# **Supplementary Information**

# Algorithm-aided engineering of aliphatic halogenase WelO5\* for the asymmetric late-stage functionalization of soraphens

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# I. Supplementary Tables

Natural enzymes			
Name	Source organism		
Bmp5 <sup>1</sup>	P. luteoviolacea 2ta16		
PyrH <sup>2</sup>	Streptomyces ru- gosporus LL-42D005		
KtzR <sup>3</sup>	Kutzneria sp. 744		
Th-Hal <sup>4</sup>	Streptomyces violaceusniger SPC6		
SttH⁵	Streptomyces toxytricini NRRL 15443		
PrnA <sup>6</sup>	Pseudomonas fluorescens		
KtzQ <sup>3</sup>	Kutzneria sp. 744		
RebH <sup>7</sup>	Lechevalieria aerocolonigenes (strain 39243)		
ThaL <sup>8</sup>	Streptomyces albogriseolus		
PrnC <sup>6</sup>	Pseudomonas fluorescens		
RadH <sup>9</sup>	Chaetomium chiversii		
MalA' <sup>10</sup>	Malbranchea graminicola (086937A)		
Rdc2 <sup>11</sup>	Pochonia chlamydosporia		
ChIA <sup>12</sup>	Dictyostelium discoideum		
	Engineered enzymes		
Name	Name		
PrnA_F103A <sup>13</sup>	RebH_3SS <sup>14</sup>		
PrnA_E450K_F454K <sup>13</sup>	RebH_4V <sup>14</sup>		
SttH_Triple	RebH_5LS <sup>15</sup>		
RebH_0S <sup>15</sup>	RebH_6TL <sup>15</sup>		
RebH_1PVM <sup>14</sup>	RebH_8F <sup>15</sup>		
RebH_2T <sup>14</sup>	RebH_10S <sup>15</sup>		
RebH_3S <sup>14</sup>	RebH_Thermo <sup>16</sup>		

# Supplementary Table 1. FI-Hal panel.

#### **Supplementary Table 2.** αKGHs panel.

Natural enzymes			
Name	Source organism		
WelO5 <sup>17</sup>	Hapalosiphon welwitschii UTEX B1830		
WelO5 <sup>*18</sup>	Hapalosiphon welwitschii IC-52-3		
AmbO5 <sup>19</sup>	Fischerella ambigua UTEX1903		
Engineered enzymes			
Name	Name		
WelO5*_N74L	WelO5*_V81L		
WelO5*_V81T	WelO5*_184F		
WelO5*_A88G	WelO5*_A88S		
WelO5*_A88T	WelO5*_V90P		
WelO5*_P153F	WelO5*_P153K		
WelO5*_161A	WelO5*_I161D		
WelO5*_I161G	WelO5*_I161R		
WelO5*_I161S	WelO5*_I161T		
WelO5*_I161E	WelO5*_I225M		
WelO5*_E76V_V81L	WelO5*_V81G_I161G		
WelO5*_ V81G_I161P	WelO5*_ V81L_A88T		
WelO5*_ V81L_l161D	WelO5*_ V81L_I161M		
WelO5*_ V81L_I161V	WelO5*_ V81R_I161D		
WelO5*_ V81R_I161G	WelO5*_ V81R_I161S		
SadA_D157G <sup>20</sup>			

**Supplementary Table 3.** Ranking of the predicted variants using machine learning. Variants were predicted towards increase in activity and towards increase in selectivity. The produced "activity" variants were chosen on the highest activity predictions. The "selectivity for 1b" variants were chosen on high predicted selectivity towards product **1b** with the additional threshold that the activity predictions had to be higher than 0.6. The activity label (*A*) was calculated using the formula A = tot. *Cl conversion WelO5\* variant / tot. Cl conversion WelO5\* GAP (tot. Cl conversion* = (*SIM*<sub>1a</sub> + *SIM*<sub>1b</sub>) / (*SIM*<sub>1a</sub> + *SIM*<sub>1b</sub> + *SIM*<sub>1c</sub> + *SIM*<sub>1</sub>)). The selectivity label (*S*) was calculated using the formula  $S = (SIM_{1a} - SIM_{1b}) / (SIM_{1a} + SIM_{1b})$ .

Activity				Selectivity 1b	)	
Mutant	Ranking activity	Activity p/m <sup>#</sup>	Selectivity p/m <sup>#</sup>	Mutant	Selectivity p/m <sup>#</sup>	Activity p/m <sup>#</sup>
SIP	1	10.2/10.4	0.32/0.08	AHS	-0.77/-0.75	0.7/2.7
VIA	2	9.3/11.4	0.32/0.23	MHS	-0.68/-0.77	1.2/0.9
AIP	3	9.2/11.2	0.07/0.08	AMS	-0.67/*	0.8/*
ALP	4	8.2/11.3	0.56/0.70	VHS	-0.60/-0.75	1.5/1.9
CIA	5	8.0/*	0.25/*	LHT	-0.55/-1.0	0.7/0.5
CIP	6	7.84/*	0.30/*	LHA	-0.53/-0.67	1.8/1.9
SVP	7	7.6/8.5	0.63/0.52	AHG	-0.52/-1.0	1.2/4.2
SIA	8	7.4/5.9	0.24/0.28	АНА	-0.52/*	1.2/*
CLP	9	7.2/*	0.74/*	LMS	-0.51/0.52	0.6/0.6
VLA	10	7.2/11.0	0.77/0.72	LHG	-0.51/-1.0	1.7/3.8

\* these variants were not measured, <sup>#</sup>predicted/measured

**Supplementary Table 4**. Activity of WelO5\* variants for the functionalization of soraphen derivatives. The halogenase variants were capable to produce multiple products (as reported). Each of the reported products were derivatized once only as observed by selected ion monitoring.

Compound	Observed products	WelO5* variant
Soraphen C, <b>2</b>	<ul> <li>2 chlorinated products</li> <li>2 hydroxylated products</li> </ul>	- VAA⁺ - ILV°
о с с с с с с с с с с с с с с с с с с с	<ul> <li>4 chlorinated products</li> <li>3 hydroxylated products</li> </ul>	- SLP⁺ - SLP°
	<ul> <li>- 5 chlorinated products</li> <li>- 2 hydroxylated products</li> </ul>	- SLP⁺ - SHP°

<sup>+</sup> Variant showing the highest amount of total chlorination; <sup>°</sup> Variant showing the highest amount of total hydroxylation

Year	Enzyme	Target	Number of variants	Perc. Covered /%	Reference
2007	halohydrin	activity	30-150 at each	~(3*N/2 <sup>N</sup> ) * 100	Fox et al. <sup>21</sup>
	dehalogenase		round for 18 rounds	N= mutation sites	
2012	epoxide	enantioselectivity	95	23.8	Feng et al. <sup>22</sup>
	hydrolase				
2018	green flourescent	color change	218	0.14	Saito et al. <sup>23</sup>
	protein				
2018	epoxide	enantioselectivity	37	7.4	Cadet et al. <sup>24</sup>
	hydrolase				
2019	nitric oxide	stereodivergence	445 over 2 rounds	0.6 - 8.9	Wu et al. <sup>25</sup>
	dioxygenase				
2021	artificial	activity	400	80.0	Vornholt et al. <sup>26</sup>
	metalloenzymes				

**Supplementary Table 5.** Selected examples of the application of machine learning used for the engineering of enzymes.

Supplementary Table 6. Enzyme and substrate concentrations used for the kinetic experiments.

Variants	Enzyme conc. / µM	Substrate conc. / µM
GAP	20	40, 70, 100, 150, 200, 250,500, 750
VLA	2	40, 70, 100, 150, 200, 250,500, 750
SLP	2	40, 70, 100, 150, 200, 250,500, 750

Supplementary Equation 1. Substrate inhibition model.

$$v_0 = \frac{v_{max} * [S]_0}{K_m + [S]_0 + \frac{[S]_0^2}{K_i}}$$

Supplementary Table 7. Comparison of <sup>13</sup>C-NMR chemical shifts between soraphen C (reported<sup>2</sup> and

synthesized) and synthesized epi-soraphen C.

<sup>13</sup> C-NMR shifts of	<sup>13</sup> C-NMR shifts of		<sup>13</sup> C-NMR shifts of	
fermented soraphen C	synthesized soraphen C	Δ1 (ppm)	synthesized epi-soraphen C	Δ2 (ppm)
(δ in ppm) <sup>2</sup>	(δ in ppm)		(δ in ppm)	
170.6	170.78	-0.2	172.82	-2.2
141.0	141.12	-0.1	139.69	1.3
137.3	137.45	-0.1	137.96	-0.7
128.6	128.72	-0.1	128.76	-0.2
128.2	128.29	-0.1	128.70	-0.5
126.2	126.35	-0.1	128.46	-2.3
125.0	125.18	-0.2	127.09	-2.1
99.4	99.60	-0.2	99.91	-0.5
83.7	83.89	-0.2	82.86	0.8
76.1	76.28	-0.2	77.57	-1.5
74.9	75.04	-0.1	76.33	-1.4
74.6	74.81	-0.2	73.58	1.0
72.5	72.66	-0.2	71.74	0.8
68.8	68.98	-0.2	68.93	-0.1
57.6	57.78	-0.2	57.49	0.1
57.3	57.47	-0.2	57.45	-0.2
46.2	46.35	-0.1	45.42	0.8
35.8	36.01	-0.2	36.84	-1.0
35.6	35.79	-0.2	35.46	0.1
35.2	35.32	-0.1	35.02	0.2
29.4	29.57	-0.2	27.58	1.8
26.0	26.14	-0.1	25.12	0.9
23.0	23.17	-0.2	22.57	0.4
12.5	12.64	-0.1	16.57	-4.1
11.7	11.83	-0.1	12.55	-0.9
10.3	10.49	-0.2	10.55	-0.3

# II. Supplementary Figures



Supplementary Figure 1. Crystal structure of soraphen A bound to the BC domain of yeast acetyl-coenzyme A carboxylases. The crystal structure (green) reveals the active conformation of the macrocycle (PDB ID: 1W96). Right: Visualization of the interactions between the soraphen A (wheat sticks) and the residues of the BC domain (green sticks).



