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Supplementary file 1

Designing a new generation of expression toolkits for engineering of green microalgae; robust production of human interleukin-2

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Table S1. The results of the ELISA assay from the total extracted soluble proteins from the transformed and untransformed *C. reinhardtii* cells.

	Blank			Sample 1			Sample 2			Sample 3		
	B-1	B-2	B-3	S1-1	S-1-2	S1-3	S2-1	S2-2	S2-3	S3-1	S3-2	S3-3
OD (nm)	0.079	0.093	0.086	0.113	0.11	0.114	0.155	0.154	0.156	0.198	0.203	0.202
IL-2 Conentration (pg/mL)	0	0	0	5.0526	4.2631	5.3157	16.1052	15.8421	16.3684	27.421	28.7368	28.4736

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C. reinhardtii chloramphenicol sensitivity Control $10 \mu g/mL$ 50 μg/mL 100 μg/mL 120 μg/mL 150 μg/mL C. vulgaris chloramphenicol sensitivity D. salina chloramphenicol sensitivity 50 μg/mL Control Control 10 μg/mL 60 μg/mL 100 μg/mL 130 μg/mL $40 \mu g/mL$

Fig. S1. Evaluating the sensitivity of the *C. reinhardtii*, *C. vulgaris*, and *D. salina* to the chloramphenicol for the selection of the transformants.

>LB TGGCAGGATATATTGTGGTGTAAACAGCAAATTACATATGTCTGCGTGACGGCGGGGA GCTCGCTGAGGCTTGACATGATTGGTGCGTATGTTTGTATGAAGCTACAGGACTGATTT CCAGAAGGCGCCATACGGCCCGCTGGCGGCACCCATCCGGTATAAAAGCCCGCGACCCC >hsp70 GCTATACATAACCACTCAGCTAGCTTAAGAGTCAGTAAACGTGAGTCGACGAGCAAGCC CGGCGGATCAGGCAGCGTGCTTGCAGATTTGACTTGCAACGCCCGCATTGTGTCGACGA AGGCTTTTGGCTCCTCTGTCGCTGTCTCAAGCAGCATCTAACCCTGCGTCGCCGTTTCCAT TTGCAGTCCCATCAAGCTTGCATGCCGGGCGCGCCAGAAGGAGCGCAGCCAAACCAGG ATGATGTTTGATGGGGTATTTGAGCACTTGCAACCCTTATCCGGAAGCCCCCTGGCCCAC AAAGGCTAGGCGCCAATGCAAGCAGTTCGCATGCAGCCCCTGGAGCGGTGCCCTCCTGA >rbcS2 TAAACCGGCCAGGGGCCTATGTTCTTTACTTTTTTACAAGAGAAGTCACTCAACATCTT >5-UTR

Fig. S2. The sequence of the promoter cassette composed of hsp70 and rbcS2 regulatory elements, Int-1, and 5-UTR of the rbcS2 gene. The elements were determined in different colors.

>Sc >Kozak **ATGGCCGCCGAGAAGAAGATCACCGGCTACACCACCGTGGACATCTCCCAGTGGCACCG** CAAGGAGCACTTCGAGGCGTTCCAGTCCGTGGCCCAGTGCACCTACAACCAGACCGTGC AGCTGGACATCACCGCCTTCTTGAAGACCGTGAAGAACAAGCACAAGTTCTACCCC GCCTTCATCCACATCCTGGCCCGCCTGATGAACGCCCACCCCGAGTTCCGCATGGCCATG AAGGACGGCGAGCTGGTGATCTGGGACAGCGTGCACCCCTGCTACACCGTGTTCCACGA GCAGACCGAGACCTTCTCCTCGCTGTGGAGCGAGTACCACGACGACTTCCGCCAGTTCCT GCACATCTACTCGCAGGACGTGGCCTGCTACGGCGAGAACCTGGCCTACTTCCCCAAGG GCTTCATCGAGAACATGTTCTTCGTGTCGGCCAACCCCTGGGTGAGCTTCACCAGCTTCG ACTTGAACGTGGCCAACATGGACAACTTCTTCGCCCCCGTGTTCACCATGGGCAAGTACT ACACCCAGGGCGACAAGGTGCTGATGCCCCTGGCCATCCAGGTGCACCATGCCGTGTGC GACGGCTTCCACGTGGGCAGGATGCTGAACGAGTTGCAGCAGTACTGCGACGAGTGGC >CmR AGGGCGCCGCCCGCCCCCCACCTCCAGCCTGCAAGCAGGCCGGCGACGT GGACGAGAACCCCGGCCCCGAGGGCAGGGGCAGCCTGCTGACCTGCGGCGACGTGGA >Hybrid 2A **GGAGAACCCCGGCCCC**

