

Supporting information

Light-controlled cell factories – Employing photocaged IPTG for light-mediated optimization of *lac*-based gene expression and (+)-valencene biosynthesis in *Corynebacterium glutamicum*

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DNA-Sequence of the synthetic codon-usage optimized CnVS gene (oCnVS):

ATGGCGGAAATGTTAACGGCAACAGCAGCAACGATGGCAGCAGCTGCATGCCGGTCAAAGATGCGCTGC
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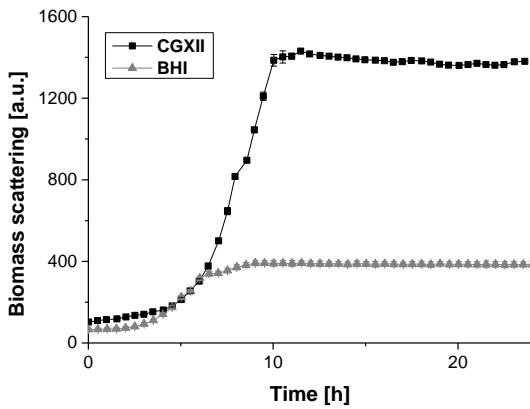


FIG S1 Growth curves of *C. glutamicum* cultures in triplicates in BHI complex (grey triangles) and CGXII minimal medium (black squares).

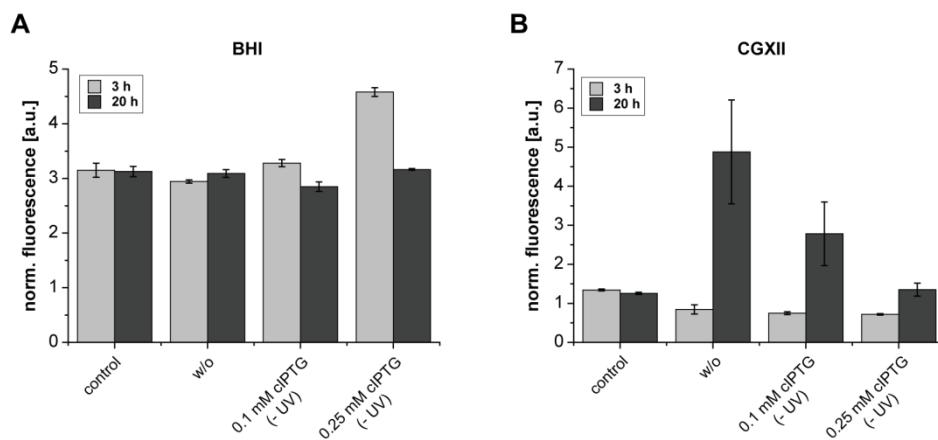


FIG S2 Basal expression background of clPTG supplemented cultures in BHI (A) and CGXII medium (B) in the dark. Normalized fluorescence values originate from biomass-normalized triplicates analog to values for induced gene expression depicted in Fig. 2B,D. Control: Wildtype control strain without the pEKEx-2-EYFP plasmid.

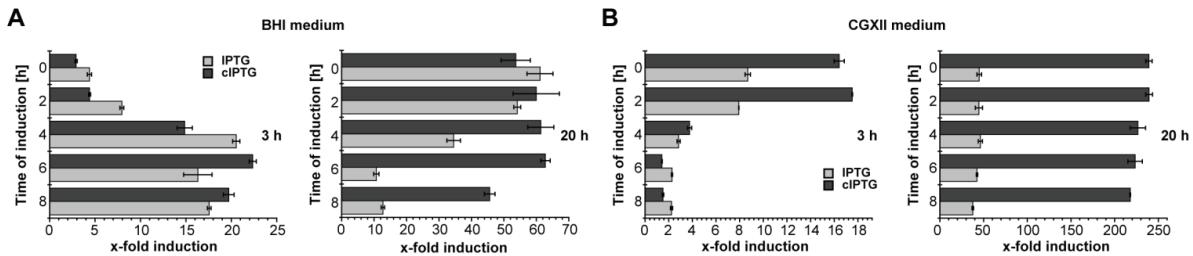


FIG S3 Dynamic range of induction for IPTG (light grey) and clIPTG-based (dark grey) induction after 3 (left) and 20 h (right) of expression in BHI (**A**) and CGXII medium (**B**) using *C. glutamicum* ATCC13032 (pEKEx-2-EYFP). Calculations originate from data depicted in Fig.2 B,C (biomass-normalized fluorescence) and Fig.S2 (basal fluorescence levels).

