

Supplementary Materials for
Screening for Type II L-Asparaginases: Lessons from the Genus *Halomonas*

Zeinab Sharafi, Mahmood Barati, Mohammad Reza Khoshayand and Sina
Adrangi*

*To whom correspondence should be addressed. E-mail: s.adrangi@sbmu.ac.ir

Volume 16, Issue 4 (Autumn 2017)

This PDF file includes:

Figures S1 to S6

Tables S1 and S2

huangheensis	1	CGAGGATCATGCCAAAAGCCATGGGGTCCACCAGCGGATCACCAGGTTCTTGCGAATGA
KO116	1	CCAGGATCATCCCGAACGCCATTGGATCGACGAGCGGTGTGCCGGGTTCCGCGCGAATCA
R57-5	1	-----ATCATTCCAAATGCCATCGGGTCGACGAGTGGCGTGCCTGGTTCGCGCGCAATCA
huangheensis	61	TGGTGCCCTGCCCGCCTTCACTTCGACAATGCCGATGGCC <u>TGCAGCATCTTGATGGCTT</u>
KO116	61	TGGTGCCTTGGCCTCGTTTGACTTCGACAATCCCTATCGCC <u>TGCAGCATCTTGATGGCTT</u>
R57-5	56	TGGTGCCTTGGCCGCTTGGACTTCCACAATGCCAATGGCT <u>TGCAGCATCTTGATGGCTT</u>
		ANSF →
huangheensis	121	CTCGCACACTTGACTTTCGGATTCCCAGGCTTTGGGTGAGCTCTGTTTCCGAGGCAGGT
KO116	121	CGCGCACACTGGATTACCAATGCCAGGGTGTGAGTGAAGCTCCGTTTCTGAGGGTAGGT
R57-5	116	CACGCACGCTCGATTTGCCAATCCCCAACTGTGGGTCAATTGAGTTCTGAGGTTAGAT
huangheensis	181	AGTCACCTGGCTGAAGCTCGCCGCTGATCAGCGCGCTCTTGATACGCTCGAGCAGTTGAA
KO116	181	AATCCCCTGGCTTAAGATCGCCTCGTATTAGGGCGCTTTAATACGCTCTAGCAGCTGCA
R57-5	176	AATCCCCTGGTTTGGAGTCTCCACGTATCAAGGCGCTTTAATGCGCTCTAGCAGTTGCA
huangheensis	241	GGGCAACGGAACACTACGGTCAATCGAACTCCCGAATGACTCCATGGATGACGTCCTTGGCT
KO116	241	GCGCAACTGAACCTGCGATCCATCGAATCTCCAAAAGAGGTTGACTGGTTCGAGTTAATCT
R57-5	236	GCGCAACGAGCTGCGCTCCATCGAATCCCCAAAAGACGTTGTAGTGGTTCGAGGTAATCT
huangheensis	301	GGCGTTTTACTGGCATGGTCCGGTCTTGCTCGACTGCCGTCAG-----ACTCT
KO116	301	CGGGCTGGCGTTTAAAACCTCATGTGGGCTCGCGTGGCTCATCGGTGTTTTCTGCATTTT
R57-5	296	CGGGCTGGCGTTTAAAACCTCATGGGGGCTCGCGTGGCTCATCGGTGTTTTTTGCATCTT
huangheensis	350	GGAAAGCCGCC-GGAAATCAGGCTTCCACTCTATCCATGATAACGATCTCCAGCCGACC
KO116	361	ATCAATGTCCTAAGCCTAGCGGCTTAAG-CCATTCGATTGCATATA-TTTAAAAGAAATA
R57-5	356	ATCAATGCCATAAGCCTAGCGGCTTAAG-ACATCTGATGGCATATA-TTTAAAAGTAATT
huangheensis	409	TTGTTGTGCTGTTTCAACTGGGACGC-----CTGGCGCTTCTGCAAAATCGAGCCTAGCA
KO116	419	TAGTTATGTTTCATTGGTTGGTGAAGTCTGGCTTGCTAACAGGTTGTTCTTTAGTCTAGTC
R57-5	414	TGTTTATTTGTTTCGGTTTATATGCCCGACTTGCTAATGGTTGTGCTTTAGTCTAGTC
huangheensis	464	TGTGACAGGACATCAGACATCAGATGTCTGATTAC-----AAGAATAAGGACCATG
KO116	479	TGAAATCAGACATCTGATGTCTGTGTTTGTGTT-CAGAAAGATATAAAACACGAGAACCTG
R57-5	474	TGAAATCAGACATCTGATGTCTGGTGTGTTGTATTAGCTAGACATAAAACAAGAGAATCTG
huangheensis	515	AATGTCAGACTCCAGAATCATTGTTCATGACCACCGGTGGAACCATAGCCAGCGAACTAGA
KO116	538	AATGACCAAAAAGCACATCGTGGTATTGACCACGGGCGGCACGATTGCCAGCAAGCCAG
R57-5	534	CATGAGCCAAAAGCACATCGTGGTGTGACCACGGGCGGCACATTGCCAGCAAGCCAG
huangheensis	575	TATTTCTGGCCGAGTCTTCCGGCGCGCTGTCCGGCGCCGACCTGCTGTCTCGTGCCAA
KO116	598	TGACTCAGGGCGAAGCCAGTCAGGGGCTTTAAGCGGCGAGCAGTTGCTGGATCAGGTGGC
R57-5	594	TGATTCAGGGCGTAGCCAGTCAGGGGCATTAAGCGGGGAGCAGCTACTTGTATCGGGTGGC
huangheensis	635	CGTCCCTGAACGTTTTTCGGTCCAGTGTGGAGGTACATTCCGGTACTGCAGAAACCCAGCAA
KO116	658	ACTGCCGAAGGGCGTCGACGTTACGTTAGAAGTGATATCGATCCTGCAGAAACCCAGCAA
R57-5	654	GCTGCCCCAGGGCGTCGACGTTACGTTAGAGGTGATGTCGATCTGCAGAAACCCAGTAA
		← ANSR
huangheensis	695	CGCCATCACTCTCGATGACTTGTGACAGTGCGC-
KO116	718	TGCGGTCAACCTCGCCGATCTTGTGAACTCCAT-
R57-5	714	TGCCGTCTCCCTCGCCGATCTTGTGAGCTCCATC

