
Supplementary information

A biosensor for the direct visualization of auxin

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SI Guide

A biosensor for direct visualisation of auxin

Ole Herud-Sikimić^{1*}, Andre C. Stiel^{1,2*}, Martina Kolb^{1*}, Sooruban Shanmugaratnam^{1,3},
Kenneth W. Berendzen⁴, Christian Feldhaus¹, Birte Höcker^{1,3^}, Gerd Jürgens^{1^}

Suppl. Table S1.

Table S1a. Data collection and refinement statistics (molecular replacement) for TrpR-Wt with ligand IAA

Table S1b. Data collection and refinement statistics (molecular replacement) for TrpR-S88Y with ligand IAA

Table S1c. Data collection and refinement statistics (molecular replacement) for TrpR-T44L-S88Y with ligand IAA

Table S1d. Data collection and refinement statistics (molecular replacement) for TrpR-T44L-T81M-S88Y with ligand IAA

Table S1e. Data collection and refinement statistics (molecular replacement) for TrpR-T44L-T81M-N87G-S88Y with ligand IAA

Table S1f. Data collection and refinement statistics (molecular replacement) for TrpR-M42F-T44L-T81M-N87G-S88Y with ligand IAA

Table S1g. Data collection and refinement statistics (molecular replacement) for TrpR-M42F-T44L-T81I-S88Y with ligand IAA

Table S1h. Data collection and refinement statistics (molecular replacement) for TrpR-M42F-T44L-T81I-S88Y with ligand IAN

Suppl. Table S2.

Table S2. Flow cytometry statistics

Table S1a Data collection and refinement statistics (molecular replacement) for TrpR-Wt with ligand IAA

TrpR-Wt IAA: 6EJW	
Data collection	
Space group	P 4 ₃
Cell dimensions	
<i>a</i> , <i>b</i> , <i>c</i> (Å)	81.31 81.31 72.00
α , β , γ (°)	90 90 90
Resolution (Å)	44.93 - 1.99 (2.06 - 1.99) ^a
<i>R</i> _{sym} or <i>R</i> _{merge}	0.094 (1.962)
<i>I</i> / σ <i>I</i>	11.55 (0.91)
Completeness (%)	99.45 (96.48)
Redundancy	6.5 (6.0)
Refinement	
Resolution (Å)	44.93 - 1.99 (2.06 - 1.99)
No. reflections	32088 (3099)
<i>R</i> _{work} / <i>R</i> _{free}	0.195 (0.309) / 0.243 (0.364)
No. atoms	
Protein	3293
Ligand/ion	130
Water	130
<i>B</i> -factors	
Protein	50.2
Ligand/ion	46.4
Water	51.5
R.m.s. deviations	
Bond lengths (Å)	0.008
Bond angles (°)	0.90

^a Numbers in brackets correspond to values in the highest resolution shell.

Table S1b Data collection and refinement statistics (molecular replacement) for TrpR-S88Y with ligand IAA

TrpR-S88Y IAA: 6EJZ	
Data collection	
Space group	P 4 ₃
Cell dimensions	
<i>a</i> , <i>b</i> , <i>c</i> (Å)	82.44 82.44 75.65
α , β , γ (°)	90 90 90
Resolution (Å)	41.22 - 1.90 (1.97 - 1.90) ^a
<i>R</i> _{sym} or <i>R</i> _{merge}	0.081 (1.816)
<i>I</i> / σ <i>I</i>	20.28 (1.47)
Completeness (%)	99.79 (98.67)
Redundancy	12.8 (12.6)
Refinement	
Resolution (Å)	41.22 - 1.90 (1.97 - 1.90)
No. reflections	40085 (3931)
<i>R</i> _{work} / <i>R</i> _{free}	0.184 (0.305) / 0.218 (0.337)
No. atoms	
Protein	3337
Ligand/ion	118
Water	220
<i>B</i> -factors	
Protein	43.8
Ligand/ion	49.4
Water	46.8
R.m.s. deviations	
Bond lengths (Å)	0.006
Bond angles (°)	0.69

^a Numbers in brackets correspond to values in the highest resolution shell.

