Supporting Information

Solid-Phase Synthesis of BODIPY Dyes and Development of an

Immunoglobulin Fluorescent Sensor

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<u>1. General procedures (synthesis, screening, SPR and Job plot analysis) and</u> characterization for 1-3.

Synthesis. General. All the chemicals (building block aldehydes plus others) and solvents were purchased from Sigma Aldrich, Alfa Aesar, Fluka, MERCK or Acros, and used without further purification. 2-chlorotrityl chloride polystyrene resin (100-200 mesh, 1% DVB cross-linking) was purchased from BeadTech (Korea). Human immunoglobulins (G, A and M), immunoglobulin-depleted human serum and other reagents used for the biological characterization were supplied by Sigma Aldrich. Normal phase purifications were carried out using Merck Silica Gel 60 (particle size: 0.040-0.063) mm, 230-400 mesh). Analytical characterization was performed on a HPLC-MS (Agilent-1200 series) with a DAD detector and a single quadrupole mass spectrometer (6130 series) with an ESI probe. Analytical method, unless indicated: eluents: A: H_2O (0.1% HCOOH), B: ACN (0.1% HCOOH), gradient from 5 to 95% B in 10 min; C₁₈(2) Luna column (4.6 \times 50 mm², 5 µm particle size). ¹H-NMR, ¹⁹F-NMR and ¹³C-NMR spectra were recorded on Bruker Avance 300 NMR spectrometer, and chemical shifts are expressed in parts per million (ppm). For ¹⁹F-NMR, trifluoroacetic acid was used as a reference ($\delta = 0.00$ ppm). Spectroscopic and quantum yield data were measured on a SpectraMax M2 spectrophotometer (Molecular Devices), and the data analysis was performed using GraphPrism 5.0.

(9H-fluoren-9-yl)methyl 3-oxo-3-(1H-pyrrol-2-yl)propylcarbamate (1). Fmoc- β -Ala-OH (2.5 g, 8.0 mmol) was dissolved in dry THF. 2,2'-dipyridyl disulfide (2.7 g, 12.0 mmol) and PPh₃ (4.2 g, 16.0 mmol) were added, and the mixture was stirred under N₂ atmosphere at r.t for 24 h. In a separate flask, methylmagnesium bromide (12 mL of a 3.0

M solution in THF, 36.0 mmol) was added dropwise at -78 °C to a solution of pyrrole (3.3 mL, 48.0 mmol) in dry THF (50 mL). The mixture was stirred at -78 °C for 30 min and at -20 °C for another 30 min. Then the thioester crude mixture (after 24 h reaction) was added drop wise at -78 °C, and the whole solution was stirred at -78 °C for 30 min and another 30 min at r.t. The reaction was quenched with a saturated solution of NH₄Cl. After dilution with diethyl ether, the organic layer was washed with water $(3 \times 100 \text{ mL})$ and brine (100 mL), dried over anhydrous sodium sulfate and evaporated under vacuum. Column chromatography (elution with hexane-ethyl acetate 1:1) rendered 2.53 g of 1 as a white solid (yield: 88%). ESI m/z (C₂₂H₂₀N₂O₃), calc: 359.2; found (M+Na⁺): 382.9. ¹H-NMR (300 MHz, CDCl₃): 9.68 (bs, 1H), 7.75 (d, 2H, J = 7.3Hz), 7.57 (d, 2H, J = 7.8Hz), 7.38 (t, 2H, J = 7.6Hz), 7.28 (t, 2H, J = 7.3Hz), 7.04 (bs, 1H), 6.94 (bs, 1H), 6.28 (d, 1H, J = 2.9Hz), 5.48 (bt, 1H, J = 5.9Hz), 4.38 (d, 2H, J = 7.0Hz), 4.20 (t, 1H, J = 7.0Hz), 3.61 (dt, 2H, J = 5.9Hz, 6.1Hz), 3.06 (t, 2H, J = 5.9Hz). ¹³C-NMR (75 MHz, CDCl₃): 189.0, 156.4, 143.9, 141.3, 131.7, 127.6, 127.0, 125.1, 124.9, 119.9, 116.8, 110.9, 66.7, 47.2, 37.6, 36.3.

10 - (2 - (((9H-fluoren-9-yl)methoxy) carbonylamino) ethyl) - 5, 5 - difluoro - 1, 3 - dimethyl - 1,

5H-dipyrrolo[**1,2-c:1',2'-f**][**1,3,2**]**diazaborinin-4-ium-5-uide** (**2**). **1** (1.5 g, 4.2 mmol) and 2,4-dimethylpyrrole (692 μ L, 6.7 mmol) were dissolved in CH₂Cl₂ at 0 °C. After 10 min stirring, the mixture was treated drop wise with POCl₃ (1.3 g, 8.4 mmol), and the resulting solution was stirred at 0 °C for 1 h, and at 35 °C for 20 h. Afterwards, DIEA (2.9 mL, 16.8 mmol) and BF₃·OEt₂ (2.3 mL, 16.8 mmol) were added and the crude mixture was stirred for 4 h at r.t. Column chromatography (elution with hexane-ethyl acetate 4:1) rendered 1.2 g of **2** as a reddish solid (yield: 57%). ESI *m/z* (C₂₈H₂₆BF₂N₃O₂),

