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2 APPENDIXES

3 **Appendix A**

4
 5 The rate equations of the model of photosynthetic carbon metabolism. The subscripts of
 6 $v_1, v_2 \dots, v_{131}$ correspond to the numbers in Fig. 1. The kinetic parameters in the rate
 7 equations are listed in Appendices C and D. See Appendix G for definitions of
 8 abbreviations.

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$$v_1 = \frac{\text{RuBP} \times W_c \times \min(1, \frac{\text{RuBP}}{E_t})}{(\text{RuBP} + K_r(1 + \frac{\text{PGA}}{K_{I11}} + \frac{\text{FBP}}{K_{I12}} + \frac{\text{SBP}}{K_{I13}} + \frac{\text{Pi}}{K_{I14}} + \frac{\text{NADPH}}{K_{I15}}))}$$

$$W_C = \frac{V_{C\max} \times CO_2}{CO_2 + K_{M11}(1 + \frac{O_2}{K_{M12}})}$$

$$v_2 = \frac{V_2 \times (\text{PGA} \times ATP - \frac{DPGA \times ADP}{K_{E2}})}{(\text{PGA} + K_{M21})(ATP + K_{M22}(1 + \frac{ADP}{K_{M23}}))}$$

$$v_3 = \frac{V_3 \times DPGA \times NADPH}{(DPGA + K_{M31})(NADPH + K_{M32})}$$

$$v_5 = \frac{V_5 \times (GAP \times DHAP - \frac{FBP}{K_{E5}})}{K_{M51}K_{M52}(1 + \frac{GAP}{K_{M51}} + \frac{DHAP}{K_{M52}} + \frac{FBP}{K_{M53}} + \frac{GAP \times DHAP}{K_{M51}K_{M52}})}$$

$$v_6 = \frac{V_6 \times (FBP - \frac{F6P \times Pi}{K_{E6}})}{FBP + K_{M61}(1 + \frac{F6P}{K_{I61}} + \frac{Pi}{K_{I62}})}$$

$$TEMP1 = Xu5P \times (1 + \frac{E4P \times Ri5P}{Km71}) + E4P + Ri5P$$

$$Den = 1 + (1 + \frac{GAP}{K_{M72}}) \times (\frac{F6P}{K_{M104}} + \frac{S7P}{K_{M103}}) + \frac{GAP}{K_{M102}} + \frac{TEMP1}{K_{M101}}$$

$$v_7 = \frac{V_7 \times (F6P \times GAP \times K_{E7} - Xu5P \times E4P)}{(K_{M71} \times K_{M101} \times Den)}$$

$$v_8 = \frac{V_8 \times (DHAP \times E4P - \frac{SBP}{K_{E8}})}{(E4P + K_{M82})(DHAP + K_{M81})}$$

$$v_9 = \frac{V_9 \times (SBP - \frac{Pi \times S7P}{K_{E9}})}{SBP + K_{M9}(1 + \frac{Pi}{K_{I9}})}$$

$$v_{10} = \frac{V_{10} \times (GAP \times S7P \times K_{E10} - Ri5P \times Xu5P)}{(K_{M71} \times K_{M101} \times Den)}$$

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- 2 Alternate simplified equations for v_7 and v_{10} can also be used following Poolman et al
 3 (2000).

$$v_{13} = \frac{V_{13} \times (ATP \times Ru5P - \frac{ADP \times RuBP}{K_{E13}})}{(ATP \times (1 + \frac{ADP}{K_{I134}}) + K_{M132} (1 + \frac{ADP}{K_{I135}})) (Ru5P + K_{M131} (1 + \frac{GAP}{K_{I131}} + \frac{RuBP}{K_{I132}} + \frac{Pi}{K_{I133}}))}$$

$$v_{16} = \frac{V_{16} \times (ADP \times Pi - \frac{ATP}{K_{E13}})}{(ADP + K_{M161}) (Pi + K_{M162})}$$

$$v_{23} = \frac{V_{23} \times G1P \times ATP}{(G1P + K_{M231}) ((1 + \frac{ADP}{K_{I23}}) (ATP + K_{M232}) + \frac{K_{M232} \times Pi}{K_{A231} \times PGA + K_{A232} \times F6P + K_{A233} \times FBP})}$$

$$N = 1 + (1 + \frac{K_{M313}}{P_{ext}}) (\frac{Pi}{K_{M312}} + \frac{PGA}{K_{M32}} + \frac{GAP}{K_{M33}} + \frac{DHAP}{K_{M311}})$$