Table S1c Data collection and refinement statistics (molecular replacement) for TrpR-T44L-S88Y with ligand IAA

TrpR-T44L-S88Y IAA: 6ENI	
Data collection	
Space group	P 2 ₁ 2 ₁ 2 ₁
Cell dimensions	
<i>a</i> , <i>b</i> , <i>c</i> (Å)	54.78 63.41 64.77
α , β , γ (°)	90 90 90
Resolution (Å)	34.91 - 1.10 (1.14 - 1.10) ^a
<i>R</i> _{sym} or <i>R</i> _{merge}	0.052 (0.701)
<i>I</i> / σ <i>I</i>	26.67 (3.85)
Completeness (%)	94.44 (80.40)
Redundancy	12.7 (11.4)
Refinement	
Resolution (Å)	34.91 - 1.10 (1.14 - 1.10)
No. reflections	87117 (7337)
<i>R</i> _{work} / <i>R</i> _{free}	0.158 (0.263) / 0.176 (0.263)
No. atoms	
Protein	1805
Ligand/ion	30
Water	469
<i>B</i> -factors	
Protein	14.1
Ligand/ion	13.1
Water	23.7
R.m.s. deviations	
Bond lengths (Å)	0.010
Bond angles (°)	1.08

^a Numbers in brackets correspond to values in the highest resolution shell.

Table S1d Data collection and refinement statistics (molecular replacement) for TrpR-T44L-T81M-S88Y with ligand IAA

TrpR-T44L-T81M-S88Y IAA: 6EKP	
Data collection	
Space group	P 2 ₁ 2 ₁ 2 ₁
Cell dimensions	
<i>a</i> , <i>b</i> , <i>c</i> (Å)	55.04 63.12 64.64
α , β , γ (°)	90 90 90
Resolution (Å)	41.48 - 1.46 (1.51 - 1.46) ^a
<i>R</i> _{sym} or <i>R</i> _{merge}	0.034 (0.770)
<i>I</i> / σ <i>I</i>	20.30 (1.97)
Completeness (%)	98.54 (94.76)
Redundancy	4.3 (4.2)
Refinement	
Resolution (Å)	41.48 - 1.46 (1.51 - 1.46)
No. reflections	39383 (3744)
<i>R</i> _{work} / <i>R</i> _{free}	0.172 (0.306) / 0.190 (0.332)
No. atoms	
Protein	1689
Ligand/ion	41
Water	250
<i>B</i> -factors	
Protein	35.0
Ligand/ion	30.8
Water	45.4
R.m.s. deviations	
Bond lengths (Å)	0.002
Bond angles (°)	0.50

^a Numbers in brackets correspond to values in the highest resolution shell.

Table S1e Data collection and refinement statistics (molecular replacement) for TrpR-T44L-T81M-N87G-S88Y with ligand IAA

TrpR-T44L-T81M-N87G-S88Y IAA: 6ENN	
Data collection	
Space group	P 2 ₁ 2 ₁ 2 ₁
Cell dimensions	
<i>a</i> , <i>b</i> , <i>c</i> (Å)	54.58 63.19 64.90
α , β , γ (°)	90 90 90
Resolution (Å)	41.77 - 1.17 (1.21 - 1.17) ^a
<i>R</i> _{sym} or <i>R</i> _{merge}	0.052 (0.659)
<i>I</i> / σ <i>I</i>	16.16 (1.90)
Completeness (%)	98.04 (87.61)
Redundancy	5.8 (4.1)
Refinement	
Resolution (Å)	41.77 - 1.17 (1.21 - 1.17)
No. reflections	74760 (6616)
<i>R</i> _{work} / <i>R</i> _{free}	0.148 (0.225) / 0.180 (0.264)
No. atoms	
Protein	1837
Ligand/ion	26
Water	347
<i>B</i> -factors	
Protein	15.7
Ligand/ion	12.8
Water	30.9
R.m.s. deviations	
Bond lengths (Å)	0.010
Bond angles (°)	1.10

^a Numbers in brackets correspond to values in the highest resolution shell.