calc: 485.2; found (M-F): 466.3. ¹H-NMR (300 MHz, CDCl₃): 7.77 (d, 2H, J = 7.6Hz), 7.61 (bs, 1H), 7.56 (d, 2H, J = 7.0Hz), 7.41 (t, 2H, J = 7.6Hz), 7.31 (t, 2H, J = 7.3Hz), 7.11 (d, 1H, J = 3.5Hz), 6.42 (d, 1H, J = 3.2Hz), 6.17 (bs, 1H), 5.10 (bt, 1H, J = 6.2Hz), 4.43 (d, 2H, J = 6.7Hz), 4.19 (t, 1H, J = 6.7Hz), 3.50 (dt, 2H, J = 6.7 Hz, 7.2Hz), 3.18 (t, 2H, J = 7.0Hz), 2.58 (s, 3H), 2.46 (s, 3H). ¹³C-NMR (75 MHz, CDCl₃): 161.2, 156.3, 145.2, 145.1, 143.7, 142.3, 141.3, 138.2, 134.0, 127.7, 127.0, 125.0, 124.0, 123.8, 120.0, 116.1, 66.8, 47.2, 42.8, 30.1, 16.2, 15.0. ¹⁹F-NMR (282 MHz, CDCl₃): -70.31, -70.53 (dd, J = 31Hz, J = 62Hz, BF₂).^{1.2}

10-(2-aminoethyl)-5,5-difluoro-1,3-dimethyl-5H-dipyrrolo[1,2-c:1',2'-

f][1,3,2]diazaborinin-4-ium-5-uide hydrochloride (3). A solution of 2 (1.0 g, 2.06 mmol) in CH₂Cl₂ was treated with DBU every 15 min for a total of 4 times ($4 \times 73 \mu$ L, 4 \times 0.51 mmol) and a total reaction time of 1 h. Afterwards, the reaction was quenched with a aqueous solution of 0.5 N HCl and stirred for 15 min at r.t. The resulting red solid was filtered off, and thoroughly washed with diethyl ether to render 350 mg of **3** as the hydrochloride salt (yield: 65%). ESI *m/z* (C₁₃H₁₆BF₂N₃), calc: 263.1; found (M-F): 244.1. ¹H-NMR (300 MHz, DMSO-d₆): 8.51 (bs, 2H), 7.65 (s, 1H), 7.63 (s, 1H), 6.52 (s, 1H), 6.46 (s, 1H), 3.37 (m, 2H), 3.16 (s, 3H), 3.03 (m, 2H), 2.50 (s, 3H). ¹³C-NMR (75 MHz, DMSO-d₆): 161.6, 146.3, 141.0, 138.0, 133.5, 133.3, 124.7, 124.0, 116.1, 48.5, 47.8, 15.9, 14.6. ¹⁹F-NMR (282 MHz, DMSO-d₆): -66.97, -67.19 (dd, *J* = 29Hz, *J* = 61Hz, BF₂).

Synthesis of the BDM Library. A solution of **3** (350 mg, 1.3 mmol) and DIEA (1.1 mL, 6.5 mmol) was dissolved in *N*-methylpyrrolidone (NMP) and added to 2-chlorotrityl chloride polystyrene resin (loading: 1.2 mmol/g, 3.3 g, 3.9 mmol). The reaction was

shaken at r.t. for 16 h, after which the resin was capped with MeOH (0.8 mL/g resin) for 4 h, and finally filtered off and washed with NMP, DMF and DCM (× 4 each solvent). For every compound, 50 mg of the loaded resin (corresponding to 0.06 mmol of 3) were re-suspended in 2 mL DMSO-ACN (1:1), and treated with pyrrolidine (75 µL, 0.9 mmol), AcOH (54 µL, 0.9 mmol) and the corresponding aldehydes (0.9 mmol). The resulting suspension was heated at 85 °C for 5 min, cooled down to r.t., filtered off, and washed with DMF and DCM (× 4 each solvent). Afterwards, the resins were treated with a solution of TFA-DCM (0.5:99.5) (2×10 min), and the resulting filtrates were combined and evaporated under pressure. **BDM** products were isolated as the free-amine compounds after silica-based SPE elution with DCM-MeOH (98:2) containing 1% NH₃ _{conc} in MeOH (full characterization data on Table S1). Ig Orange (BDM-69): ¹H-NMR $(300 \text{ MHz}, \text{CDCl}_3)$: 7.57-7.61-7.69 (m, 8H), 7.39 (d, 4H, J = 7.6Hz), 7.17 (d, 4H, J = 7.6\text{Hz}), 7.17 (d, 4H, J = 7.6\text{Hz}), 7.17 (d, 4H, J = 7.6\text{Hz}), 7.17 (d, 7.6\text{Hz}), 7.17 (d, 7.6\text{Hz})), 7.17 (d, 7.6\text{Hz}), 7.17 (d, 7.6\text{Hz})), 7.18 (d, 7.6\text{Hz})), 7.18 (d, 7.6\text{Hz})), 8.8Hz), 6.80 (s, 1H), 6.50 (dd, 1H, J = 2.1Hz, J = 4.0Hz), 3.18 (d, 2H, J = 6.4Hz), 3.12 (d, 2H, J = 6.4Hz), 2.54 (s, 3H), 1.63 (bs, 2H). ¹³C-NMR (75 MHz, CDCl₃): 157.0, 143.9, 140.6, 139.2, 138.9, 137.8, 136.9, 135.0, 134.4, 129.8, 128.6, 128.2, 127.8, 127.8, 126.9, 126.5, 123.4, 122.1, 119.7, 118.0, 116.0, 77.2, 43.4, 31.8, 29.5, 16.2. ¹⁹F-NMR (282 MHz, CDCl₃): -65.82, -66.03 (dd, J = 30Hz, J = 59Hz, BF₂).

