

A Review of Enhancement of Biohydrogen Productions by Chemical Addition Using a Supervised Machine Learning Method

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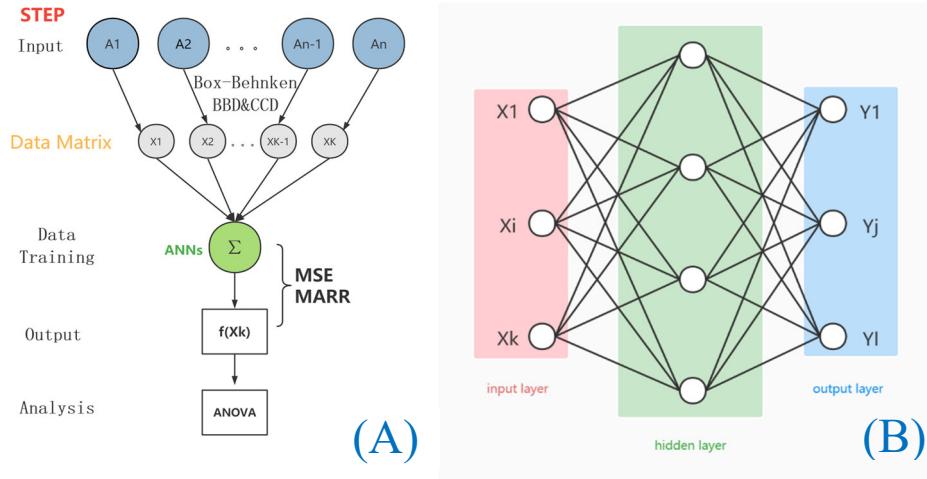


Figure S1. Schematic diagram of methodology: (A) The procedures flowchart, (B) ANNs construction: feed forward three layers networks.

Table S1. Ions comparisons upon BioH₂ generation, - refers to all data missing as, for convenience of calculation, the missing value was replaced by the averaged value during artificial neuron network learning process.

Mg ²⁺ /mmol/L	Cu ²⁺ /mmol/L	Na ⁺ /mmol/L	NH ₄ ⁺ /mmol/L	K ⁺ /mmol/L	HY/mmol/g	HER/mmol/L.h ⁻¹	Reference
1.05	0.002	10.21	17.57	1.38	2.9	5.4	[45]
0.99	-	47	-	1.1	3	13.07	[46]
1.23	-	-	9.09	-	20	0.4	[47]
67.9	-	49.9	420.8	122.5	13.3	2	[48]
5.2	-	32.5	-	25.1	0.9	2.4	[49]
1.754	-	9.58	1.77	2.29	15.9	10.1	[50]
0.35	-	0.939	0.39	0.46	12.9	5.69	[51]
1.05	0.002	10.21	17.57	1.38	5	13.07	[46]
1.23	-	26.37	9.17	-	24.2	0.8	[47]
1.02	-	14.6	3.17	-	1.92	2.5	[52]
0.62	-	37.27	-	-	16.75	102.5	[25]
2.15	-	-	2	-	7.85	62.4	[24]
0.4	-	12.7	2.5	0.7	1.9	49.4	[53]
0.11	0.06	0.1	6.6	0.4	8.4	0.6	[42]
3.596	-	26.37	45.05	-	3.875	1.92	[54]
3.596	-	-	45.05	-	9.2	3.1	[54]
1.04	-	14.6	3.17	-	1.92	2.5	[52]
3.596	-	26.37	10	2	10.1	0.23	[55]
2.25	-	100.02	49.94	49.97	11.66	3.2	[30]
0.348	0.097	5.23	3.74	0.735	1.58	0.066	[56]
0.035	-	0.01	42.2	0.38	6.7	0.23	[57]
0.985	-	75.47	9.49	2.87	10.4	6	[58]
1.5	0.01	-	9.7	3	14.1	11.5	[59]
1.02	0.003	43	-	1.5	2.4	10.3	[46]
0.2	0.02	3.9	7.8	0.8	1.11	1.5	[20]
1.1	0.06	42.7	9.7	36.1	9.5	30	[27]
1.04	-	-	10	2	11.7	0.28	[55]
1.4	0.05	-	9.9	3.2	12.7	10.4	[59]
0.99	-	47	-	1.1	10.8	1.3	[60]
0.99	-	47	-	1.1	2.7	11.5	[46]
0.11	0.07	0.09	521	9.2	1.21	0.22	[27]
0.11	0.06	0.1	6.6	0.4	7.25	0.5	[42]
14.8	-	-	10.4	-	15.7	44.9	[25]

Table S2. ANOVA analysis for the effect of ions concentration upon HY, where r2 0.94, Adjust r2 0.93, Predicted r2 0.93, adequate precision (AP) 15.

Source	Sum of squares	DF	Mean square	F-value	p-value
Model	2309.24	20	115.46	1.32	0.2554
A-Mg ²⁺	19.50	1	19.50	0.2223	0.6414
B-Cu ²⁺	31.18	1	31.18	0.3553	0.5565
C-Na ⁺	51.66	1	51.66	0.5887	0.4501
D-NH ⁴⁺	32.96	1	32.96	0.3756	0.5455
E-K ⁺	33.49	1	33.49	0.3816	0.5423
AB	178.00	1	178.00	2.03	0.1667
AC	0.0263	1	0.0263	0.0003	0.9863
AD	456.62	1	456.62	5.20	0.0313
AE	2.38	1	2.38	0.0271	0.8706
BC	48.12	1	48.12	0.5483	0.4659
BD	69.86	1	69.86	0.7960	0.3808
BE	144.30	1	144.30	1.64	0.2115
CD	2.01	1	2.01	0.0229	0.8810
CE	1.52	1	1.52	0.0173	0.8963
DE	54.59	1	54.59	0.6221	0.4377
A ²	484.11	1	484.11	5.52	0.0270
B ²	244.16	1	244.16	2.78	0.1078
C ²	565.76	1	565.76	6.45	0.0177
D ²	681.36	1	681.36	7.76	0.0100
E ²	375.15	1	375.15	4.28	0.0492
Residue	2193.86	25	87.75		
Lack of fit	2193.86	20	109.69		
Pure Error	0.0000	5	0.0000		
Cor total	4503.10	45			