Figure S1. Multiple sequence alignment of the L-asparaginase gene promoter region of entries NZ_CP013106.1 (*H. Huangheensis*), NZ_CP011052.1 (*Halomonas*. sp. KO116) and LN813019.1 (*Halomonas*. sp. R57-5). The positions of the primers are underlined and conserved bases in these regions are highlighted.

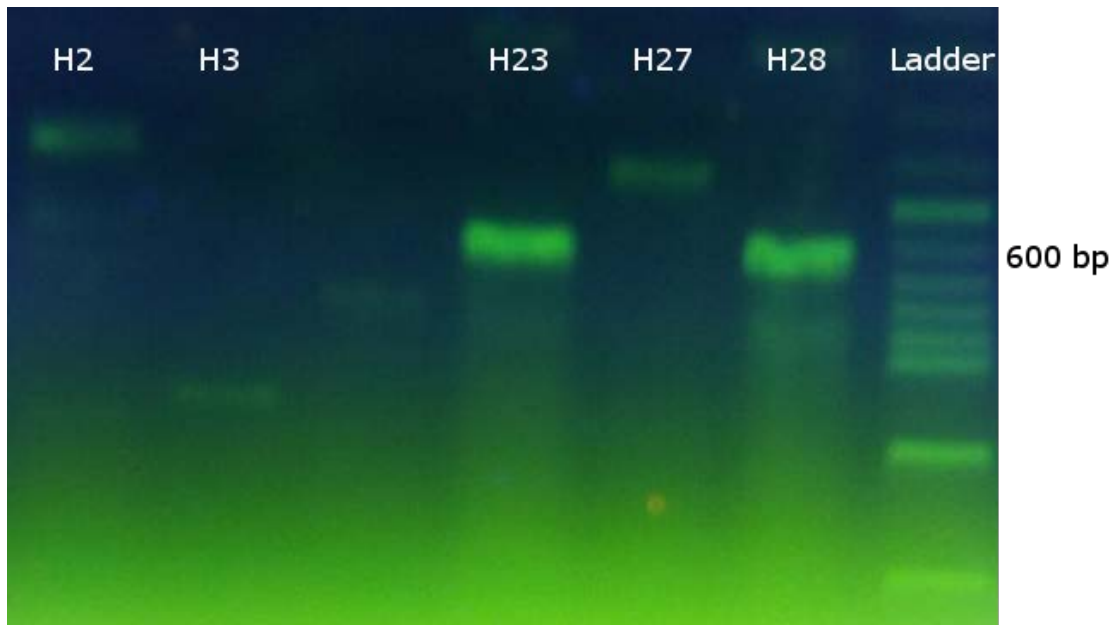


Figure S2. Amplification of genomic DNA from isolates H2, H3, H23, H27 and H28 using primers ANSF and ANSR. A band of the expected size (600 bp) can be seen with H23 and H28.