$$v_{31} = \frac{V_{31} \times DHAP}{K_{M311} N}$$

$$v_{32} = \frac{V_{32} \times PGA}{K_{M32} N}$$

$$v_{33} = \frac{V_{33} \times GAP}{K_{M33} N}$$

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$$3 \quad v_{55} = \frac{V_{55} \times (UTPc \times G1Pc - \frac{UDPGc \times OPOPc}{K_{E551}})}{K_{m551} K_{m552} (1 + \frac{UTPc}{K_{m551}} + \frac{G1Pc}{K_{m552}} + \frac{UDPGc}{K_{m553}} + \frac{OPOPc}{K_{m554}} + \frac{UTPc \times G1Pc}{K_{m551} K_{m552}} + \frac{UDPGc \times OPOPc}{K_{m553} K_{m554}})}$$
$$4 \quad v_{56} = \frac{V_{56} \times (F6Pc \times UDPGc - \frac{SUCPc \times UDPc}{K_{E56}})}{(F6Pc + K_{m561} (1 + \frac{FBPc}{K_{I562}})) (UDPGc + K_{m562} (1 + \frac{UDPc}{K_{I561}}) (1 + \frac{SUCPc}{K_{I563}}) (1 + \frac{SUCC}{K_{I565}}) (1 + \frac{Pic}{K_{I564}}))}$$
$$5 \quad v_{57} = \frac{V_{57} \times (SUCPc - \frac{SUCC \times Pic}{K_{E57}})}{SUCPc + K_{m571} \times (1 + \frac{SUCC}{K_{m572}})}$$
$$6 \quad v_{58} = \frac{V_{58} \times F26BPc}{K_{m581} \times (1 + \frac{F26BPc}{K_{m581}}) (1 + \frac{Pic}{K_{I582}}) (1 + \frac{F6Pc}{K_{I581}})}$$
$$7 \quad v_{59} = \frac{V_{59} \times (ATPc \times F6Pc - \frac{ADPc \times F26BPc}{K_{E59}})}{(F6Pc + K_{m593} \times (1 + \frac{F26BPc}{K_{m592}}) (1 + \frac{DHAPc}{K_{I592}})) ((ATPc + K_{m591} \times (1 + \frac{ADPc}{K_{I591}})))}$$
$$8 \quad v60 = \frac{V_{60} \times (ATPc \times UDPc - ADPc \times UTPc / KE60)}{K_{m602} K_{m603} \times (1 + \frac{ATPc}{K_{m602}} + \frac{UDPc}{K_{m603}} + \frac{ATPc \times UDPc}{K_{m602} \times K_{m603}} + \frac{ADPc}{K_{m601}} + \frac{UTPc}{K_{m604}} + \frac{ADPc \times UTPc}{K_{m601} K_{m604}})}$$

$$v_{111} = \frac{RuBP \times W_o \times \min(1, \frac{RuBP}{E_t})}{(RuBP + K_r(1 + \frac{PGA}{K_{I11}} + \frac{FBP}{K_{I12}} + \frac{SBP}{K_{I13}} + \frac{PI}{K_{I14}} + \frac{NADPH}{K_{I15}}))}$$

$$W_o = \frac{V_{111} \times O_2}{O_2 + K_o(1 + \frac{CO_2}{k_C})}$$

$$v_{112} = \frac{V_{112} \times PGCA}{PGCA + K_{m112}(1 + GCA/K_{i1121})(1 + Pi/K_{I1122})}$$

$$v_{113} = \frac{V_{113} \times (ATP \times GCEA - \frac{ADP \times PGA}{K_{E113}})}{(ATP + Km_{1131}(1 + \frac{PGA}{K_{i113}}))(GCEA + K_{m1132})}$$

$$v_{121} = \frac{V_{121} \times GCAC}{GCAC + K_{m121}}$$

$$v_{122} = \frac{V_{122} \times (GOAc \times SERc - \frac{HPRc \times GLYc}{K_{E122}})}{(GOAc + K_{m1221})(SERc + K_{m1222}(1 + \frac{GLYc}{K_{i1221}}))}$$

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$$v_{123} = \frac{V_{123} \times (HPRc \times NADHc - \frac{NADc \times Gcea}{K_{E123}})}{HPRc + K_{m1231}(1 + HPRc/K_{I123})}$$

$$v_{124} = \frac{V_{124} \times (GOAc \times GLUc - \frac{KGc \times GLYc}{K_{E124}})}{(GOAc + K_{m1241})(GLUc + K_{m1242}(1 + GLYc/K_{I124}))}$$

$$v_{131} = \frac{V_{131} \times GLYc}{GLYc + K_{m1311}(1 + SERc/K_{I1311})}$$

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$$v_{1in} = V_{1T} \frac{GCEAc}{GCEAc + K_{M1011}(1 + \frac{GCAc}{K_{I1011}})}$$

$$v_{1out} = V_{1T} \frac{GCEA}{GCEA + K_{M1011}(1 + \frac{GCA}{K_{I1011}})}$$

$$v_{2out} = V_{2T} \frac{GCA}{GCA + K_{M1012}(1 + \frac{GCEA}{K_{I1012}})}$$

$$v_{2in} = V_{2T} \frac{GCAC}{GCAC + K_{M1012}(1 + \frac{GCEAc}{K_{I1012}})}$$

$$G6P = \frac{\text{HexP}}{\frac{1}{K_{E21}} + K_{E22} + 1}$$

$$F6P = \frac{\frac{1}{K_{E21}} \times \text{HexP}}{\frac{1}{K_{E21}} + K_{E22} + 1}$$

$$G1P = \frac{K_{E22} \times \text{HexP}}{\frac{1}{K_{E21}} + K_{E22} + 1}$$

$$T3P = GAP + DHAP$$

$$Pent = Ru5P + Ri5P + Xu5P$$

$$HexP = G6P + F6P + G1P$$

$$GAP = \frac{T3P}{1 + K_{e4}}$$

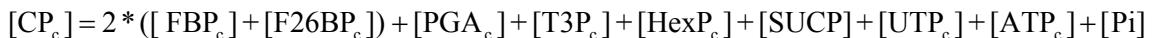
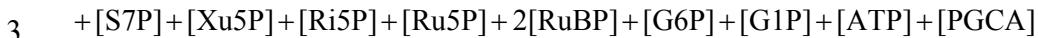
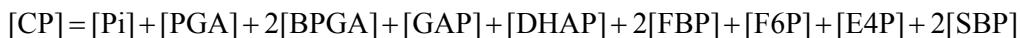
$$DHAP = \frac{K_{e3} \times T3P}{1 + K_{e4}}$$

$$Ru5P = \frac{Pent}{\frac{1}{K_{E11}} + \frac{1}{K_{E12}} + 1}$$

$$Ri5P = \frac{\frac{1}{K_{E11}} \times Pent}{\frac{1}{K_{E11}} + \frac{1}{K_{E12}} + 1}$$

$$Xu5P = \frac{\frac{1}{K_{E12}} \times Pent}{\frac{1}{K_{E11}} + \frac{1}{K_{E12}} + 1}$$

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1 Appendix B

- 2 Differential equations to describe rates of change in each intermediate of carbon
 3 metabolism. See Appendix A for definition of $v_1, v_2 \dots, v_{131}$.

