

Optimizing high-density phototrophic cultivation of *C. reinhardtii* for chloroplast recombinant protein production

Gordon B. Wellman, Amna A. Abdullah, Malak Abdallah and Kyle J. Lauersen

Biological and Environmental Sciences and Engineering Division, King Abdullah University of Science and Technology (KAUST), Thuwal 23955-6900, Kingdom of Saudi Arabia.

INTRODUCTION

Recent advances in transgene design and strain domestication have demonstrated the use of *C. reinhardtii* as a photosynthetic cell chassis for sustainable heterologous production of a range of natural products. A domesticated strain (UVM4) with high nuclear genome transgene expression capability was optimized for the capacity to grow on inorganic phosphite and nitrate as sole sources of P and N, respectively, to enable high-cell density cultivation with reduced risk of contamination^[1]. This strain (UPN) serves as a platform to produce heterologous metabolites in our group^[2]. A further goal in the use of this strain was to add value to spent biomass by leveraging *C. reinhardtii*'s generally regarded as safe (GRAS) status to sustainably produce food-grade recombinant proteins (RPs). A range of proteins of interest to the food industry were selected and coding sequences were optimized for chloroplast expression. A Strep-II tag sequence was added for detection by Western blot and down-stream purification.

The chloroplast of UPN was engineered to encode these RPs by *PsbH* knock-out/restoration transformation^[3] to enable antibiotic-free selection. We present here successful chloroplast genome expression of the ice binding protein *LpIBP* (PDB: 3ULT) from perennial ryegrass *Lolium perenne*.