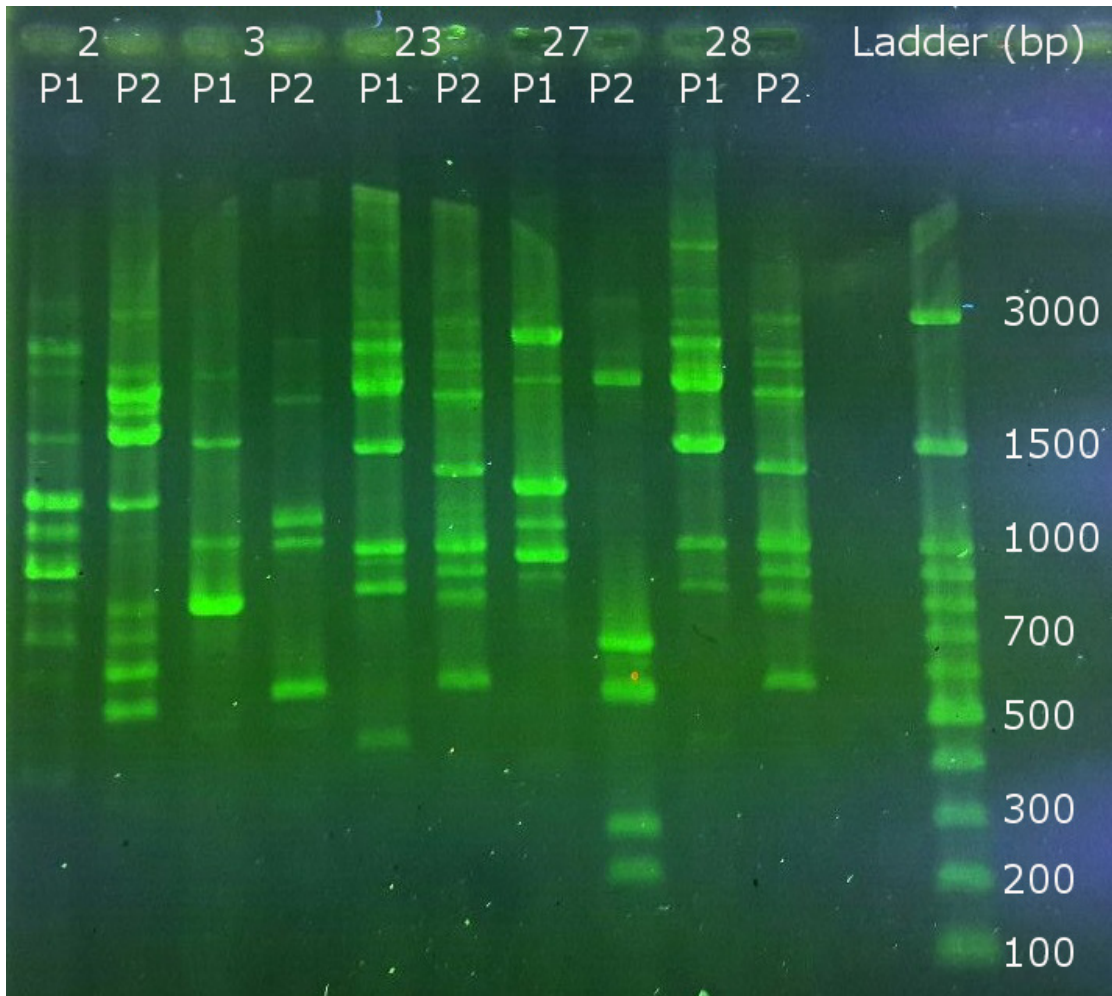


Figure S3. RAPD-PCR profiles of isolates H2, H3, H23, H27 and H28. Genomic DNA samples from each isolate were amplified using primers 5'-CCGCAGCCAA-3' (P1) and 5'-GATGACCGCC-3' (P2).

