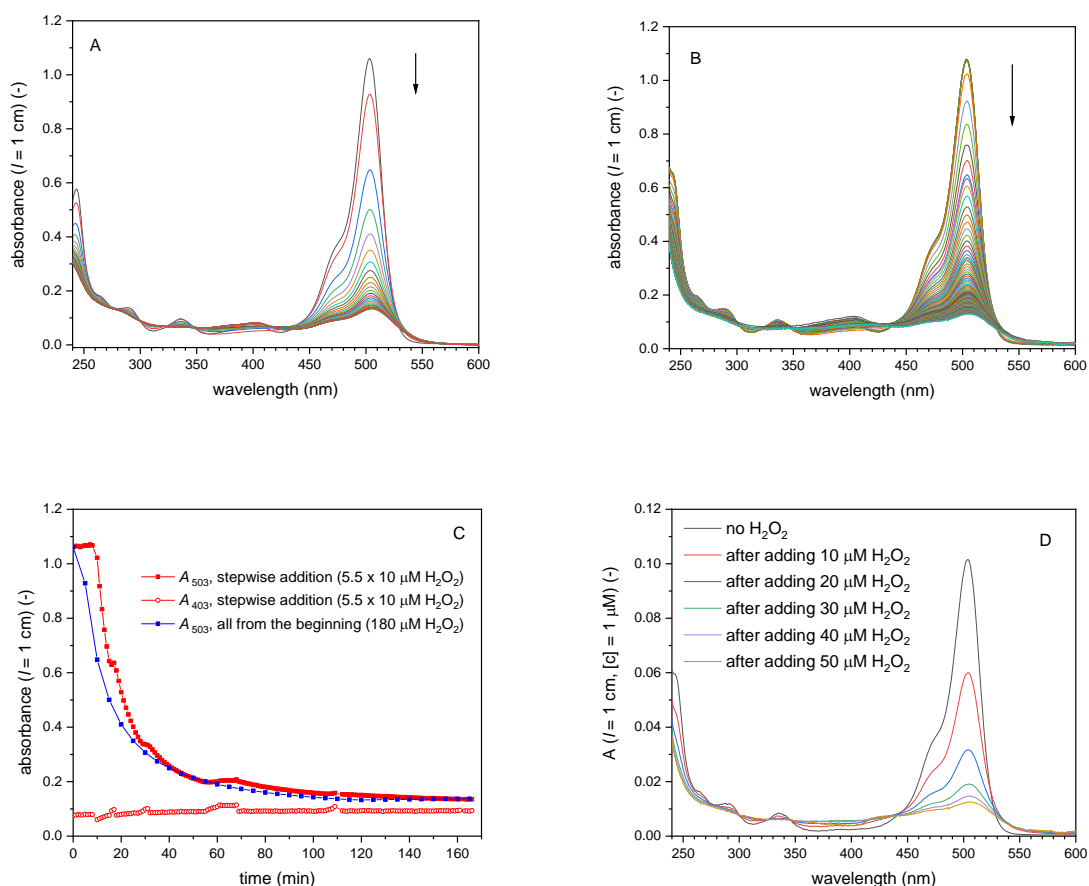


Figure S-8. Oxidation of DCF in PB (10 mM sodium phosphate buffer solution, pH = 7.2) at $T = 25\text{ }^{\circ}\text{C}$ with HRP and H_2O_2 .

(A) $[\text{DCF}]_0 = 10\text{ }\mu\text{M}$, $[\text{HRP}] = 0.5\text{ }\mu\text{M}$, $[\text{H}_2\text{O}_2]_0 = 180\text{ }\mu\text{M}$ *added at once*. The spectrum was recorded every 5 min for 180 min. A_{503} decreased with time. Addition of more H_2O_2 did not lead to a further decrease in A_{503} .

(B) $[\text{DCF}]_0 = 10\text{ }\mu\text{M}$, $[\text{HRP}] = 0.5\text{ }\mu\text{M}$, with *stepwise addition* of H_2O_2 (10 μM) up to 55 μM . The spectrum was recorded every 10 min. After each H_2O_2 addition, the spectrum was recorded until it did not change anymore. Then the next portion of H_2O_2 was added, up to a total of 5.5 equivalents with respect to $[\text{DCF}]_0$. After addition of 5 equivalents H_2O_2 (50 μM), the spectrum did not change anymore.

(C) Comparison of the two DCF oxidation experiments shown in **(A)** and **(B)**. Changes in A_{503} and A_{403} . The data for A_{403} reflect transient changes in the oxidation state of the heme group in HRP.^{S9, S10}

(D) Stable absorption spectra after stepwise addition of H_2O_2 , from top to bottom: DCF only (10 μM , no H_2O_2), after addition of 10, 20, 30, 40, 50 or 55 μM H_2O_2 (and waiting until equilibrium was reached). From the recorded spectra, the spectrum of HRP (0.5 μM , see **Figure S-6**) was subtracted and they were normalized to 1 μM with respect to the concentration and volume of the initial DCF solution. The last two spectra were identical and taken as the spectrum of DCF_{ox} . Oxidation of DCF is indicated by the increase in absorbance at $\lambda = 530\text{-}550\text{ nm}$, see Rota et al. (1999).^{S11}

9. Catalase-like activity of HRP under similar conditions as used for reactions with DCF derivatives in bulk solution

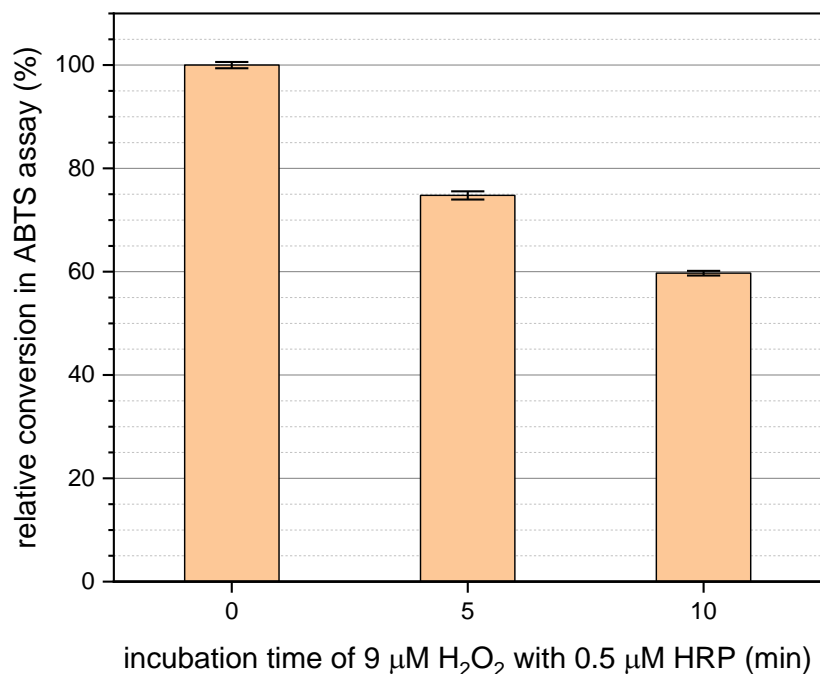


Figure S-9. Decreasing H₂O₂ concentrations observed with the ABTS assay. HRP (0.5 μM) and H₂O₂ (9 μM) were mixed in PB (10 mM phosphate buffer solution, pH 7.2). Either at the same time or after 5 or 10 min incubation time in light-shielded polypropylene tubes, ABTS²⁻ (1 mM) was added to the solutions that were placed in a 1 cm quartz cell. After mixing by pipette, UV/vis spectra were recorded until A_{414} became stable (A_{414} indicated formation of ABTS^{•-}, the reaction was finished within the first 10 seconds). After subtracting A_{414} contributions from HRP and the ABTS²⁻ substrate solution in PB, the obtained ΔA_{414} was attributed relatively to the amount of H₂O₂ present at the time of ABTS²⁻ addition (that was converted to ABTS^{•-} proportionally). It was clearly shown that under the conditions used (HRP 0.5 μM in PB), the concentration of H₂O₂ significantly decreased over time. This can be attributed to the HRP-catalysed decomposition of H₂O₂ (“catalytic” activity of HRP).^{S12}

