

## Supplementary Materials

### S1. Absorption model

Figure S1 is a mutiple-tube mass balance model during the absorption of CO<sub>2</sub>. Assume  $S$  is the cross section area of the tube,  $\Delta z$  is the differential length, the  $u$  is the linear velocity of gas in the tube, the  $-r_A$  is the reaction rate, the  $d$  is the diameter of the tube, and  $C_A$  is the concentration of CO<sub>2</sub>. According to mass balance at steady-state condition, the balance for multiple tube is showed as follow:

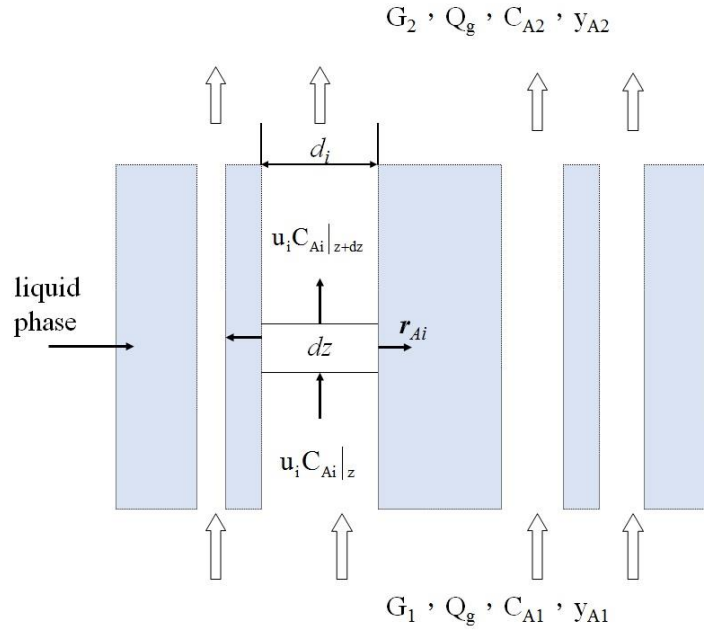


Figure S1. A mutiple-tube mass balance model

$$\sum_{i=1}^N S_i u_i (C_{Ai}|_z - C_{Ai}|_{z+\Delta z}) - \sum_{i=1}^N (-r_{Ai})(\pi d_i) \Delta z = 0 \quad (s1)$$

Eq. (s1) is divided by  $\Delta z$  and take a limit  $\Delta z \rightarrow 0$ , the equation becomes:

$$-\frac{\sum d(u_i S_i C_{Ai})}{dz} - \sum_{i=1}^N (-r_{Ai})(\pi d_i) = 0 \quad (s2)$$

Because  $G_{Ai} = u_i S_i C_{Ai}$ , Eq. (s2) can be rewritten as follow:

$$-\frac{dG_A}{dz} - \sum_{i=1}^N (-r_{Ai})(\pi d_i) = 0 \quad (s3)$$

where  $G_A = \sum_{i=1}^N G_{Ai}$ . Eq.(s3) multiplies by  $dz$  and takes an integration:

$$\int_1^2 dG_A + \int_1^2 \sum_{i=1}^N (-r_{Ai}) dA_i = 0 \quad (s4)$$

or

$$G_{A2} - G_{A1} = \int_1^2 \sum_{i=1}^N (-r_{Ai}) dA_i \quad (s5)$$

The average absorption rate can be defined as follow:

$$-\bar{r}_A = \frac{\sum_1^2 \int (-r_{Ai}) dA_i}{\sum_1^2 \int dA_i} = \frac{\sum_1^2 \int (-r_{Ai}) dA_i}{A} \quad (s6)$$

Therefore, Eq.(s6) becomes:

$$G_{A2} - G_{A1} = \int_1^2 \sum_{i=1}^N (-r_{Ai}) dA_i = (-\bar{r}_A) A \quad (s7)$$

If Eq. (s7) is the first stage in the column, all the other stage could be showed below:

$$\text{First stage} \quad G_{A2} - G_{A1} = (-\bar{r}_{A1}) A_1 \quad (s8)$$

$$\text{Second stage} \quad G_{A3} - G_{A2} = (-\bar{r}_{A2}) A_2 \quad (s9)$$

$$\text{Third stage} \quad G_{A4} - G_{A3} = (-\bar{r}_{A3}) A_3 \quad (s10)$$

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$$\text{Mth stage} \quad G_{AM+1} - G_{AM} = (-\bar{r}_{AM}) A_M \quad (s11)$$

Take summation for all stages, the result is:

$$G_{AM+1} - G_{A1} = \sum_{j=1}^M (-\bar{r}_{Aj}) A_j = (-R_A) V_L \quad (s12)$$

where  $-\bar{r}_{Aj}$  means average rate for jth stage and  $-R_A$  is:

$$-R_A = \frac{\sum_{j=1}^M (-\bar{r}_{Aj}) A_j}{V_L} \quad (s13)$$

where  $V_L$  is the liquid volume in the column; therefore, Eq.(s12) becomes:

$$-R_A = \frac{G_{AM+1} - G_{A1}}{V_L} \quad (s14)$$

In Fig. 3,  $G_{AM+1}$  and  $G_{A1}$  are represented molar flow rate for inlet and outlet, respectively. In here, assume only two components,  $\text{CO}_2(\text{A}) + \text{N}_2(\text{B})$ , in which B is the tie element. Due to this, the molar flow rate of B can be estimated below:

$$G_{BM+1} = G_{AM+1} \left[ \frac{1 - y_{AM+1}}{y_{AM+1}} \right] \quad (\text{s15})$$