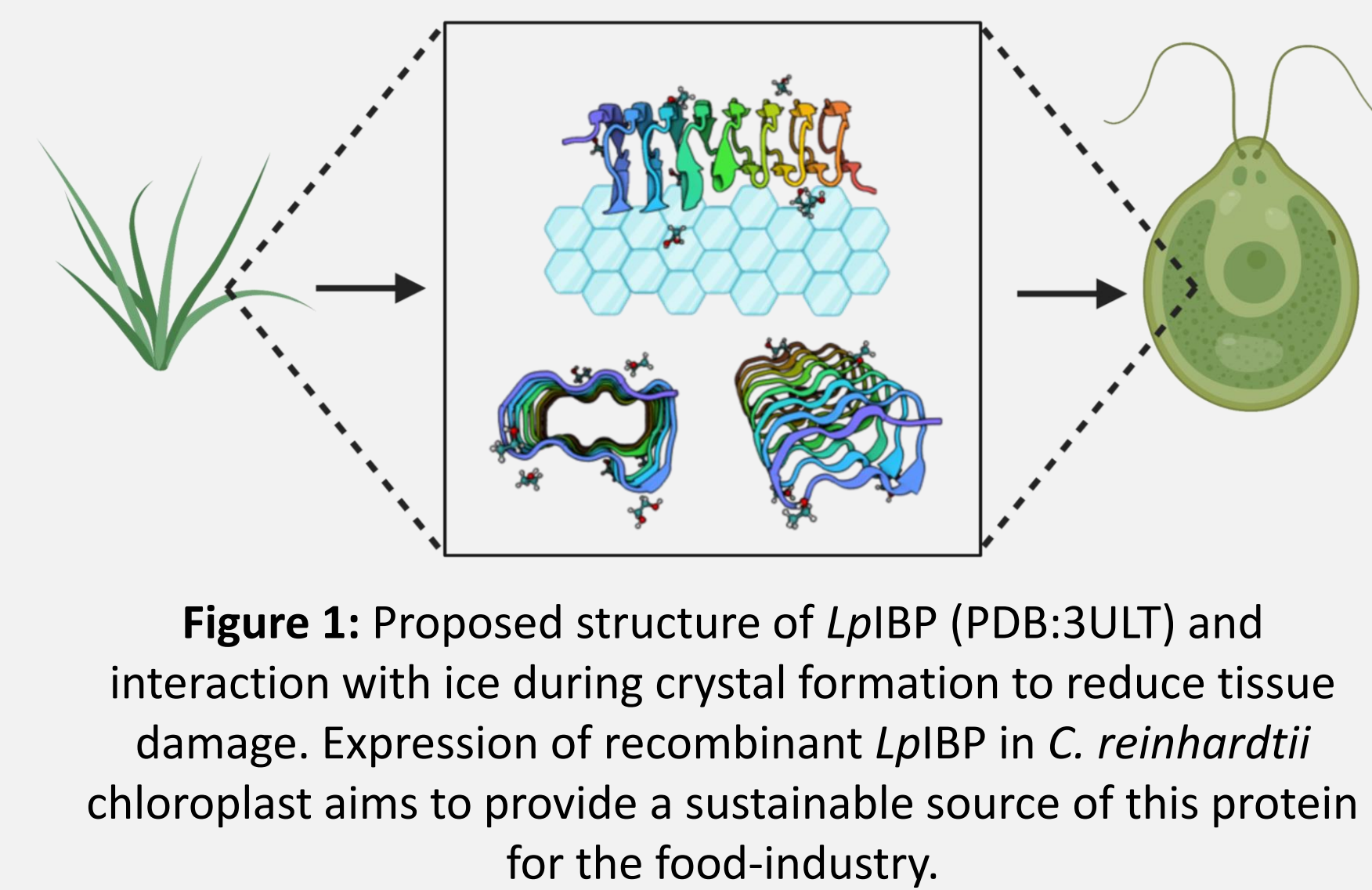


Figure 1: Proposed structure of *LpIBP* (PDB:3ULT) and interaction with ice during crystal formation to reduce tissue damage. Expression of recombinant *LpIBP* in *C. reinhardtii* chloroplast aims to provide a sustainable source of this protein for the food-industry.

DEVELOPMENT OF 'UPN'

The high-density capable strain 'UPN' was generated from cell-wall deficient UVM4 by chloroplast genome transformation with WM88 phosphite NAD⁺ oxidoreductase from *Pseudomonas stutzeri* (*ptxD*)^[4] allowing conversion of inorganic phosphite (PO_3) into bio-available phosphate (PO_4). Subsequently, NO_3 metabolic capacity was restored by co-transformation of NIT1/NIT2^[5]. Modified versions of TAP (tris-acetate-phosphate medium) with equal-molar replacement of phosphate and NH_4Cl with phosphite and NaNO_3 were developed for reduced contamination cultivation.

For high-density cultivation experiments, 6xP medium^[5] was similarly modified by replacing phosphate with equal-molar phosphite solutions. This medium (6xPhi) supports high density growth of UPN in CellDEG photobioreactors.

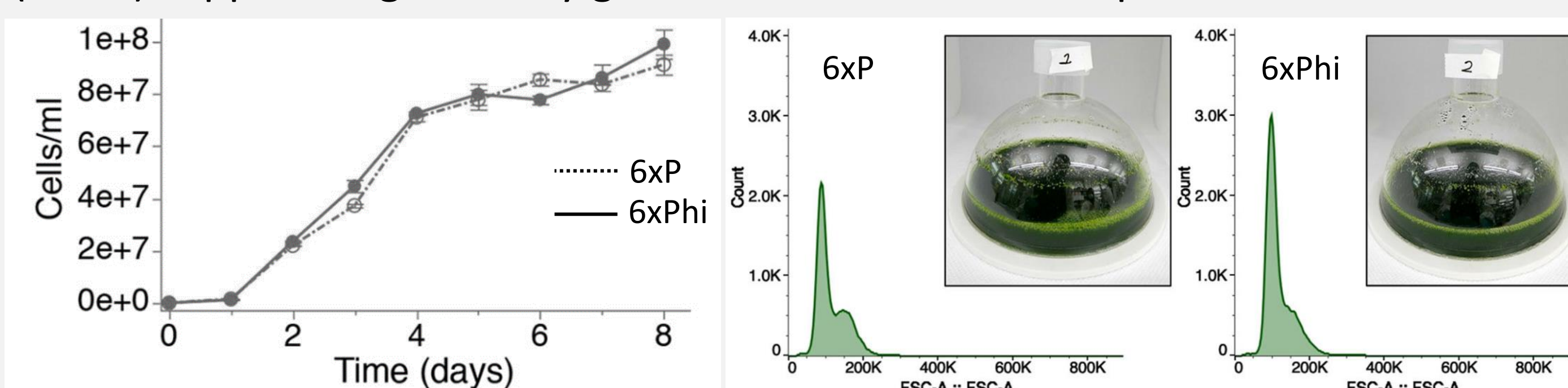


Figure 3: Growth of UPN in CellDEG photobioreactors in 6xP and 6xPhi media. Adapted from^[1].