10. Hydrolysis of DCFH₂-DA catalysed by BCA

Control measurements 1

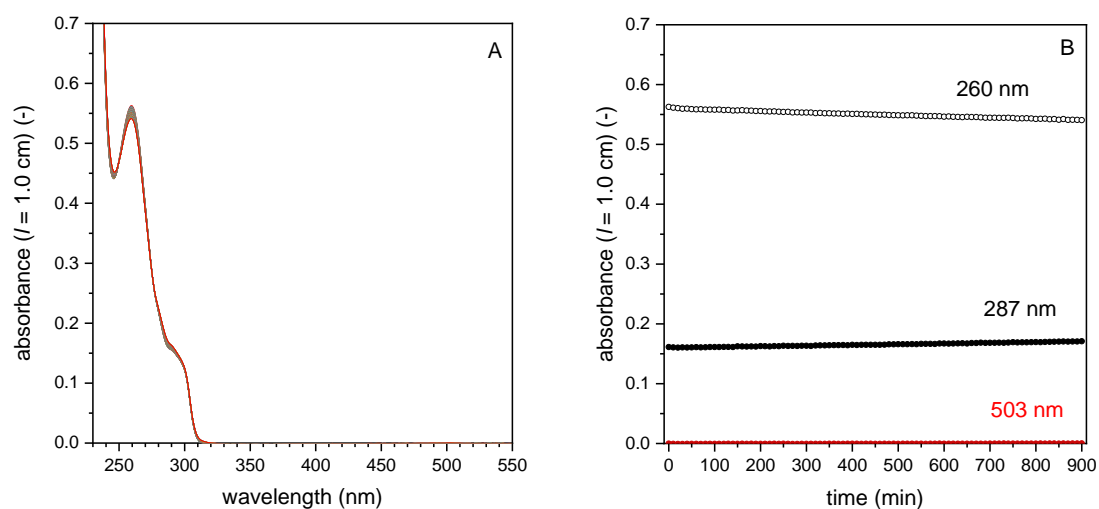


Figure S-10. Stability of DCFH₂-DA (50 μ M, 1 vol% DMSO) in PB (10 mM sodium phosphate buffer solution, pH = 7.2) at 25 $^{\circ}$ C, determined by recording the UV/vis spectrum as a function of time. **(A)** Change in the spectrum, as recorded every 10 min for 15 h. **(B)** Changes of A_{260} , A_{287} , and A_{503} .

Control measurements 2

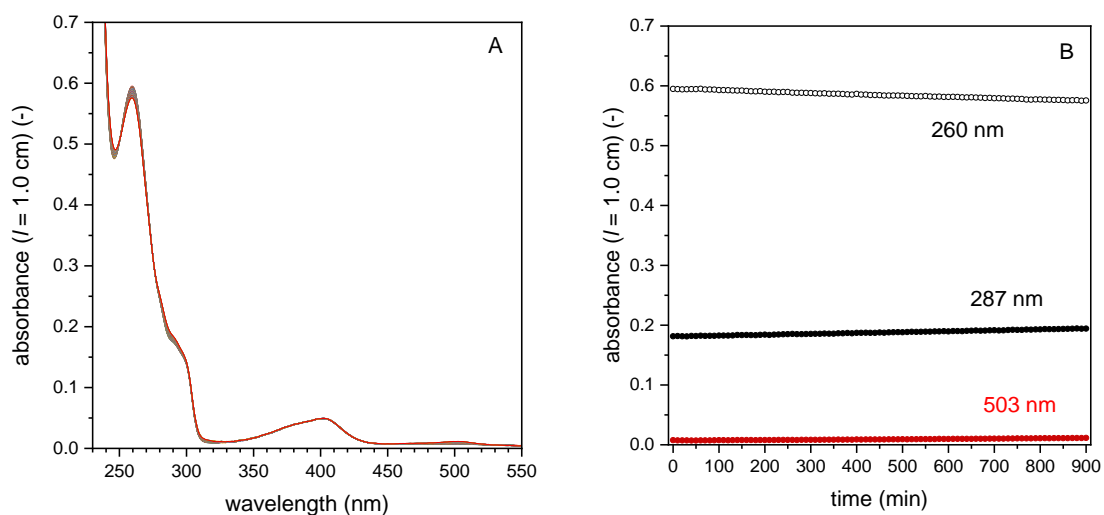


Figure S-11. Stability of DCFH₂-DA (50 μ M, 1 vol% DMSO) in PB (10 mM sodium phosphate buffer solution, pH = 7.2) at 25 $^{\circ}$ C in the presence of HRP (0.5 μ M). **(A)** Change in the UV/vis absorption spectrum, recorded every 10 min for 15 h. **(B)** Changes of A_{260} , A_{287} , and A_{503} . The absorption band with maximal absorption at 403 nm originated from the heme group of HRP.^{S8, S13}

Control measurements 3

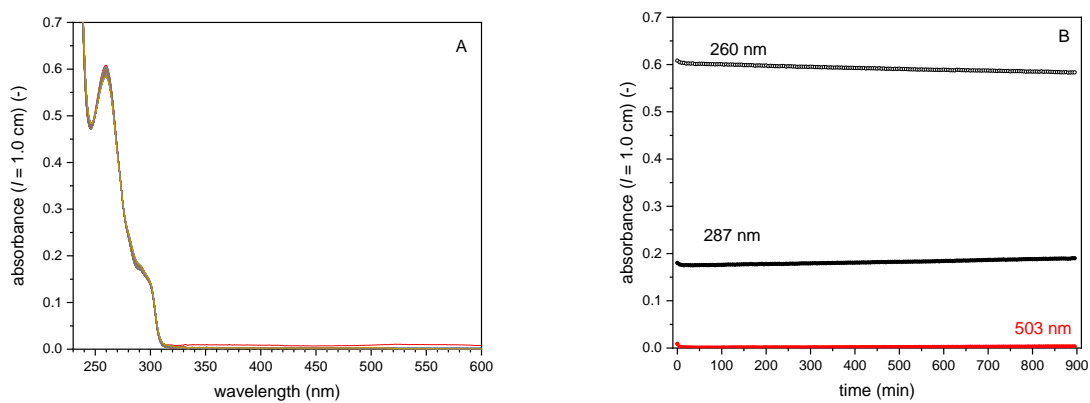


Figure S-12. Stability of DCFH₂-DA (50 μ M, 1 vol% DMSO) in PB (10 mM sodium phosphate buffer solution, pH = 7.2) at 25 $^{\circ}$ C in the presence of H₂O₂ (9 μ M). **(A)** Change in the UV/vis absorption spectrum, recorded every 5 min for 15 h. **(B)** Changes of A_{260} , A_{287} , and A_{503} .