Table S1f Data collection and refinement statistics (molecular replacement) for TrpR-M42F-T44L-T81M-N87G-S88Y with ligand IAA

TrpR-M42F-T44L-T81M-N87G-S88Y IAA: 6ELB	
Data collection	
Space group	P 2 ₁ 2 ₁ 2 ₁
Cell dimensions	
<i>a</i> , <i>b</i> , <i>c</i> (Å)	55.09 63.31 65.05
α , β , γ (°)	90 90 90
Resolution (Å)	45.37 - 1.44 (1.49 - 1.44) ^a
<i>R</i> _{sym} or <i>R</i> _{merge}	0.048 (1.086)
<i>I</i> / σ <i>I</i>	19.27 (1.37)
Completeness (%)	98.25 (93.32)
Redundancy	7.2 (6.9)
Refinement	
Resolution (Å)	45.37 - 1.44 (1.49 - 1.44)
No. reflections	41354 (3869)
<i>R</i> _{work} / <i>R</i> _{free}	0.154 (0.227) / 0.193 (0.316)
No. atoms	
Protein	1771
Ligand/ion	26
Water	309
<i>B</i> -factors	
Protein	24.3
Ligand/ion	18.7
Water	36.4
R.m.s. deviations	
Bond lengths (Å)	0.006
Bond angles (°)	0.82

^a Numbers in brackets correspond to values in the highest resolution shell.

Table S1g Data collection and refinement statistics (molecular replacement) for TrpR-M42F-T44L-T81I-S88Y with ligand IAA

TrpR- M42F-T44L-T81I-S88Y IAA: 6ELF	
Data collection	
Space group	P 2 ₁ 2 ₁ 2 ₁
Cell dimensions	
a, b, c (Å)	54.96 63.41 64.82
α, β, γ (°)	90 90 90
Resolution (Å)	31.70 - 1.83 (1.90 - 1.83) ^a
R_{sym} or R_{merge}	0.094 (1.985)
$I / \sigma I$	18.08 (1.22)
Completeness (%)	99.50 (96.29)
Redundancy	12.5 (11.2)
Refinement	
Resolution (Å)	31.70 - 1.83 (1.90 - 1.83)
No. reflections	20416 (1945)
$R_{\text{work}} / R_{\text{free}}$	0.177 (0.409) / 0.217 (0.421)
No. atoms	
Protein	1701
Ligand/ion	26
Water	155
<i>B</i> -factors	
Protein	35.6
Ligand/ion	31.3
Water	42.4
R.m.s. deviations	
Bond lengths (Å)	0.013
Bond angles (°)	1.20

^a Numbers in brackets correspond to values in the highest resolution shell.

Table S1h Data collection and refinement statistics (molecular replacement) for TrpR-M42F-T44L-T81I-S88Y with ligand IAN

TrpR- M42F-T44L-T81I-S88Y IAN: 6ELG	
Data collection	
Space group	P 2 ₁ 2 ₁ 2 ₁
Cell dimensions	
<i>a</i> , <i>b</i> , <i>c</i> (Å)	55.01 63.47 65.36
α , β , γ (°)	90 90 90
Resolution (Å)	42.09 - 1.38 (1.43 - 1.38) ^a
<i>R</i> _{sym} or <i>R</i> _{merge}	0.063 (0.772)
<i>I</i> / σ <i>I</i>	16.65 (1.83)
Completeness (%)	99.55 (96.20)
Redundancy	6.3 (5.8)
Refinement	
Resolution (Å)	42.09 - 1.38 (1.43 - 1.38)
No. reflections	47907 (4557)
<i>R</i> _{work} / <i>R</i> _{free}	0.143 (0.239) / 0.177 (0.295)
No. atoms	
Protein	1790
Ligand/ion	24
Water	387
<i>B</i> -factors	
Protein	16.8
Ligand/ion	12.8
Water	29.8
R.m.s. deviations	
Bond lengths (Å)	0.010
Bond angles (°)	0.99

^a Numbers in brackets correspond to values in the highest resolution shell.

