

Article

Production of Satratoxin G and H is Tightly Linked to Sporulation in *Stachybotrys chartarum*

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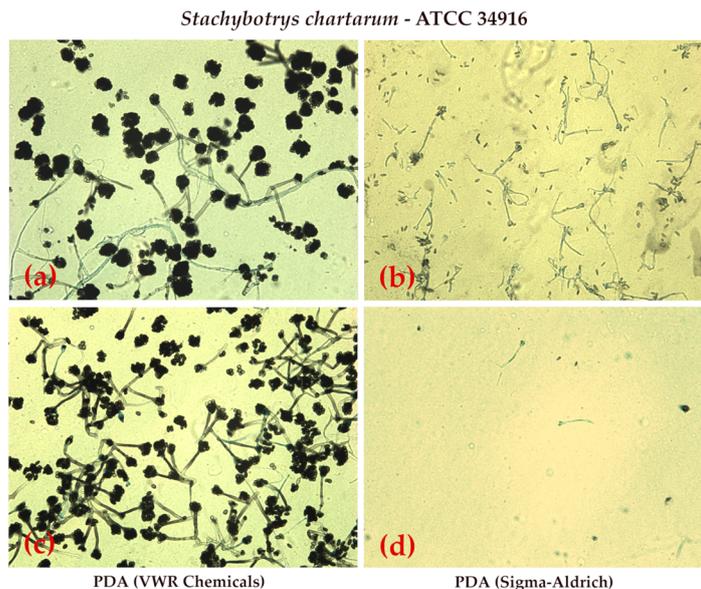


Figure S1. Microscopy (light microscopy, 200x, lactophenol blue staining) of *S. chartarum* genotype S strain ATCC 34916 as three-point (**a and b**) and one-point (**c and d**) culture shows the significantly reduced spore production on PDA-S (**b and d**) and thinner mycelium compared to the high rate of sporulation with well-grown mycelium on PDA-V (**a and c**).

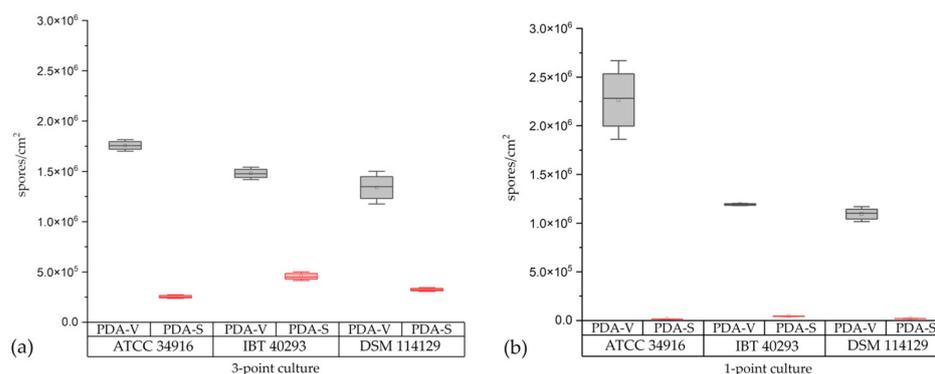


Figure S2. Spore count per cm² of *S. chartarum* genotype S strains (ATCC 34916, IBT 40293, and DSM 114129) on PDA-V (gray box plots) and PDA-S (red box plots) harvested from the nutrition media shown in Figure 1 (**a**: three-point cultures, **b**: one-point cultures).