Control measurements 4

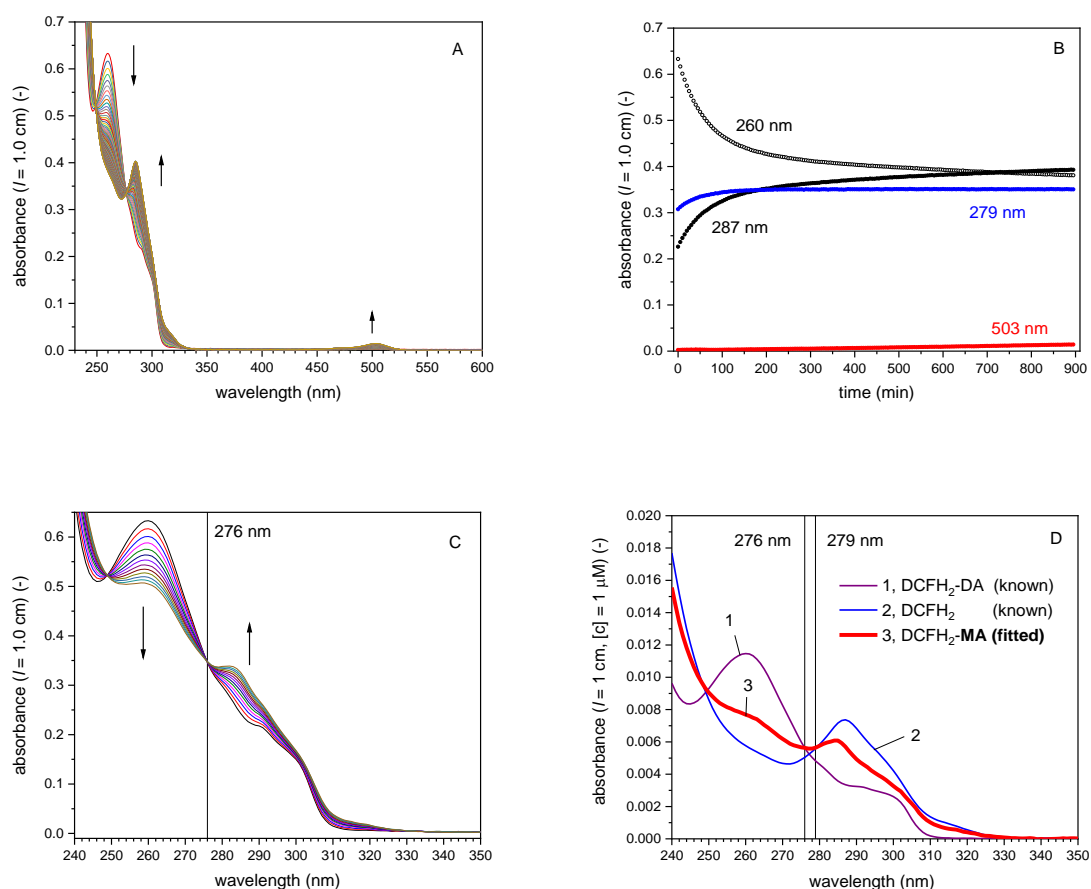


Figure S-13. BCA-catalysed hydrolysis of DCFH₂-DA (50 μM, 1 vol% DMSO) in PB (10 mM sodium phosphate buffer solution, pH = 7.2) at 25 °C. [BCA] = 1.0 μM. **(A)** Change in the UV/vis absorption spectrum, recorded every 5 min for 15 h. **(B)** Changes of A_{260} , A_{287} , A_{279} and A_{503} . After $t = 900$ min, A_{503} was 0.014, corresponding to 0.14 μM DCF (ϵ_{503} (DCF) = $1.02 \cdot 10^5$ M⁻¹cm⁻¹, see **Figure S-3**). At $\lambda = 279$ nm, the isosbestic point between DCFH₂ and DCFH₂-MA becomes visible with stable absorption the latest after $t = 300$ min (no DCFH₂-DA left). **(C)** Change in the UV/vis absorption spectrum, recorded every 5 min within the first 60 min of reaction. At $\lambda = 276$ nm, the isosbestic point between DCFH₂-DA and DCFH₂-MA becomes visible, as the small early DCFH₂ formation can be neglected. **(D)** Absorption spectra of DCFH₂-DA (1), DCFH₂ (2) (both known, see **Figure 2**) and DCFH₂-MA (3) (fitted). With help of the two isosbestic points of DCFH₂-MA (at 276 nm with DCFH₂-DA and at 279 nm with DCFH₂), the molar absorption spectrum could be fitted from the time evolution of the reaction between $t = 300$ -900 min (see **B**). Thereby, DCFH₂-MA hydrolysis was assigned to the differential spectrum ($A_{900\text{min}} - A_{300\text{min}}$) in such a concentration that the resulting molar fit for DCFH₂-MA (adding the molar differential spectrum to the known spectrum of DCFH₂) complied with both isosbestic points (both isosbestic points are from different reaction phases and thereby independent).

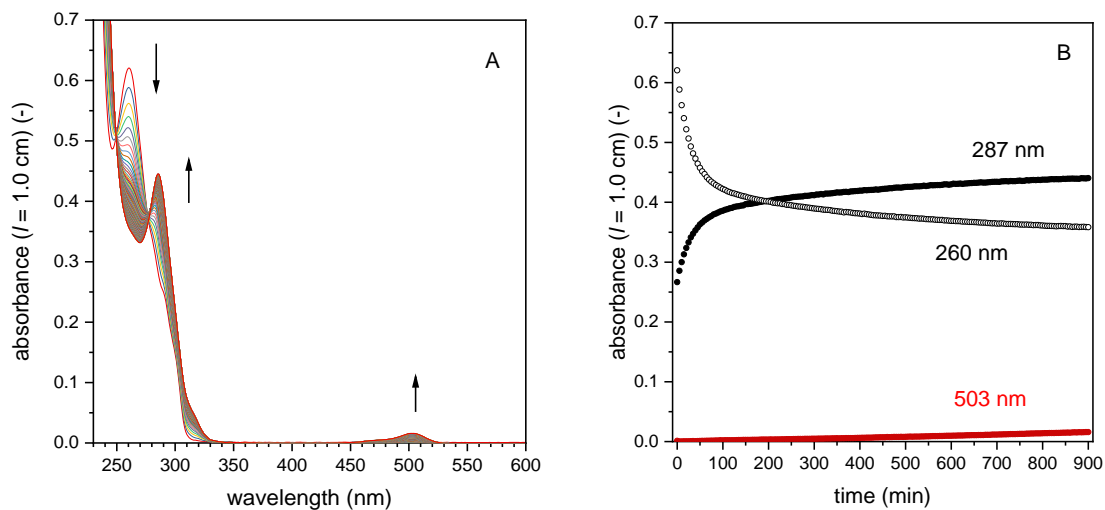


Figure S-14. BCA-catalysed hydrolysis of DCFH₂-DA (50 μ M, 1 vol% DMSO) in PB (10 mM sodium phosphate buffer solution, pH = 7.2) at 25 $^{\circ}$ C. [BCA] = 2.0 μ M. **(A)** Change in the UV/vis absorption spectrum, recorded every 5 min for 15 h. **(B):** Changes of A_{260} , A_{287} , and A_{503} . After $t = 900$ min, A_{503} was 0.016, corresponding to 0.16 μ M DCF (ϵ_{503} (DCF) = $1.02 \cdot 10^5$ M⁻¹cm⁻¹, see **Figure S-3**).

Control measurements 5

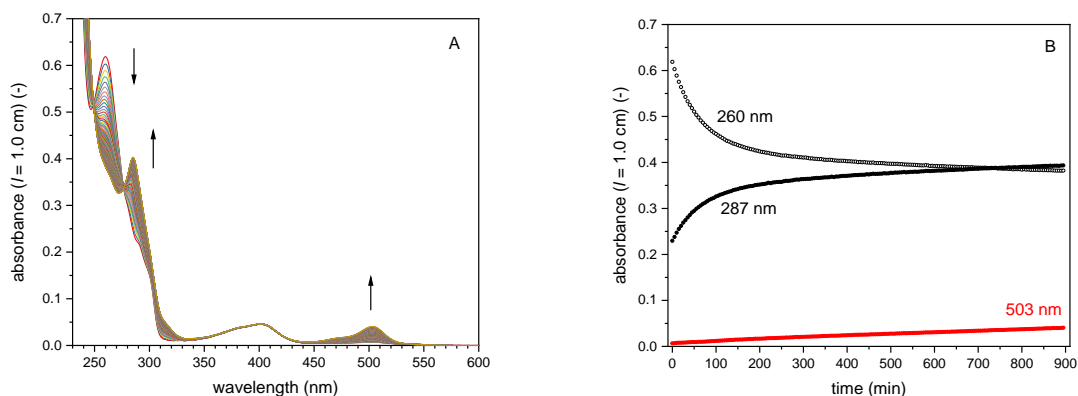


Figure S-15. BCA-catalysed hydrolysis of DCFH₂-DA (50 μM, 1 vol% DMSO) in PB (10 mM sodium phosphate buffer solution, pH = 7.2) at 25 °C in the presence of HRP (0.5 μM). [BCA] = 1.0 μM, *no* added H₂O₂. **(A)** Change in the UV/vis absorption spectrum, recorded every 5 min for 15 h. **(B)** Changes of A₂₆₀, A₂₈₇, and A₅₀₃. After *t* = 900 min, A₅₀₃ was 0.04, corresponding to 0.4 μM DCF (ϵ_{503} (DCF) = $1.02 \cdot 10^5 \text{ M}^{-1}\text{cm}^{-1}$, see **Figure S-3**).

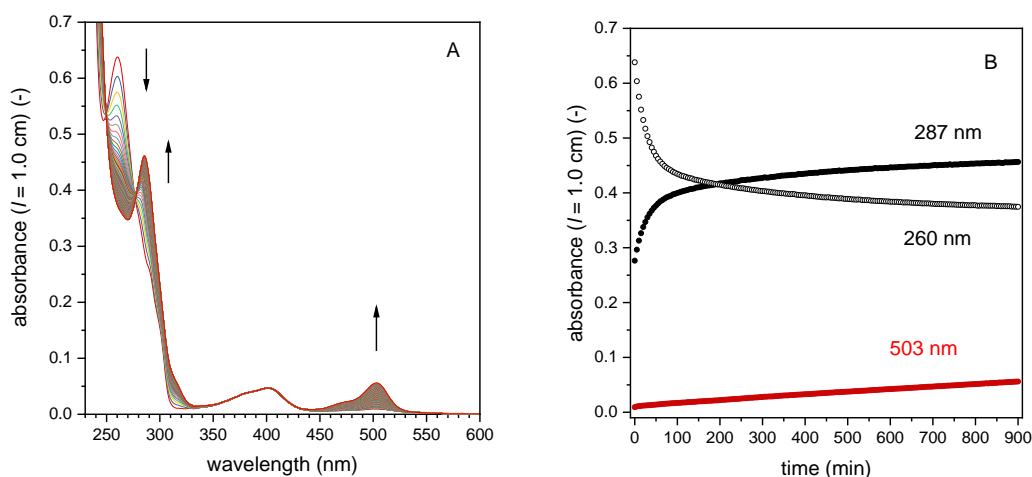


Figure S-16. BCA-catalysed hydrolysis of DCFH₂-DA (50 μM, 1 vol% DMSO) in PB (10 mM sodium phosphate buffer solution, pH = 7.2) at 25 °C in the presence of HRP (0.5 μM). [BCA] = 2.0 μM, *no* added H₂O₂. **(A)** Change in the UV/vis absorption spectrum, recorded every 5 min for 15 h. **(B)** Changes of A₂₆₀, A₂₈₇, and A₅₀₃. After *t* = 900 min, A₅₀₃ was 0.056, corresponding to 0.55 μM DCF (ϵ_{503} (DCF) = $1.02 \cdot 10^5 \text{ M}^{-1}\text{cm}^{-1}$, see **Figure S-3**).