Table S2. Flow cytometry statistics

Sample description	Y Param	Gate	Mean Y	Median Y	Gate2	% Hist	Sample #	Sample replicate	Dilution	Conc [fM]	FRET-ratio meanY	Conc [fM]	% Hist	Conc [fM]	FRET-ratio meanY	Conc [fM]	FRET-ratio %
Sample 189_H2O	(FL10-Log_Height/FL9-Log_Height)	Total	0	0	FRET-RESPONSE	0	189	x	0	0	0	0	0	0	0	0	0.00
Sample 190_AQUA	(FL10-Log_Height/FL9-Log_Height)	Total	290.88	295.52	FRET-RESPONSE	66.67	190	x	0	0	290.88	0	66.67	0	0	0	2.94
Sample 191_NEON	(FL10-Log_Height/FL9-Log_Height)	Total	394.34	338.59	FRET-RESPONSE	4.18	191	x	0	0	394.34	0	4.18	0	0	0	0.18
Sample 192_GFP	(FL10-Log_Height/FL9-Log_Height)	Total	987.72	666.08	FRET-RESPONSE	17.03	192	x	0	0	987.72	0	17.03	0	0	0	0.75
Sample 193_AUX_0.0	(FL10-Log_Height/FL9-Log_Height)	Total	312.95	309.17	FRET-RESPONSE	21.38	193	x	0	0	312.95	0	21.38	0	0	0	0.94
Sample 194_AUX_a.1	(FL10-Log_Height/FL9-Log_Height)	Total	749.86	636.68	FRET-RESPONSE	27.26	194	x	1	0.01	749.86	0.01	27.26	0.01	0.01	0.01	1.20
Sample 195_AUX_a.2	(FL10-Log_Height/FL9-Log_Height)	Total	670.34	608.57	FRET-RESPONSE	28.34	195	a	2	0.0025	670.34	0.0025	28.34	0.0025	2.11	0.0025	1.25
Sample 196_AUX_a.3	(FL10-Log_Height/FL9-Log_Height)	Total	597.79	556.03	FRET-RESPONSE	26.88	196	a	3	0.000625	597.79	0.000625	26.88	0.000625	1.88	0.000625	1.18
Sample 197_AUX_a.4	(FL10-Log_Height/FL9-Log_Height)	Total	452.96	424.08	FRET-RESPONSE	25.66	197	a	4	0.00015625	452.96	0.00015625	25.66	0.00015625	1.42	0.00015625	1.13
Sample 198_AUX_a.5	(FL10-Log_Height/FL9-Log_Height)	Total	377.88	370.36	FRET-RESPONSE	23.66	198	a	5	3.90625E-05	377.88	3.90625E-05	23.66	3.90625E-05	1.19	3.90625E-05	1.04
Sample 199_AUX_a.6	(FL10-Log_Height/FL9-Log_Height)	Total	340.12	338.39	FRET-RESPONSE	23.64	199	a	6	9.76563E-06	340.12	9.76563E-06	23.64	9.76563E-06	1.07	9.76563E-06	1.04
Sample 200_AUX_a.7	(FL10-Log_Height/FL9-Log_Height)	Total	330.7	323.45	FRET-RESPONSE	23.39	200	a	7	2.44141E-06	330.7	2.44141E-06	23.39	2.44141E-06	1.04	2.44141E-06	1.03
Sample 201_AUX_a.8	(FL10-Log_Height/FL9-Log_Height)	Total	325.35	323.45	FRET-RESPONSE	23.35	201	a	8	6.10352E-07	325.35	6.10352E-07	23.35	6.10352E-07	1.02	6.10352E-07	1.00
Sample 202_AUX_a.9	(FL10-Log_Height/FL9-Log_Height)	Total	320.86	323.45	FRET-RESPONSE	22.72	202	a	9	1.52588E-07	320.86	1.52588E-07	22.72	1.52588E-07	1.01	1.52588E-07	1.00