**Supplementary Figure 2. Synthesis scheme to obtain soraphen C and soraphen analogues.** Individual reaction steps are described in the Supplementary Methods.



Supplementary Figure 3. LC-MS analysis of the biotransformation of soraphen A. a Selected ion chromatograms (SIM) of the m/z values of interest. Biotransformation using negative control (blue), WT WelO5\* (orange) and WelO5\*\_V81G\_I161P (green) are compared. The top chromatogram shows the trace of soraphen A (543.2 m/z =  $1+Na+H^+$ ), the middle chromatogram shows two species corresponding to chlorinated soraphen A (577.2 m/z =  $1+Na+H^++Cl^{35}$ ) and the bottom chromatogram shows one hydroxylated soraphen A species (577.2 m/z =  $1+Na+H^++OH$ ). b MS chart of the chlorinated species showing the characteristic M: M + 2 = 3 : 1 isotopic pattern of a chlorinated compound.



**Supplementary Figure 4. Biotransformation products of soraphen A.** Observed products of the biotransformation reactions of soraphen A with WelO5\* variants.



**Supplementary Figure 5. Illustration of the algorithm aided approach to predict improved variants.** Activity (or selectivity) data obtained by LC-MS analysis was used as a label for the machine learning algorithm. Amino acids properties were represented as a 17-dimensional vector. The feature vector of a sequence was defined by joining the vector representation of its individual amino acids at sites V81X, A88X, I161X and aggregated into the *504 x 51*-dimensional training matrix. This was used to train a machine learning model. To avoid overfitting and to better gauge the generalizability of our model, we cross-validated over ten splits, and model performance was evaluated on the coefficient of determination (R^2).



**Supplementary Figure 6. Docking studies of soraphen A into WelO5\* homology models.** Enzyme models were prepared with SWISS-MODEL<sup>27</sup> (enzyme model and soraphen A in wheat, engineered residues in red) or AlphaFold<sup>28</sup> (enzyme model and soraphen A in palegreen, engineered residues in orange) and soraphen A was docked using AutoDock Vina<sup>29</sup>. In the active site the histidines coordinating to the iron (orange) are shown in grey, the chlorine in green and the α-ketoglutarate in pale cyan. **a-c** View into the active site of the WelO5\* variants GAP, VLA and AHG, respectively. The active site of WelO5\* including the engineered residues at position 81, 88 and 161 are nearly identical in both homology models (SWISS-MODEL (wheat) and AlphaFold (palegreen)). **d-f** Soraphen A docked into the WelO5\* variants GAP, VLA and AHG. Distances were measured from the iron and chloride to C14 of soraphen A (grey dotted lines) and to C16 of soraphen A (yellow dotted lines). The two models of **d** WelO5\* GAP and **f** WelO5\* AHG show a similar positioning of the engineered amino acid residues as well as of the docked soraphen A. In the AlphaFold model of **e** WelO5\* VLA shorter distances between the iron and chloride to C14 of soraphen A (grey dotted lines) than to C16 of the macrolide (yellow dotted lines) suggest the structural reason for the predominant formation of regioisomer **1a**. **g** Overall structural homology of the enzyme models prepared with SWISS-MODEL (wheat) or AlphaFold (palegreen): Main structural differences lay in the two α-helices marked in the black circle.



**Supplementary Figure 7. Structure of soraphen A**. Numbering of the carbon atoms of the cyclic polyketide backbone of soraphen A.



Supplementary Figure 8. *In vitro* activity assays using mono-chlorinated products (1a, 1b and 2a) as substrates for the engineered WelO5\* variants GAP, SLP, VLA and WVS. Depicted is the estimated conversion to chlorinated or hydroxylated product (*SIM area of product / SIM area of all products and starting material* \* 100). Reactions were performed according to the method described in the main paper using an enzyme concentration of 5  $\mu$ M for variants GAP, SLP, VLA and WVS and a substrate concentration of 60  $\mu$ M (1a, 1b and 2a). The conversion values to the chlorinated or hydroxylated products by the engineered halogenase variants were determined in triplicates (N = 3 independent experiments). The depicted boxes correspond to the interquartile range and end at the quartiles Q<sub>1</sub> and Q<sub>3</sub>, respectively. The statistical median is depicted as a horizontal line in the box. The whiskers comprise the farthest points that are not outliers (i.e., that are within 1.5x of the interquartile range of Q<sub>1</sub> and Q<sub>3</sub>, respectively).



Supplementary Figure 9. *In vitro* activity assays showing the conversion of soraphen A to the chlorinated product 1a by the engineered WelO5\* variants GAP, SLP and VLA. The conversion was determined at a substrate concentration of 60  $\mu$ M and the reaction was quenched at stable product concentration using the procedure described in the method section of the main paper. To account for the different enzyme concentrations used (GAP = 5  $\mu$ M, SLP = 0.5  $\mu$ M and VLA = 0.5  $\mu$ M), the observed product concentration in the GAP reactions was divided by ten. The conversion values of soraphen A to the chlorinated product 1a by the engineered halogenase variants were determined in quadruplicates in each case (N = 4 independent experiments). The depicted boxes correspond to the interquartile range and end at the quartiles Q<sub>1</sub> and Q<sub>3</sub>. The statistical median is depicted as a horizontal line in the box. The whiskers comprise the farthest points that are not outliers (i.e., that are within 1.5x the interquartile range of Q<sub>1</sub> and Q<sub>3</sub>, respectively).



**Supplementary Figure 10. LC-MS analysis of the anion promiscuity of selected WelO5\* variants**. The variants GAP (blue), SLP (orange), WVS (green) and negative control (no enzyme, red) were analysed in the presence of 500 mM of NaF, NaCl, NaBr, NaI, NaN<sub>3</sub> and NaNO<sub>2</sub>, respectively, by selected ion monitoring. Masses corresponding to the introduction of the anions into soraphen A could be observed for chloride, bromide, azide and nitrite salts.



Supplementary Figure 11. Calibration curve of 1a used for product quantification.



**Supplementary Figure 12. Michaelis Menten kinetics for WelO5\* variants GAP, VLA and SLP.** The formation of product **1a** for the WelO5\* variants GAP, VLA and SLP was measured in triplicate at each substrate concentration (N=3 independent experiments). All data points belonging to individual Michaelis Menten measurement series are marked by triangles, squares or circles. Substrate inhibition can be observed in all cases.



**Supplementary Figure 13. Out of fold predicted vs measured values. a** The measured activity values of the training set were predicted using Gaussian processes (y-axis) and compared to the measured activity (x-axis) **b** The measured selectivity of the training set were predicted using a random forest algorithm (y-axis) and compared to the measured selectivity (x-axis). A linear regression is shown for these values.



Supplementary Figure 14. Overview of the AlphaFold models (green) of WelO5\* and the variants GAP, SLP and VLA showing a view of the entrance to the active site. The engineered amino acid residues are shown in orange, while the co-factor  $\alpha$ -ketoglutarate is shown in light cyan. To comparatively evaluate substrate access to the active sites of all enzyme variants, the bottleneck radii were calculated using CAVER Web 1.0 with default parameters<sup>30</sup>. This investigation highlighted that the employed enzyme engineering approach has led to a widening of the access tunnel from 2.1 Å (wildtype enzyme) to 3.2 Å (variant GAP), 2.6 Å (variant SLP) and 2.4 Å (variant VLA). The resulting improved access to the active site might explain why the wildtype enzyme cannot convert the macrolide soraphen A, while variants GAP, SLP and VLA accept the bulky substrate.



Supplementary Figure 15. Docking experiments of soraphen A, 12-epi-fischerindole U and chlorinated soraphen A into wildtype WelO5\* and its engineered variants. Overview of the docking scores of soraphen A (blue, 1), the natural substrate 12-epi-fischerindole U (red) and chlorinated soraphen A (green, 1a) into AlphaFold models of WelO5\* wild type (WT) and the variants GAP, SLP and VLA, respectively. The docking was performed as described in the method section of the main paper. All ligands, irrespective of them being native or non-native substrates, showed similar docking scores for the enzyme variants. Scores were obtained through the AutoDock Vina scoring function<sup>[32]</sup> which consists of the weighted sum of steric interactions (gauss<sub>1</sub>, gauss<sub>2</sub> and repulsion, identical for all atom pairs), hydrophobic interaction between hydrophobic atoms and hydrogen bonding (where applicable). A lower score indicates higher affinity of the ligand towards the receptor. In each case, nine docking solutions were obtained from one docking experiment (n = 1 individual experiment). Results were visually inspected using PyMOL software and only solutions in which the ligand docked close to the active site were considered in the depicted analysis. The depicted boxes correspond to the interquartile range and end at the quartiles Q<sub>1</sub> and Q<sub>3</sub>. The median is depicted as a horizontal line in the box. The whiskers comprise the farthest points that are not outliers (i.e., that are within 1.5x the interquartile range of Q<sub>1</sub> and Q<sub>3</sub>, respectively).

# III. Supplementary Methods

#### **Chemical synthesis methods**

Synthesis: Soraphen A was obtained by fermentation at Syngenta (former Novartis) using the published procedure<sup>31</sup>. The reagents for synthesis were obtained from commercial sources and used without further purification unless otherwise stated. The solvents for synthesis were obtained from commercial sources and stored over molecular sieves.

Purification: purification over silica gel were performed on a Combi*Flash* Rf 200i instrument using standard commercial pre-packed silica gel cartridges.

NMR: NMRs were recorded either on a Bruker 400 MHz spectrometer or a Bruker 600 MHz spectrometer. <sup>1</sup>H-NMR chemical shifts are reported relative to TMS and are referenced based on the residual proton resonances of the corresponding deuterated solvent (CDCl<sub>3</sub>: 7.26 ppm) whereas <sup>13</sup>C NMR spectra are reported relative to TMS using the carbon signals of the deuterated solvent (CDCl<sub>3</sub>: 77.16 ppm). Assignments were made on the basis of chemical shifts, coupling constants, COSY, HSQC, HMBC, ROESY data. Resonances are described using the following abbreviations; s (singlet), d (doublet), t (triplet), q (quartet), quin. (quintet), sext. (sextet), sept. (septet), m (multiplet), br. (broad), app. (apparent), dd (double doublet) and so on. Coupling constants (*J*) are given in Hz and are rounded to the nearest 0.1 Hz.