Control measurements 6

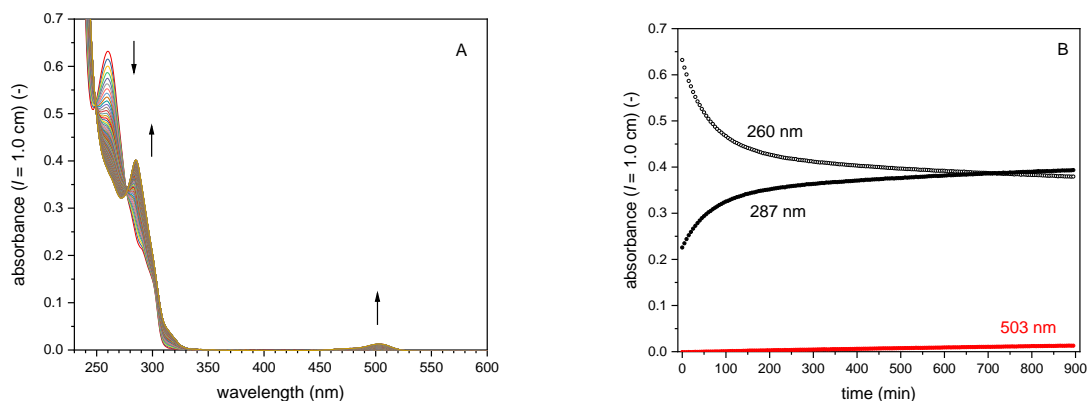


Figure S-17. BCA-catalysed hydrolysis of DCFH₂-DA (50 μ M, 1 vol% DMSO) in PB (10 mM sodium phosphate buffer solution, pH = 7.2) at 25 $^{\circ}$ C in the presence of H₂O₂ (9 μ M) *without* HRP. [BCA] = 1.0 μ M. **(A)** Change in the UV/vis absorption spectrum, recorded every 5 min for 15 h. **(B)** Changes of A_{260} , A_{287} , and A_{503} . After $t = 900$ min, A_{503} was 0.013, corresponding to 0.13 μ M DCF (ϵ_{503} (DCF) = $1.02 \cdot 10^5$ M⁻¹cm⁻¹, see **Figure S-3**).

Control measurements 7

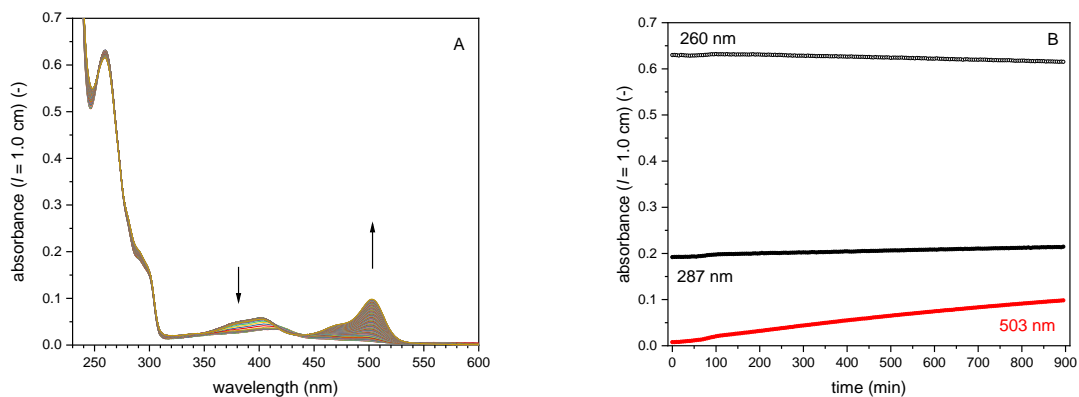


Figure S-18. Non-enzymatic hydrolysis of DCFH₂-DA (50 μ M, 1 vol% DMS) in PB (10 mM sodium phosphate buffer solution, pH = 7.2) at 25 $^{\circ}$ C in the presence of HRP (0.5 μ M) and H₂O₂ (9 μ M) *without* BCA. **(A)** Change in the UV/vis absorption spectrum, recorded every 5 min for 15 h. **(B)** Changes of A_{260} , A_{287} , and A_{503} . After $t = 900$ min, A_{503} was 0.1, corresponding to 0.98 μ M DCF (ϵ_{503} (DCF) = $1.02 \cdot 10^5$ M⁻¹cm⁻¹, see **Figure S-3**).

11. Specificity constants (k_{cat}/K_M) for the BCA-catalysed hydrolysis of DCFH₂-DA, DCFH₂-MA and DCF-MA in bulk solution

Table S-2. Specificity constants (or kinetic efficiencies), k_{cat}/K_M (Michaelis-Menten kinetics), for the hydrolysis of DCFH₂-DA, DCFH₂-MA and DCF-MA catalysed by BCA. The experimentally measured initial rates of hydrolysis were linearly dependent on the concentration of BCA used ($[E]_0$) and on the current substrate concentration ($[S]$). Due to this linear dependency and BCA being rather inefficient in catalysing the hydrolysis of these substrates, Michaelis-Menten kinetics for $[S] \ll K_M$ was assumed to apply. Under these conditions, the reaction rate is linearly dependent on $[S]$ and $[E]_0$ and the rate constant is k_{cat}/K_M , such that: $d[S]/dt = (k_{cat}/K_M) \cdot [E]_0 \cdot [S]$. Thus, first order kinetics applies, such that $[S]_t = [S]_0 \cdot e^{-kt}$, whereas $k = (k_{cat}/K_M) \cdot [E]_0$. The experimentally determined decrease of $[S]_t$ with time was fitted as exponential decay with time and the obtained value for k was divided by $[E]_0$ to get k_{cat}/K_M . 10 mM sodium phosphate buffer solution, pH = 7.2, 25 °C.

Substrate (Reaction Step in Scheme 2)	k_{cat}/K_M ($\mu\text{M}^{-1} \text{min}^{-1}$)	Initial experimental conditions	Fitting conditions (Respective figure with data)
DCFH ₂ -DA ^a (Hyd_1)	0.0187	50 μM DCFH ₂ -DA 1.0 μM BCA	$t = 0-300$ min, $R^2=0.99891$ (Figure 3A, purple data points)
	0.0209	50 μM DCFH ₂ -DA 2.0 μM BCA	$t = 0-200$ min, $R^2=0.99912$ (Figure 3B, purple data points)
DCFH ₂ -MA ^b (Hyd_2)	0.0012	50 μM DCFH ₂ -DA 2.0 μM BCA	$t = 200-900$ min, $R^2=0.99948$ (Figure 3B, red data points)
DCF-MA ^b (Hyd_3)	0.0043	20 μM DCF-DA 1.0 μM BCA	$t = 200-500$ min, $R^2=0.99928$ (Figure S-19C, red data points)
DCF-MA within cascade reaction, thus coexisting DCFH ₂ -DA and DCFH ₂ -MA ^c (Hyd_3)	0.0050	50 μM DCFH ₂ -DA 2.0 μM BCA 0.5 μM HRP 9 μM H ₂ O ₂	$t = 50-350$ min, $R^2=0.99992$ (DCF-MA concentration calculated from A_{503} and A_{460} in Figure 5, blue data points)
	0.0048	50 μM DCFH ₂ -DA 3.0 μM BCA 0.5 μM HRP 9 μM H ₂ O ₂	$t = 50-350$ min, $R^2=0.99990$ (DCF-MA concentration calculated from A_{503} and A_{460} in Figure 5, red data points)

^a The increasing amount of DCFH₂-MA (resulting from the hydrolysis of DCFH₂-DA) did not have a significant impact on $d[\text{DCFH}_2\text{-DA}]/dt$. No product inhibition was observed, although the hydrolysis of DCFH₂-MA was also catalysed by BCA. The reason could be a low ES-complex concentration which is usual for $[S] \ll K_M$, and thus not many BCA enzyme molecules were blocked by other substrates.

^b The DCFH₂-MA and DCF-MA concentrations were taken into account after the concentrations of DCFH₂-DA and DCF-DA, respectively, were approaching zero.