Sample 203_AUX_a.10	(FL10-Log_Height/FL9-Log_Height)	Total	326.59	323.45	FRET-RESPONSE	22.69	203	a	10	3.8147E-08	326.59	3.8147E-08	22.69	3.8147E-08	1.03	3.8147E-08	1.00
Sample 204_AUX_a.11	(FL10-Log_Height/FL9-Log_Height)	Total	321.15	323.45	FRET-RESPONSE	24.06	204	a	11	9.53674E-09	321.15	9.53674E-09	24.06	9.53674E-09	1.01	9.53674E-09	1.06
Sample 205_AUX_a.12	(FL10-Log_Height/FL9-Log_Height)	Total	324.87	323.45	FRET-RESPONSE	22.95	205	a	12	2.38419E-09	324.87	2.38419E-09	22.95	2.38419E-09	1.02	2.38419E-09	1.01
Sample 206_AUX_b.1	(FL10-Log_Height/FL9-Log_Height)	Total	737.1	636.68	FRET-RESPONSE	28.63	206	b	1	0.01	737.1	0.01	28.63	0.01	2.32	0.01	1.26
Sample 207_AUX_b.2	(FL10-Log_Height/FL9-Log_Height)	Total	673.65	608.57	FRET-RESPONSE	27.49	207	b	2	0.0025	673.65	0.0025	27.49	0.0025	2.12	0.0025	1.21
Sample 208_AUX_b.3	(FL10-Log_Height/FL9-Log_Height)	Total	593.06	556.03	FRET-RESPONSE	28.33	208	b	3	0.000625	593.06	0.000625	28.33	0.000625	1.86	0.000625	1.25
Sample 209_AUX_b.4	(FL10-Log_Height/FL9-Log_Height)	Total	461.27	443.67	FRET-RESPONSE	26.26	209	b	4	0.00015625	461.27	0.00015625	26.26	0.00015625	1.48	0.00015625	1.16
Sample 210_AUX_b.5	(FL10-Log_Height/FL9-Log_Height)	Total	376.13	370.36	FRET-RESPONSE	24.93	210	b	5	3.90625E-05	376.13	3.90625E-05	24.93	3.90625E-05	1.18	3.90625E-05	1.10
Sample 211_AUX_b.6	(FL10-Log_Height/FL9-Log_Height)	Total	343.42	338.39	FRET-RESPONSE	23.94	211	b	6	9.76563E-06	343.42	9.76563E-06	23.94	9.76563E-06	1.08	9.76563E-06	1.06
Sample 212_AUX_b.7	(FL10-Log_Height/FL9-Log_Height)	Total	327.81	323.45	FRET-RESPONSE	23.5	212	b	7	2.44141E-06	327.81	2.44141E-06	23.5	2.44141E-06	1.03	2.44141E-06	1.04
Sample 213_AUX_b.8	(FL10-Log_Height/FL9-Log_Height)	Total	320.02	323.45	FRET-RESPONSE	22.93	213	b	8	6.10352E-07	320.02	6.10352E-07	22.93	6.10352E-07	1.01	6.10352E-07	1.01
Sample 214_AUX_b.9	(FL10-Log_Height/FL9-Log_Height)	Total	320.66	323.45	FRET-RESPONSE	22.96	214	b	9	1.52588E-07	320.66	1.52588E-07	22.96	1.52588E-07	1.01	1.52588E-07	1.01
Sample 215_AUX_b.10	(FL10-Log_Height/FL9-Log_Height)	Total	322.21	323.45	FRET-RESPONSE	23.1	215	b	10	3.8147E-08	322.21	3.8147E-08	23.1	3.8147E-08	1.01	3.8147E-08	1.02
Sample 216_AUX_b.11	(FL10-Log_Height/FL9-Log_Height)	Total	321.47	323.45	FRET-RESPONSE	23.33	216	b	11	9.53674E-09	321.47	9.53674E-09	23.33	9.53674E-09	1.01	9.53674E-09	1.03