RESULTS

After recovering photoautotrophic capable transformants and confirming integration of *LpIBP* coding sequence into the chloroplast genome, selected transformants were grown in 6-well plates under different media and CO_2 supply to assess growth compared to empty vector (EV) control (Fig 4).

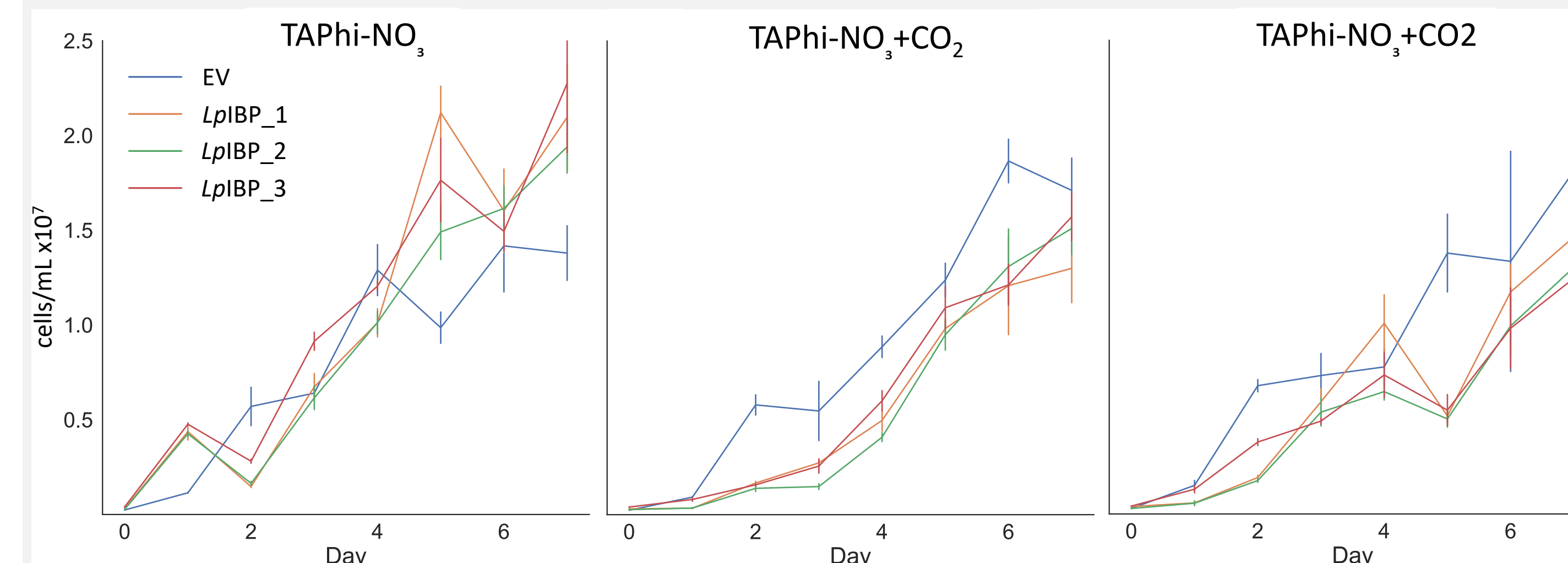


Figure 4: Growth of 3 independent UPN:*LpIBP* transformants and empty vector control grown in TAPhi (left), TAPhi + 5% CO_2 in air (mid), and acetate free media (TPhi) + 5% CO_2 in air (right). Mean cells/mL \pm SEM plotted of 3 biological replicates of each transformant measured (n=3).