^c The respective k_{cat}/K_M was found to be similar for different enzyme concentrations within the cascade reaction as well as for an “undisturbed” DCF-MA/DCF mixture (entry above), including higher DCF-species concentration (50 μM as opposed to 20 μM). Although [DCF-MA] within the time frame used for the fitting was smaller than coexisting [DCFH₂-DA] + [DCFH₂-MA], no significant impact on $d[\text{DCF-MA}]/dt$ was found within the cascade reaction. No competitive inhibition by the reduced DCF-species was observed, although the hydrolysis of DCFH₂-DA and DCFH₂-MA was also catalysed by BCA at the same time. The reason could be a low ES-Complex concentration, which is usual for $[S] \ll K_M$, and thus not many BCA enzymes being blocked by other substrates.

12. Hydrolysis of DCF-DA catalysed by BCA

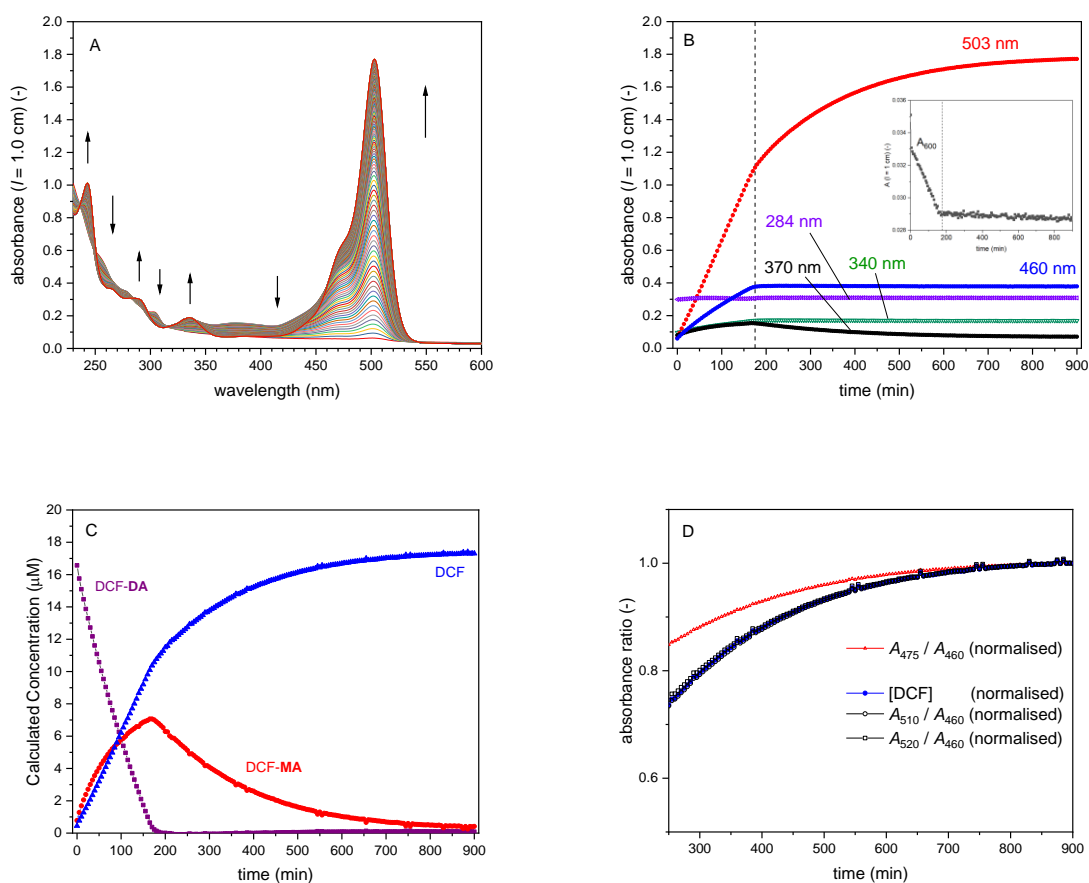


Figure S-19. BCA-catalysed hydrolysis of DCF-DA (20 μM , 1 vol% DMSO) in PB (10 mM sodium phosphate buffer solution, pH = 7.2) at 25 $^{\circ}\text{C}$. [BCA] = 1.0 μM . **(A)** Change in the UV/vis absorption spectrum, recorded every 5 min for 15 h. **(B)** Changes of A_{503} , A_{460} , A_{370} , A_{340} and A_{284} . A_{503} can be mainly attributed to the formation of DCF, and A_{370} to DCF-MA. After an initial reaction phase (up to the dashed vertical line at $t = 175$ min), isobestic points were at $\lambda = 284$, 340 and 460 nm. After $t = 175$ min, all DCF-DA molecules were at least partially hydrolysed (presence of only DCF-MA and DCF). As the samples were turbid at the beginning of the reaction (see inset), no quantitative analysis was performed during the initial phase of the reaction (initial hydrolysis of DCF-DA). **(C)** Concentration determination during the reaction, using the known absorbance of DCF at $\lambda = 520$ nm ($\epsilon_{520}(\text{DCF}) = 2.7 \cdot 10^4 \text{ M}^{-1}\text{cm}^{-1}$) and the isobestic point between DCF-MA/DCF at $\lambda = 460$ nm ($\epsilon_{460}(\text{DCF-MA/DCF}) = 1.98 \cdot 10^4 \text{ M}^{-1}\text{cm}^{-1}$). **(D)** Comparison of the time course of A_{520} , A_{510} and A_{475} with the rising DCF concentration. All values were normalised to the value observed at $t = 900$ min. Whereas the ratio for A_{520} and A_{510} aligned well with the calculated concentration of DCF, A_{475} (as an arbitrary example at a wavelength at which DCF-MA absorbs light) did deviate. This shows that DCF-MA does not absorb at $\lambda > 510$ nm, which was used to fit the molar absorption spectrum of DCF-MA (see 3.1.3 and **Figure 2**). For the BCA-catalysed hydrolysis of DCF-MA, the specificity constant ($k_{\text{cat}}/K_{\text{M}}$) was determined (see **Table S-2**).

13. Oxidation of DCFH₂ catalysed by HRP with H₂O₂ as oxidant

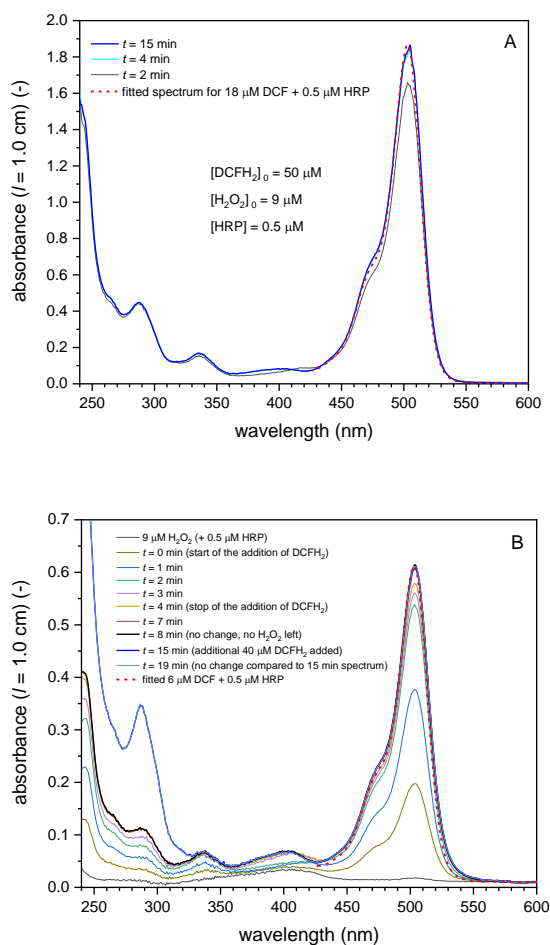


Figure S-20. HRP-catalysed oxidation of DCFH₂ (50 μM) in PB (10 mM sodium phosphate buffer solution, pH = 7.2) at 25 °C, [HRP] = 0.5 μM, [H₂O₂]₀ = 9 μM. **(A)** Solutions of DCFH₂, HRP, and H₂O₂ were mixed at the beginning, and the UV/vis spectrum of the reaction mixture was measured after $t = 2, 4,$ and 15 min. The latter spectrum (blue curve) was identical with the spectrum of a 18 μM solution of DCF (red line) between $\lambda = 430$ and 600 nm, as calculated from the reference spectrum of DCF given in **Figure 2B**. As a result, the two-electron reduction of H₂O₂ yielded quantitative conversion of DCFH₂ to DCF according to **Scheme 1**: One-electron oxidations of two molecules of DCFH₂ to DCFH[•] (Oxi_1'a and Oxi_1'b), followed by one-electron oxidations of two molecules of DCFH[•] to two molecules of DCF with dissolved O₂ (Oxi_1''). **(B)** DCFH₂ was added in portions to the reaction mixture. 10 μM DCFH₂ were added slowly during the first 4 min. After the UV/vis spectrum did not change anymore with time ($t = 8$ min), the remaining 40 μM DCFH₂ were added ($t = 15$ min). This final addition did not lead to a further change of the absorption spectrum above $\lambda = 330$ nm. Spectral changes below $\lambda = 330$ nm were due to DCFH₂. This clearly indicates that H₂O₂ (9 μM initially added) reacted only with the initially added DCFH₂ (10 μM), resulting in the formation of DCF and DCF_{ox}. Moreover, another part of the added H₂O₂ was probably oxidized to O₂ *via* the catalytic activity of HRP (see **Figure S-9**).^{S12, S14} In any case, the final spectrum could not be fitted well with the spectrum of DCF only (red dashed line), see the deviations at $\lambda = 425 - 450$ nm and at $\lambda = 525 - 550$ nm. They are clear indications of DCF overoxidation. In any case, the amount of DCF obtained was much lower than in the case of the experiments in **(A)**, $A_{503} = 0.61$ **(B)** vs $A_{503} = 1.83$ **(A)**.