Length: 1357
 Identity: 1271/1357 (93.7%)
 Similarity: 1271/1357 (93.7%)
 Gaps: 35/1357 (2.6%)
 Score: 5981.5

```

H23 17 GGTAGCTTGCTACCCGCTGACGAGCGGGACGGGTGAGTAATGCATAGG
H28 5 GGAAGCTTGCTTCCAGGCGTCGAGCGGGACGGGTGAGTAATGCATAGG

H23 67 AATCTGCCCGGTAGTGGGGGATAACCTGGGGAAACCCAGGCTAATACCGC
H28 55 AATCTGCCCGGTAGTGGGGGATAACCTGGGGAAACTCAGGCTAATACCGC

H23 117 ATACGTCTACGGGAGAAAAG--GGGG--CTCCGG--CTCCCGTATTGGA
H28 105 ATACGTCTACGGGAGAAAAGCAGGGGATCTTCGGACCTTCGCTATCGGA

H23 161 TGAGCCTATGTCGGATTAGCTAGTTGGTAAGGTAATGGCTTACCAAGGCA
H28 155 TGAGCCTATGTCGGATTAGCTAGTTGGTGAGGTAACGGCTACCAAGGCG

H23 211 ACGATCCGTAGCTGGTCTGAGAGGATGATCAGCCACATCGGGACTGAGAC
H28 205 ACGATCCGTAGCTGGTCTGAGAGGATGATCAGCCACACTGGGACTGAGAC

H23 261 ACGGCCCAACTCCTACGGGAGGCAGCAGTGGGGAATATTGGACAATGGG
H28 255 ACGGCCCAACTCCTACGGGAGGCAGCAGTGGGGAATATTGGACAATGGG

H23 311 GGGAAACCTGATCCAGCCATGCCCGGTGTGTGAAGAAGGCCCTCGGGTTG
H28 305 GGGAAACCTGATCCAGCCATGCCCGGTGTGTGAAGAAGGCCCTCGGGTTG

H23 361 TAAAGCACTTTCAGCGAGGAAGAAGCCCT-AGTGGTTAATACCATTAGG
H28 355 TAAAGCACTTTCAGCGAGGAAGAAGCCCTGAG-GGCTAATACCCTTCAGG

H23 410 AAAGACATCACTCGCAGAAGAAGCACC GGCTAACTCCGTGCCAGCAGCCG
H28 404 AAGGACATCACTCGCAGAAGAAGCACC GGCTAACTCCGTGCCAGCAGCCG

H23 460 CGGTAATACGGAGGGTGCAAGCGTTAATCGGAATTACTGGCGTAAAGCG
H28 454 CGGTAATACGGAGGGTGCGAGCGTTAATCGGAATTACTGGCGTAAAGCG

H23 510 CGCGTAGGTGGCTTGATAAGCCGGTTGTGAAAGCCCGGGCTCAACCTGG
H28 504 CGCGTAGGTGGCTTGATAAGCCGGTTGTGAAAGCCCGGGCTCAACCTGG

H23 560 GAACGGCATCCGGAAGTGTGAGGCTAGAGTGCAGGAGAGGAAGGTAGAAT
H28 554 GAACGGCATCCGGAAGTGTGAGGCTAGAGTGCAGGAGAGGAAGGTAGAAT

H23 610 TCCCGGTGTAGCGGTGAAATGCGTAGAGATCGGGAGGAATACCAGTGGCG
H28 604 TCCCGGTGTAGCGGTGAAATGCGTAGAGATCGGGAGGAATACCAGTGGCG

H23 660 AAGGCGCCTTCTGACTGACACTGACACTGAGGTGCGAAAGCGTGGGTA
H28 654 AAGGCGCCTTCTGACTGACACTGACACTGAGGTGCGAAAGCGTGGGTA

H23 710 GCAAACAGGATTAGATACCCCTGGTAGTCCACGCCGTAACAGATGTCGACC