HPLC-MS: HPLC traces were obtained on an Acquity UPLC from Waters: Binary pump, heated column compartment, diode-array detector and ELSD detector. Column: Waters UPLC HSS T3, 1.8  $\mu$ m, 30 x 2.1 mm, Temp: 60 °C, DAD Wavelength range: 210 to 500 nm, Solvent Gradient: A = water + 5% MeOH + 0.05 % HCOOH, B= Acetonitrile + 0.05 % HCOOH, gradient: 10-100% B in 2.7 min; Flow: 0.85 mL/min. Low resolution mass spectra were recorded on a mass spectrometer from Waters (SQD, SQDII Single quadrupole mass spectrometer) equipped with an electrospray source (Polarity: positive and negative ions); Capillary: 3.00 kV, Cone range: 30V, Extractor: 2.00 V, Source Temperature: 150 °C, Desolvation Temperature: 350 °C, Cone Gas Flow: 50 L/h, Desolvation Gas Flow: 650 L/h, Mass range: 100 to 900 Da.

#### Synthesis and characterization of soraphen C and soraphen analogues

For a scheme of the synthesis route refer to Supplementary Figure 2.

(1*R*,2*S*,5*S*,10*S*,11*R*,12*E*,14*S*,15*S*,16*R*,17*S*,18*R*)-17-[tert-butyl(dimethyl)silyl]oxy-1-hydroxy-10,11,18-trimethoxy-2,14,16-trimethyl-5-phenyl-4,19-dioxabicyclo[13.3.1]nonadec-12-en-3-one (**S1**)



Soraphen A (100 mg, 0.19 mmol), DMAP (25 mg, 0.20 mmol), imidazole (52 mg, 0.77 mmol) and TBSCI (58 mg, 0.38 mmol) were dissolved in DMF (2.0 mL). The slightly yellow solution was stirred at room temperature for 48 h. Then the reaction mixture was diluted with EtOAc and washed with aq HCl (1M), water and brine, dried over  $Na_2SO_4$ , filtered and concentrated under reduced pressure.

The crude was purified by flash chromatography on silica gel (gradient: cHex/EtOAc 95:5 to 55:45, 30 mL/min, 15 min) to afford the title compound (60 mg, 49%) as well as some recovered starting material (50 mg).

<sup>1</sup>H-NMR (400 MHz, CDCl<sub>3</sub>):  $\delta$  (ppm) = 7.42–7.23 (m, 5H), 6.40 (dd, *J* = 16.2, 3.8 Hz, 1H), 6.12 (dd, *J* = 12.0, 2.5 Hz, 1H), 5.41 (ddd, *J*=16.0, 9.5, 1.8 Hz, 1H), 5.17 (d, *J*=1.8 Hz, 1H), 4.17 (t, *J*=2.5 Hz, 1H), 3.78 (dd, *J*=9.5, 2.2 Hz, 1H), 3.69 (dd, *J*=10.5, 2.5 Hz, 1H), 3.47 (s, 3H), 3.42 (dt, *J*=11.1, 2.5 Hz, 1H), 3.38 (s, 3H), 3.31 (s, 3H), 3.05 (qd, *J*=7.0, 1.1 Hz, 1H), 3.00 (dd, *J*=2.7, 0.9 Hz, 1H), 2.56–2.49 (m, 1H), 2.13–2.04 (m, 1H), 1.81–1.72 (m, 2H), 1.65–1.54 (m, 3H), 1.34–1.25 (m, 2H), 1.16 (d, *J*=7.3 Hz, 3H), 1.09 (d, *J*=7.3 Hz, 3H), 1.05–1.01 (m, 1H), 1.00 (d, *J*=6.5 Hz, 3H), 0.94 (s, 9H), 0.17 (d, *J*=4.0 Hz, 6H);

<sup>13</sup>C-NMR (101 MHz, CDCl<sub>3</sub>): δ (ppm) = 171.25, 142.77, 140.74, 128.43 (2C), 127.52, 126.29 (2C), 121.80, 99.45, 84.99, 83.65, 77.39, 72.08, 71.90, 70.86, 58.32, 57.41, 56.31, 46.32, 37.30, 35.62, 35.22, 31.02, 25.95 (3C), 25.23, 24.18, 18.26, 12.42, 11.49, 10.32, -4.82, -4.85.

(1*R*,2*S*,5*S*,10*S*,12*E*,14*S*,15*S*,16*R*,17*S*,18*R*)-17-[tert-butyl(dimethyl)silyl]oxy-1-hydroxy-10,18-dimethoxy-2,14,16-trimethyl-5-phenyl-4,19-dioxabicyclo[13.3.1]nonadec-12-ene-3,11-dione (**S2**)



To a solution of **S1** (750 mg, 1.18 mmol) in 1,2-dichloroethane (14.8 mL) at room temperature was added DDQ (621 mg, 2.60 mmol). The yellow suspension was stirred for 4 days. Then the reaction mixture was diluted with EtOAc, washed twice with aq.  $Na_2S_2O_3$  (10%), then brine, dried over  $Na_2SO_4$ , filtered and concentrated under reduced pressure.

The crude was purified by flash chromatography on silica gel (gradient: cHex/EtOAc 90:10 to 80:20, 40 mL/min, 16 min) to afford the title compound (522.8 mg, 72%).

<sup>1</sup>H-NMR (400 MHz, CDCl<sub>3</sub>):  $\delta$  (ppm) = 7.45 (dd, *J*=16.9, 4.8 Hz, 1H), 7.39–7.28 (m, 4H), 6.24 (dd, *J*=16.9, 1.5 Hz, 1H), 5.91 (dd, *J*=8.6, 5.3 Hz, 1H), 5.05 (d, *J*=1.5 Hz, 1H), 4.29 (dd, *J*=7.7, 5.5 Hz, 1H), 4.16 (t, *J*=2.6 Hz, 1H), 3.86 (dd, *J*=10.3, 2.6 Hz, 1H), 3.37 (s, 6H), 3.05 (qd, *J*=7.0, 1.47 Hz, 1H), 3.00 (dd, *J*=2.8, 0.9 Hz, 1H), 2.70–2.64 (m, 1H), 2.03–1.94 (m, 1H), 1.82–1.71 (m, 2H), 1.70–1.58 (m, 4H), 1.49–1.39 (m, 3H), 1.13 (d, *J*=7.0 Hz, 3H), 1.10 (d, *J*=7.3 Hz, 3H), 1.00 (d, *J*=6.6 Hz, 3H), 0.94 (s, 9H), 0.16 (d, *J*=5.1 Hz, 6H);

<sup>13</sup>C-NMR (101 MHz, CDCl<sub>3</sub>): δ (ppm) = 202.74, 171.44, 152.37, 141.78, 128.45 (2C), 127.67, 127.22, 126.64 (2C), 99.90, 83.63, 74.30, 70.76, 70.61, 57.92, 57.50, 46.31, 36.52, 36.01, 35.73, 32.33, 25.92 (3C), 23.98, 23.49, 18.23, 13.20, 11.55, 10.24, -4.82, -4.86;

HPLC-MS: rt = 2.69 min, m/z = 504  $[M-C_6H_{14}Si]^-$ ; 618  $[M-H]^-$ 

2*S*,5*S*,10*S*,12*E*,14*S*,15*S*,16*S*,17*S*,18*R*)-1,17-dihydroxy-10,18-dimethoxy-2,14,16-trimethyl-5-phenyl-4,19-dioxabicyclo[13.3.1]nonadec-12-ene-3,11-dione (**3**)



To a solution of **S2** (30.0 mg, 0.485 mmol) in THF (0.2 mL) were added at 0°C AcOH (8.8  $\mu$ L, 0.15 mmol) and TBAF (0.10 mL, 1 M in THF, 0.10 mmol). The reaction mixture was stirred at 0 °C for 1.5 h. The reaction mixture was poured into sat. aq. NH<sub>4</sub>Cl, then extracted with DCM/MeOH (9:1) The combined organic layers were dried over Na<sub>2</sub>SO<sub>4</sub>, filtered and concentrated under reduced pressure.

The crude product was purified by flash chromatography on silica gel (gradient: cHex/EtOAc 85:15 to 50:50, 18 mL/min, 12 min) to afford the title compound (21 mg, 86%).

<sup>1</sup>H-NMR (400 MHz, CDCl<sub>3</sub>):  $\delta$  (ppm) = 7.35–7.30 (m, 6H), 6.51 (dd, *J*=16.1, 1.5 Hz, 1H), 5.61 (t, *J*=7.3 Hz, 1H), 4.51 (s, 1H), 4.04–4.01 (m, 2H), 3.80 (dd, *J*=9.2, 4.4 Hz, 1H), 3.38 (s, 3H), 3.35 (s, 3H), 3.17–3.12 (m, 2H), 2.69–2.60 (m, 1H), 1.99–1.93 (m, 1H), 1.90–1.84 (m, 2H), 1.80–1.60 (m, 3H), 1.51–1.41 (m, 2H), 1.36–1.29 (m, 1H), 1.25–1.17 (m, 1H), 1.07 (d, *J*=7.3 Hz, 3H), 1.06 (d, *J*=7.3 Hz, 3H), 1.04 (d, *J*=7.0 Hz, 3H);

<sup>13</sup>C-NMR (101 MHz, CDCl<sub>3</sub>): δ (ppm) = 202.60, 170.99, 152.95, 140.15, 128.67 (2C), 128.31, 126.68 (2C), 124.57, 99.85, 86.02, 76.62, 76.16, 72.63, 68.99, 57.76, 57.62, 45.76, 36.53, 35.65, 34.44, 30.18, 23.74, 22.77, 13.94, 11.79, 10.49;