Quantum Yield Measurements. Quantum yields were calculated by measuring the integrated emission area of the fluorescent spectra, and referring them to the area measured for Rhodamine B in EtOH after excitation at 510 nm ($\Phi_{\text{Rho-B}} = 0.70$). Quantum yields for the **BDM** products were then calculated using equation below, where *F* represents the area of fluorescent emission, *n* is reflective index of the solvent, and *Abs* is

absorbance at excitation wavelength selected for standards and samples. Emission was integrated from 540 nm to 800 nm.

$$\Phi_{flu}^{sample} = \Phi_{fl}^{reference} \left(\frac{F^{sample}}{F^{reference}} \right) \left(\frac{\eta^{sample}}{\eta^{reference}} \right) \left(\frac{Abs^{reference}}{Abs^{sample}} \right)$$

Screening of the BDM Library. Fluorescence intensities were measured using a Spectra Max Gemini XSF plate reader in a 384-well plate. BDM compounds were dissolved to a final concentration of 10 μ M (20 mM HEPES buffer, pH 7.4, containing 1% DMSO) and incubated with different proteins and peptides at different serial concentrations in 20 mM HEPES buffer (pH 7.4). The excitation wavelength was set at 510 nm, and the emission spectra were recorded from 560 to 700 nm. Fluorescence fold increase ratios were determined by referring the maximum fluorescence intensity emission of BDM compounds in the presence of the screened proteins and peptides to the maximum fluorescence intensity emission of BDM compounds in 20 mM HEPES buffer (pH 7.4).

Surface Plasmon Resonance. Human IgG (5.0 mg, 0.033 μ mol) and biotin-OSu (0.11 mg, 0.33 μ mol) were dissolved in 0.1 M NaHCO₃ (pH 8.5) and shaken for 1 h at 25 °C. Excess of biotinylating reagent was removed by centrifugation with Microcon 30K filters (3 rounds at 14,000 rpm for 20 min at 4 °C). The purified biotinylated IgG was resuspended in PBS, and characterized by SDS-PAGE and Western blotting with HRP-conjugated streptavidin (Figure S6 in SI). The concentration of biotinylated IgG before fluorescence and SPR experiments was determined by Bradford assay. SPR measurements were performed on a T-200 instrument (Biacore AB, GE Healthcare) equipped with Series S sensor chip CM7 (Biacore AB, GE Healthcare). All experiments were performed at 25 °C as previously described.³ Briefly, the carboxymethylated sensor surface was activated for 7 min with 1:1 of 0.2 M EDC (*N*-ethyl-*N'*-[3-(diethylamino)

propyl] carbodiimide) and 50 mM NHS (*N*-hydroxysuccinimide), then human IgG was coupled to 12,000 RU and finally the surface was deactivated for 7 min with 1 M ethanolamine–HCl (pH 8.5). The immobilized immunoglobulin was probed with one positive- and one negative-binding fragment diluted into the running buffer (10 mM sodium phosphate buffer, 150 mM NaCl, 1% DMSO, pH 7.4) to ascertain its functionality. **Ig Orange** (0.07, 0.21, 0.62, 1.86, 5.57 and 16.7 μ M, in duplicates) was injected across a control surface and the immobilized surface serially. Sensorgrams obtained were DMSO calibrated and buffer- and reference-subtracted.⁴ Responses at equilibrium were plotted against analyte concentration and fit to a simple isotherm to obtain the affinity constant using Scrubber 2 software (BioLogics Inc. Australia).

Job Plot Analysis. Ig Orange (final concentrations: 0, 1.6, 3.2, 4.8, 6.4, 8.0, 9.6, 11.2, 12.8, 14.4, 16 μ M) and human IgG (final concentrations: 16, 14.4, 12.8, 11.2, 9.6, 8.0, 6.4, 4.8, 3.2, 1.6, 0 μ M) were incubated in PBS buffer (pH 7.3) containing 1% DMSO, and the fluorescence intensities of the different solutions were recorded on a SpectraMax M2 plate reader (excitation: 530 nm; emission: 590 nm). The Job plot is represented as [fluorescence fold increase × ratio Ig Orange] *vs.* ratio Ig Orange, with values as means (*n*=3) and error bars as standard deviations.

2. Chemical structures and characterization data for the BDM library.

Code	Structure	Mcalc.	Mexp.	tR (min)	purity	λ _{max} (ABS)	λ _{max} (EM)	φ	mg
BDM-1	NH2 NBN FF	441.3	422.0	3.74	87%	561	586	0.21	5.1
BDM-2	NH ₂ N _B N FF	394.3	375.0	3.96	88%	607	685	0.01	5.8
BDM-3		433.3	414.0	4.02	87%	600	683	0.05	5.9
BDM-4	NH2 PFF	563.4	543.9	5.48	95%	566	590	0.42	6.1
BDM-5	NH2 NBN FFF	427.3	408.0	5.16	96%	561	575	0.50	7.7
BDM-6	NH2 NBN FF	451.3	432.0	4.97	89%	536	682	0.12	9.0

Table S1. Chemical structures and characterization data for the **BDM** library.

	NH-	105.3		4.50	00~	F / /	501	0.07	
BDM-7		487.3	467.9	4.68	89%	566	591	0.37	6.6
BDM-8	ZZ Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z	417.3	418.0	2.31	95%	561	575	0.34	3.9
BDM-9	NH2 NBN FF	365.3	346.0	4.21	88%	554	567	0.33	3.5
BDM-12	NH2 FF Q	385.3	365.9	4.46	90%	556	567	0.32	6.2
BDM-14	NH2 NBN FF	518.3	498.9	6.18	96%	593	624	0.02	7.8
BDM-16	NH2 FFFF	485.4	466.0	5.46	94%	561	574	0.14	4.9
BDM-18	NH ₂ N _B N FF	397.3	378.0	4.61	100%	567	588	0.33	6.7