H28 704 GCAAACAGGATTAGATACCCCTGGTAGTCCACGCCGTAACAGATGTCGACC

H23 760 AGCCGTTGGGTGCCTAGAGCACTTTGTGGCGAAGTTAACCGGATAAGTCCG
H28 754 AGCCGTTGGGTGCCTAGAGCACTTTGTGGCGCAGTTAACCGGATAAGTCCG

H23 810 ACCGCCTGGGGAGTACGGCCGCAAGGTTAAAACCTCAAATGAATTGACGGG
H28 804 ACCGCCTGGGGAGTACGGCCGCAAGGTTAAAACCTCAAATGAATTGACGGG

H23 860 GGCCCGCACAAGCGGTGGAGCATGTGGTTTAATTCGATGCAACCGGAAGA
H28 854 GGCCCGCACAAGCGGTGGAGCATGTGGTTTAATTCGATGCAACCGGAAGA

H23 910 ACCTTACCTACCTTGACATC-----TACAGAAGCCGGAAGAGAT
H28 904 ACCTTACCTACCTTGACATC-----TACAGAAGCCGGAAGAGAT

H23 950 TCTGGTGTGCCTTCGGGAAGTGTGTA-AGACAGGTGTCGATGGCTGTCGTC
H28 948 T-----GGTGCCTTCGGGAAGTGTGTA-AGACAGGTGTCGATGGCTGTCGTC

H23 999 AGCTCGTGTGTGAAATGTTGGGTTAAGTCCCGTAAACGAGCGCAACCCCTT
H28 993 AGCTCGTGTGTGAAATGTTGGGTTAAGTCCCGTAAACGAGCGCAACCCCTT

H23 1049 GTCCTTATTTGCCAGCGAGTAATGTCGGGAAGTCTAAGGAGACTGCGCGT
H28 1043 GTCCTTATTTGCCAGCGA-TTCGGTTCGGGAAGTCTAAGGAGACTGCGCGT

H23 1099 GACAAACCGGAGGAAGTGGGGACGACGTCAGTCAATCATGTCGCCCTTACG
H28 1092 GACAAACCGGAGGAAGTGGGGACGACGTCAGTCAATCATGTCGCCCTTACG

H23 1149 GGTAGGGCTACACACGTGCTACAATGGCCGGTACAAGGGCGGCGA-GCT
H28 1142 GGTAGGGCTACACACGTGCTACAATGGTTCGGTACAAGGGTTCGAATGC-

H23 1198 CGCGAGAGT-CAGCGAATCCCTTAAAGCCGGTCTCAGTCCGGATCGGAGT
H28 1191 CGCGAG-GTGGAGCTAATCCCATAAAGCCGGTCTCAGTCCGGATCGGAGT

H23 1247 CTGCAACTCGACTCCGTGAAGTCGGAATCGCTAGTAATCGTGAATCAGAA
H28 1240 CTGCAACTCGACTCCGTGAAGTCGGAATCGCTAGTAATCGTGAATCAGAA

H23 1297 TGTACGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTACACCA
H28 1290 TGTACGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTACACCA

H23 1347 TGGGAGT
H28 1340 TGGGAGT

```

Figure S4. Pairwise alignment of 16S rDNA sequences of isolates H23 and H28. The two sequences were aligned using EMBOSS Water (<http://www.ebi.ac.uk/Tools/emboss/>) with gap opening and extension penalties of 10 and 0.5 respectively and the DNAbfull scoring matrix.