$$\frac{d[RuBP]}{dt} = v_{13} - v_1 - v_{111};$$

$$\frac{d[PGA]}{dt} = 2v_1 - v_2 - v_{32} + v_{111} + v_{113};$$

$$\frac{d[DPGA]}{dt} = v_2 - v_3;$$

$$\frac{d[T3P]}{dt} = v_3 - 2v_5 - v_7 - v_8 - v_{10} - v_{31} - v_{33};$$

$$\frac{d[FBP]}{dt} = v_5 - v_6;$$

$$\frac{d[E4P]}{dt} = v_7 - v_8;$$

$$\frac{d[S7P]}{dt} = v_9 - v_{10};$$

$$\frac{d[SBP]}{dt} = v_8 - v_9;$$

$$\frac{d[ATP]}{dt} = v_{16} - v_2 - v_{23} - v_{13} - v_{113};$$

$$\frac{d[HexP]}{dt} = v_6 - v_7 - v_{23};$$

$$\frac{d[PenP]}{dt} = v_7 + 2v_{10} - v_{13};$$

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$$\frac{d[GOAc]}{dt} = v_{121} - v_{122} - v_{124}$$

$$\frac{d[SERc]}{dt} = v_{131} - v_{122}$$

$$\frac{d[GLYc]}{dt} = v_{122} + v_{124} - 2v_{131}$$

$$\frac{d[HPRc]}{dt} = v_{122} - v_{123}$$

$$\frac{d[GCEAc]}{dt} = v_{123} - v_{1in} + v_{1out}$$

$$\frac{d[GCEA]}{dt} = v_{1in} - v_{113} - v_{1out}$$

$$\frac{d[GCA]}{dt} = v_{112} - v_{2out} + v_{2in}$$

$$\frac{d[PGCA]}{dt} = v_{111} - v_{112}$$

$$\frac{d[GCAC]}{dt} = v_{2out} - v_{121} - v_{2in}$$

$$\frac{d[OPOPc]}{dt} = v_{55} - v_{61}$$

$$\frac{d[UTPc]}{dt} = v_{60} - v_{55}$$

$$1 \quad \frac{d[SUCPc]}{dt} = v_{56} - v_{57}$$

$$\frac{d[SUCc]}{dt} = v_{57} - v_{62}$$

$$\frac{d[PGAc]}{dt} = v_{32} - v_{pga_use}$$

$$\frac{d(T3Pc)}{dt} = v_{31} + v_{33} - 2v_{51}$$

$$\frac{d(FBPc)}{dt} = v_{51} - v_{52}$$

$$\frac{d(HexPc)}{dt} = v_{52} - v_{55} - v_{59} + v_{58} - v_{56}$$

$$2 \quad \frac{d[F26BPc]}{dt} = v_{59} - v_{58}$$

$$\frac{d[UDPGc]}{dt} = v_{55} - v_{56}$$

$$\frac{d[ATPc]}{dt} = v_{atpf} - v_{59} - v_{60}$$

1 Appendix C

2 Kinetic parameters of the enzymes of the Calvin cycle, starch synthesis and triose
3 phosphate export illustrated in Fig. 1.

4 Table C1

5 The Michaelis-Menten constants, inhibition constants and activation constants of the enzymes in
6 the Calvin cycle, starch synthesis and triose phosphate export. The response coefficient of each
7 parameter was calculated for the initial nitrogen distribution (See text).

RN ^a	Reaction	Par ^b	Value (mM) ^c	Description n ^d	CC ^e	Reference
1	RuBP+CO ₂ →2PGA	K _{M11}	0.0115	CO ₂	-0.60	Jordan and Ogren (1981), Tcherkez <i>et al.</i> (Tcherkez et al., 2006)
1	RuBP+CO ₂ →2PGA	K _{M12}	0.222	O ₂	0.54	Jordan and Ogren (1981), von Caemmerer (2000)
1	RuBP+CO ₂ →2PGA	K _{M13}	0.020	RuBP	-0.07	Farquhar (1979)
1	RuBP+CO ₂ →2PGA	K _{I11}	0.84	PGA	8×10 ⁻⁴	Badger and Lorimer (1981)
1	RuBP+CO ₂ →2PGA	K _{I12}	0.04	FBP	1×10 ⁻³	Badger and Lorimer (1981)
1	RuBP+CO ₂ →2PGA	K _{I13}	0.075	SBP	0.02	Badger and Lorimer (1981)
1	RuBP+CO ₂ →2PGA	K _{I14}	0.9	Pi	2×10 ⁻⁴	Badger and Lorimer (1981)
1	RuBP+CO ₂ →2PGA	K _{I15}	0.07	NADPH	0.004	Badger and Lorimer (1981)
2	PGA+ATP ↔ ADP + DPGA	K _{M21}	0.240	PGA	6×10 ⁻⁴	Larsson-Raznickiewicz (1967), Kopkesecundo et al. (1990)
2	PGA+ATP ↔ ADP + DPGA	K _{M22}	0.390	ATP	2×10 ⁻³	Larsson-Raznickiewicz (1967), Kopkesecundo et al. (1990)
2	PGA+ATP ↔ ADP + DPGA	K _{M23}	0.23	ADP	-1×10 ⁻³	Lee (1982)
2	PGA+ATP↔ADP + DPGA	K _{E2}	7.6	Equ. ×10 ⁻⁴	Const.	Laisk <i>et al.</i> (1989); Dietz et al (1984); Heber et al (1986)
3	DPGA+NADPH+H ⁺ ↔GAP + Pi+NADP	K _{M31}	0.004	BPGA	-5×10 ⁻⁵	Trost <i>et al.</i> (1993)
3	DPGA+NADPH +H ⁺ ↔GAP + Pi+NADP	K _{M32}	0.100	NADPH	-4×10 ⁻⁵	Cerf (1978), Ferri <i>et al</i> (1978), Trost (1993), Macioszek and Anderson (1987)
4	DHAP ↔GAP	K _{E4}	0.05	Equ. Const.	6×10 ⁻³	Bassham and Krause (1969)
5	GAP+DHAP ↔FBP	K _{M51}	0.3	GAP	-2×10 ⁻³	Iwaki <i>et al</i> (1991)
5	GAP+DHAP ↔FBP	K _{M52}	0.4	DHAP	-9×10 ⁻⁴	Iwaki <i>et al</i> (1991)
5	GAP+DHAP ↔FBP	K _{M53}	0.02	FBP	3×10 ⁻⁴	Brooks and Criddle (1966), Schnarrenberger and Kruger (1986)
5	GAP+DHAP ↔FBP	K _{E5}	7.1	Equ. Const.	5×10 ⁻⁵	Bassham and Krause (1969), Iwaki <i>et al</i> (1991)
6	FBP→F6P+Pi	K _{M61}	0.033	FBP	-2×10 ⁻³	Charles and Halliwell (1981)
6	FBP→F6P+Pi	K _{I61}	0.7	F6P	4×10 ⁻⁴	Heldt (1983)
6	FBP→F6P+Pi	K _{I62}	12	Pi	1×10 ⁻⁵	Charles and Halliwell (1981)
6	FBP→F6P+Pi	K _{E6}	6.7×10 ⁵	Equ	4×10 ⁻⁹	Bassham and Krause (1969) , Laisk <i>et al.</i> (1989)