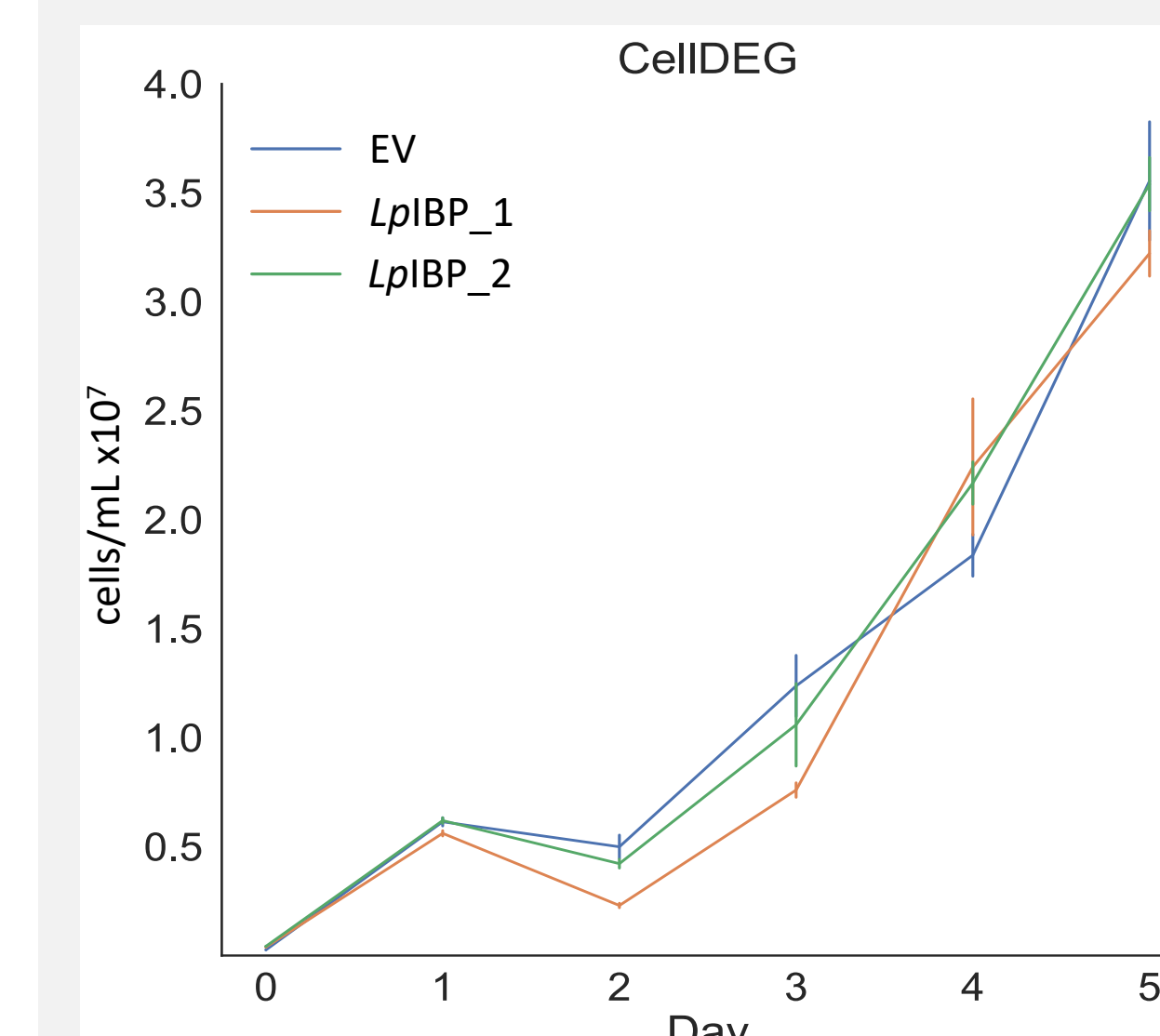


Figure 5: Growth of 2 independent UPN:*LpIBP* transformants and empty vector control grown 6xPhi+5% CO_2 in CellDEG photobioreactors. Mean cells/mL \pm SEM plotted of 2 biological replicates of each transformant.