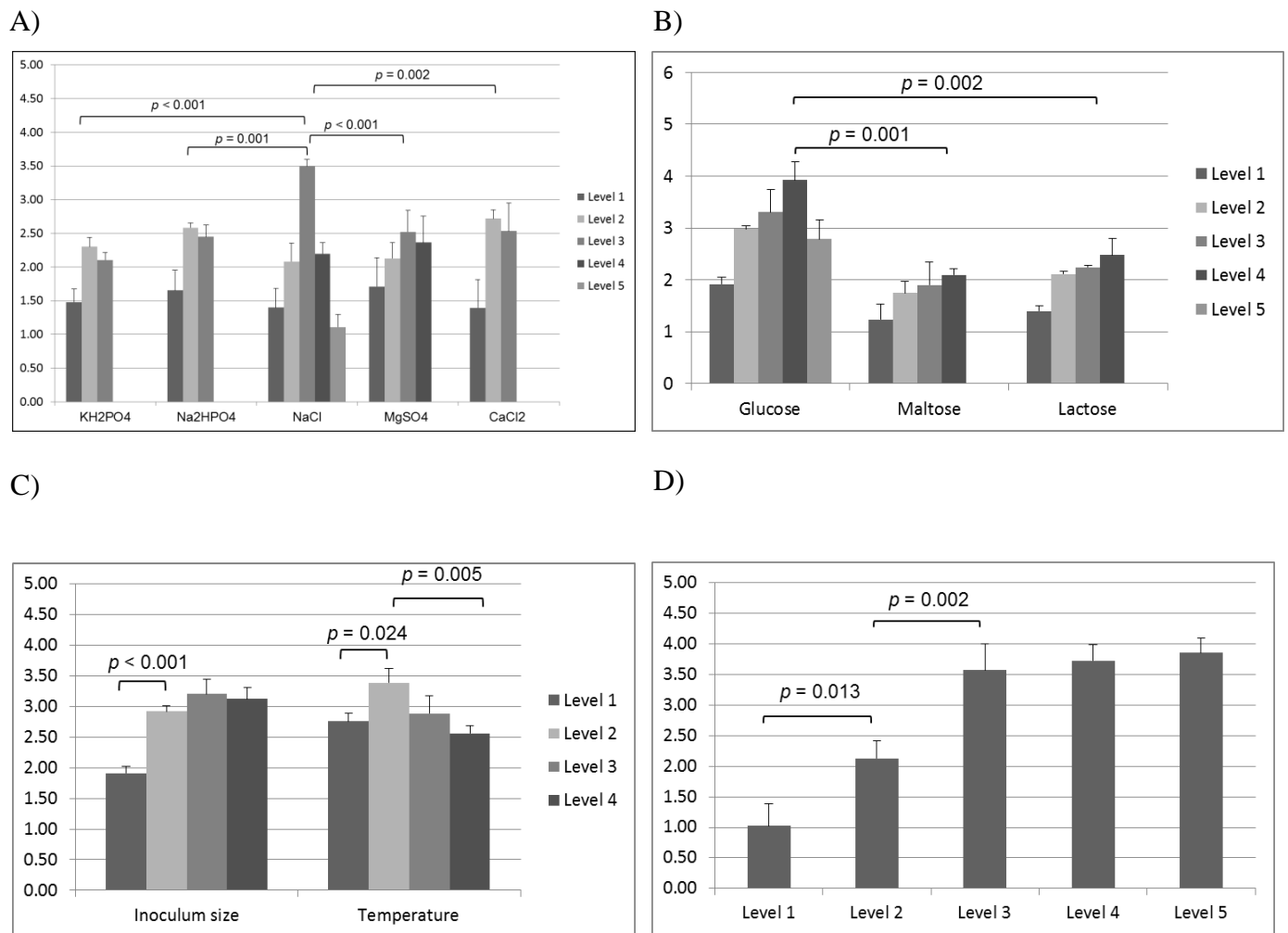


Figure S5. Effect of different salts (A), carbohydrates (B), temperatures and inoculum sizes (C) and L-asparagine concentrations (D) on L-asparaginase production by isolate H28. Vertical axes show enzyme activity in U/mL. In each case the effect was evaluated at least at 3 levels as indicated in Table S1. All p -values were calculated using ANOVA and Tukey's HSD test. (A) The concentration of each salt was increased until either no significant increase or a significant decrease in enzyme activity was observed (p -values for this stage are not shown). Subsequently, the salt resulting in the highest enzyme activity was chosen for further study (shown p -values refer to this stage). (B) The concentration of each carbohydrate was increased until either no significant increase or a significant decrease in enzyme activity was observed (p -values for this stage are not shown). Subsequently, the carbohydrate resulting in the highest enzyme activity

was chosen for further study (shown *p*-values refer to this stage). (C) Inoculum size (%) and temperature were increased until either no significant increase or a significant decrease in enzyme activity was observed (*p*-values for significant differences are shown) and the levels resulting in highest enzyme activity were used throughout the rest of the study. (D) L-asparagine concentration was increased until no significant increase in enzyme activity was observed (*p*-values for significant differences are shown) and results were used to choose the range of L-asparagine concentration for the next step of optimization.