	NH ₂	393.3	374.0	4.91	90%	555	569	0.71	8.0
BDM-19									
BDM-22	NH2 NBN FF	341.3	322.0	3.61	85%	567	584	0.14	5.8
BDM-30	NH2 NBN FF	439.3	419.9	4.99	86%	596	634	0.11	4.4
BDM-32	NH2 NBN FF	431.3	412.0	4.62	91%	576	624	0.25	5.6
BDM-33	H ₂ H _B N FF	381.3	362.0	3.69	88%	564	580	0.14	9.7
BDM-34	NH ₂ NB FF	415.3	396.0	4.98	92%	547	581	0.65	7.4
BDM-36	NH ₂ N _B N FF	415.3	396.0	5.02	94%	568	595	0.23	6.9
BDM-38	HH2 HBN FF	432.3	411.9	3.71	87%	565	579	0.06	9.8

	NH ₂	427.3	408.0	4.94	88%	554	567	0.53	5.6
BDM-40	FF GG								
BDM-42	NH2 NBF FF	473.3	454.0	5.06	93%	554	563	0.25	4.3
BDM-43	NH2 HAN FF	423.3	404.0	4.55	97%	564	579	0.21	4.5
BDM-45	NH2 SNBN FFF	451.3	432.0	5.47	93%	564	583	0.74	4.5
BDM-46	NH2 NBN FF	423.3	404.0	5.30	94%	564	583	0.53	6.0
BDM-48	NH2 NBN FF	407.3	388.0	4.46	98%	563	583	0.83	4.0
BDM-49	NH ₂ N _B N FF	379.2	360.0	3.76	92%	562	572	0.24	5.7

	NH ₂	439.3	420.0	4.07	87%	568	600	0.46	5.9
BDM-53	N,BN FF	439.3	420.0	4.07	0170	508	000	0.40	3.9
BDM-54	NH2 NBR FF	460.3	439.9 441.9	4.39	88%	559	573	0.52	6.2
BDM-61	NH2 NBN FF	431.3	411.9	4.73	91%	576	610	0.22	8.7
BDM-62	Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z	411.3	392.0	3.58	89%	567	594	0.43	5.2
BDM-63	NH ₂ N _F F F	425.3	406.0	4.01	94%	568	597	0.54	6.5
BDM-65	S S S S S S S S S S S S S S S S S S S	433.3	413.9	5.21	84%	589	615	0.15	7.2
BDM-68	NH2 P, BF FF	435.3	415.9	4.90	87%	551	561	0.63	5.5

	NH ₂	452.2	424.0	5.01	94%	571	594	0.22	06
BDM-69		453.3	434.2			574		0.32	8.6
BDM-70	Ź J Z L O Z L O J J J J J J J J J J J J J J J J J J J	461.3	442.0	4.82	93%	582	642	0.13	7.1
BDM-73	NH2 NH2 N BR F H OH	446.3	425.8 427.8	3.89	84%	561	575	0.28	4.4
BDM-75	NH2 NH2 FF	423.3	404.0	4.64	88%	560	585	0.53	4.4
BDM-76	NH ₂ Y, B, N, F, F	439.3	420.0	5.20	84%	572	588	0.25	4.0
BDM-77	NH ₂ Y _B N FF	393.3	374.0	5.03	84%	557	569	0.54	5.2
BDM-78	NH ₂ N _B N FF	439.3	420.0	4.55	87%	591	n.d	0.01	5.0

	NH ₂	428.3	409.0	3.57	95%	565	576	0.21	4.9
BDM-88	N N N N N N N N N N N N N N N N N N N	12010	429.0		<i></i>	202		0.21	,
BDM-90	NH2 NBN FF	457.3	438.0	5.18	97%	565	582	0.26	7.3
BDM-91	NH2 NBN FF	424.3	405.0	3.59	95%	614	699	0.02	7.0
BDM-93	NH2 NBN FF	420.3	401.0	3.95	91%	592	653	0.06	4.0
BDM-97	NH ₂ N _B N FF OH	397.3	378.0	3.68	89%	561	572	0.04	6.8
BDM-107	NH2 NBN FF	422.3	403.0	4.22	97%	617	681	0.05	4.7
BDM-108	NH ₂ N _B N FF	393.3	374.1	4.86	85%	549	573	0.35	6.2

	NH ₂	201.2	262.0	A 11	0107	5(2	500	0.95	25
BDM-110	NBN FF	381.3	362.0	4.11	91%	563	583	0.85	3.5
BDM-111	EF2 PEFF FF FF	408.3	389.0	3.11	100%	566	589	0.79	4.7
BDM-132	HO O	397.3	378.0	3.36	100%	568	594	0.05	4.5
BDM-135	N.B.F.F	355.3	336.0	3.98	93%	577	601	0.10	4.4
BDM-137	NH2 NH2 FF	401.3	382.0	4.84	87%	563	575	0.28	7.1
BDM-139	NH2 NBN FF	395.2	376.0	3.94	92%	564	584	0.55	3.7
BDM-140	NH2 N.B.N. F.F.	409.3	390.0	4.97	93%	564	585	0.57	5.6
BDM-143	NH2 NF2 PFF	365.2	346.0	4.51	84%	554	566	0.42	6.1

	NH2	441.2	422.0	267	0.407	570	612	0.05	51
BDM-144		441.3	422.0	3.67	94%	578		0.05	5.4
BDM-177		407.3	388.0	5.03	96%	556	567	0.78	4.1
BDM-178	9-0-0 9-0-0	473.2	454.0	5.06	96%	561	577	0.37	5.0
BDM-186	NH2 N.B.N. F.F.	381.2	362.0	4.15	97%	553	564	0.97	5.4
BDM-192	HO HO HO HO HO HO HO HO HO HO HO HO HO H	397.2	378.0	3.75	86%	574	605	0.20	4.0
BDM-195	NH2 NH2 NH2 FFF	481.7	461.9	5.18	100%	600	663	0.03	6.5
BDM-101	tr	443.3	423.9	5.23	97%	559	576	0.66	5.3
BDM-100	NH2 FFF	436.1	415.8 417.8	4.49	86%	571	582	0.43	5.0