				Const.		
7	F6P+GAP→E4P+Xu5P	K _{M71}	0.1	Xu5P	4×10 ⁻⁴	Murphy and Walker (1982), Laisk <i>et al</i> (1989)
7	F6P+GAP→E4P+Xu5P	K _{M72}	0.1	GAP	2×10 ⁻³	Sprenger et al (1995), Schenk et al (1998)
7	F6P+GAP→E4P+Xu5P	K _{E7}	10	Equ.	9×10 ⁻⁴	Bassham and Krause (1969), Laisk <i>et al.</i> (1989)
				Const.		
8	E4P+DHAP→SBP	K _{M8}	0.02	SBP	0	Brooks and Criddle (1966)
8	E4P+DHAP→SBP	K _{M81}	0.4	DHAP	-3×10 ⁻⁴	Iwaki et al (1991)
8	E4P+DHAP→SBP	K _{M82}	0.2	E4P	-6×10 ⁻⁴	Estimate
8	E4P+DHAP→SBP	K _{E8}	1.07	Equ.	4×10 ⁻³⁴	Bassham and Krause (1969) , Laisk <i>et al.</i> (1989)
				Const.		
9	SBP→S7P+Pi	K _{M9}	0.05	SBP	-0.03	Woodrow et al (1983), Cadet and Meunier (1988)
9	SBP→S7P+Pi	K _{I9}	12	Pi	3×10 ⁻⁴	Woodrow et al(1983)
9	SBP→S7P+Pi	K _{E9}	6.7×10 ⁵	Equ.	4×10 ⁻⁸	Bassham and Krause (1969), Laisk <i>et al.</i> (1989)
				Const.		
10	S7P+GAP→Ri5P+Xu5P	K _{M101}	0.118	P5P	2×10 ⁻⁴	Racher (1961), Laisk <i>et al</i> (1989)
10	S7P+GAP→Ri5P+Xu5P	K _{M102}	0.072	GAP	-4×10 ⁻⁴	Albe (1991); Laisk <i>et al</i> (1989)
10	S7P+GAP→Ri5P+Xu5P	K _{M103}	0.46	S7P	-5×10 ⁻⁴	Albe (1991); Laisk <i>et al</i> (1989)
10	S7P+GAP→Ri5P+Xu5P	K _{M104}	1.54	F6P	3×10 ⁻⁵	Albe (1991); Laisk <i>et al</i> (1989)
10	S7P+GAP→Ri5P+Xu5P	K _{E10}	1.17	Equ.	8×10 ⁻⁴	Bassham and Krause (1969), Laisk <i>et al.</i> (1989)
				Const.		
11	Ri5P↔Ru5P	K _{E11}	0.4	Equ.	2×10 ⁻³	Bassham and Krause (1969)
				Const.		
12	Xu5P↔Ru5P	K _{E12}	0.67	Equ.	3×10 ⁻³	Bassham and Krause, (1969)
				Const.		
13	Ru5P+ATP→RuBP+ADP	K _{M131}	0.05	Ru5P	-4×10 ⁻³	Gardemann et al (1983), Omnaas et al (1985)
13	Ru5P+ATP→RuBP+ADP	K _{M132}	0.059	ATP	-2×10 ⁻³	Laing et al (1981), Gardemann et al (1983), Omnaas et al (1985)
13	Ru5P+ATP→RuBP+ADP	K _{I131}	2	PGA	2×10 ⁻⁴	Gardemann et al (1983)
13	Ru5P+ATP→RuBP+ADP	K _{I132}	0.7	RuBP	4×10 ⁻³	Gardemann et al (1983)
13	Ru5P+ATP→RuBP+ADP	K _{I133}	4	Pi	2×10 ⁻⁵	Gardemann et al (1983)
13	Ru5P+ATP→RuBP+ADP	K _{I134}	2.5	ADP	1×10 ⁻³	Gardemann et al (1983)
13	Ru5P+ATP→RuBP+ADP	K _{E13}	0.4	ADP	2×10 ⁻⁴	Gardemann et al (1983)
13	Ru5P+ATP→RuBP+ADP	K _{I135}	6846	Equ.	2×10 ⁻³	Bassham and Krause (1969), Laisk <i>et al.</i> (1989)
				Const.		
16	ADP+Pi→ATP	K _{M161}	0.014	ADP	-1×10 ⁻⁵	Davenport and Mccarty (1986)
16	ADP+Pi→ATP	K _{M162}	0.3	Pi	-6×10 ⁻⁴	Aflalo and Shavit (1983)
16	ADP+Pi→ATP	K _{M163}	0.3	ATP	5×10 ⁻⁶	r.f. Laisk <i>et al.</i> (1989)
16	ADP+Pi→ATP	K _{E16}	5.7	Equ.	0	Bassham and Krause (1969), Laisk <i>et al.</i> (1989)
				Const.		
21	F6P↔G6P	K _{E21}	2.3	Equ.	-2×10 ⁻³	Bassham and Krause (1969)
				Const.		
22	G6P↔G1P	K _{E22}	0.058	Equ.	-5×10 ⁻⁵	Colowick and Sutherland (1942)
				Const.		
23	G1P+ATP+Gn→PPi+ADP+	K _{M231}	0.08	G1P	-6×10 ⁻⁵	Pettersson and Ryde-Pettersson (1988)
	G _{n+1}					
23	G1P+ATP+Gn→PPi+ADP+	K _{M232}	0.08	ATP	-4×10 ⁻⁵	Pettersson and Ryde-Pettersson (1988)