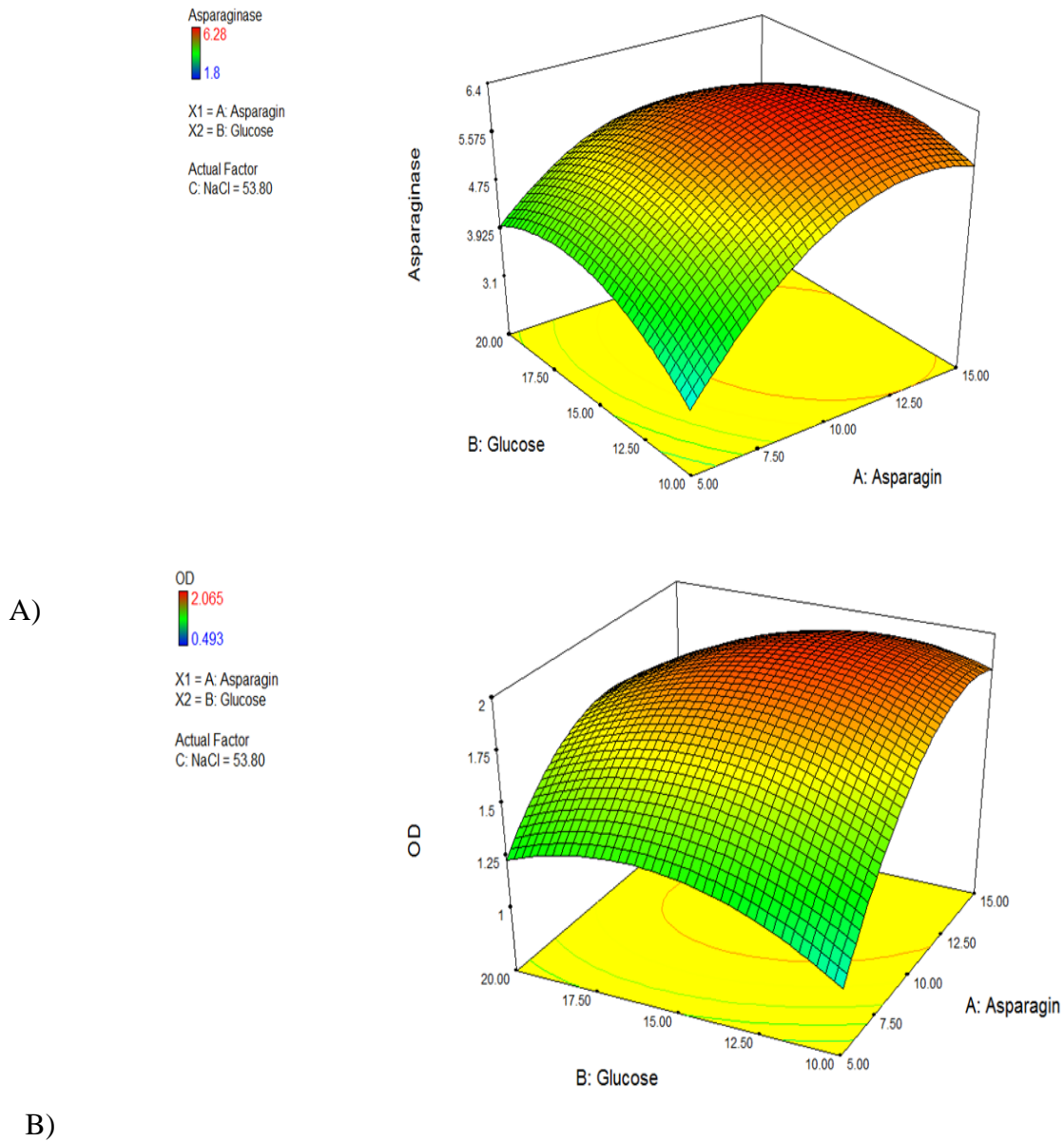


Figure S6. Representative response surface plots showing the effect of L-asparagine and glucose concentrations on (A) enzyme activity and (B) cell density at an optimal NaCl concentration of 54 g/L. Details of the design are provided in Table S2.

Table S1. Different levels of factors evaluated in the first step of medium optimization.

Factor	Unit	Level				
		1	2	3	4	5
KH ₂ PO ₄	g /L	2.5	5	7.5	-	-
Na ₂ HPO ₄	g /L	2.5	5	7.5	-	-
NaCl	g /L	5	20	50	100	150
MgSO ₄ .7H ₂ O (1M)	mL	0.2	0.4	0.6	0.8	-
CaCl ₂ (0.1M)	mL	0.1	0.2	0.4	-	-
Glucose	g /L	2.5	5	10	15	20
Maltose	g /L	2.5	5	10	15	--
Lactose	g /L	2.5	5	10	15	--
L-asparagine	g /L	2.5	5	10	15	20
Inoculum Size	%	1	5	10	15	
Temperature	°C	27	30	33	36	-

Table S2. Central composite design and corresponding results for enzyme activity and cell density.

Run	Variable levels (g/L)			Activity (U/mL)*		Cell density (OD ₆₀₀)**	
	A: L-asparagine	B: Glucose	C: NaCl	Predicted	Actual	Predicted	Actual
1	5	10	25	2.939320	3.0	0.924556	0.97