14. Quantitative analysis of different cascade reaction mixtures run in bulk solution with dissolved enzymes

Table S-3. Summary of the quantitative analyses of various bulk cascade reaction mixtures, prepared in PB (pH = 7.2) at $[\text{DCFH}_2\text{-DA}]_0 = 50 \mu\text{M}$ and different concentrations of BCA, HRP, and H_2O_2 , kept for $t = 15 \text{ h}$ at room temperature. The concentrations given for the DCF derivatives in the final spectrum ($t = 15 \text{ h}$) were determined by spectral fitting on the basis of isosbestic points (λ_{iso} , see **Table 1**) and characteristic absorptions (e.g. A_{503} for DCF or A_{550} for DCF_{ox}), using the reference spectra given in **Figure 2** and **Table S-1**. For examples, see **Figure 4C** or **Figure 13**.

Starting conditions: $[\text{DCFH}_2\text{-DA}]_0 = 50 \mu\text{M}$, PB: 10 mM sodium phosphate buffer solution, pH = 7.2			Determined concentrations in the reaction mixtures after a reaction time of 15 h							
[BCA] (μM)	[HRP] (μM)	$[\text{H}_2\text{O}_2]_0$ (μM)	$[\text{DCFH}_2\text{-DA}]$ (μM)	$[\text{DCFH}_2\text{-MA}]$ (μM)	$[\text{DCFH}_2]$ (μM)	$[\text{DCF-MA}]$ (μM)	[DCF] (μM)	$[\text{DCF}_{\text{ox}}]$ (μM)	Extent of hydrolysis (%) ^a	Extent of over- oxidation (%) ^b
1	0.5	9	0	11.5	30.9	0	7.6 (42%) ^c	1.0	77	12
2	0.5	9	0	5.0	36.0	0	9.3 (52%) ^c	0.6	90	6
3	0.5	9	0	2.9	38.1	0	9.6 (52%) ^c	0.3	94	3
1	0.5	4.5	0	15.3	31.7	0	4.9 (54%) ^c	0.5	71	9
1	0.5	9	0	11.5	30.9	0	7.6 (42%) ^c	1.0	77	12
1	0.5	18	0	9.0	27.3	0	12.4 (34%) ^c	2.0	82	14
1	0.5	27	0	7.0	24.0	0	15.7 (29%) ^c	3.5	86	18
1	0.5	180	0	n/a ^d	n/a ^d	n/a ^d	24.0 ^d (7%) ^c	26.0 ^d	n/a ^d	52 ^d
1	0.0050	9	0	18.2	17.3	0.4	14.2 (80%) ^c	0	63	0
1	0.0075	9	0	18.9	16.8	0.4	14.4 (80%) ^c	0	62	0
1	0.5	9	0	11.5	30.9	0	7.6 (42%) ^c	1.0	77	12
1	1	9	0	9.7	33.3	0	7.1 (39%) ^c	1.1	81	13
1	1.5	9	0	10.4	33.6	0	6.7 (37%) ^c	1.1	80	14
1	0	0	0	24.6	28.0	n/a	n/a	n/a	53	n/a
2	0	0	0	6.2	43.8	n/a	n/a	n/a	88	n/a

n/a: not applicable

Colour code: yellow, data to **Figure 5**; blue, data to **Figure 6**; red, data to **Figure 7**, green, data to **Figure 3**

^a The extent of hydrolysis was determined by subtracting the total concentration of the remaining mono-acetate species ($[\text{DCFH}_2\text{-MA}]_{15 \text{ h}} + [\text{DCF-MA}]_{15 \text{ h}}$) from $[\text{DCFH}_2\text{-DA}]_0$.

^b Defined as $([\text{DCF}_{\text{ox}}]_{15 \text{ h}} / ([\text{DCF}_{\text{ox}}]_{15 \text{ h}} + [\text{DCF}]_{15 \text{ h}})) \cdot 100$ (%).

^c Yield compared to a maximal theoretic conversion/consumption of 1 H_2O_2 for yielding 2 DCF.

^d The complete final spectrum of the reaction mixture initially containing a large excess of H_2O_2 could not be fitted convincingly. However, from the spectral region between $\lambda = 460$ and 600 nm , the ratio between DCF and DCF_{ox} was estimated.