	G_{n+1}						
23	$G1P + ATP + Gn \rightarrow PPi + ADP + G_{n+1}$	K_{A231}	0.1	PGA	2×10^{-5}	Pettersson and Ryde-Pettersson (1988)	
23	$G1P + ATP + Gn \rightarrow PPi + ADP + G_{n+1}$	K_{A232}	0.02	F6P	2×10^{-6}	Pettersson and Ryde-Pettersson (1988)	
23	$G1P + ATP + Gn \rightarrow PPi + ADP + G_{n+1}$	K_{A233}	0.02	FBP	3×10^{-7}	Pettersson and Ryde-Pettersson (1988)	
23	$G1P + ATP + Gn \rightarrow PPi + ADP + G_{n+1}$	K_{I23}	10	ADP	7×10^{-6}	Pettersson and Ryde-Pettersson (1988)	
31	$P_{ext} + DHAP_i \rightarrow Pi + DHAP_o$	K_{M311}	0.077	DHAP	-6×10^{-5}	Fliege et al (1978), Portis Jr. (1983)	
31	$P_{ext} + DHAP_i \rightarrow Pi + DHAP_o$	K_{M312}	0.63	Pi	1×10^{-5}	Fliege et al (1978), Portis Jr. (1983)	
31	$P_{ext} + DHAP_i \rightarrow Pi + DHAP_o$	K_{M313}	0.74	P_{ext}	-4×10^{-4}	Fliege et al (1978), Portis Jr. (1983)	
32	$P_{ext} + PGA_i \rightarrow Pi + PGA_o$	K_{M32}	0.25	PGA	-9×10^{-6}	Fliege et al (1978), Portis Jr. (1983)	
33	$P_{ext} + GAP_i \rightarrow Pi + GAP_o$	K_{M33}	0.075	GAP	-3×10^{-6}	Fliege et al (1978), Portis Jr. (1983)	

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2 ^aRN: Reaction number corresponding to the number in Fig. 1.3 ^b Parameters beginning K_M represent the Michaelis-Menten constant of the metabolite listed in the *description* column4 Parameters beginning with K_I represent the inhibition constant of the inhibitor listed in the *description* column.5 Parameters beginning with K_A represent the activation constant of the activator listed in the *description* column.6 ^cThe units of the equilibrium constant are mM (which is equivalent to mmol l⁻¹) for all constants, except K_e . The unit of K_e for
7 reactions with one substrate and one product, or reactions with two substrates and two products is dimensionless. The unit of K_e for
8 reactions with one substrate and two products is mM. The unit for reactions with two substrates and one product is (mM)⁻¹.9 ^dThe description column lists the compounds to which the kinetic constant applies.10 ^e Response coefficients for different parameters used in the model

11

1 **Table C2**

2 The maximum rate of each enzyme (V_m) normalized on maximum Rubisco carboxylation activity
 3 (V_1)

Maximum Velocity	Enzyme	Reaction	V_m/V_1	Reference
V_1	Rubisco	$\text{RuBP} + \text{CO}_2 \rightarrow 2\text{PGA}$	1	Latzko et al (1981), Woodrow (1986), Geiger et al. (1999), Haake et al.(1999), Chen et al. (2005), Strand et al. (2000), Strand et al (1999), Tamoi et al (2006), Peterkofsky and Racher (1961), Chen et al (2005)
V_2	PGA Kinase	$\text{PGA} + \text{ATP} \rightarrow \text{ADP} + \text{DPGA}$	10.3	Same as above
V_3	GAP dehydrogenase	$\text{DPGA} + \text{NADPH} \rightarrow \text{GAP} + \text{Pi} + \text{NADP}$	1.39	Same as above
V_5	FBP Aldolase	$\text{GAP} + \text{DHAP} \rightarrow \text{FBP}$	0.42	Same as above
V_6	FBPase	$\text{FBP} \rightarrow \text{F6P} + \text{Pi}$	0.25	Same as above
V_7	Transketolase	$\text{F6P} + \text{GAP} \rightarrow \text{E4P} + \text{Xu5P}$	1.07	Same as above
V_8	Aldolase	$\text{E4P} + \text{DHAP} \rightarrow \text{SBP}$	0.42	Same as above
V_9	SBPase	$\text{SBP} \rightarrow \text{S7P} + \text{Pi}$	0.11	Same as above
V_{10}	Transketolase	$\text{S7P} + \text{GAP} \rightarrow \text{Ri5P} + \text{Xu5P}$	1.07	Same as above
V_{13}	Ribulosebiphosphate kinase	$\text{Ru5P} + \text{ATP} \rightarrow \text{RuBP} + \text{ADP}$	3.71	Same as above
V_{16}	ATP synthase	$\text{ADP} + \text{Pi} \rightarrow \text{ATP}$	5.5	Same as above
V_{23}	ADP-glucose pyrophosphorylase and Starch Synthase	$\text{ADPG} + \text{G}_n \rightarrow \text{G}_{(n+1)} + \text{ADP}$	0.1	Same as above
V_{31}	Phosphate translocator	$\text{DHAP}_i \rightarrow \text{DHAP}_o$	0.3	Lilley et al (1977)
V_{32}	Phosphate translocator	$\text{PGA}_i \rightarrow \text{PGA}_o$	0.3	Lilley et al (1977)
V_{33}	Phosphate translocator	$\text{GAP}_i \rightarrow \text{GAP}_o$	0.3	Lilley et al (1977)

1

2 **Table C3** The concentrations of the metabolites used in the model. The range of values found
 3 in the literature are given in the adjacent column, together with their source.