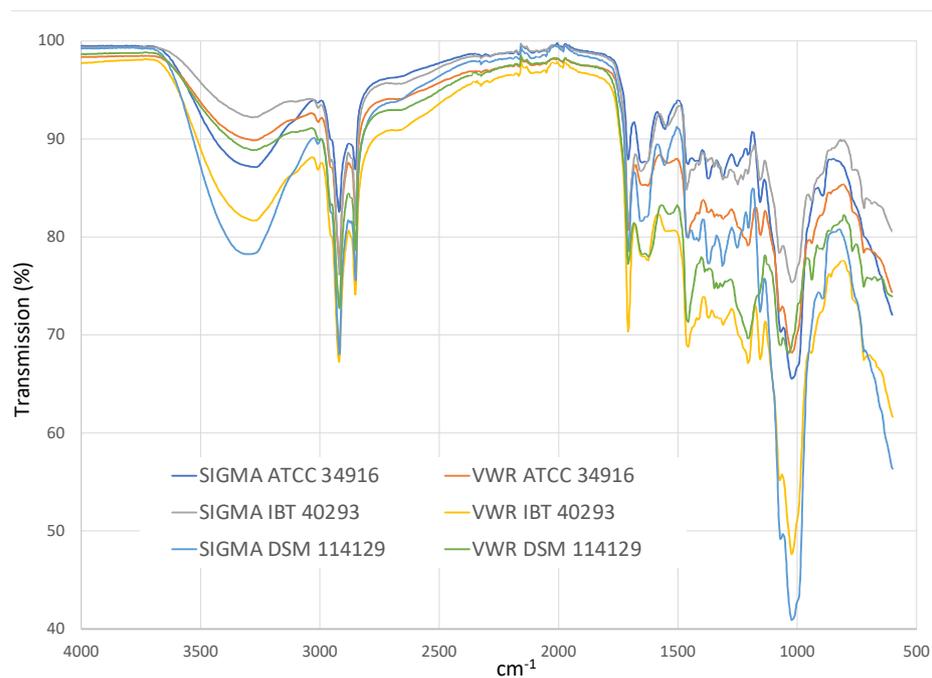


Figure S3. ATR-IR spectra of melanin extracted from *S. chartarum* genotype S strains (ATCC 34916: dark blue and orange, IBT 40293: gray and yellow, DSM 114129: light blue and green) grown on PDA-V and PDA-S (raw data can be found in Table S1).

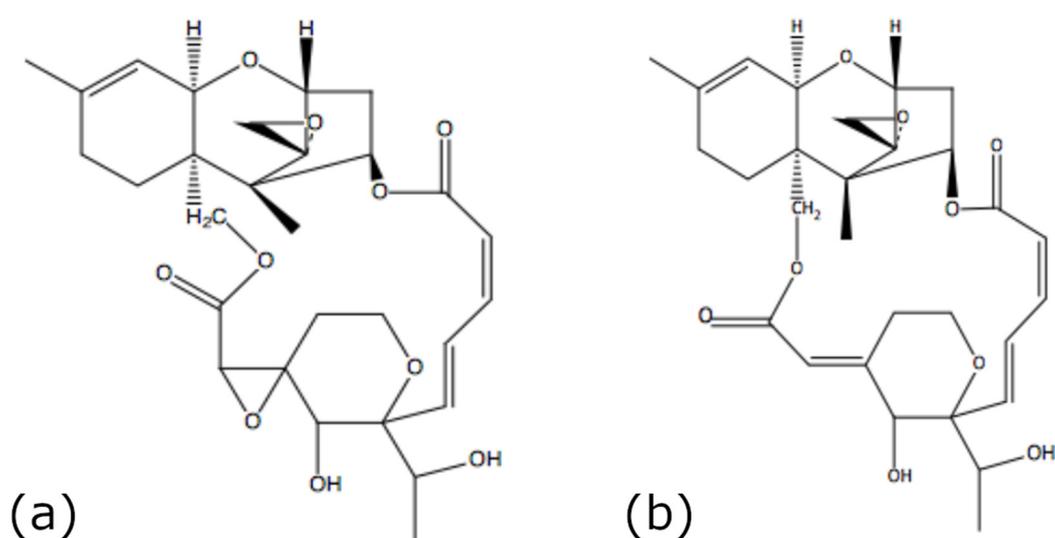


Figure S4. The chemical structures of the macrocyclic trichothecenes satratoxin G (a) and satratoxin H (b) produced by *S. chartarum* genotype S strains (Ulrich 2016).