	NH ₂	443.3	424.0	5.09	94%	554	564	0.47	4.4
BDM-20	Solution of the second	443.3	424.0	5.09	94 70	554	J0 4	0.47	4.4
BDM-202		407.3	388.0	4.64	85%	559	575	0.52	4.1
BDM-231	N-F-F-F-F-F-F-F-F-F-F-F-F-F-F-F-F-F-F-F	395.2	376.0	3.94	93%	558	568	0.17	5.9
BDM-198		381.2	362.0	4.08	95%	559	574	0.59	5.2
BDM-218	NH2 NBM	457.3	438.0	5.18	100%	559	574	0.37	5.8
BDM-179	NH2 NH2 P F F F F	485.3	466.0	5.09	96%	566	581	0.26	5.5
BDM-199	NH2 FF	429.3	410.0	3.79	94%	562	574	0.18	7.6

	NH ₂	385.7	366.0	4.47	88%	553	566	0.24	3.9
BDM-207		365.7	300.0	4.47	00 //	333	500	0.24	5.7
BDM-209	NH2 H, BN FF	411.3	392.0	4.09	93%	555	567	0.38	4.1
BDM-237	NH2 FFF	477.1	457.8	4.69	97%	558	572	0.35	5.2
BDM-228	Br ()	490.2	469.8 471.9	4.48	92%	557	567	0.24	8.8
BDM-236	NH2 NH2 F F	371.3	352.0	4.29	90%	574	594	0.22	4.9
BDM-208	NH2 N.B.N. F.F.	365.2	346.1	4.54	100%	558	572	0.27	3.8
BDM-27	NH2 NB/N/FF	351.2	332.0	4.01	87%	553	566	0.39	4.2
BDM-37	NH2 N ig/N F F	407.3	387.9	4.81	84%	566	588	0.15	4.5

	NH2	452.4	452.0	270	0001	566	502	0.41	54
BDM-163	Jan Harris	452.4	453.0	2.78	88%	566	586	0.41	5.6
BDM-239	Z Z Z Z Z Z B B F F F F F F F F F F F F	399.2	380.0	4.42	100%	559	572	0.28	5.5
BDM-240	NH2 NH2 NH2 NH2 NH2 F	417.2	397.9	4.48	87%	552	565	0.23	5.5
BDM-241	N- N- P- Br Br	539.0	517.8	4.88	84%	560	576	0.32	6.5
BDM-243	H- Z- Z- F F F	379.3	360.0	4.72	89%	543	565	0.87	3.0
BDM-245	NHE - D - P - P - P - P - P	399.2	380.0	4.32	87%	555	563	0.32	4.4
BDM-246	NH2 N + B F F C	417.7	397.9	4.93	100%	554	566	0.17	5.8
BDM-247	NH2 H2 FFF FFF	490.2	469.9 471.9	4.48 4.67	90%	571	599	0.54	5.6

	NH ₂	202.2	264.0	4 70	1000	557	5(0	0.24	4.0
BDM-249	F F F	383.2	364.0	4.72	100%	557	568	0.24	4.8
BDM-251	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	379.3	360.0	4.84	100%	556	572	0.22	5.9
BDM-252		424.3	405.0	4.86	91%	556	569	0.08	4.6
BDM-259	Br - O O	490.2	469.8 471.8	4.70	87%	559	573	0.35	5.0
BDM-260	NH2 NF2 Br	460.1	439.8 441.8	4.55	93%,	564	582	0.05	4.0
BDM-263		407.3	388.0	5.69	85%	559	571	0.26	3.9
BDM-267	Bry F F	444.1	423.9 425.9	4.96	92%	558	573	0.21	4.4

	NH2	1157	205.0	1 10	0.207	560	577	0.46	2.0
BDM-274		415.7	395.9	4.48	93%	562	577	0.46	3.9
BDM-275		431.7	411.9	4.30	85%	562	n.d	0.05	5.9
BDM-277		477.7	457.9	5.71	87%	559	574	0.59	4.9
BDM-281		415.7	395.9	4.13	92%	554	567	0.28	5.1
BDM-282		445.7	425.9	4.74	88%	558	565	0.19	4.5
BDM-283	NH₂ + ġ.N. + F F	401.7	381.9	3.68	100%	556	567	0.15	4.3
BDM-284		451.7	431.9	5.23	84%	594	622	0.03	4.6

	NH ₂	399.7	379.9	4.74	86%	544	561	0.51	6.0
BDM-285									
BDM-289		399.7	380.0	4.97	84%	559	570	0.26	4.9
BDM-290		477.7	457.9	5.61	85%	558	574	0.36	4.5
BDM-298	¢	473.3	454.0	5.30	89%	559	568	0.32	6.8
BDM-299	a o	512.2	491.8 493.8	5.89	91%	555	563	0.27	7.1
BDM-300	NH2 H BNN F F	457.3	438.0	5.65	89%	562	579	0.44	5.3
BDM-301	NH2 H-B-N H-B-F H-F-F	461.3	442.0	4.76	100%	563	586	0.41	5.6