HPLC-MS: rt = 1.83 min, m/z = 504 [M-H]<sup>-</sup>; 528 [M+H+Na]<sup>2+</sup>

(1*R*,2*S*,5*S*,10*S*,11*R*,12*E*,14*S*,15*S*,16*R*,17*S*,18*R*)-17-[tert-butyl(dimethyl)silyl]oxy-1,11-dihydroxy-10,18-dimethoxy-2,14,16-trimethyl-5-phenyl-4,19-dioxabicyclo[13.3.1]nonadec-12-en-3-one (S3)

(1*R*,2*S*,5*S*,10*S*,11*S*,12*E*,14*S*,15*S*,16*R*,17*S*,18*R*)-17-[tert-butyl(dimethyl)silyl]oxy-1,11-dihydroxy-10,18-dimethoxy-2,14,16-trimethyl-5-phenyl-4,19-dioxabicyclo[13.3.1]nonadec-12-en-3-one (S4)

(1*R*,2*S*,5*S*,10*S*,14*S*,15*S*,16*R*,17*S*,18*R*)-17-[tert-butyl(dimethyl)silyl]oxy-1,11-dihydroxy-10,18-dimethoxy-2,14,16-trimethyl-5-phenyl-4,19-dioxabicyclo[13.3.1]nonadecan-3-one (**S5**)



To a solution of **S2** (380 mg, 0.610 mmol) in 1,2-dimethoxyethane (3 mL) was added at 0°C a solution of ZnCl<sub>2</sub> (1 M in Et<sub>2</sub>O, 9.0 mL, 0.92 mmol) followed by addition of NaBH<sub>4</sub> (64.5 mg, 1.53 mmol). The reaction mixture was stirred at 0 °C for 1h40. It was then quenched with sat. aq. NH<sub>4</sub>Cl and extracted with EtOAc. The combined organic phases were dried over Na<sub>2</sub>SO<sub>4</sub>, filtered and concentrated under reduced pressure.

The crude product was purified by flash chromatography on silica gel (gradient: cHex/EtOAc 95:05 to 70:30, 40 mL/min, 22 min) to afford three products: **S3** (186.1 mg, 49%), **S4** (78.4 mg, 21%), and **S5** (23.9 mg, 6%) as a mixture of epimers.

Data for S3:

<sup>1</sup>H-NMR (400 MHz, CDCl<sub>3</sub>):  $\delta$  (ppm) = 7.37–7.22 (m, 5H), 6.33 (dd, *J*=16.1, 4.0 Hz, 1H), 6.07 (dd, *J*=12.1, 2.6 Hz, 1H), 5.39 (ddd, *J*=16.1, 9.4, 1.7 Hz, 1H), 5.17 (d, *J*=1.8 Hz, 1H), 4.24 (dd, *J*=9.2, 2.6 Hz, 1H), 4.14 (t, *J*=2.8 Hz, 1H), 3.64 (dd, *J*=10.6, 2.6 Hz, 1H), 3.44 (s, 3H), 3.36 (s, 3H), 3.34–3.33 (m, 1H), 3.03 (qd, *J*=7.0, 1.5 Hz, 1H), 2.98 (dd, *J*=2.9, 1.1 Hz, 1H), 2.52–2.43 (m, 1H), 2.29 (br m, 1H), 2.13–2.07 (m, 1H), 1.79–1.56 (m, 5H), 1.32–1.24 (m, 2H), 1.21–1.17 (m, 1H), 1.13 (d, *J*=7.3 Hz, 3H), 1.06 (d, *J*=7.7 Hz, 3H), 0.95 (d, *J*=6.6 Hz, 3H), 0.93 (s, 9H), 0.15 (s, 3H), 0.14 (s, 3H);

HPLC-MS: rt = 2.58 min, m/z = 644  $[M+H+Na]^{2+}$ ; 506  $[M-C_6H_{14}Si]^{-1}$ 

The relative stereochemistry of **S3** was determined retrospectively from soraphen C after deprotection. The relative stereochemistry of **S4** was determined by comparison with **S3**.

Data for S4:

<sup>1</sup>H-NMR (400 MHz, CDCl<sub>3</sub>):  $\delta$  (ppm) = 7.38–7.24 (m, 5H), 6.12 (ddd, *J*=16.3, 6.1, 1.1 Hz, 1H), 5.90 (dd, *J*=8.6, 5.7 Hz, 1H), 5.55 (ddd, *J*=16.2, 5.8, 1.1 Hz, 1H), 5.08 (d, *J*=1.1 Hz, 1H), 4.29 (t, *J*=5.7 Hz, 1H), 4.11 (t, *J*=2.6 Hz, 1H), 3.84 (dd, *J*=9.9, 2.6 Hz, 1H), 3.43 (s, 3H), 3.35 (s, 3H), 3.35–3.30 (m, 1H), 3.03 (qd, *J*=7.1, 1.1 Hz, 1H), 2.97 (dd, *J*=2.9, 0.7 Hz, 1H), 2.70–2.58 (br s, 1H), 2.53–2.46 (m, 1H), 2.09–2.00 (m, 1H), 1.75–1.62 (m, 5H), 1.56–1.47 (m, 2H), 1.38–1.29 (m, 2H), 1.11 (d, *J*=7.0 Hz, 3H), 1.07 (d, *J*=7.3 Hz, 3H), 0.97 (d, *J*=7.0 Hz, 3H), 0.93 (m, 9H), 0.15 (s, 3H), 0.14 (s, 3H);

<sup>13</sup>C-NMR (101 MHz, CDCl<sub>3</sub>): δ (ppm) = 171.91, 141.78, 136.68, 128.43 (2C), 127.65, 126.96, 126.71 (2C), 99.54, 83.58, 77.68, 74.18, 73.18, 71.62, 70.95, 57.69, 57.52, 46.66, 36.69, 36.22, 35.79, 28.91, 25.92 (3C), 24.47, 23.97, 18.23, 15.63, 11.72, 10.74, -4.83, -4.85;

HPLC-MS: rt = 2.66 min, m/z = 644 [M+H+Na]<sup>2+</sup>

Data for S5:

<sup>1</sup>H-NMR (400 MHz, CDCl<sub>3</sub>): δ (ppm) = 7.39–7.28 (m, 4H), 7.25–7.21 (m, 1H), 5.98 (dd, *J*=10.8, 3.1 Hz, 1H), 5.15 (d, *J*=1.8 Hz, 1H), 4.15 (t, *J*=2.6 Hz, 1H), 3.88 (dd, *J*=10.6, 2.6 Hz, 1H), 3.74 (dt, *J*=9.1, 3.0 Hz, 1H), 3.42 (s, 3H), 3.39–3.36 (m, 1H), 3.34 (s, 3H), 3.01–2.94 (m, 2H), 2.12–1.98 (m, 3H), 1.93–1.75 (m, 2H), 1.68–1.56 (m, 6H), 1.53–1.38 (m, 4H), 1.12 (d, *J*=7.0 Hz, 3H), 1.03 (d, *J*=7.3 Hz, 3H), 0.94 (s, 9H), 0.81 (d, *J*=7.0 Hz, 3H), 0.16 (s, 3H), 0.15 (s, 3H); <sup>13</sup>C-NMR (101 MHz, CDCl<sub>3</sub>): δ (ppm) = 171.14, 142.53, 128.43 (2C), 127.58, 126.49 (2C), 99.61, 83.92, 72.81, 72.24, 70.93, 67.36, 57.76, 57.28, 46.52, 36.52, 35.11, 32.06, 28.98, 27.79, 25.94 (3C), 25.15, 23.09, 22.92, 18.25, 14.67, 11.56, 10.45, –4.82 (2C);

HPLC-MS: rt = 2.63 min, m/z = 646 [M+H+Na]<sup>2+</sup>; 508 [M-C<sub>6</sub>H<sub>14</sub>Si]<sup>-</sup>

(1*R*,2*S*,5*S*,10*S*,11*R*,12*E*,14*S*,15*S*,16*S*,17*S*,18*R*)-1,11,17-trihydroxy-10,18-dimethoxy-2,14,16-trimethyl-5-phenyl-4,19-dioxabicyclo[13.3.1]nonadec-12-en-3-one (soraphen C, **2**)



To a solution of **S3** (271 mg, 0.535 mmol) in THF (4 mL) were added at 0°C AcOH (75.8  $\mu$ L, 1.31 mmol) and TBAF (0.87 mL, 1 M in THF, 0.87 mmol). The reaction mixture was stirred at 0°C for 2 h then at room temperature for 18 h. The reaction mixture was poured into sat. aq. NH<sub>4</sub>Cl, then extracted with EtOAc. The combined organic layers were dried over Na<sub>2</sub>SO<sub>4</sub>, filtered and concentrated under reduced pressure.

The crude product was purified by flash chromatography on silica gel (gradient: cHex/EtOAc 90:10 to 40:60, 35 mL/min, 18 min) to afford the title compound (219 mg, 99%).

Comparison of the spectral data with the ones of the isolated natural product<sup>32,33</sup> confirmed the identity of the product. A table of compared <sup>13</sup>C-NMR shifts is presented in Supplementary Table 7.

<sup>1</sup>H-NMR (400 MHz, CDCl<sub>3</sub>):  $\delta$  (ppm) = 7.37–7.28 (m, 5H), 6.15 (dd, *J*=16.0, 3.9 Hz, 1H), 5.82 (dd, *J*=11.2, 3.5 Hz, 1H), 5.48 (ddd, *J*=16.0, 9.4, 1.8 Hz, 1H), 4.36 (s, 1H), 4.17 (td, *J*=9.0, 2.6 Hz, 1H), 4.01 (br d, *J*=8.0 Hz, 1H), 3.81 (dd, *J*=10.4, 2.8 Hz, 1H), 3.62 (d, *J*=9.9 Hz, 1H), 3.43 (s, 3H), 3.38 (s, 3H), 3.37–3.34 (m, 1H), 3.18 (dd, *J*=2.6, 1.1 Hz, 1H), 3.14 (q, *J*=7.3 Hz, 1H), 2.50–2.42 (m, 2H), 2.15–2.07 (m, 1H), 1.93 (q, *J*=7.0 Hz, 1H), 1.82–1.75 (m, 1H), 1.72–1.64 (m, 1H), 1.51–1.45 (m, 2H), 1.38–1.34 (m, 1H), 1.22–1.13 (m, 2H), 1.09 (d, *J*=7.0 Hz, 3H), 1.05 (d, *J*=7.7 Hz, 3H), 1.00 (d, *J*=6.6 Hz, 3H);

<sup>13</sup>C-NMR (101 MHz, CDCl<sub>3</sub>): δ (ppm) = 170.78, 141.12, 137.45, 128.72 (2C), 128.29, 126.35 (2C), 125.18, 99.60, 83.89, 76.28, 75.04, 74.81, 72.66, 68.98, 57.78, 57.47, 46.35, 36.01, 35.79, 35.32, 29.57, 26.14, 23.17, 12.64, 11.83, 10.49;

HPLC-MS: rt = 1.65 min, m/z = 505 [M-H]<sup>-</sup>

HR-MS: m/z calculated for C<sub>28</sub>H<sub>43</sub>O<sub>8</sub> [(M+H)<sup>+</sup>]: 507.2952, found 507.2962.