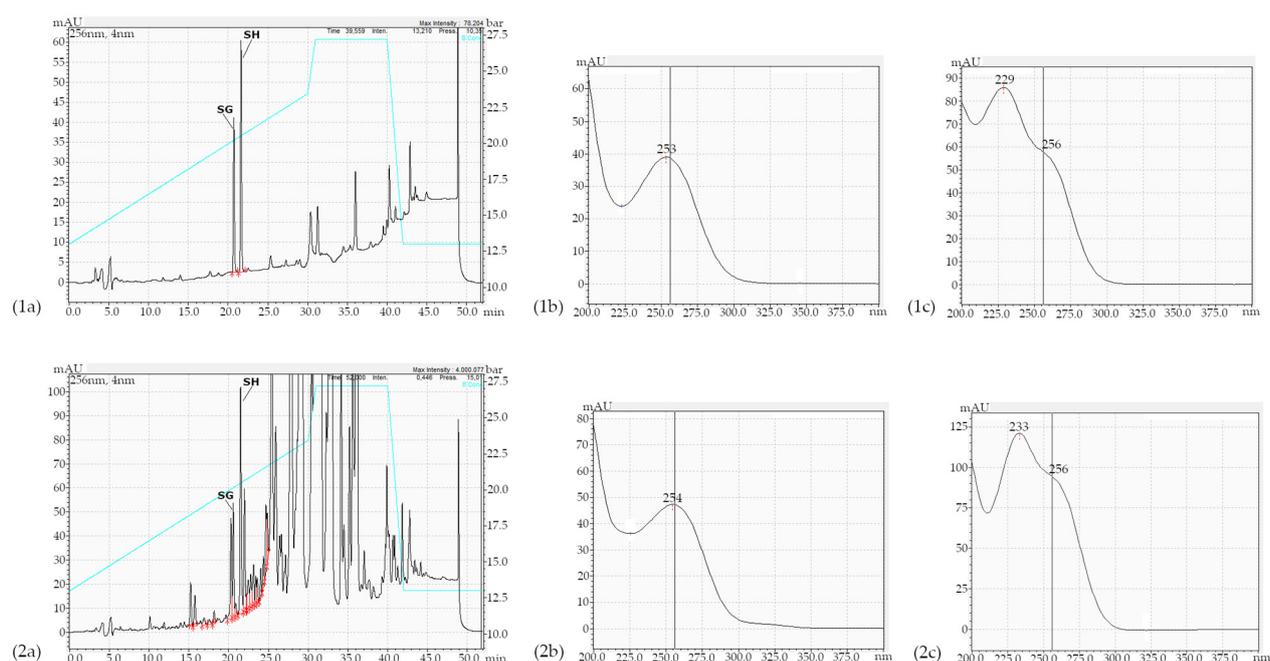


Figure S5. The HPLC chromatograms recorded at 256 nm of satratoxin G (SG) with a retention time (RT) of 20.73 min and satratoxin H (SH) with a RT of 21.62 min diluted in methanol (standard solution) at a concentration of 10.0 $\mu\text{g/mL}$ (1a) with the corresponding UV spectra for SG (1b) and SH (1c) and the detected satratoxin G (15.6 $\mu\text{g/mL}$, RT: 20.60 min) and satratoxin H (22.9 $\mu\text{g/mL}$, RT: 21.60 min) in a toxin extract of *S. chartarum* genotype S strain ATCC 34916 grown on PDA-V as a three-point culture (2a) with the corresponding UV spectra for SG (2b) and SH (2c).

Table S2. Parameters describing colonies of ATCC 34916, IBT 40293, and DSM 114129 on PDA-V and PDA-S were used to determine the spore count after 21 days of growth (3-point cultures & 1-point cultures).

	Colony area		Spore count	
	cm^2	spores/Agar Plate	spores/ cm^2	
ATCC 34916				
PDA-V	51.1 ± 0.5	$8.992 \times 10^7 \pm 2.765 \times 10^6$	$1.758 \times 10^6 \pm 3.817 \times 10^4$	
3-point culture				
IBT 40293				
PDA-V	49.1 ± 2.1	$7.275 \times 10^7 \pm 5.019 \times 10^6$	$1.480 \times 10^6 \pm 4.135 \times 10^4$	
3-point culture				
DSM 114129				
PDA-V	46.1 ± 0.8	$6.183 \times 10^7 \pm 6.010 \times 10^6$	$1.339 \times 10^6 \pm 1.085 \times 10^5$	
3-point culture				
ATCC 34916				
PDA-S	36.1 ± 1.1	$9.167 \times 10^6 \pm 2.566 \times 10^5$	$2.544 \times 10^5 \pm 1.244 \times 10^4$	
3-point culture				
IBT 40293				
PDA-S	38.0 ± 2.7	$1.735 \times 10^7 \pm 1.083 \times 10^6$	$4.572 \times 10^5 \pm 2.892 \times 10^4$	
3-point culture				
DSM 114129				
PDA-S	34.4 ± 0.2	$1.118 \times 10^7 \pm 4.752 \times 10^5$	$3.248 \times 10^5 \pm 1.278 \times 10^4$	
3-point culture				
ATCC 34916				
PDA-V	17.3 ± 0.4	$3.933 \times 10^7 \pm 5.150 \times 10^6$	$2.265 \times 10^6 \pm 2.689 \times 10^5$	
1-point culture				
IBT 40293	20.8 ± 0.4	$2.483 \times 10^7 \pm 6.292 \times 10^5$	$1.192 \times 10^6 \pm 7.351 \times 10^3$	

PDA-V			
1-point culture			
DSM 114129			
PDA-V	16.0 ± 0.2	1.750 × 10 ⁷ ± 1.000 × 10 ⁶	1.092 × 10 ⁶ ± 5.099 × 10 ⁴
1-point culture			
ATCC 34916			
PDA-S	9.2 ± 0.3	1.344 × 10 ⁵ ± 5.079 × 10 ³	1.466 × 10 ⁴ ± 6.075 × 10 ²
1-point culture			
IBT 40293			
PDA-S	6.5 ± 0.3	2.808 × 10 ⁵ ± 2.504 × 10 ⁴	4.297 × 10 ⁴ ± 1.658 × 10 ³
1-point culture			
DSM 114129			
PDA-S	8.8 ± 0.3	1.755 × 10 ⁵ ± 7.013 × 10 ³	1.991 × 10 ⁴ ± 6.816 × 10 ²
1-point culture			