[NH2	177 1	1570	4.02	1000	555	577	0.21	56
BDM-303		477.1	457.8	4.92	100%	555	566	0.31	5.6
BDM-305		429.2	410.0	3.99	86%	567	587	0.25	8.1
BDM-307		427.3	408.0	3.50	84%	564	580	0.07	4.0
BDM-308	NH2 H BN F F	512.2	491.8	6.15	86%	556	567	0.19	4.9
BDM-310	N F F	409.3	390.0	4.90	84%	566	600	0.42	6.5
BDM-311	NF2 NF2 NF F F F F	461.3	441.9	5.28	84%	556	565	0.21	5.1
BDM-312	N − N − N − N − N − N − N − N − N − N −	399.2	380.0	4.36	83%	557	568	0.18	3.0

	NH ₂	399.2	380.0	4.26	100%	556	570	0.10	4.7
BDM-314	F F								
BDM-316		439.3	420.0	4.81	87%, 4.81	572	610	0.46	5.9
BDM-318	OH	385.2	366.0	3.83	85%, 3.83	563	576	0.11	7.4
BDM-320	NH2 NH2 NHBN FFF	425.3	406.0	4.46	84%	558	569	0.21	4.3
BDM-322	Nte 	461.3	441.9	5.37	85%	559	577	0.15	5.5
BDM-323	NH2 H-B-M F-F	395.3	376.0	4.61	87%	556	570	0.44	5.0
BDM-325	NH2 N.B.N. FF	493.4	474.0	6.59	85%	557	573	0.47	5.3
BDM-326		417.2	398.0	4.40	91%	554	565	0.71	5.9

[NH ₂	562 4	542.0	6.05	0107	551	566	0.65	0.2
BDM-331	Co-Co-Co-Co-Co-Co-Co-Co-Co-Co-Co-Co-Co-C	563.4	543.9	6.05	84%	554	566	0.65	9.3
BDM-332	NH2 NH2 F F F	431.2	411.9	4.83	91%	552	562	0.61	5.3
BDM-333		397.3	378.0	4.51	95%	557	575	0.30	5.6
BDM-335	NH2 X- Z- F- F- F- F	395.3	376.0	4.57	95%	565	591	0.78	6.2
BDM-336	P. B.N. F.F	395.3	376.0	4.53	91%	560	577	0.58	3.4
BDM-340	NH2 FF	465.3	445.9	5.43 5.57	96%	534	541	0.11	5.0
BDM-341	F-C)-O-C-C-C-C-C-C-C-C-C-C-C-C-C-C-C-C-C-	461.3	441.9	5.09	90%	556	570	0.37	12. 4

	NH ₂	202.2	274.0	5.00	0107	550	577	0.29	0.6
BDM-343		393.3	374.0	5.08	91%	559	577	0.28	9.6
BDM-345	The second secon	467.2	447.9	4.92	88%	551	564	0.22	7.4
BDM-347	N- , is F F F	425.3	406.0	4.67	96%	566	591	0.57	5.3
BDM-350	N- , BF F NO2	426.2	406.9	3.93	98%	556	571	0.14	5.9
BDM-356	NH ₂ N ₁ N ₅ N F F	479.3	460.0	5.16	92%	573	682	0.03	4.2
BDM-357	NH2 N, j, N, F, F NO2 N	465.3	446.0	4.72	90%	585	687	0.04	5.0
BDM-359	NH ₂ N: B'N F F	399.2	380.0	4.33	100%	559	574	0.83	6.6

	NH ₂	451.2	421.0	4.00	0607	550	564	0.51	61
BDM-363	H B F F F F F F F F F F F F F F F F F F	451.3	431.8	4.98	96%	553	564	0.51	6.1
BDM-365	F F F	435.2	415.9	4.77	96%	549	563	0.24	5.2
BDM-369	N-B- P- F- F- S	357.2	337.9	3.96	93%	554	570	0.30	3.6
BDM-371		451.7	431.9	5.35	90%	590	630	0.03	4.8
BDM-372		445.7	425.9	4.34	87%	561	578	0.55	8.5
BDM-377	NH2 N, B, N F F	444.3	445.0	4.76	98%	586	n.d	0.04	6.9
BDM-378	H2 H2 H2 H2 H H H H H H H H H H H H H H	475.3	456.0	5.14	94%	563	585	0.41	3.8

	NH ₂	379.2	360.0	3.86	91%	554	563	0.31	5.3
BDM-391									
BDM-397	NH ₂ N _{*B} N FF	411.2	392.0	3.67	92%	570	614	0.10	4.9
BDM-401	A C C C C C C C C C C C C C C C C C C C	469.3	449.9	4.28	100%	565	590	0.85	5.6
BDM-409	NH ₂ N _F F F Br	430.1	409.9 411.8	4.93	85%	558	570	0.12	4.0
BDM-425	NH2 H, BR F F	445.3	425.9	5.27	89%	562	576	0.12	4.4
BDM-429	NH2 H, B, N, F F	395.3	376.0	4.49	100%	564	586	0.23	4.6
BDM-430	NH2 N× B F F	383.3	364.0	3.09	100%	565	592	0.99	3.3

	NH ₂	499.4	480.0	6.14	94%	555	565	0.39	4.9
BDM-435	HO-0								
BDM-437	NH2 NH2 FFF	403.6	383.9	4.62	95%	555	566	0.21	4.1
BDM-440	NH2 N.B.N. F.F.	451.3	432.0	5.42	93%	563	589	0.37	6.1
BDM-89	NH2 YBN FF	427.3	408.0	5.15	85%	576	612	0.30	6.1
BDM-384	NH2 NH2 NFF	477.1	457.8	4.60	92%	551	565	0.32	3.1
BDM-105	NH2 FFF	367.2	348.0	3.48	93%	556	567	0.18	4.4
BDM-82	N the point of the	441.3	422.0	4.36	87%	579	621	0.40	6.6

	NH ₂	478.4	459.0	6.17	89%	622	695	0.03	4.0
BDM-17	N HEF		479.0						

HPLC conditions: A: H₂O-HCOOH: 99.9:0.1. B: ACN-HCOOH: 99.9:0.1; gradient 5% B to 95% B (10 min), isocratic 95% B (2 min). Reverse-phase Agilent C₁₈ Zorbax column (2.1 x 30 mm²) 3.5 μ m, flow rate: 1 mL/min. Purity determined by integration of the absorbance peaks at 350 nm. ESI (+) *m/z* signals mostly correspond to the [M-F] fragmentation.