(1*R*,2*S*,5*S*,10*S*,11*S*,12*E*,14*S*,15*S*,16*S*,17*S*,18*R*)-1,11,17-trihydroxy-10,18-dimethoxy-2,14,16-trimethyl-5-phenyl-4,19-dioxabicyclo[13.3.1]nonadec-12-en-3-one (*epi*-soraphen C, **S7**)



To a solution of **S4** (10 mg, 0.016 mmol) in THF (0.4 mL) were added at 0°C AcOH (2.8  $\mu$ L, 0.048 mmol) and TBAF (0.03 mL, 1 M in THF, 0.32 mmol). The reaction mixture was stirred at 0°C for 2 h then at room temperature for 18 h. The reaction mixture was poured into sat. aq. NH<sub>4</sub>Cl, then extracted with a mixture of DCM/MeOH (9:1). The combined organic layers were dried over Na<sub>2</sub>SO<sub>4</sub>, filtered and concentrated under reduced pressure. The crude product was purified by flash chromatography on silica gel (gradient: cHex/EtOAc 70:30 to 50:50, 18 mL/min, 9 min) to afford the title compound (5 mg, 61%) containing some impurities.

<sup>1</sup>H-NMR (600 MHz, CDCl<sub>3</sub>):  $\delta$  (ppm) = 7.37–7.29 (m, 5H), 5.89 (dd, *J*=15.9, 6.3 Hz, 1H), 5.67 (t, *J*=7.3 Hz, 1H), 5.53 (ddd, *J*=15.8, 7.1, 1.1 Hz, 1H), 4.69 (s, 1H), 4.20 (t, *J*=6.6 Hz, 1H), 3.99 (dd, *J*=10.4, 2.5 Hz, 1H), 3.98 (br. s, 1H), 3.69 (br. s, 1H), 3.42 (s, 3H), 3.37 (s, 3H), 3.30–3.26 (m, 1H), 3.14 (dd, *J*=2.5, 0.9 Hz, 1H), 3.11 (q, *J*=7.2 Hz, 1H), 2.53 (br s, 1H), 2.47–2.43 (m, 1H), 2.24–2.18 (m, 1H), 1.94–1.90 (m, 1H), 1.85–1.78 (m, 2H), 1.66–1.61 (m, 1H), 1.56–1.51 (m, 1H), 1.46–1.38 (m, 3H), 1.04 (d, *J*=7.5 Hz, 3H), 1.03 (d, *J*=7.1 Hz, 3H), 0.99 (d, *J*=6.6 Hz, 3H);

<sup>13</sup>C-NMR (151 MHz, CDCl<sub>3</sub>): δ (ppm) = 172.82, 139.69, 137.96, 128.76 (2C), 128.70, 128.46 (2C), 127.09, 99.91, 82.86, 77.57, 76.33, 73.58, 71.74, 68.93, 57.49, 57.45, 45.42, 36.84, 35.46, 35.02, 27.58, 25.12, 22.57, 16.57, 12.55, 10.55;

HPLC-MS: rt = 1.77 min, m/z = 505 [M-H]<sup>-</sup>

(1*S*,2*R*,3*R*,5*S*,6*S*,7*S*,12*S*,15*S*,16*R*,17*R*,18*S*,19*R*)-18-[tert-butyl(dimethyl)silyl]oxy-16-hydroxy-6,7,17-trimethoxy-2,15,19-trimethyl-12-phenyl-4,13,20-trioxatricyclo[14.3.1.03,5]icosan-14-one (**S6**)



To a solution of **S3** (400 mg, 0.630 mmol) in DCM (6.3 mL) at rt was added mCPBA (706 mg, 3.150 mmol). The reaction mixture was stirred for 17 h. The reaction mixture was poured into sat. aq. NaHCO<sub>3</sub>. The phases were separated, the aqueous layer was extracted with EtOAc. The combined organic layers were washed with brine, dried over Na<sub>2</sub>SO<sub>4</sub>, filtered and concentrated under reduced pressure.

The crude product was purified by flash chromatography on silica gel (gradient: cHex/EtOAc 100:0 to 80:20, 40 mL/min, 18min) to afford the title compound (330 mg, 80%).

<sup>1</sup>H-NMR (400 MHz, CDCl<sub>3</sub>):  $\delta$  (ppm) = 7.37–7.32 (m, 4H), 7.26–7.23 (m, 1H), 6.06 (dd, *J*=12.3, 2.8 Hz, 1H), 5.43 (d, *J*=1.8 Hz, 1H), 4.18 (t, *J*=1.8 Hz, 1H), 4.00 (dd, *J*=11.0, 2.6 Hz, 1H), 3.55 (s, 3H), 3.49 (t, *J*=2.0 Hz, 1H), 3.41 (s, 3H), 3.38 (s, 3H), 3.37–3.35 (m, 1H), 3.16 (dd, *J*=8.1, 2.2 Hz, 1H), 3.06–3.01 (m, 2H), 2.95 (dd, *J*=8.3, 1.7 Hz, 1H), 2.37–2.29 (m, 1H), 2.06–1.97 (m, 2H), 1.72–1.66 (m, 1H), 1.64–1.55 (m, 2H), 1.47–1.44 (m, 1H), 1.40–1.28 (m, 2H), 1.18 (d, *J*=7.3 Hz, 3H), 1.08 (d, *J*=7.3 Hz, 3H), 1.04–0.97 (m, 1H), 0.94 (s, 9H), 0.60 (d, *J*=7.0 Hz, 3H), 0.17 (s, 3H), 0.15 (s, 3H);

<sup>13</sup>C-NMR (101 MHz, CDCl<sub>3</sub>): δ (ppm) = 170.83, 142.61, 128.48 (2C), 127.55, 126.20 (2C), 99.68, 83.66, 83.23, 71.87, 70.52, 68.14, 58.58, 58.07, 57.39, 55.13, 53.63, 46.39, 36.89, 35.49, 34.05, 30.39, 27.06, 25.80 (3C), 24.83, 24.18, 18.05, 11.26, 10.09, 8.92, -4.86, -4.99;

HPLC-MS: rt = 2.70 min, m/z = 674 [M+Na]<sup>+</sup>

HR-MS: m/z calculated for C<sub>35</sub>H<sub>58</sub>NaO<sub>9</sub>Si [(M+Na)<sup>+</sup>]: 673.3742, found 673.3727.

(1*R*,2*R*,3*R*,5*S*,6*S*,7*S*,12*S*,15*S*,16*R*,17*R*,18*S*,19*S*)-16,18-dihydroxy-6,7,17-trimethoxy-2,15,19-trimethyl-12-phenyl-4,13,20-trioxatricyclo[14.3.1.03,5]icosan-14-one (**4**)



To a solution of **S6** (660 mg, 1.01 mmol) in THF (10 mL) were added at 0°C AcOH (176  $\mu$ L, 3.04 mmol) and TBAF (2.00 mL, 1 M in THF, 2.03 mmol). The reaction mixture was stirred at 0°C for 2.5 h. The reaction mixture was poured into sat. aq. NH<sub>4</sub>Cl, then extracted with EtOAc. The combined organic layers were dried over Na<sub>2</sub>SO<sub>4</sub>, filtered and concentrated under reduced pressure.

The crude product was purified by flash chromatography on silica gel (gradient: cHex/EtOAc 85:15 to 60:40, 18 mL/min, 11 min) to afford the title compound (500 mg, 92%).

<sup>1</sup>H-NMR (400 MHz, CDCl<sub>3</sub>):  $\delta$  (ppm) = 7.35–7.24 (m, 5H), 5.86 (dd, *J*=12.3, 1.8 Hz, 1H), 4.47 (s, 1H), 4.11–4.08 (m, 2H), 3.59 (d, *J*=8.0 Hz, 1H), 3.54 (s, 3H), 3.43 (s, 3H), 3.40–3.38 (m, 1H), 3.37 (s, 3H), 3.30 (t, *J*=2.0 Hz, 1H), 3.17 (d, *J*=1.8 Hz, 1H), 3.13–3.07 (m, 2H), 2.97 (dd, *J*=8.4, 3.6 Hz, 1H), 2.32–2.25 (m, 1H), 2.10–2.04 (m, 1H), 1.87–1.81 (m, 1H), 1.73–1.62 (m, 3H), 1.55–1.44 (m, 3H), 1.25–1.20 (m, 1H), 1.14 (d, *J*=6.9 Hz, 3H), 1.05 (d, *J*=7.6 Hz, 3H), 0.64 (d, *J*=6.9 Hz, 3H);

<sup>13</sup>C-NMR (101 MHz, CDCl<sub>3</sub>): δ (ppm) = 170.72, 141.64, 128.59 (2C), 127.99, 126.14 (2C), 99.78, 82.98, 76.34, 74.08, 68.96, 68.87, 58.75, 58.40, 57.35, 55.78, 54.43, 53.53, 46.67, 36.47, 35.40, 34.02, 29.47, 25.78, 23.73, 11.58, 10.25, 9.01;

HPLC-MS: rt = 1.86 min, m/z = 535 [M-H]<sup>-</sup>

HR-MS: m/z calculated for C<sub>29</sub>H<sub>44</sub>NaO<sub>9</sub> [(M+Na)<sup>+</sup>]: 559.2878, found 559.2882.