4

Metabolite name	Model default (mmol l ⁻¹)	Concentration range (mmol l ⁻¹)	Reference
RuBP	2.000	0.6-6	Bassham and Krause (1969), Dietz and Heber (1984), Schimkat et al (1990), Woodrow and Mott (1993)
PGA	2.400	1.4-12	Schimkat et al (1990), Woodrow and Mott (1993)
DPGA	0.0011	0.8-1.4	Dietz and Heber (1984), Woodrow and Mott (1993)
GAP	0.02	0.032-0.04	Bassham and Krause (1969), Dietz and Heber (1984), Woodrow and Mott (1993)
DHAP	0.48	0.37-0.7	Bassham and Krause (1969), Dietz and Heber (1984), Schimkat et al (1990)
FBP	0.670	0.067-0.52	Bassham and Krause (1969), Dietz and Heber (1984), Schimkat et al (1990), Woodrow and Mott (1993)
E4P	0.050	0.02-0.05	Bassham and Krause (1969), Woodrow and Mott (1993)
S7P	2.0	0.248- 0.4	Bassham and Krause (1969), Woodrow and Mott (1993)
SBP	0.30	0.114-0.3	Bassham and Krause (1969), Schimkat et al (1990), Woodrow and Mott (1993)
ATP	0.68	0.2-2.5	Bassham and Krause (1969), Woodrow and Mott (1993), Igamberdiev et al (2001)
NADPH	0.21	0.21	Giersch et al (1980), Woodrow and Mott (1993)
CO ₂	0.012	0.006	Dietz and Heber (1984)
O ₂	0.26	0.25	Model estimate
HexP	2.2	3.2-4.85	Schimkat et al (1990), Woodrow and Mott (1993), Winter et al (1994)
PenP	0.25	0.05-0.3	Bassham and Krause (1969), Schimkat et al (1990), Woodrow and Mott (1993)
Pi	5	4-10	Dietz and Heber (1984), Schimkat et al (1990), Woodrow and Mott (1993)
CP	15	15	Lilley et al (1977)
CA	1.5	1.5	Igamberdiev et al (2001)
CN	0.5	0.5	Giersch et al (1980)
P _{ext}	0.5	5	Bligny et al (1990), Woodrow and Mott (1993)

5

1 Appendix D

2 Kinetic parameters of the enzymes of photorespiration (PCOP). The source of each value
3 is given in the final column.

4 Table D1

5 The Michaelis-Menten constants and inhibition constants for enzymes in PCOP

RN ^a	Reaction	Param. ^b	Value (mM) ^c	Description ^d	CC ^e	Reference
112	2-PGCA +H ₂ O → GCA + Pi	K _{M112}	0.026	PGCA	-5×10 ⁻⁷	Christeller and Tolbert (1978)
112	2-PGCA +H ₂ O → GCA + Pi	K _{I1121}	94	GCA, competitive with PGCA	5×10 ⁻⁹	Christeller and Tolbert (1978)
112	2-PGCA +H ₂ O → GCA + Pi	K _{I1122}	2.55	Pi, competitive with PGCA	7×10 ⁻⁹	Christeller and Tolbert (1978)
113	GCEA + ATP→ PGA + ADP	K _{M1131}	0.21	ATP	1×10 ⁻⁸	Kleczkowski et al (1985)
113	GCEA + ATP→ PGA + ADP	K _{M1132}	0.25	GCEA	-3×10 ⁻¹⁰	Kleczkowski et al (1985)
113	GCEA + ATP→ PGA + ADP	K _{I113}	0.36	PGA, competitive with ATP	-2×10 ⁻⁸	Kleczkowski and Randall (1988)
113	GCEA + ATP→ PGA + ADP	K _{E113}	300	Equil. Const.	6×10 ⁻⁸	Kleczkowski et al (1985)
121	GCA _C +O ₂ →H ₂ O ₂ +GOAc	K _{M121}	0.1	GCAC	-8×10 ⁻⁹	Tolbert (1981)
122	GOAc + SERc→ HPRc + GLYc	K _{M1221}	0.15	GOAc	-1×10 ⁻⁸	Nakamura and Tolbert (1983)
122	GOAc + SERc→ HPRc + GLYc	K _{M1222}	2.7	SERc	4×10 ⁻¹²	Nakamura and Tolbert (1983)
122	GOAc + SERc→ HPRc + GLYc	K _{I1221}	33	GLYc, competitive with SERc	2×10 ⁻¹⁴	Nakamura and Tolbert (1983)
122	GOAc + SERc→ HPRc + GLYc	K _{E122}	0.24	Equil. Const.	-2×10 ⁻⁸	Guynn (1982)
123	HPRc + NADc→ NADHc + GCEAc	K _{M123}	0.09	HPRc	-2×10 ⁻¹¹	Kleczkowski and Edwards (1989)
123	HPRc + NADc→ NADHc + GCEAc	K _{I123}	12	HPRc, self inhibition	-1.5×10 ⁻⁹	Kleczkowski and Edwards (1989)
123	HPRc + NADc→ NADHc + GCEAc	K _{E123}	2.5 ×10 ⁵	Equil. Const.	-6×10 ⁻¹⁰	Guynn (1982)
124	GOAc + GLUc→ KGc + GLYc	K _{M1241}	0.15	GOAc	1×10 ⁻⁷	Nakamura and Tolbert (1983)
124	GOAc + GLUc→ KGc + GLYc	K _{M1242}	1.7	GLUc	-2×10 ⁻⁸	Nakamura and Tolbert (1983)
124	GOAc + GLUc→ KGc + GLYc	K _{I124}	2	GLYc competitive	-2×10 ⁻¹³	Calibrated