Sample 217_AUX_b.12	(FL10-Log_Height/FL9-Log_Height)	Total	323.82	323.45	FRET-RESPONSE	23.43	217	b	12	2.38419E-09	323.82	2.38419E-09	23.43	2.38419E-09	1.02	2.38419E-09	1.03
Sample 218_AUX_c.1	(FL10-Log_Height/FL9-Log_Height)	Total	731.46	636.68	FRET-RESPONSE	28.08	218	c	1	0.01	731.46	0.01	28.08	0.01	2.30	0.01	1.24
Sample 219_AUX_c.2	(FL10-Log_Height/FL9-Log_Height)	Total	669.51	608.57	FRET-RESPONSE	28.21	219	c	2	0.0025	669.51	0.0025	28.21	0.0025	2.10	0.0025	1.24
Sample 220_AUX_c.3	(FL10-Log_Height/FL9-Log_Height)	Total	599.75	556.03	FRET-RESPONSE	28.05	220	c	3	0.000625	599.75	0.000625	28.05	0.000625	1.89	0.000625	1.24
Sample 221_AUX_c.4	(FL10-Log_Height/FL9-Log_Height)	Total	473.14	443.67	FRET-RESPONSE	26.17	221	c	4	0.00015625	473.14	0.00015625	26.17	0.00015625	1.49	0.00015625	1.15
Sample 222_AUX_c.5	(FL10-Log_Height/FL9-Log_Height)	Total	382.84	370.36	FRET-RESPONSE	25.36	222	c	5	3.90625E-05	382.84	3.90625E-05	25.36	3.90625E-05	1.20	3.90625E-05	1.12
Sample 223_AUX_c.6	(FL10-Log_Height/FL9-Log_Height)	Total	341.03	338.39	FRET-RESPONSE	24.15	223	c	6	9.76563E-06	341.03	9.76563E-06	24.15	9.76563E-06	1.07	9.76563E-06	1.06
Sample 224_AUX_c.7	(FL10-Log_Height/FL9-Log_Height)	Total	326.56	323.45	FRET-RESPONSE	23.68	224	c	7	2.44141E-06	326.56	2.44141E-06	23.68	2.44141E-06	1.03	2.44141E-06	1.04
Sample 225_AUX_c.8	(FL10-Log_Height/FL9-Log_Height)	Total	318.71	309.17	FRET-RESPONSE	23.02	225	c	8	6.10352E-07	318.71	6.10352E-07	23.02	6.10352E-07	1.00	6.10352E-07	1.01
Sample 226_AUX_c.9	(FL10-Log_Height/FL9-Log_Height)	Total	319.9	323.45	FRET-RESPONSE	23.54	226	c	9	1.52588E-07	319.9	1.52588E-07	23.54	1.52588E-07	1.01	1.52588E-07	1.04
Sample 227_AUX_c.10	(FL10-Log_Height/FL9-Log_Height)	Total	322.59	323.45	FRET-RESPONSE	23.83	227	c	10	3.8147E-08	322.59	3.8147E-08	23.83	3.8147E-08	1.01	3.8147E-08	1.05
Sample 228_AUX_c.11	(FL10-Log_Height/FL9-Log_Height)	Total	318.1	309.17	FRET-RESPONSE	23.82	228	c	11	9.53674E-09	318.1	9.53674E-09	23.82	9.53674E-09	1.00	9.53674E-09	1.05
Sample 229_AUX_c.12	(FL10-Log_Height/FL9-Log_Height)	Total	319.87	309.17	FRET-RESPONSE	23.6	229	c	12	2.38419E-09	319.87	2.38419E-09	23.6	2.38419E-09	1.01	2.38419E-09	1.04

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