#### **Purification of bio-extracts**

Starting material	erial Structure of isolated compounds			
Soraphen A				
	1a	1c	1b	
Soraphen C				
	2a			

Supplementary Table 8. Structures of isolated modified soraphen compounds.

All bioextracts were purified by preparative reverse-phase HPLC. In each case, the bioextract was dissolved in 1 mL of DMSO and injected on a Fraction Lynx Prep HPLC equipped with a Column Hichrom C18 ODS-2 5 250 mm x 2.1mm i.d. The mobile phase consisted of a mixture of  $H_2O+0.1\%$  HCOOH and ACN+0.1% HCOOH. The flow was set at 20 mL min<sup>-1</sup>. The gradients used for the purification of the different extracts are presented below.

Supplementary lable 9. Prep. HPLC gradient for 1a and
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Time (min)	% H <sub>2</sub> O + 0.1% HCOOH	% ACN + 0.1% HCOOH
0	60	40
2	60	40
37	0	100
40	1	100
41	50	50
45	50	50

(1a) rt = 23.6 min; (1c) rt = 11.7 min;

#### Supplementary Table 10. Prep. HPLC gradient for 1b.

First column:			Second o	olumn:	
Time	% H <sub>2</sub> O	% ACN	Time	% H₂O	% ACN
(min)	+ 0.1% HCOOH	+ 0.1% HCOOH	(min)	+ 0.1% HCOOH	+ 0.1% HCOOH
0	60	40	0	50	50
2	60	40	2	50	50
35	0	100	35	0	100
40	0	100	40	0	100
41	60	40	41	50	50
45	60	40	45	50	50

(1b) rt (second column)= 20.6 min.

#### Supplementary Table 11. Prep. HPLC gradient for 2a.

Time (min)	% H <sub>2</sub> O + 0.1% HCOOH	% ACN + 0.1% HCOOH
0	60	40
2	60	40
37	0	100
40	0	100
41	60	40
45	60	40

#### Analytical data of the halogenated and hydroxylated products

(1*R*,2*S*,5*S*,10*S*,11*R*,12*E*,14*S*,15*S*,16*S*,17*S*,18*R*)-8-chloro-1,17-dihydroxy-10,11,18-trimethoxy-2,14,16-trimethyl-5-phenyl-4,19-dioxabicyclo[13.3.1]nonadec-12-en-3-one (**1a**)



Isolated yield: 10% (11.0 mg)

<sup>1</sup>H-NMR (600 MHz, CDCl<sub>3</sub>) δ (ppm) = 7.36–7.34 (m, 4H), 7.30 (dq, *J*=8.5, 4.3 Hz, 1H), 6.21 (dd, *J*=16.3, 3.7 Hz, 1H), 5.92 (dd, *J*=11.8, 2.5 Hz, 1H), 5.49 (ddd, *J*=16.2, 9.5, 1.7 Hz, 1H), 4.57 (s, 1H), 4.12–4.06 (m, 1H), 4.00 (tt, *J*=11.7, 2.5 Hz, 1H), 3.88 (dt, *J*=10.7, 2.7 Hz, 1H), 3.77 (dd, *J*=10.5, 2.5 Hz, 1H), 3.68 (dd, *J*=9.5, 1.9 Hz, 1H), 3.48 (s, 3H), 3.38 (s, 3H), 3.32 (s, 3H), 3.19-3.15 (m, 2H), 3.12 (q, *J*=7.0 Hz, 1H), 2.53–2.49 (m, 1H), 2.21–2.15 (m, 1H), 2.06–2.02 (m, 1H), 2.01–1.96 (m, 1H), 1.92–1.89 (m, 1H), 1.88–1.85 (m, 1H), 1.77–1.73 (m, 1H), 1.69 (ddd, *J*=13.9, 11.5, 2.1 Hz, 1H), 1.56 (br s, 1H), 1.13 (d, *J*=7.1 Hz, 3H), 1.07 (d, *J*=7.6 Hz, 3H), 1.02 (d, *J*=6.7 Hz, 3H);

<sup>13</sup>C-NMR (151 MHz, CDCl<sub>3</sub>) δ = 170.97, 141.00, 140.12, 128.75 (2C), 128.30, 126.15 (2C), 122.97, 99.54, 84.03, 79.98, 76.33, 73.15, 72.40, 69.22, 58.48, 57.47, 56.89, 56.51, 46.49, 39.67, 35.85, 35.42, 34.33, 32.90, 12.58, 11.71, 10.43 ppm;

HPLC-MS: rt = 2.07 min, m/z = 578/580 [M+H+Na]<sup>2+</sup>, 553/555 [M–H]<sup>-</sup> chloro isotopic pattern

(1*R*,2*S*,5*S*,10*S*,11*R*,12*E*,14*S*,15*S*,16*S*,17*S*,18*R*)-1,8,17-trihydroxy-10,11,18-trimethoxy-2,14,16-trimethyl-5-phenyl-4,19-dioxabicyclo[13.3.1]nonadec-12-en-3-one (**1***c*)



Isolated yield: 3% (3.7 mg)

<sup>1</sup>H-NMR (600 MHz, CDCl<sub>3</sub>) δ = 7.36–7.33 (m, 4H), 7.30 (dq, *J*=8.7, 4.2 Hz, 1H), 6.16 (dd, *J*=16.3, 3.7 Hz, 1H), 5.87 (dd, *J*=10.2, 4.7 Hz, 1H), 5.49 (ddd, *J*=16.2, 9.3, 1.8 Hz, 1H), 4.48 (s, 1H), 4.05 (br d, *J*=8.7 Hz, 1H), 3.86 (br s, 1H), 3.81 (br dd, *J*=10.5, 2.5 Hz, 1H), 3.76–3.73 (m, 1 H), 3.72 (dd, *J*=9.3, 1.8 Hz, 1H), 3.47 (s, 3H), 3.42 (br d, *J*=9.3 Hz, 1H), 3.38 (s, 3H), 3.30 (s, 3H), 3.18 (d, *J*=1.6 Hz, 1H), 3.13 (q, *J*=7.0 Hz, 1H), 2.51 (m, 1H), 2.17 (s, 1H), 2.05–1.96 (m, 2H), 1.97–1.91 (m, 1H), 1.69 (ddd, *J*=14.0, 10.4, 1H), 1.55–1.48 (m, 5H), 1.11 (d, *J*=7.1 Hz, 3H), 1.06 (d, *J*=7.5 Hz, 3H), 1.03 (d, *J*=6.7 Hz, 3H) ppm

<sup>13</sup>C-NMR (151 MHz, CDCl<sub>3</sub>) δ = 170.93, 140.83, 140.18, 128.75 (2C), 128.37, 126.41 (2C), 122.99, 99.60, 84.83, 80.05, 76.30, 74.32, 72.39, 69.11, 65.78, 58.36, 57.49, 56.44, 46.38, 38.78, 35.80, 35.46, 33.53, 31.78, 31.08, 12.69, 11.71, 10.51 ppm

MS: m/z = 559 [M+Na]<sup>+</sup>

(2*S*,5*R*,10*S*,11*R*,12*E*,14*S*,15*S*,16*S*,17*S*,18*R*)-6-chloro-1,17-dihydroxy-10,11,18-trimethoxy-2,14,16-trimethyl-5-phenyl-4,19-dioxabicyclo[13.3.1]nonadec-12-en-3-one (**1b**)



Isolated yield: 4% (4.1 mg)

<sup>1</sup>H-NMR (600 MHz, CDCl<sub>3</sub>)  $\delta$  = 7.41–7.35 (m, 4H), 7.35–7.32 (m, 1H), 6.22 (dd, *J*=16.2, 3.9 Hz, 1H), 5.88 (d, *J*=9.1Hz, 1H), 5.48 (ddd, *J*=16.2, 9.5, 1.8 Hz, 1H), 4.39 (s, 1H), 4.38–4.35 (m, 1H), 4.12–4.06 (m, 1H), 3.79 (dd, *J*=10.5, 2.7 Hz, 1H), 3.63 (dd, *J*=9.5, 2.1 Hz, 1H), 3.47–3.43 (m, 4H), 3.37 (s, 3H), 3.30 (s, 3H), 3.16–3.14 (m, 1H), 3.14–3.09 (m, 2H), 2.56–2.46 (m, 1H), 2.17 (s, 1H) 2.03 (qd, *J*=9.9, 4.5 Hz, 1H) 1.90 (q, *J*=7.5 Hz, 1H) 1.87–1.80 (m, 1H), 1.66–1.61 (m, 2H), 1.45–1.36 (m, 1H), 1.35–1.25 (m, 2H), 1.07 (d, *J*=6.3 Hz, 3H), 1.06 (d, *J*=6.0 Hz, 3H), 1.02 (d, *J*=6.8 Hz, 3H) ppm;

<sup>13</sup>C-NMR (151 MHz, CDCl<sub>3</sub>) δ = 170.01, 140.07, 137.28, 128.99, 128.40 (2C), 128.05 (2C), 122.64, 99.66, 84.90, 83.23, 76.26, 75.65, 72.68, 69.27, 62.15, 58.25, 57.52, 56.39, 46.37, 35.82, 35.30, 33.83, 29.53, 18.98, 12.61, 11.55, 10.39 ppm.