3. Characterization data for Ig Orange (BDM-69).

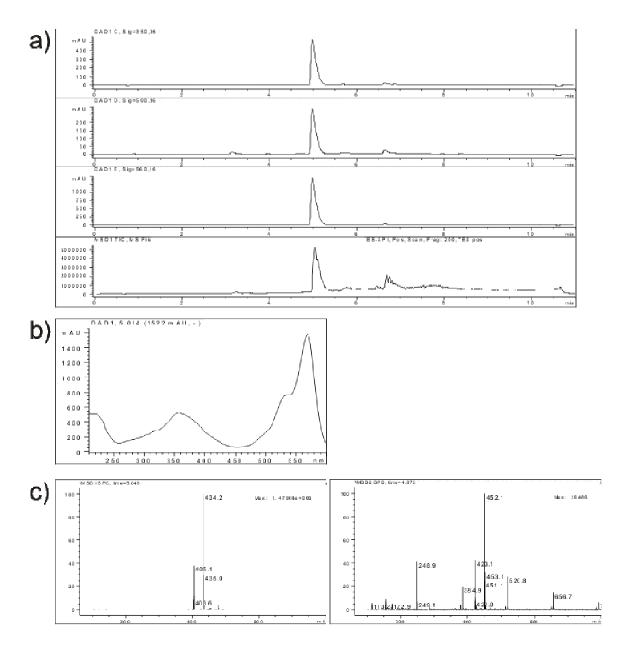


Figure S1. HPLC-MS characterization of Ig Orange (BDM-69). a) Chromatograms (*descending order*) at 350 nm, 500 nm, 560 nm and MS-TIC; b) absorbance spectra; c) *left, right*: ESI-positive and ESI-negative MS spectra, respectively.

4. Extended screening of Ig Orange (BDM-69).

Fluorescence intensities were measured using a Spectra Max Gemini XSF plate reader in a 384-well plate. **IgG orange** was dissolved to a final concentration of 10 μ M (20 mM HEPES buffer, pH 7.4, containing 1% DMSO) and incubated with different biomolecules at 4 serial concentrations (*see below*) in 20 mM HEPES buffer (pH 7.4). The excitation wavelength was set at 510 nm, and the emission spectra were recorded from 560 to 700 nm. Fluorescence increase ratios were determined by referring the fluorescence intensity emission (590 nm) of **Ig Orange** in the presence of the screened analytes to the fluorescence intensity emission of **Ig Orange** in the absence of the screened analytes.

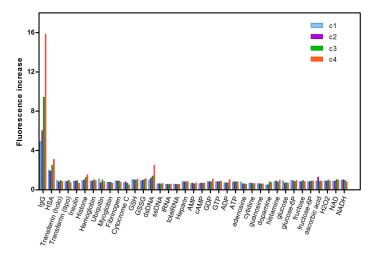


Figure S2. Fluorescence response of **Ig Orange** against an extended set of macromolecules and small molecules in 20 mM HEPES buffer, pH 7.4 (fold increase determined at 590 nm). Analyte serial concentrations (c1-c4, ascending concentration order): proteins, peptides, dsDNA, ssDNA, tRNA, total RNA and heparin: 0.125, 0.25, 0.5 and 1 mg/mL; AMP, cAMP, GDP, GTP, ADP, ATP, adenosine, cytidine, guanosine: 12.5, 25, 50, 100 μ M; dopamine, histamine: 0.04, 0.2, 1 and 5 mg/mL; glucose, glucose-6P, fructose, fructose-6P: 0.08, 0.4, 2, 10 mM; ascorbic acid, H₂O₂, NAD, NADH: 0.04, 0.2, 1, 5 mM.

IgG Orange was dissolved to a final concentration of 10 μ M (PBS buffer, pH 7.3, containing 1% DMSO) and incubated with human IgG, IgA and IgM at different concentrations (*see below*) in PBS buffer (pH 7.3). The excitation wavelength was set at 530 nm, and fluorescence fold increase ratios were determined by referring the maximum fluorescence intensity emission (590 nm) of **Ig Orange** in the presence of immunoglobulins to the maximum fluorescence intensity emission of **Ig Orange** in buffer.

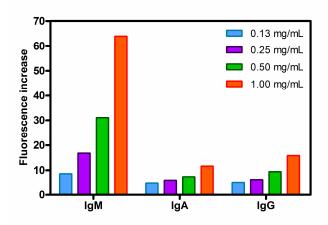


Figure S3. Fluorescence response of **IgG orange** upon incubation with different concentrations of human IgM, IgA and IgG.

5. Ig Orange response against IgG in human serum samples.

Known amounts of human IgG covering the whole physiological range (3.3, 6.2, 9.4 and 12.5 mg/mL) were added to immunoglobulin(G, A, M)-depleted human serum (Sigma Aldrich). The resulting samples were diluted 10 times in PBS buffer (pH 7.3) to reach a final volume of 100 μ L and **Ig Orange** (1 μ L, 1 mM) was added (final **Ig Orange** concentration: 10 μ M). Fluorescence intensities of the samples were recorded on a SpectraMax M2 plate reader (exc: 530 nm; emission: 590 nm). The experimental data was fitted to a linear regression using Graphpad Prism 5.0 software.