with GLU						
124	GOAc + GLUc \rightarrow KGc + GLYc	K_{E124}	607	Equi. Const.	-3×10^{-14}	Cooper and Meister (1972)
131	GLYc + NADc \rightarrow CO2 + NH3 + SERc +NADHc	K_{M1311}	6	GLYc	-2×10^{-13}	Douce et al (2001)
131	GLYc + NADc \rightarrow CO2 + NH3 + SERc +NADHc	K_{I1311}	4	SERc, competitive with GLYc	6×10^{-9}	Douce et al (2001)
101a	GCEAc \rightarrow GCEA	K_{M1011}	0.39	GCEA	-4×10^{-8}	Howitz and McCarty (1986)
101a	GCEAc \rightarrow GCEA	K_{I1011}	0.28	GCA, competitive with GCEA	-2×10^{-11}	Howitz and McCarty (1986)
101b	GCA \rightarrow GCAC	K_{M1012}	0.2	GCA	-4×10^{-9}	Howitz and McCarty (1985)
101b	GCA \rightarrow GCAC	K_{I1012}	0.22	GCEA, competitive with GCA	-1×10^{-10}	Howitz and McCarty (1985)

1 ^{a b c d e}The description is same as in the Table C1.

1

2 **Table D2** The maximum rate of each enzyme (V_m) of the PCOP as given in Table 1, but here
 3 normalized on the maximum Rubisco carboxylation activity (V_1).
 4

Max. Velocity	Enzyme name	Reaction	V_m/V_1	Reference
V_{111}	Rubisco	$\text{RuBP} + \text{O}_2 \rightarrow \text{PGA} + \text{PGCA}$	0.24	Ueno et al (2005); Marek and Sspalding (1991); Ku et al (1991), Devi and Raghavendra (1993), Devi et al. (Devi et al., 1995)
V_{112}	Phosphoglycolate phosphatase	$2\text{-PGCA} + \text{H}_2\text{O} \rightarrow \text{GCA} + \text{Pi}$	18.0	Same as above
V_{113}	Glycerate kinase	$\text{GCEA} + \text{ATP} \rightarrow \text{PGA} + \text{ADP}$	1.96	Same as above
V_{121}	Glycolate oxidase	$\text{GCAc} + \text{O}_2 \rightarrow \text{H}_2\text{O}_2 + \text{GOAc}$	0.45	Same as above
V_{122}	Serine glyoxylate aminotransferase	$\text{GOAc} + \text{SERc} \rightarrow \text{HPRc} + \text{GLYc}$	1.13	Same as above
V_{123}	NADH- hydroxypyruvate reductase	$\text{HPRc} + \text{NADc} \rightarrow \text{NADHc} + \text{GCEAc}$	3.44	Same as above
V_{124}	Glutamate glyoxylate aminotransferase (GGAT)	$\text{GOAc} + \text{GLUc} \rightarrow \text{KGc} + \text{GLYc}$	0.94	Same as above
V_{131}	Glycine decarboxylase	$\text{GLYc} + \text{NADc} \rightarrow \text{CO}_2 + \text{NH}_3 + \text{SERc} + \text{NADHc}$	0.86	Same as above
V_{1T}	Glycerate/glycolate transporter	$\text{GCEAc} \leftrightarrow \text{GCEA}$	0.4	Howitz and McCarty (1986)
V_{2T}	Glycerate/glycolate transporter	$\text{GCAC} \leftrightarrow \text{GCA}$	0.4	Howitz and McCarty (1985)

5

6 ^a see table C2

1 **Table D3**

2 The concentrations of the metabolites of photorespiration (PCOP). The range of values found in
 3 the literature are given in the adjacent column, together with their source.

4

Metabolite	Locat-ion ^a	Concentration (mM)	Assumptions	Reference
NADH	Chl	0.22	This was kept constant in the preliminary model	Igamberdiev et al (2001)
NADHc	Cyto	0.47	As for NADH	Igamberdiev et al (2001)
NAD	Chl	0.08	As for NADH	Igamberdiev et al (2001)
NADc	Cyt	0.02	As for NADH	Igamberdiev et al (2001)
ATP	Chl	0.68		Igamberdiev et al (2001)
ATPc	Cyt	0.36	As for NADH	Igamberdiev et al (2001)
ADP	Chl	0.82		Igamberdiev et al (2001)
ADPc	Cyt	0.64	As for NADH	Igamberdiev et al (2001)
GLUC	Cyt	24	As for NADH	Winter et al (1993)
KGc	Cyt	0.4	As for NADH	Winter et al (1993)
Pic	Chl	5	As for NADH	Pieters et al (2001)
SERC	Cyt	7.5		Winter et al (1993)
GLYc	Cyt	1.8		Winter et al (1993)
PGA	Chl	4.3		Winter et al (1994)
GOAc	Cyt	0.028	30 µl cytosol per mg chlorophyll	Winter et al (1993), Wingler et al (1997)
GCA	Chl	0.36		Coombs and Whittingham (1966)
GCAc	Cyt	0.36		Coombs and Whittingham (1966)
PGCA	Chl	0.003	Based on the Michaelis-Menton equation for reaction 112 ^b	
HPRc	Cyt	0.004	Based on the Michaelis-Menton equation for reaction 123 ^b	
GCEA	Chl	0.18	Based on the Michaelis-Menton equation for reaction 113 ^b	
GCEAc	Cyt	0.18	Assume equilibrium with stromal Glycerate concentration	

5

6 Comment: ^a Chl: chloroplast stroma; Mit: Mitochondrion; Cyt: Cytosol;

1 **Appendix E**
 2 Kinetic parameters of enzymes in the metabolic pathway leading to sucrose synthesis in
 3 the cytosol.