HPLC-MS: rt = 1.80 min, m/z = 564/566 [M+H+Na]<sup>2+</sup>, 539/541 [M–H]<sup>-</sup> chloro isotopic pattern

(1*R*,2*S*,5*S*,10*S*,11*R*,12*E*,14*S*,15*S*,16*S*,17*S*,18*R*)-8-chloro-1,11,17-trihydroxy-10,18-dimethoxy-2,14,16-trimethyl-5-phenyl-4,19-dioxabicyclo[13.3.1]nonadec-12-en-3-one (**2a**)



Isolated yield: 10% (16.1 mg)

<sup>1</sup>H-NMR (600 MHz, CDCl<sub>3</sub>)  $\delta$  = 7.38–7.33 (m, 4H), 7.33–7.28 (m, 1H), 6.16 (dd, *J*=16.1, 3.8 Hz, 1H), 5.89 (dd, *J*=11.7, 2.7 Hz, 1H), 5.50 (ddd, *J*=16.1, 9.4, 1.8 Hz, 1H), 4.58 (s, 1H), 4.19 (dd, *J*=9.3, 2.1 Hz, 1H), 4.08 (br s, 1H), 4.00 (tt, *J*=11.5, 2.7 Hz, 1H), 3.81 (dt, *J*=11.0, 2.9 Hz, 1H), 3.76 (dd, *J*=10.6, 2.6 Hz, 1H), 3.46 (s, 3H), 3.38 (s, 3H), 3.30 (br d, *J*=7.3 Hz, 1H), 3.17 (d, *J*=1.8 Hz, 1H), 3.12 (q, *J*=7.2 Hz, 1H), 2.51–2.43 (m, 1H), 2.39–2.29 (m, 1H), 2.17 (ddt, *J*=15.0, 11.6, 3.5, 3.5 Hz, 1H), 2.08–2.01 (m, 2H), 1.93–1.88 (m, 1H), 1.84 (ddt, *J*=14.8, 11.8, 3.4, 3.4 Hz, 1H), 1.75 (ddt, *J*=14.9, 11.2, 4.0, 4.0 Hz, 1H), 1.69–1.64 (m, 2H), 1.11 (d, *J*=7.0 Hz, 3H), 1.06 (d, *J*=7.5 Hz, 3H), 1.00 (d, *J*=6.8 Hz, 3H) ppm

 $^{13}\text{C-NMR}$  (151 MHz, CDCl<sub>3</sub>)  $\delta$  = 170.83, 140.85, 137.99, 128.77 (2C), 128.34, 126.17 (2C), 125.19, 99.54, 81.27, 76.29, 74.54, 73.31, 72.42, 69.18, 58.09, 57.47, 56.58, 46.39, 39.15, 35.86, 35.35, 34.29, 32.84, 12.60, 11.71, 10.47 ppm

MS:  $m/z = 563/565 [M+Na]^+$  chloro isotopic pattern

#### Supplementary Table 12. Assignment Table <sup>1</sup>H-NMR.

Chemical shifts (in ppm) for multiplets were taken at the center of the multiplet for clarity. The assignment of the peaks was done based on analysis of 2D spectra: COSY, HSQC, HMBC, ROESY.



Carbon		(1a)	(1b)	(1c)	Soraphen C	(2a)
Carbon	Soraphen A	R=Me;	R=Me;	R=Me;	R=H;	R=H;
number		X1=CI, X2=H	X <sub>1</sub> =H, X <sub>2</sub> =Cl	X <sub>1</sub> =OH, X <sub>2</sub> =H	X1=H, X2=H	X <sub>1</sub> =Cl, X <sub>2</sub> =H
2	3.14	3.15	3.12	3.16	3.15	3.12
4	3.18	3.19	3.15	3.20	3.18	3.17
5	4.02	4.12	4.09	4.07	4.02	4.00
6	1.94	1.95	1.90	1.95	1.93	1.90
7	3.83	3.79	3.79	3.84	3.82	3.76
8	2.49	2.54	2.51	2.54	2.45	2.47
9	6.17	6.23	6.22	6.19	6.15	6.16
10	5.48	5.51	5.48	5.51	5.49	5.50
11	3.68	3.71	3.63	3.75	4.17	4.19
12	3.41	3.91	3.45	3.77	3.36	3.81
12	1.24	1.72	1.34	1.52	1.17	1.66
15	1.69	2.01	1.64	1.71	1.77	2.05
1/	1.16	4.02	1.43	3 86	1.17	4.00
14	1.46	4.02	1.64	5.80	1.51	4.00
15	1.34	1.77	1.86	1.58	1.35	1.75
	1.46	1.89	2.03	1.58	1.48	1.85
16	1.67	2.07	1 37	2.02	1.67	2.05
	2.10	2.20	4.57	2.05	2.10	2.18
17	5.84	5.95	5.88	5.90	5.82	5.89
18	1.11	1.15	1.07	1.14	1.09	1.11
19	3.38	3.41	3.37	3.41	3.38	3.38
20	1.05	1.10	1.06	1.19	1.05	1.06
21	1.03	1.05	1.02	1.05	1.00	1.00
22	3.29	3.35	3.30	3.33	/	/
23	3.44	3.50	3.45	3.49	3.43	3.46
2′	7.32	7.38	7.38	7.37	7.35	7.35
3′	7.32	7.38	7.38	7.38	7.35	7.35
4'	7.32	7.32	7.34	7.33	7.27	7.30
5'	7.32	7.38	7.38	7.38	7.35	7.35
6'	7.32	7.38	7.38	7.37	7.35	7.35

# NMR spectra

Supplementary Figure 16. NMR of compound 1.



Supplementary Figure 17. NMR of compound S1.



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Supplementary Figure 18. NMR of compound S2.



Supplementary Figure 19. NMR of compound 3.



Supplementary Figure 20. NMR of compound S3.



Supplementary Figure 21. NMR of compound S4.



Supplementary Figure 22. NMR of compound S5.



Supplementary Figure 23. NMR of compound soraphen C, 2.







Supplementary Figure 24. NMR of compound *epi*-soraphen C, S7.



Supplementary Figure 25. NMR of compound S6.



Supplementary Figure 26. NMR of compound 4.



Supplementary Figure 27. NMR of compound 1a.







Supplementary Figure 28. NMR of compound 1c.







Supplementary Figure 29. NMR of compound 1b.







Supplementary Figure 30. NMR of compound 2a.







#### Methods for biological testing and BP80 determination

The assays were performed at Syngenta's high-throughput screening facilities, using standardized assays and the necessary standards.

#### Leaf disk assays:

*Erysiphe graminis f.sp. tritici* (Wheat powdery mildew): preventive application

Barley leaf segments were placed on agar in multiwell plates (24-well format) and sprayed with test solutions. After drying, the leaf disks were inoculated with spores of the fungus. After appropriate incubation the activity of a compound was assessed 7 dpi (days post inoculation) as preventive fungicidal activity.

#### Puccinia recondita (Brown rust): curative application

Wheat leaf segments are placed on agar in multiwell plates (24-well format). The leaf disks are then inoculated with a spore suspension of the fungus. One day after inoculation the test solution is applied. After appropriate incubation the activity of a compound is assessed 8 dpi (days post inoculation) as curative fungicidal activity.

#### Liquid culture assays:

#### Botrytis cinerea (Gray mould):

Conidia of the fungus from cryogenic storage were directly mixed into nutrient broth (Vogel's minimal media). A DMSO solution of the test compounds was placed into a microtiter plate (96-well format) and the nutrient broth containing the fungal spores was added to it. The test plates were incubated at 24 °C and the inhibition of growth was determined photometrically after 72 hours at 620 nm.

#### Mycosphaerella arachidis (Brown leaf spot of peanut):

Conidia of the fungus from cryogenic storage were directly mixed into nutrient broth (PDB potato dextrose broth). A DMSO solution of the test compounds was placed into a microtiter plate (96-well format) and the nutrient broth containing the fungal spores was added to it. The test plates were incubated at 24 °C and the inhibition of growth was determined photometrically after approximately 5-6 days at 620 nm.

#### Septoria tritici (leaf blotch):

Conidia of the fungus from cryogenic storage were directly mixed into nutrient broth (PDB potato dextrose broth). A DMSO solution of the test compounds was placed into a microtiter plate (96-well format) and the nutrient broth containing the fungal spores was added to it. The test plates were incubated at 24 °C and the inhibition of growth was determined photometrically after 72 hours at 620 nm.

#### Monographella nivalis (snow mould, foot rot of cereals):

Conidia of the fungus from cryogenic storage were directly mixed into nutrient broth (PDB potato dextrose broth). A DMSO solution of the test compounds was placed into a microtiter plate (96-well format) and the nutrient broth containing the fungal spores was added to it. The test plates were incubated at 24 °C and the inhibition of growth was determined photometrically after 72 hours at 620 nm.

#### **BP80 determination:**

A dilution series was performed for each fungus/compound combination and the efficiency of the compounds evaluated. BP80 represents the breakpoint below which less than 80% efficiency is observed.

The experiments on living organisms were performed as single or triplicate experiments, against positive and negative standards (both commercial and company internal). The triplicate experiments were technical triplicates, with three samples tested in parallel during the same test session. The pest control percentage is assessed visually by a trained personal and is assigned to be 100%, 90%, 70%, 50%, 20% or 0%. The BP80 was determined as an average over all the results.

Compound 1								
	Rate of	Rate of		% Pe				
Pathogen tested	application	application	Test 1		Test 2		pBP80	BP80 (uM)
				replicate 1	replicate 2	replicate 3		
	22.2	42.6	100	100	100	100		
Enveinho graminis	7.39	14.2	100	100	100	100	5 9	0.16
Liysipne grannins	2.46	4.7	100	100	100	100	5.6	0.10
	0.819	1.6	100	100	100	100		
	22.2	42.6	100	100	100	100		
Russinia recondita	7.39	14.2	100	100	100	100	5 9	0.16
Puccinia reconulta	2.46	4.7	100	100	100	100	5.8	
	0.819	1.6	100	100	100	100		
	6.67	12.8	100	100	100	100	6.8	0.016
Potrutio cinoroo	2.22	4.3	100	100	100	100		
Bolrylis cinerea	0.739	1.4	100	100	100	100		
	0.246	0.5	100	100	100	100		
	6.67	12.8	100	100	100	100		0.050
Mycosphaerella arachidis	2.22	4.3	100	100	100	100	6.3	
	0.739	1.4	100	100	100	100		
	6.67	12.8	100	100	100	100		
Oo ato sia tsitiai	2.22	4.3	100	100	100	100	0.0	
Septoria tritici	0.739	1.4	100	100	100	100	6.3	0.050
	0.246	0.5	100	70	70	70		
	6.67	12.8	100	100	100	100		
Monographella	2.22	4.3	100	100	100	100		0.040
nivalis	0.739	1.4	100	100	100	100	٥.४	0.016
	0.246	0.5	100	100	100	100		