15. Kinetic reaction pathway analysis for the cascade reaction run in bulk solution depending on the HRP concentration

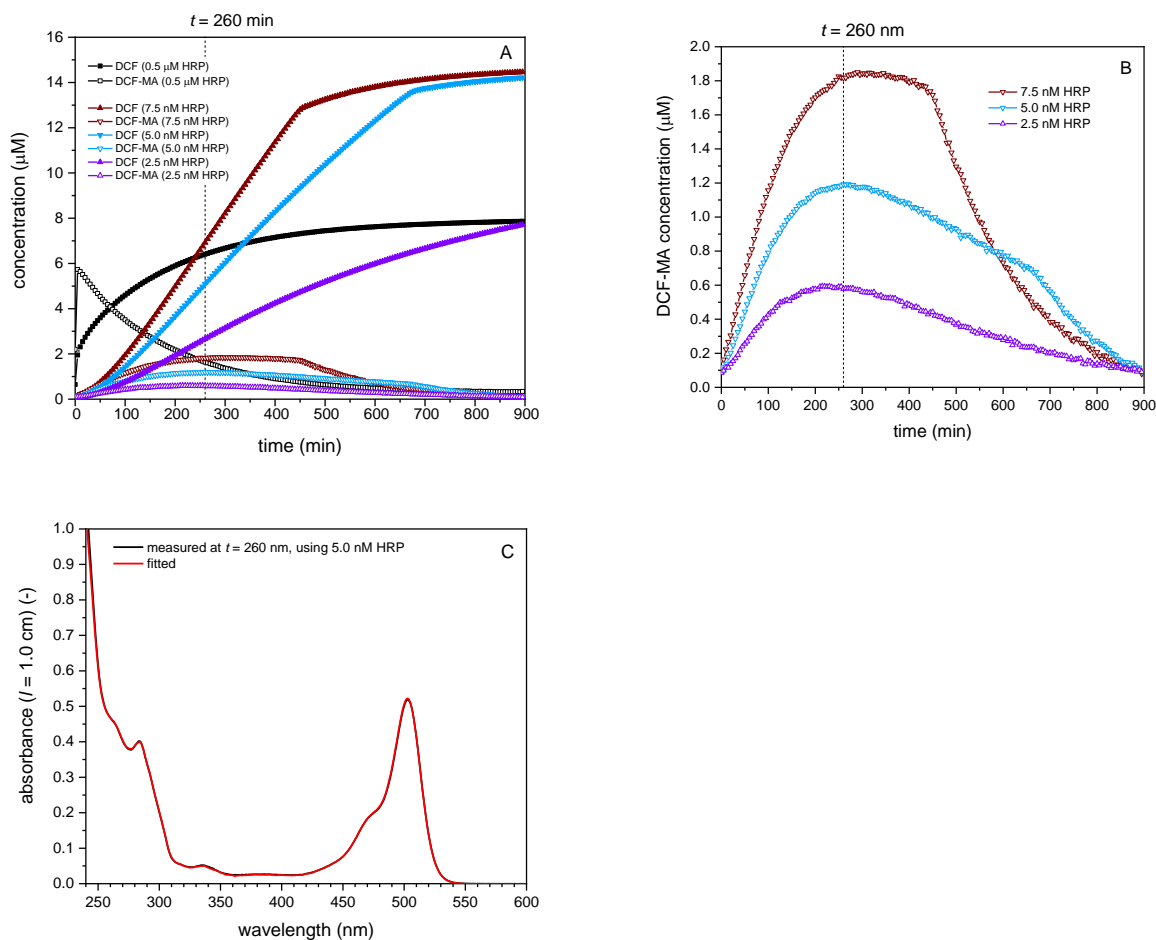


Figure S-21. (A) Concentration of DCF and DCF-MA against time in cascade reactions using 50 μM DCFH₂-DA, 9 μM H₂O₂, 1.0 μM BCA as well as 0.5 μM or 7.5 nM, 5.0 nM or 2.5 nM HRP (in 10 mM phosphate buffer solution, pH = 7.2, at 25 °C). The concentrations were calculated from A_{503} and A_{460} , as shown in Figure 7. **(B)** DCF-MA concentration for those reactions in (A) which were run with 7.5 nM, 5.0 nM, or 2.5 nM HRP. **(C)** Comparison of the spectrum measured after $t = 260$ min for the reaction shown in (A) using 5.0 nM HRP (black line) with the sum of the spectra obtained by fitting (red line). The concentrations of the individual species present in the reaction mixture after $t = 260$ min were determined by spectral fitting, as described in the caption of Table S-3 (except for $t = 260$ min instead of 15 h). The following concentrations resulted from fitting: [BCA] = 1.0 μM ; [HRP] = 5.0 nM; [DCFH₂-DA] = 0 μM ; [DCFH₂-MA] = 40.1 μM ; [DCFH₂] = 4.5 μM ; [DCF-MA] = 1.2 μM ; [DCF] = 5.0 μM ; [DCF_{ox}] = 0 μM .

Reaction pathway analysis for a “hydrolysis-limited” reaction ($[HRP] = 0.5 \mu\text{M}$ in **Figure S-21A**):

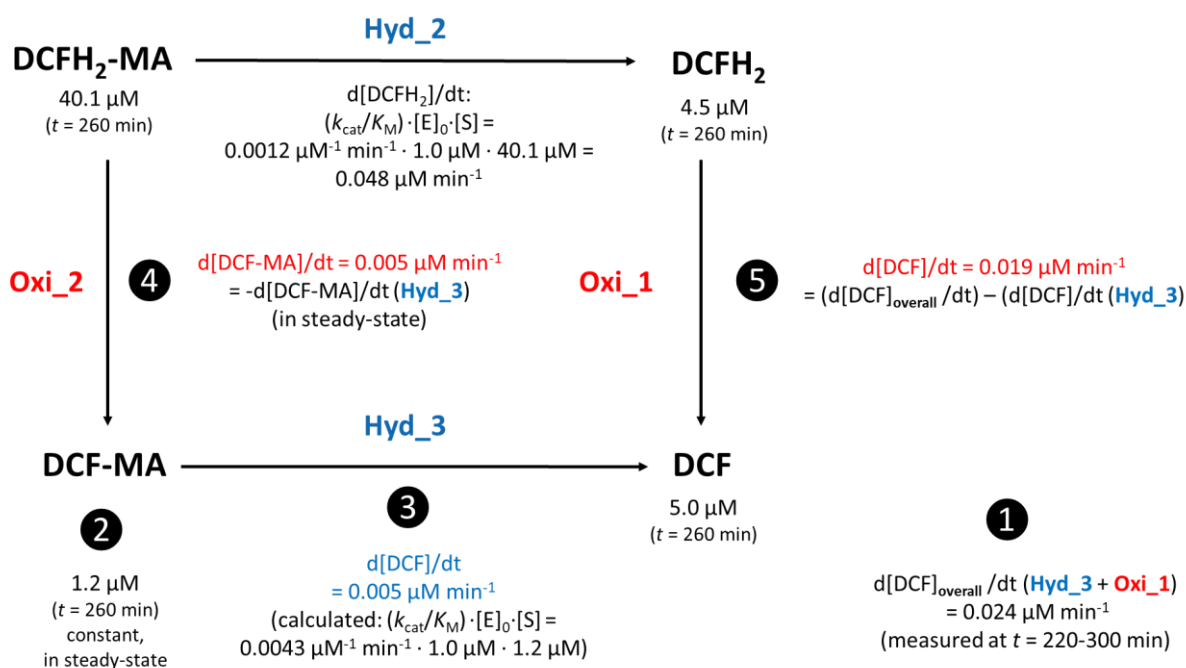
Most of the oxidation reactions were completed after $t = 5 \text{ min}$ where H_2O_2 was nearly used up (**Figure S-21A**). At that time, most of the oxidised species was still DCF-MA ($1 \mu\text{M DCF}_{\text{ox}}$ neglected). This indicates that the reaction proceeded predominantly along *pathway 2* (see **Scheme 2**). The concentration change after 5 min was solely caused by the hydrolysis of DCF-MA to DCF (also part of *pathway 2*). Since the hydrolysis reactions were rate limiting for DCF+DCF-MA formation, a substantial part of H_2O_2 was finally not used for DCF formation. This could be attributed to some overoxidised DCF found (DCF_{ox}) and “catalytic” activity of “excess” HRP (see section 3.1.4 and **Figure S-9**).

Reaction pathway analysis for “oxidation-limited” reactions ($[HRP] = 2.5, 5.0$ and 7.5 nM in **Figure S-21A**):

The oxidation reactions took much longer (H_2O_2 was used up at $t = 450 \text{ min}$ for 7.5 nM HRP , $t = 650 \text{ min}$ for 5.0 nM HRP and not yet within the used experiment time for 2.5 nM HRP , **Figure S-21A**). The formation of DCF-MA (*via* oxidation of $\text{DCFH}_2\text{-MA}$) and its hydrolysis occurred in parallel as long as H_2O_2 was still present (see **Scheme 2**). At the beginning of the reaction, similar DCF and DCF-MA concentrations indicate that *pathway 2* was predominant. With ongoing time, *pathway 1* was gaining more and more share in DCF formation. The oxidation became rate limiting for the formation of DCF + DCF-MA. With this the concentrations of $\text{DCFH}_2\text{-MA}$ and DCFH_2 increased (although H_2O_2 was still present). Since the oxidation was rate limiting, side reactions were largely suppressed (most of the initially added H_2O_2 was used for DCF formation and no DCF_{ox} was found). At the beginning, mainly $\text{DCFH}_2\text{-MA}$ was abundant (favouring **Oxi_2**, *pathway 2*). Over time a significant concentration of DCFH_2 was slowly building up (slowly, on expense of $\text{DCFH}_2\text{-MA}$, see **Figure 3**), speeding up **Oxi_1** of *pathway 1*. Thereby, **Oxi_1** got more and more favoured over **Oxi_2**, therefore favouring *pathway 1* with increasing DCFH_2 to $\text{DCFH}_2\text{-MA}$ ratio. For all three HRP concentrations, the DCF-MA

concentration remained similar at the beginning of the reaction for quite a while and then stayed almost constant for at least 20 min (at $t \approx 260$ min, see **Figure S-21B**).

A comparison of **Oxi_1** with **Oxi_2** was made using steady-state kinetic data (see **Scheme S-1**). The DCF-MA concentration remained stable at $t = 260$ min ($d[\text{DCF-MA}]/dt = 0$). The comparison of **Oxi_1** and **Oxi_2** in **Scheme S-1** shows that *pathway 1* was predominant at $t = 260$. Oxidation via **Oxi_1** was determined as faster (about 4-times) than **Oxi_2**, although $[\text{DCFH}_2]$ ($= 4.5 \mu\text{M}$) was about 9-times lower than $[\text{DCFH}_2\text{-MA}]$ ($= 40.1 \mu\text{M}$) at that time (see **Scheme S-1** and **Figure S-21C**). Consequently, in the following the slowly decreasing DCF-MA concentration and the stable DCF formation rate (see **Figure S-21A,B** for $t > 260$ min) showed that *pathway 1* kept being predominant for $t \approx 260\text{-}650$ min (before finally all H_2O_2 was consumed and DCF-MA hydrolysis was the last remaining DCF formation process after $t \approx 650$ min). In addition, since the DCF formation rate, dA_{460}/dt and $[\text{DCF-MA}]$ in the steady-state ($d[\text{DCF-MA}]/dt = 0$) were all linearly dependent on $[\text{HRP}]$ (see **Figure 7** and **Figure S-21A,B**), the same also applies for 2.5 and 7.5 nM HRP (just at altered time periods).



Scheme S-1. Comparison of reaction rates, as determined for the cascade reaction run with 50 μM DCFH₂-DA, 9 μM H₂O₂, 1.0 μM BCA and 5.0 nM HRP at $t = 260$ min (see **Figure S-21**). For these conditions, oxidation is rate limiting. The different hydrolysis and oxidation steps are the ones of **Scheme 2**.

The “overall” rate of formation of DCF, $\frac{d[\text{DCF}]_{\text{overall}}}{dt}$ (as a result of **Hyd_3** + **Oxi_1**) was found to be proportional to [HRP] and was obtained from **Figure S-21A**, while the given compound concentrations (at $t = 260$ min) were determined by spectral fitting (**Figure S-21C**). The specificity constant, k_{cat}/K_M , for **Hyd_3** is taken from **Table S-2**. The other reaction rates were then calculated (red and blue).