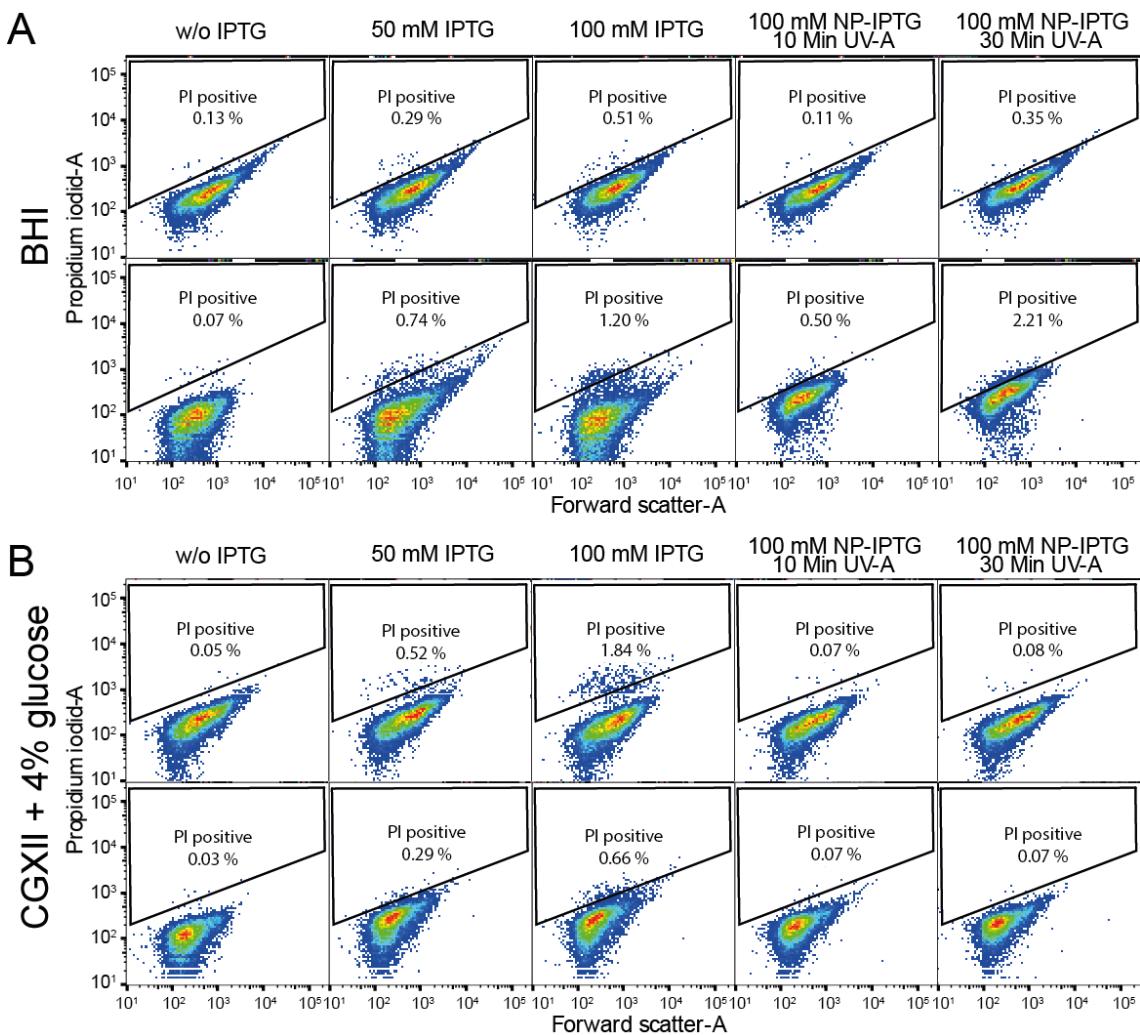


FIG S4 Propidium iodide-based live-dead-staining using flow cytometric single-cell analysis to evaluate cell viability.