2	15	10	25	5.260343	5.1	1.671235	1.56
3	5	20	25	3.578076	3.6	1.154658	1.14
4	15	20	25	4.169099	4.3	1.307837	1.40
5	5	10	75	2.963051	2.9	0.894688	0.89
6	15	10	75	5.434074	5.5	1.783868	1.84
7	5	20	75	3.671808	3.9	1.132290	1.28
8	15	20	75	4.412830	4.4	1.427970	1.43
9	1.591036	15	50	1.920546	1.8	0.575276	0.49
10	18.40896	15	50	4.495408	4.5	1.451792	1.47
11	10	6.591013	50	4.203793	4.3	1.352425	1.39
12	10	23.40896	50	3.882161	3.7	1.246643	1.15
13	10	15	7.955179	5.140523	5.1	1.621082	1.66
14	10	15	92.04482	5.365431	5.3	1.696986	1.60
15	10	15	50	6.053923	6.2	1.932282	1.83
16	10	15	50	6.053923	5.8	1.932282	1.91
17	10	15	50	6.053923	5.7	1.932282	1.89
18	10	15	50	6.053923	6.2	1.932282	1.93
19	10	15	50	6.053923	6.2	1.932282	1.98
20	10	15	50	6.053923	6.3	1.932282	2.06

* For enzyme activity the following model was obtained:

$$\text{Enzyme activity} = 6.05 + 0.77A - 0.43AB - 1.01A^2 - 0.71B^2 - 0.28C^2.$$

** For cell density the following model was obtained:

$$\text{OD} = 1.93 + 0.26A - 0.15AB - 0.32A^2 - 0.22B^2 - 0.097C^2.$$