Therefore, the outlet concentration of A becomes:

$$G_{A1} = G_{BM+1} \left[ \frac{y_{A1}}{1 - y_{A1}} \right] = G_{AM+1} \left[ \frac{1 - y_{AM+1}}{y_{AM+1}} \right] \left[ \frac{y_{A1}}{1 - y_{A1}} \right] \quad (\text{s16})$$

Substitute Eq.(s16) into Eq.(s14), the result is:

$$-R_A = \frac{G_{AM+1}}{V_L} \left( 1 - \left[ \frac{1 - y_{AM+1}}{y_{AM+1}} \right] \left[ \frac{y_{A1}}{1 - y_{A1}} \right] \right) \quad (\text{s17})$$

This means that absorption rate can be calculated when the inlet and outlet concentrations of component A are available. On the other hand, overall mass-transfer coefficient can be derived by using mass balance and two-film model as shown in Fig. 3(b). Using differential mass balance in Eq.(s11), the mass balance at  $j$ th stage becomes:

$$Q_g dC_{Aj} = dG_{Aj} = (-\bar{r}_{Aj}) dA_j = (-R_{Aj}) dV_{Lj} = (K_G a)_j (C_{Aj} - H_j C_{LA}) dV_{Lj} \quad (\text{s18})$$

where  $C_A$  is the concentration of  $\text{CO}_2$  gas in the gas phase,  $C_{LA}$  is the concentration of  $\text{CO}_2$  gas in the liquid phase, and  $H$  is the Henry's constant. In general,  $C_A$  is much higher than  $C_{LA}$  ( $C_A \gg C_{LA}$ ). Therefore, Eq.(s18) becomes:

$$Q_g dC_{Aj} = (K_G a)_j C_{Aj} dV_{Lj} \quad (\text{s19})$$

or

$$Q_g \frac{dC_{Aj}}{C_{Aj}} = (K_G a)_j dV_{Lj} \quad (\text{s20})$$

Make a summation for Eq.(s20), it becomes:

$$Q_g \sum_{j=1}^M \frac{dC_{Aj}}{C_{Aj}} = \sum_{j=1}^M (K_G a)_j dV_{Lj} \quad (\text{s21})$$

Take an integration in Eq.(s20) stage by stage,

$$\text{No.1} \quad Q_g \int_1^2 \frac{dC_A}{C_A} = \int_1^2 (K_G a)_1 dV_{L1}$$

$$\text{No.2} \quad Q_g \int_2^3 \frac{dC_A}{C_A} = \int_2^3 (K_G a)_2 dV_{L2}$$

$$\text{No.3} \quad Q_g \int_3^4 \frac{dC_{A3}}{C_{A3}} = \int_3^4 (K_G a)_3 dV_{L3}$$

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$$\text{No.M} \quad Q_g \int_{M-1}^M \frac{dC_A}{C_A} = \int_{M-1}^M (K_G a)_M dV_{LM}$$

Addition of each stage, the result becomes:

$$Q_g \int_1^M \frac{dC_A}{C_A} = K_G a V_L \quad (\text{s22})$$

where

$$K_G a = \frac{\int_1^2 (K_G a)_1 dV_{L1} + \int_2^3 (K_G a)_2 dV_{L2} + \dots + \int_{M-1}^M (K_G a)_M dV_{LM}}{V_L} \quad (\text{s23})$$

Make an integration for Eq.(s22), it becomes:

$$K_G a = \frac{Q_g}{V_L} \ln \frac{C_{AM}}{C_{A1}} \quad (\text{s24})$$

Eq.(s24) can be used to evaluate the overall mass-transfer coefficient when  $Q_g$ ,  $V_L$ ,  $C_{AM}$  and  $C_{A1}$  are available.

S2. *Regeneration Data*

Table S1. CO<sub>2</sub> loading data before and after regenerations.

1-s	$\alpha$ Before	T (°C)	$\alpha$ After
E <sub>F</sub>	0.8157	110	0.7523
		120	0.7813
		130	0.7932
R <sub>A</sub>	0.7050	110	0.6628
		120	0.6724
		130	0.6774
K <sub>Ga</sub>	0.6852	110	0.6423
		120	0.6583
		130	0.6656
$\phi$	0.7381	110	0.6816
		120	0.6956
		130	0.7013
2-s	$\alpha$ Before	T (°C)	$\alpha$ After
E <sub>F</sub>	0.7941	110	0.7477
		120	0.7537
		130	0.7616
R <sub>A</sub>	0.6614	110	0.6309
		120	0.6384
		130	0.6416
K <sub>Ga</sub>	0.6667	110	0.6357
		120	0.6385
		130	0.6432
$\phi$	0.5595	110	0.4992
		120	0.5002
		130	0.5387

3-s	$\alpha$ Before	T (°C)	$\alpha$ After
E <sub>F</sub>	0.7932	110	0.7374
		120	0.7586
		130	0.7639
R <sub>A</sub>	0.6183	110	0.5842
		120	0.5895
		130	0.5947
K <sub>Ga</sub>	0.6331	110	0.5795
		120	0.6027
		130	0.6103
$\phi$	0.4667	110	0.4271
		120	0.4325
		130	0.4381

Table S2. Heat of regeneration data obtained according to  $E_F$  optimum condition.