- 1:** The overall DCF formation rate was obtained from **Figure S-21A**.
- 2:** Steady-state concentration of DCF-MA was from **Figure S-21C**
- 3:** The reaction rate for the hydrolysis of DCF-MA was calculated using [DCF-MA], [BCA] and k_{cat}/K_M . As evident from the last three entries of **Table S-2**, k_{cat}/K_M did not change significantly when DCFH₂ and DCFH₂-MA were present, even if present at higher concentration than DCF-MA.
- 4:** Since $\frac{d[\text{DCF-MA}]}{dt} = 0$ at $t = 260$ min, DCF-MA formation by **Oxi_2** had to be equal to DCF-MA hydrolysis by **Hyd_3** ($0.005 \mu\text{M min}^{-1}$). Also note that the steady-state concentration of DCF-MA was linearly dependent on the amount of HRP used (**Figure S-21B**).
- 5:** DCF formation via **Oxi_1** from DCFH₂ could be simply calculated by subtracting the formation via **Hyd_3** from the “overall” formation of DCF at $t = 260$ min.

16. UV/vis absorption spectra and quantitative analysis of the outflow from the cascade reaction run in the glass fiber filter device with immobilised denpol-enzyme conjugates

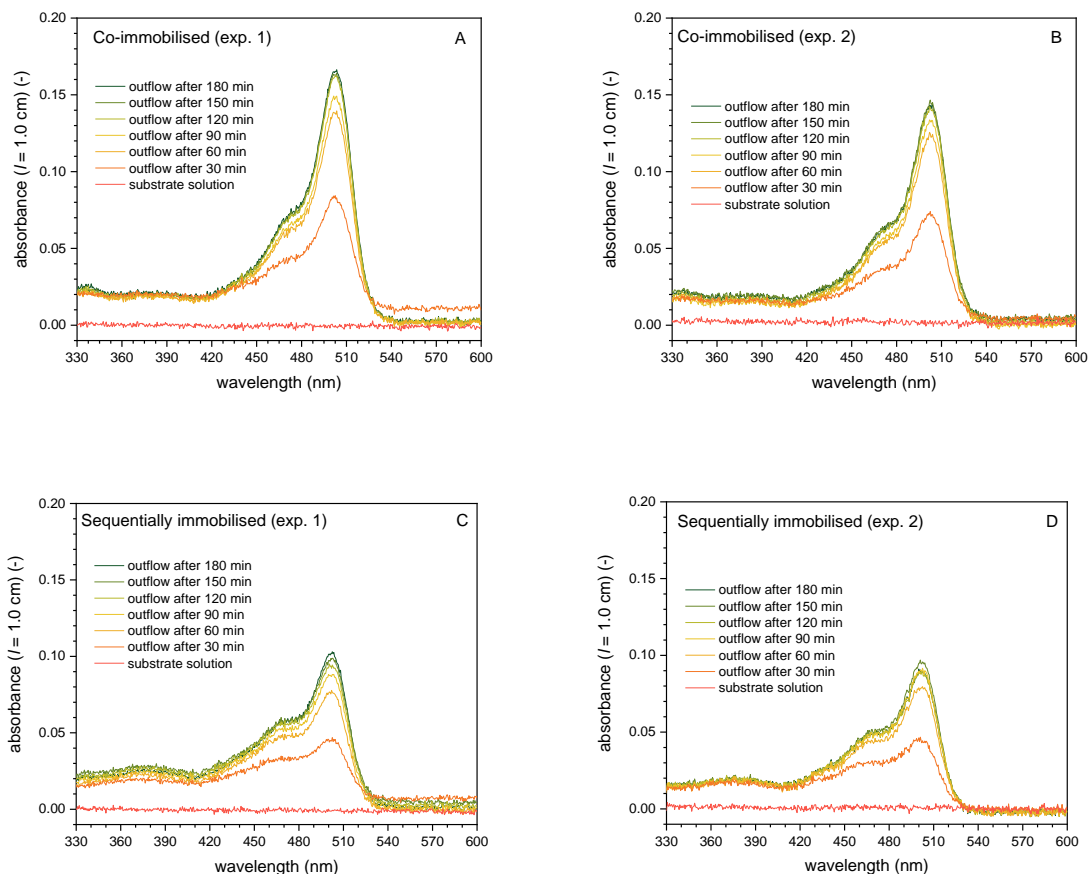


Figure S-22. UV/vis absorption spectra between $\lambda = 330$ and 600 nm of the outflow from a filter holder device consisting of two glass fiber filters with immobilised *de*-PG2₁₀₀₀-BAH₂₀₇-BCA₁₅₂ and *de*-PG2₁₀₀₀-BAH₈₆-HRP₉₀ (see ① in **Figure 11** and **Figure 12** (“exp.1” and “exp.2”). A substrate solution containing $50 \mu\text{M}$ DCFH₂-DA and $9 \mu\text{M}$ H₂O₂ was pumped through the filters and the spectra were recorded in the collected outflow after $t = 30, 60, 90, 120, 150,$ and 180 min ($t = 30$ min means that the outflow was collected until 30 min; $t = 60$ min means, collected outflow between 30 and 60 min; etc.). Between $\lambda = 330$ and 600 nm, only oxidised species absorb (DCF, DCF-MA, and DCF_{ox}, see **Figure 2**). **(A)** and **(B)**: Spectra for co-immobilised conjugates (blue data points in **Figure 12**). **(C)** and **(D)**: Spectra for the sequentially immobilised conjugates (red data points in **Figure 12**).

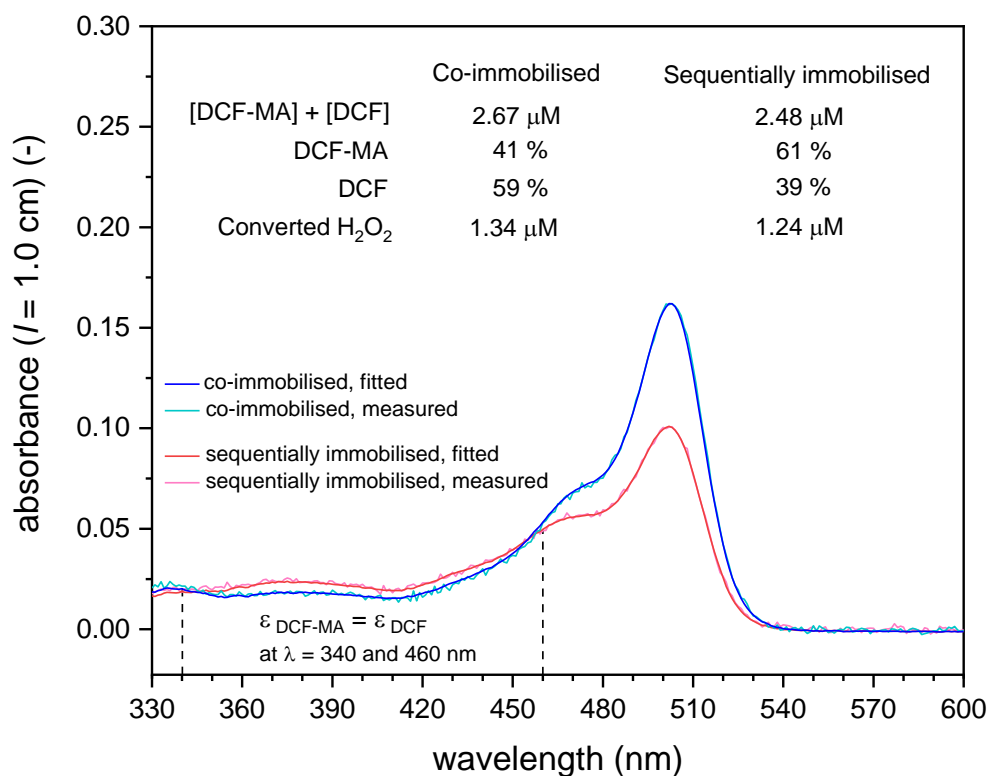


Figure S-23. UV/vis absorption spectra between $\lambda = 330$ and 600 nm of the outflow from the filter holder device for cascade reactions ① in **Figure 11** for co-immobilised (blue) and sequentially immobilised conjugates (red), collected between $t = 150$ and 180 min (“exp.1”, blue filled triangles and red empty triangles at $t = 180$ min in **Figure 12**). The measured and fitted spectra are shown with the total concentrations of DCF-MA and DCF and the relative amounts of DCF-MA and DCF present in the outflow. The fitting of the measured spectrum with the reference spectra of DCF-MA and DCF was carried out as described for the analysis of the reaction run in bulk solution (see caption of **Table S-3**).

17. References

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