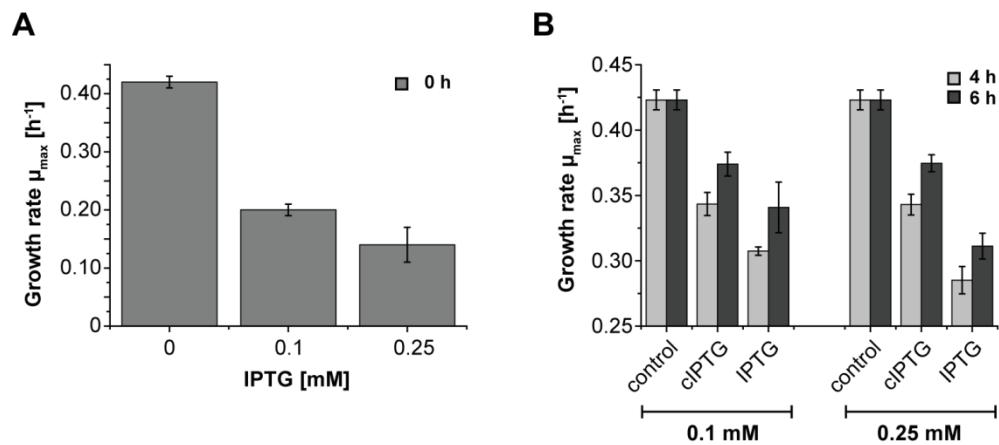


FIG S5 Reduction of growth impairment during (+)-valencene production in VLC6 *via* delayed induction and application of clIPTG-based light induction. **A)** Growth rates of VLC6 cultures for direct induction (0 h) are depicted for different IPTG inducer concentrations. **B)** Growth rates of VLC6 cultures are shown for delayed IPTG and clIPTG-induction after 4 (light grey) and 6 h (dark grey) together with un-induced cultures (control). All means and standard deviation derive from biological triplicates.