Fig. S3. The detailed sequences of the Kozak, *CmR*, and 2A peptide. The elements were determined in different colors. Sc: Start codon, *CmR*: Coding sequence of the chloramphenical acetyltransferase gene.

>XhoI CTCGAGTACAGGATGCAGCTGCTGTCCTGCATCGCCCTGAGCCTGGCCCTGGTGACCAA CAGCGCCCCACCTCGAGCTCCACCAAGAAGACCCAGCTGCAGCTGGAGCACTTGCTGC >Ex-1TGGACTTGCAGATGATCTTGAACGCCATCAACGTGAGTCGACGAGCAAGCCCGGCGGA TCAGGCAGCGTGCTTGCAGATTTGACTTGCAACGCCCGCATTGTCGACGACGACGCTTTT GGCTCCTCTGTCGCTGTCTCAAGCAGCATCTAACCCTGCGTCGCCGTTTCCATTTGCAGA ACTACAAGAACCCCAAGCTGACCAGGATGCTGACCTTCAAGTTCTACATGCCCAAGGTG AGTCGACGAGCAAGCCCGGCGGATCAGGCAGCGTGCTTGCAGATTTGACTTGCAACGC CCGCATTGTCGACGAAGGCTTTTGGCTCCTCTGTCGCTGTCTCAAGCAGCATCTAACC CTGCGTCGCCGTTTCCATTTGCAGGCCACCGAGCTGAAGCACCTGCAGTGCCTGGAGGA GGAGCTGAAGCCCCTGGAGGAGGTGCTGAACTTGGCCCAGAGCAAGAACTTCCACTTG AGGCCCAGGGACTTGATCAGCAACATCAACGTGATCGTGCTGGAGCTGAAGGGCTCCG AGACCACCTTCATGTGCGAGTACGCCGACGAGACCGCCACCATCGTGGAGTTCCTGAAC >Ex-4>Stc >BamHI
TAAGGATCCGTAAGTCTGGCGAGAGCCCGACGGGTCCACTGTGGCACTGGGTTAGCTTT TGGCACACGGGTCCACTGTGGCACTGGTTAGCTTGGCACCGGGACAGCGCCTATCTCAC CGCGGGGAACTGACGCATACCCCTGCTCGTGCTTCAGCACGGAAAAGCAAGGGGCCCA GCGCCCGCAG

Fig. S4. The detailed sequences of the microalgae expression cassette. The elements were determined in different colors. Ex-1-4: Exons 1-4 of the hIL-2, His: Histidine tag, Stc: Stop codon.

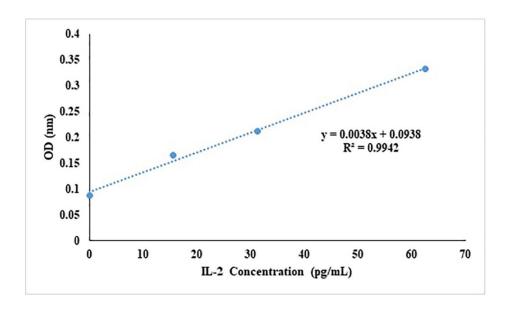


Fig. S5. The standard curve of the hIL-2 based on the four elution reactions.