#### Supplementary Table 13. BP80 determination for compound 1.

Compound 1a								
	Rate of	Rate of		% Pe	st control			
Pathogen tested	application	application	Test 1		Test 2		pBP80	BP80 (uM)
	in ppm			replicate 1	replicate 2	replicate 3		
	22.2	42.6	100	90	100	100		
Envsinhe graminis	7.39	14.2	50	20	50	70	11	3 08
Liysiphe grammis	2.46	4.7	20	0	0	20	4.4	5.50
	0.819	1.6	0	0	0	0		
	22.2	42.6	90	20	70	100		
Puccinia recondita	7.39	14.2	20	20	0	0	11	0.40
Fuccinia recondita	2.46	4.7	20	0	0	0	4.4	
	0.819	1.6	0	0	0	0		
	6.67	12.8	100	90	100	100	4.9	1.26
Dotatio cincros	2.22	4.3	50	70	70	70		
Bollylis cinerea	0.739	1.4	20	20	20	20		
	0.246	0.5	20	0	0	0		
	6.67	12.8	90	100	70	90		1.26
Mycosphaerella arachidis	2.22	4.3	0	20	20	20	4.9	
	0.739	1.4	0	0	0	0		
	6.67	12.8	20	20	70	50		
Contorio tritici	2.22	4.3	0	0	0	0	0	1000
Septona tritter	0.739	1.4	0	0	0	0	0	1000
	0.246	0.5	0	0	0	0		
	6.67	12.8	100	100	90	90		
Monographella	2.22	4.3	50	50	50	50	4.0	4.00
nivalis	0.739	1.4	20	0	0	0	4.9	1.20
	0.246	0.5	20	0	0	0		

# Supplementary Table 14. BP80 determination for compound 1a.

Compound 1b								
	Rate of	Rate of		% Pest control				
Pathogen tested	application	application	Test 1		Test 2		pBP80	BP80 (uM)
	in ppm	in uw		replicate 1	replicate 2	replicate 3		
	22.2	42.6	100	100	100	100		
Envsinhe graminis	7.39	14.2	100	100	100	100	5.8	0.16
Erysiphe grammis	2.46	4.7	100	90	100	90	5.0	0.10
	0.819	1.6	100	70	50	70		
	22.2	42.6	100	100	100	100		
Puccinia recondita	7.39	14.2	50	100	90	100	10	1.26
	2.46	4.7	0	50	90	20	4.5	
	0.819	1.6	0	0	0	20		
	6.67	12.8	100	100	100	100	5.4	0.40
Potrutio cinoroo	2.22	4.3	90	90	90	100		
Bollylis cinerea	0.739	1.4	50	90	70	70		
	0.246	0.5	0	50	0	0		
	6.67	12.8	100	100	100	100		
Mycosphaerella arachidis	2.22	4.3	50	90	50	20	4.9	1.26
	0.739	1.4	0	0	0	20		
	6.67	12.8	100	100	100	100		
Contorio tritici	2.22	4.3	70	20	50	90	4.0	4.00
Septona tritici	0.739	1.4	50	0	0	20	4.9	1.20
	0.246	0.5	0	0	0	0		
	6.67	12.8	100	100	100	100		
Monographella	2.22	4.3	100	100	90	100	5.4	0.40
nivalis	0.739	1.4	70	90	50	20	5.4	0.40
	0.246	0.5	20	0	0	20		

# Supplementary Table 15. BP80 determination for compound 1b.

Compound 1c								
	Rate of	Rate of		% Pe				
Pathogen tested	application	application	Test 1		Test 2		pBP80	BP80 (uM)
	in ppm			replicate 1	replicate 2	replicate 3		
	22.2	42.6	90	70	70	70		
Envsinhe graminis	7.39	14.2	0	0	50	20	11	3 08
Liysiphe grammis	2.46	4.7	0	0	0	20	4.4	5.90
	0.819	1.6	0	0	0	0		
	22.2	42.6	0	0	0	0		
Russinia recondita	7.39	14.2	0	0	0	0	0.05	1000
Puccinia recondita	2.46	4.7	0	0	0	0	0.05	
	0.819	1.6	0	0	0	0		
	6.67	12.8	0	0	0	0	0	
Detection aimenes	2.22	4.3	0	0	0	0		1000
Botrytis cinerea	0.739	1.4	0	0	0	0		1000
	0.246	0.5	0	0	0	0		
	6.67	12.8	0	0	0	0		1000
Mycosphaerella arachidis	2.22	4.3	0	0	0	0	0	
	0.739	1.4	0	0	0	0		
	6.67	12.8	20	0	0	0		
Contonio tritici	2.22	4.3	0	0	0	0	0	1000
Septoria tritici	0.739	1.4	0	0	0	0	0	1000
	0.246	0.5	0	0	0	0		
	6.67	12.8	20	0	0	0		
Monographella	2.22	4.3	20	0	0	0	0	1000
nivalis	0.739	1.4	0	0	0	0	U	1000
	0.246	0.5	0	0	0	0		

# Supplementary Table 16. BP80 determination for compound 1c.

Compound 2								
	Rate of	Rate of		% Pest control				
Pathogen tested	application	application	Test 1		Test 2		pBP80	BP80 (uM)
	in ppm	in uw		replicate 1	replicate 2	replicate 3		
	22.2	42.6	100	100	100	100		
Envsinhe graminis	7.39	14.2	100	100	100	100	5.8	0.16
Erysiphe grammis	2.46	4.7	100	100	100	100	5.0	0.10
	0.819	1.6	90	100	100	100		
	22.2	42.6	100	100	100	100		0.16
Puccinia recondita	7.39	14.2	100	100	100	100	5.8	
	2.46	4.7	100	100	100	100	5.0	
	0.819	1.6	100	90	100	90		
	6.67	12.8	100	100	100	100	5.8	0.16
Botrutis cinerea	2.22	4.3	100	100	100	100		
Bollyus cinerea	0.739	1.4	50	90	90	90		0.10
	0.246	0.5	20	20	20	20		
	6.67	12.8	100	100	100	100		0.4.
Mycosphaerella arachidis	2.22	4.3	90	90	90	90	5.4	
	0.739	1.4	50	0	20	50		
	6.67	12.8	90	100	100	100		
Sontorio tritici	2.22	4.3	70	70	70	70	5.4	0.40
Septona tritto	0.739	1.4	20	0	0	0	5.4	0.40
	0.246	0.5	0	0	0	0		
	6.67	12.8	100	100	100	100		
Monographella	2.22	4.3	100	90	90	90	E A	0.10
nivalis	0.739	1.4	70	50	50	50	5.4	0.40
	0.246	0.5	50	0	20	0		

# Supplementary Table 17. BP80 determination for compound 2.

Compound 2a								
	Rate of Rate of			% Pe				
Pathogen tested	application	application	Test 1		Test 2		pBP80	BP80 (uM)
	in ppm			replicate 1	replicate 2	replicate 3		
	22.2	42.6	100	100	100	100		
Envsinhe graminis	7.39	14.2	100	50	100	50	5.8	1 26
Liysiphe grammis	2.46	4.7	0	0	70	0	5.0	1.20
	0.819	1.6	0	0	0	0		
	22.2	42.6	100	100	100	100		
Russinia recondita	7.39	14.2	50	90	90	100	4.0	1.26
Puccinia recondita	2.46	4.7	0	0	90	0	4.9	
	0.819	1.6	0	0	50	0		
	6.67	12.8	50	90	90	90	4.9	1.26
Detrutia sinaraa	2.22	4.3	20	50	50	50		
Botrytis cinerea	0.739	1.4	0	0	0	0		
	0.246	0.5	0	0	0	0		
	6.67	12.8	50	20	20	20		1000
Mycosphaerella arachidis	2.22	4.3	0	0	0	0	0.05	
	0.739	1.4	0	0	0	0		
	6.67	12.8	20	0	0	20		
Contonio tritici	2.22	4.3	0	0	0	0	0	
Septoria tritici	0.739	1.4	0	0	0	0	0	1000
	0.246	0.5	0	0	0	0		
	6.67	12.8	50	50	70	70		
Monographella	2.22	4.3	20	20	20	20		4000
nivalis	0.739	1.4	0	0	0	0	0.05	1000
	0.246	0.5	0	0	0	0		

# Supplementary Table 18. BP80 determination for compound 2a.

#### **Combinatorial library WelO5\***

Codon list incorporated in the WelO5<sup>\*</sup> site-saturation library. The most abundant codons of *E. coli* were chosen in library design. By replacing the wild-type codons on positions V81, A88, and I161 with the codons in the table, the DNA sequences of the constructed variants can be obtained.

Amino acid	Codon	Amino acid	Codon
A	GCA	М	ATG
С	TGT	Ν	AAT
D	GAT	Р	CCG
E	GAA	Q	CAG
F	TTT	R	CGT
G	GGT	S	AGC
Н	CAT	Т	ACC
I	ATC	V	GTT
К	AAA	W	TGG
L	CTG	Υ	TAT

**Supplementary Table 19.** Codon list incorporated in the WelO5\* site-saturation library.

#### DNA sequence of the ordered gene fragments:

>Twist\_gene\_fragment\_of\_the\_Wel05\*\_site\_saturation\_library

green = flanking sequence (Twist) yellow = flanking sequence including restriction sites **bold** = mutation site <u>underlined</u> = WelO5\* gene sequence (ORF)

>WelO5\* amino acid sequence

MSNNTISTKPALHFLDINATEVKKYPTAIQDIIINRSFDGMIIRGVFPRDTMEQVARCLEEGNDGGMKSILNKNE EFGTK**V**AQIYGH**A**IVGQSPDLKDYFASSAIFRQACRTMFQGSPDFEEQVESIFHSLSGLPVEIPTGPEGQTYTPA TIRLLLEGRE**I**AVHVGNDFLLMPAANHLKTLLDLSDQLSYFIPLTVPEAGGELVVYNLEWNPQEVDKSADLHKYI DEVESKFKSNQSQSVAYAPGPGDMLLFNGGRYYHRVSEVIGNSPRRTIGGFLAFSKERNKIYYWS-

**bold** = mutation site

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