$E_F$	T (°C )	$q_{sen}$ (GJ/ton)	$q_{sol}$ (GJ/ton)	$q_{ads}$ (GJ/ton)	q (GJ/ton)
1-s	110	0.56 (0.148)	2.00 (0.529)	1.22 (0.323)	3.78
	120	1.03 (0.155)	4.41 (0.662)	1.22 (0.183)	6.66
	130	1.57 (0.177)	6.10 (0.686)	1.22 (0.137)	8.89
2-s	110	0.69 (0.193)	1.66 (0.465)	1.22 (0.342)	3.57
	120	0.79 (0.200)	3.32 (0.495)	1.22 (0.305)	5.33
	130	0.99 (0.154)	4.22 (0.656)	1.22 (0.190)	6.43
3-s	110	0.57 (0.147)	2.02 (0.532)	1.22 (0.321)	3.80
	120	0.92 (0.167)	3.33 (0.610)	1.22 (0.223)	5.46
	130	1.08 (0.181)	3.66 (0.614)	1.22 (0.205)	5.96

Table S3. Heat of regeneration data obtained according to  $R_A$  optimum condition.

$R_A$	T (°C)	$q_{sen}$ (GJ/ton)	$q_{sol}$ (GJ/ton)	$q_{ads}$ (GJ/ton)	q (GJ/ton)
1-s	110	0.72 (0.134)	2.92 (0.544)	1.73 (0.322)	5.37
	120	0.93 (0.135)	4.23 (0.614)	1.73 (0.251)	6.89
	130	1.10 (0.140)	4.98 (0.638)	1.73 (0.222)	7.81
2-s	110	0.89 (0.159)	2.99 (0.533)	1.73 (0.308)	5.61
	120	1.18 (0.190)	3.29 (0.531)	1.73 (0.279)	6.20
	130	1.37 (0.199)	3.81 (0.551)	1.73 (0.250)	6.91
3-s	110	0.87 (0.163)	2.78 (0.516)	1.73 (0.321)	5.39
	120	1.03 (0.182)	2.90 (0.512)	1.73 (0.306)	5.66
	130	1.26 (0.181)	3.97 (0.570)	1.73 (0.249)	6.96



Table S4. Heat of regeneration data obtained according to K<sub>Ga</sub> optimum condition.

K <sub>Ga</sub>	T (°C)	q <sub>sen</sub> (GJ/ton)	q <sub>sol</sub> (GJ/ton)	q <sub>ads</sub> (GJ/ton)	q (GJ/ton)
1-s	110	0.56 (0.101)	3.24 (0.586)	1.73 (0.313)	5.53
	120	0.87 (0.148)	3.32 (0.560)	1.73 (0.292)	5.93
	130	1.20 (0.150)	5.06 (0.633)	1.73 (0.217)	7.99
2-s	110	0.60 (0.131)	2.31 (0.497)	1.73 (0.372)	4.65
	120	0.67 (0.127)	2.85 (0.543)	1.73 (0.330)	5.25
	130	0.80 (0.134)	3.45 (0.577)	1.73 (0.289)	5.98
3-s	110	0.32 (0.090)	1.53 (0.427)	1.73 (0.483)	3.58
	120	0.56 (0.110)	2.76 (0.547)	1.73 (0.343)	5.05
	130	0.75 (0.119)	3.82 (0.606)	1.73 (0.275)	6.30

Table S5. Heat of regeneration data obtained according to  $\phi$  optimum condition.

$\phi$	T (°C)	q <sub>sen</sub> (GJ/ton)	q <sub>sol</sub> (GJ/ton)	q <sub>ads</sub> (GJ/ton)	q (GJ/ton)
1-s	110	0.69 (0.130)	3.24 (0.620)	1.31 (0.250)	5.23
	120	0.92 (0.151)	3.88 (0.635)	1.31 (0.214)	6.11
	130	1.06 (0.125)	6.13 (0.721)	1.31 (0.154)	8.50
2-s	110	0.65 (0.135)	2.86 (0.593)	1.31 (0.272)	4.82
	120	0.67 (0.121)	3.48 (0.639)	1.31 (0.240)	5.45
	130	1.89 (0.212)	5.73 (0.642)	1.31 (0.219)	8.93
3-s	110	0.99 (0.201)	2.61 (0.532)	1.31 (0.267)	4.91
	120	1.16 (0.190)	3.60 (0.594)	1.31 (0.216)	6.06
	130	1.38 (0.197)	4.32 (0.616)	1.31 (0.187)	7.01