LpIBP transformants showed similar growth to empty vector control suggesting little impact of *LpIBP* expression on growth in small scale cultures. Two transformants were selected for high-density cultivation in CellDEG photobioreactors with 6xPhi medium. Samples of cultures were taken daily for cell number (Fig 5.) and total soluble protein extraction and yield estimates (Fig 6).

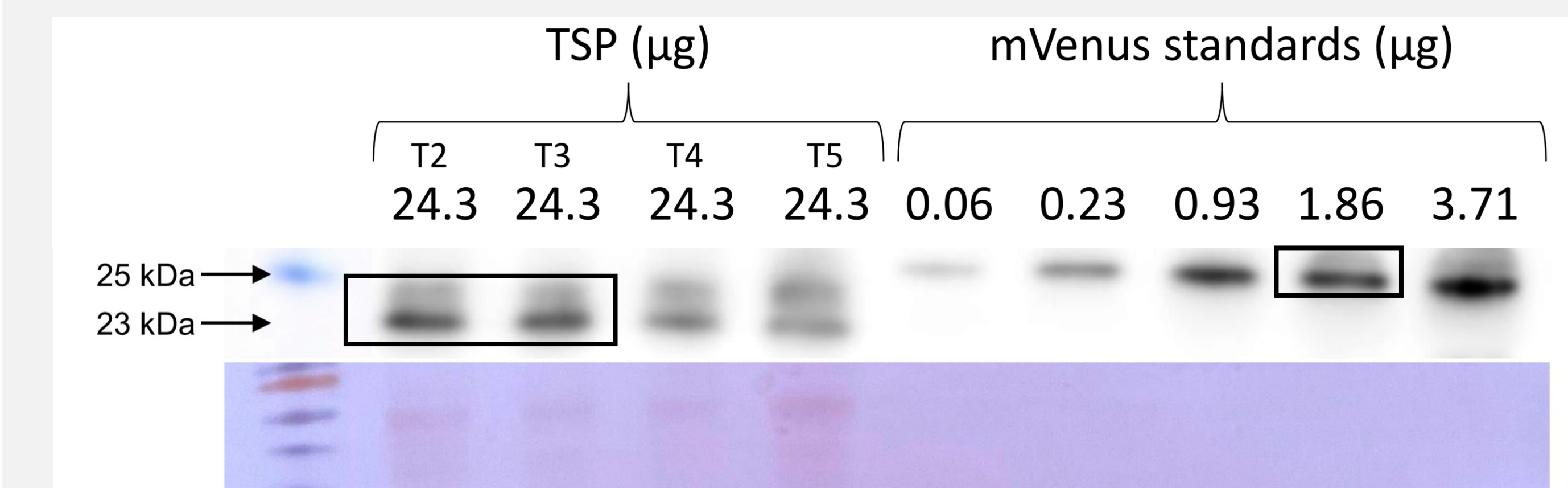


Figure 6: Estimation of *LpIBP* (23 kDa) yield as a % total soluble protein (TSP) extracted from CellDEG grown UPN:*LpIBP* cultures (Days 2-5) – compared to *E. coli* expressed fluorescent protein mVenus (25 kDa) of known concentrations. Anti-streptII-tag Western blot (top), ponceau-stained membrane (bottom). Similar chemiluminescence signals allows estimation of *LpIBP* %TSP yield of ~7.5%.

REFERENCES & CONTACT

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Contact: Gordon.Wellman@KAUST.edu.sa