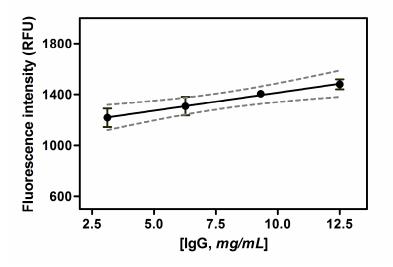


Figure S4. Fluorescence response of **Ig Orange** against known concentrations of IgG in immunoglobulin-depleted human serum. Values are represented as means (n=3) and errors bars as standard deviations. Dotted grey lines indicate a 95% reliability interval. Regression coefficient (\mathbb{R}^2): 0.99.

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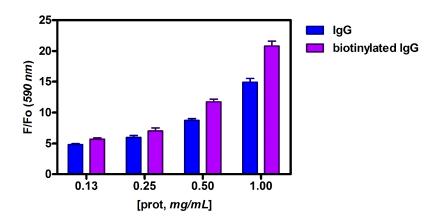


Figure S5. Comparative fluorescence increase of Ig Orange after incubation with different concentrations of non-modified and biotinylated human IgG. Values are represented as means (n=3) and error bars as standard deviations.

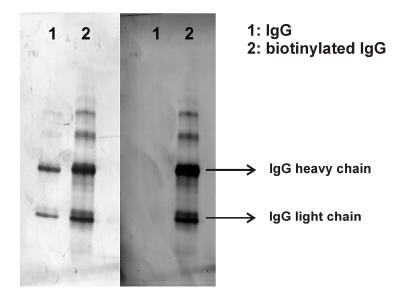


Figure S6. SDS-PAGE gel and Western blotting of human non-modified and biotinylated IgG. *Left*) samples were solved by SDS-PAGE and stained with Coomassie Blue; *right*) samples were solved by Western blotting and immunoblots were incubated with HRP-conjugated streptavidin.

7. Structure-fluorescence relationships of Ig Orange and additional SPR experiments.

Ig Orange (BDM-69) (10 μ M) and BD-69 (10 μ M) (previously synthesized as in ref. 20 in the manuscript) were incubated with different concentrations of human IgG in 20 mM HEPES buffer (pH 7.4) and the fluorescence intensities were recorded on a SpectraMax M2 plate reader (excitation: 530 nm; emission: 590 nm). Values are represented as means (*n*=3) of the fluorescence fold increase after incubation with the protein.

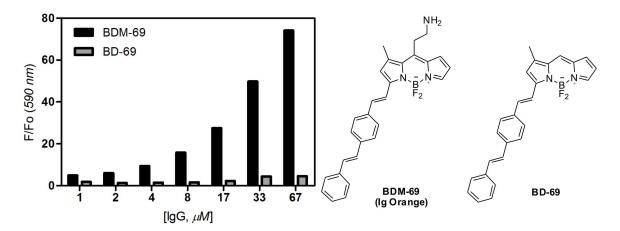


Figure S7. Fluorescence response of **Ig Orange** (**BDM-69**) and **BD-69** *vs.* human IgG. A 15-fold higher fluorescence increase was observed in the case of **Ig Orange** indicating the relevant contribution of the aminoethyl group in the interaction with human IgG.

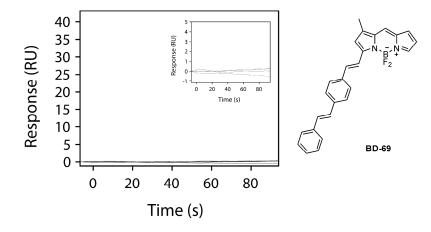


Figure S8. Binding sensorgrams of the aminoethyl-free compound (**BD-69**) upon injection across an immobilized immunoglobulin chip. Different concentrations of **BD-69** were injected across a control surface and the immobilized surface serially as described in the Experimental Section for **Ig Orange**. *Inset*) zoom image of the sensorgrams.

8. Binding of Ig Orange to polyclonal IgG and monoclonal IgG₁.

Ig Orange (10 μ M) was incubated with different concentrations of IgG from human serum (polyclonal IgG) and a monoclonal human IgG₁ in 20 mM HEPES buffer (pH 7.4). Fluorescence measurements were recorded on a SpectraMax M2 plate reader (excitation: 530 nm; emission: 590 nm).

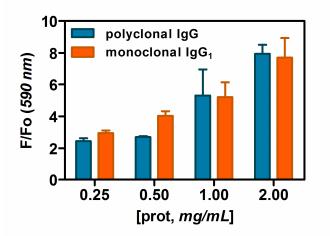


Figure S9. Fluorescence intensities of **Ig Orange** after incubation with polyclonal human IgG and monoclonal human IgG₁. Values are represented as means (n=3) and error bars as standard deviations.

9. Fractional saturation curve experiments of Ig Orange.

Ig Orange (10 μ M) was incubated with different concentrations of human IgG in 20 mM HEPES buffer (pH 7.4) and the fluorescence intensities were recorded on a SpectraMax M2 plate reader (excitation: 530 nm; emission: 590 nm).

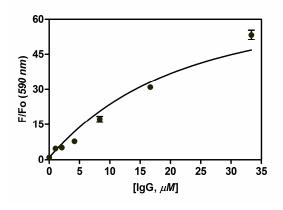


Figure S10. Fluorescence-based fractional saturation curve of **Ig Orange**. Values were represented as means (n=3) of the fluorescence fold increase after incubation with the protein, and adjusted to a binding saturation curve (GraphPad Prism 5.0) to estimate a dissociation constant K_D : 14.1 ± 0.9 µM.

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