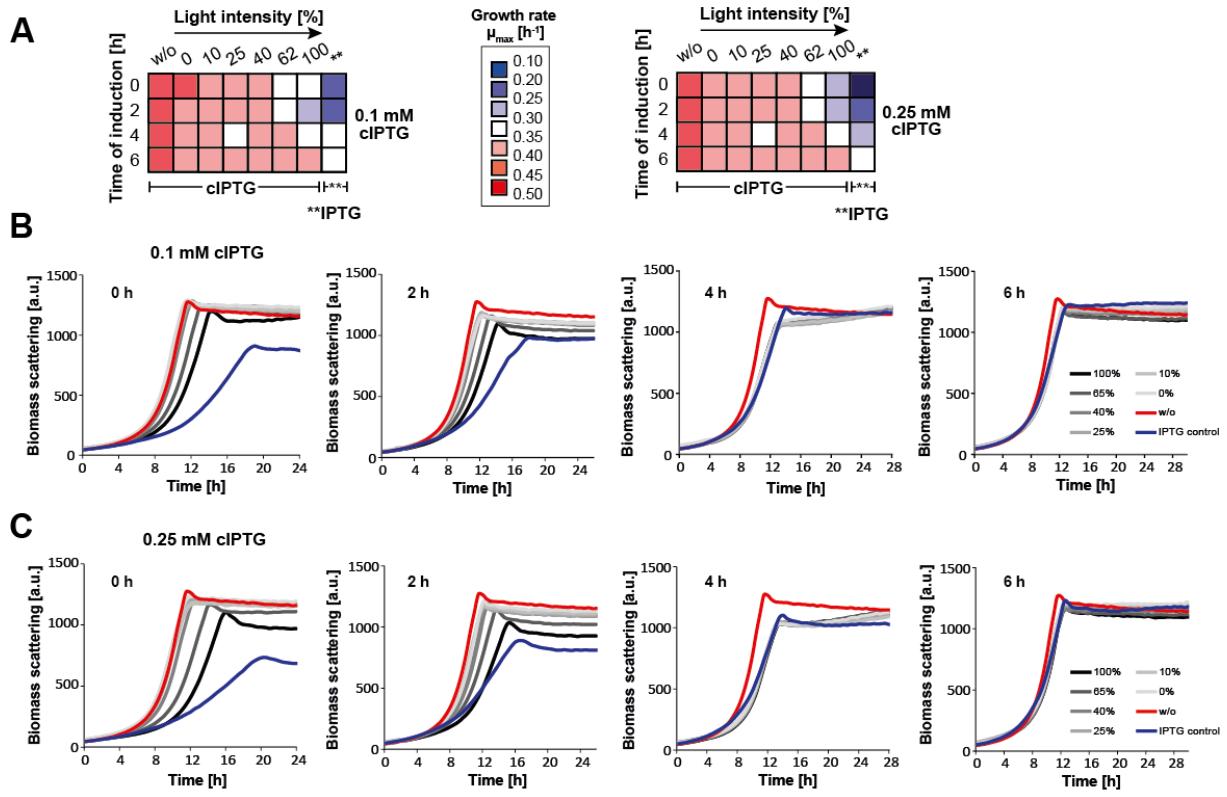


FIG S6 Growth analysis of *C. glutamicum* VLC6 during IPTG- and cIPTG-induced (+)-valencene production for different induction time points. **(A)** Maximum growth rate μ_{\max} intervals from low (blue) to high (red) in h⁻¹ after 24 h of production using different times of induction as well as different (c)IPTG concentrations (0.1 and 0.25 mM). **(B)** Growth curves of (+)-valencene production cultures supplemented with 0.1 mM (c)IPTG. Light induction (grey to black curves) is compared to IPTG induction (blue curve) after 0, 2, 4 and 6 h of cultivation and plotted against uninduced control cultures (w/o, red curve). **(C)** Growth curves of (+)-valencene production cultures analogous to Fig. S6 B supplemented with 0.25 mM (c)IPTG. All averaged data originate from at least three independent biological triplicates.

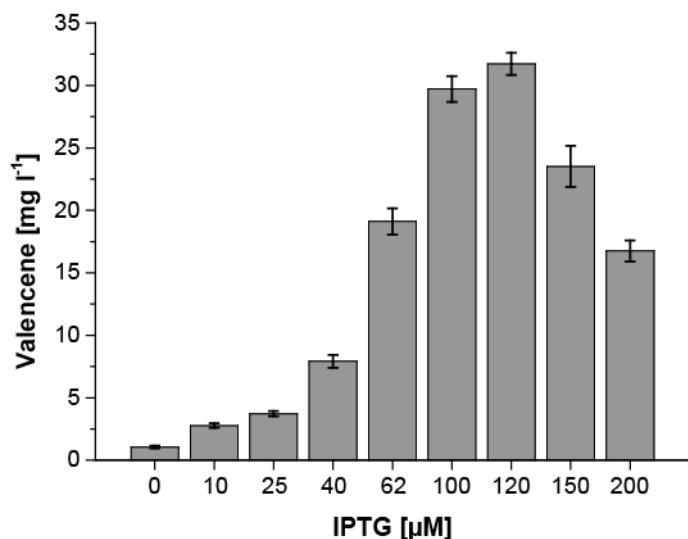


FIG S7 Optimization of IPTG concentrations for *C. glutamicum* VLC6 production cultures induced after 4 h of cultivation. All averaged data originate from at least three independent biological triplicates. Error bars indicate the respective standard deviations.

TAB S1 Summary of titers as well as volumetric and biomass-specific productivities for the conducted (+)-valencene productions in different *C. glutamicum* strains using CGXII minimal medium. Values for volumetric productivity were calculated using the overall cultivation times (28 h for induction after 4 h and 30 h for induction after 6 h).

Strain	Condition	Titer [mg l⁻¹]	Final OD ₆₀₀	Volumetric productivity [µg l⁻¹ h⁻¹]	Biomass-specific productivity [µg g CDW⁻¹ h⁻¹]
VLC3	0.1 mM IPTG, 6h, Flask	7.2±0.6	33.5±2.1	240 ± 20	29 ± 2
VLC4	0.1 mM IPTG, 6h, Flask	10.8±1.1	34.1±1.8	360 ± 37	42 ± 4
VLC5	0.1 mM IPTG, 6h, Flask	10.5±3.5	33.9±2.3	350 ± 117	41 ± 14
VLC6	0.1 mM IPTG, 6h, Flask	27.1±0.6	35.2±1.1	903 ± 20	103 ± 2
VLC6	0.1 mM IPTG, 6h, Flowerplate	29.0±0.1	59.7±3.4	967 ± 3	65 ± 0
VLC6	0.1 mM IPTG, 4h, Flask	14.8±1.1	30.3±1.2	529 ± 39	70 ± 5
VLC6	0.1 mM IPTG, 4h, Flowerplate	28.4±1.7	57.8±2.9	1014 ± 61	70 ± 4
VLC6	0.1 mM cIPTG, full light induction after 4h, Flowerplate	41.0±0.7	60.8±1.9	1464 ± 25	96 ± 2