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METHODS

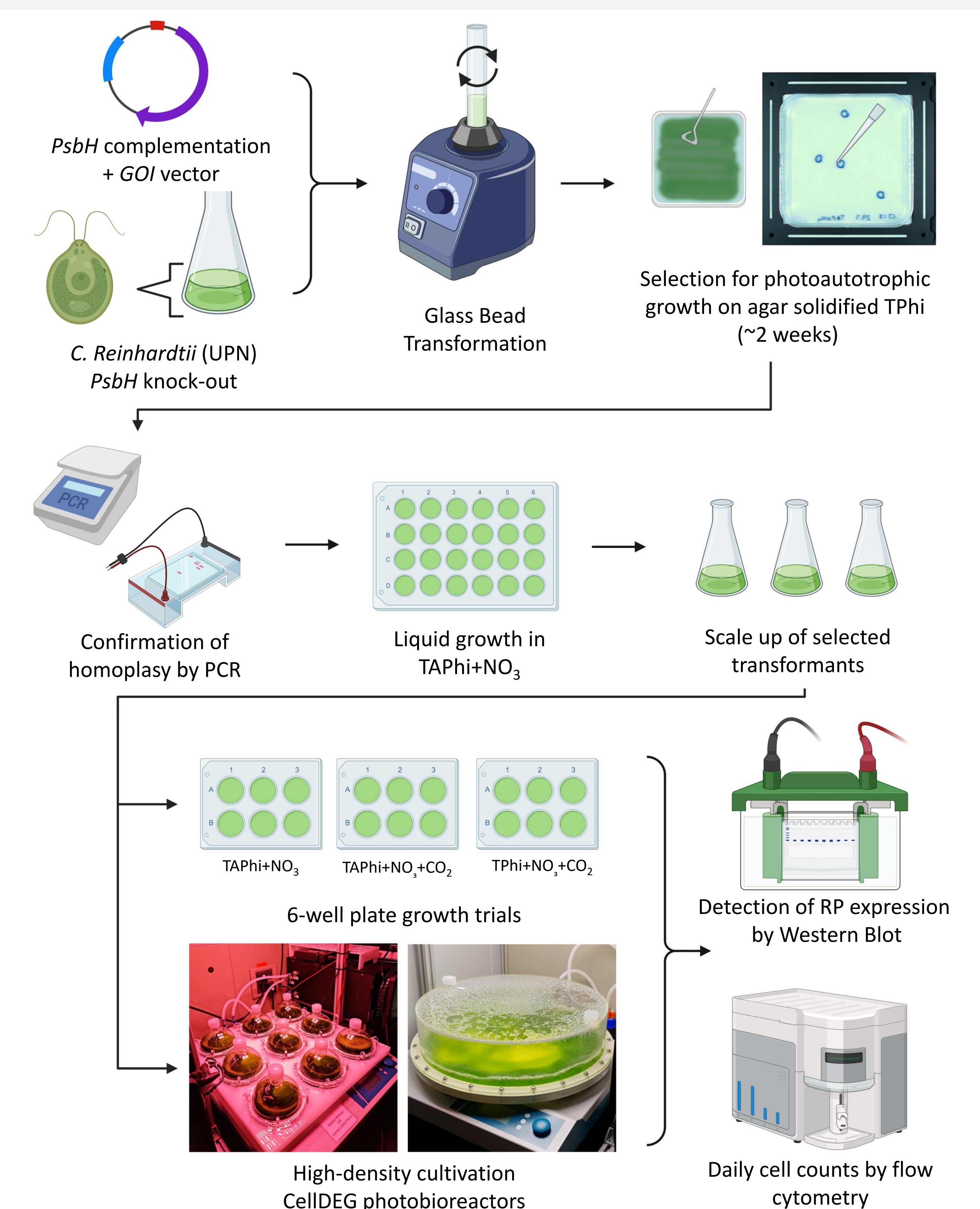


Figure 7: Overview of experimental techniques used for chloroplast genome transformation, photosynthesis restoration selection and growth trials of selected transformants

CONCLUSIONS & FUTURE WORK

The chloroplast of UPN was successfully transformed and expression of recombinant *LpIBP* detected by anti-Strep-II-tag Western blot. *psbH* complementation restores photoautotrophic growth allowing culturing in high-density photobioreactors with similar cell densities to empty vector controls. Initial estimates are that *LpIBP* accumulates to ~7.5 % of TSP.

Further optimization of high-density culture conditions (N:P ratio, light regime) for increased culture density, cell protein content and to minimize excessive nutrients supply.

Column purification of RPs to determine RP yield and feasibility of large-scale production for industry uses.

Screening of transformants with different chloroplast genome expressed RPs to find an economically attractive product compatible with other metabolic engineering projects in the SSB group.