4 **Table E1**
 5 The Michaelis-Menten constants and inhibition constants for enzymes in the metabolic pathway
 6 leading to sucrose synthesis in the cytosol.
 7

RN ^a	Reaction	Param ^b	Value (mmol l ⁻¹) ^d	Descri- ption ^c	CC ^e	Literature
51	DHAPc + PGAc ↔ FBPc	K _{m511}	.020	FBPc	4×10 ⁻⁶	Anderson et al. (1975)
51	DHAPc + PGAc ↔ FBPc	K _{m512}	.300	GAPc	-4×10 ⁻⁶	Iwaki et al. (1991)
51	DHAPc + PGAc ↔ FBPc	K _{m513}	.400	DHAPc	-4×10 ⁻⁶	Iwaki et al. (1991)
51	DHAPc + PGAc ↔ FBPc	K _{m514}	.014	SBPc	0	Harris and Koniger (1997)
51	DHAPc + PGAc ↔ FBPc	K _{E51}	12		2×10 ⁻⁶	Thomas et al. (1997)
52	FBPc ↔ F6Pc + Pic	K _{m521}	.0025	FBPc	-8×10 ⁻⁵	Jang et al. (2003)
52	FBPc ↔ F6Pc + Pic	K _{I521}	.7	F6Pc	4×10 ⁻⁵	Heldt et al. (1983)
52	FBPc ↔ F6Pc + Pic	K _{I522}	12	Pic	2×10 ⁻⁵	Charles & Halliwell (1981)
52	FBPc ↔ F6Pc + Pic	K _{I523}	7×10 ⁻⁵	F26BPc	8×10 ⁻⁵	Jang et al. (2003)
52	FBPc ↔ F6Pc + Pic	K _{E52}	6663		4×10 ⁻⁸	Bassham and Krause (1969)
55	G1Pc + UTPc ↔ GDPc + UDPGc	K _{m551}	.14	G1Pc	-1×10 ⁻⁷	Nakano et al. (1989)
55	G1Pc + UTPc ↔ GDPc + UDPGc	K _{m552}	.1	UTPc	-1×10 ⁻⁷	Nakano et al. (1989)
55	G1Pc + UTPc ↔ GDPc + UDPGc	K _{m553}	.11	OPOPc	6×10 ⁻⁸	Nakano et al. (1989)
55	G1Pc + UTPc ↔ GDPc + UDPGc	K _{m554}	.12	UDPGc	7×10 ⁻⁸	Nakano et al. (1989)
55	G1Pc + UTPc ↔ GDPc + UDPGc	K _{E55}	0.31	Equi	1×10 ⁻⁴	Lunn and Rees (1990)
56	UDPGc + F6Pc ↔ SUCPc + UDPc	K _{m561}	0.8	F6Pc	-8×10 ⁻⁵	Lunn and Rees (1990)
56	UDPGc + F6Pc ↔ SUCPc + UDPc	K _{m562}	2.4	UDPGc	-1×10 ⁻⁴	Lunn and Rees (1990)
56	UDPGc + F6Pc ↔ SUCPc + UDPc	K _{I561}	.7	UDPc	1×10 ⁻⁴	Harbron et al. (1981)
56	UDPGc + F6Pc ↔ SUCPc + UDPc	K _{I562}	.8	FBPc	5×10 ⁻⁵	Harbron et al. (1981)
56	UDPGc + F6Pc ↔ SUCPc + UDPc	K _{I563}	0.4	SUCPc	3×10 ⁻⁵	Harbron et al. (1981)
56	UDPGc + F6Pc ↔ SUCPc + UDPc	K _{I564}	11	Pic	3×10 ⁻⁵	Harbron et al. (1981)
56	UDPGc + F6Pc ↔ SUCPc + UDPc	K _{I565}	50	Sucrose	2×10 ⁻⁷	Salermo and Pontis (1978)
56	UDPGc + F6Pc ↔ SUCPc + UDPc	K _{E56}	10	Equl.	4×10 ⁻⁶	Lunn and Rees (1990)

	UDPc		Const.			
57	SUCPc \leftrightarrow Pic + SUCc	K _{m571}	.35	SUCPc	-2×10 ⁻⁵	Whitaker (1984)
57	SUCPc \leftrightarrow Pic + SUCc	K _{i572}	80	SUCc	3×10 ⁻⁸	Whitaker (1984)
57	SUCPc \leftrightarrow Pic + SUCc	K _{E57}	780	Equil.	3×10 ⁻⁷	Lunn and Rees (1990)
			Const.			
58	F26BPc \leftrightarrow F6Pc + Pic	K _{m581}	.032	F26BPc	-9×10 ⁻⁶	Macdonald et al. (1989)
58	F26BPc \leftrightarrow F6Pc + Pic	K _{i581}	.1	F6Pc	8×10 ⁻⁵	Villadsen and Nielsen (2001)
58	F26BPc \leftrightarrow F6Pc + Pic	K _{i582}	.5	Pic	8×10 ⁻⁵	Villadsen and Nielsen (2001)
59	F6Pc + ATPc \leftrightarrow F26BPc + ADPc	K _{m591}	0.5	ATPc	8×10 ⁻⁵	Walker and Huber (Walker and Huber, 1987)
59	F6Pc + ATPc \leftrightarrow F26BPc + ADPc	K _{m592}	.021	F26BPc	-7×10 ⁻⁴	Garcia de Frutos and Baanante (1995)
59	F6Pc + ATPc \leftrightarrow F26BPc + ADPc	K _{m593}	0.5	F6Pc	8×10 ⁻⁵	Walker and Huber (1987)
59	F6Pc + ATPc \leftrightarrow F26BPc + ADPc	K _{i591}	.16	ADPc	-6×10 ⁻⁵	Kretschmer and Hofmann (1984)
59	F6Pc + ATPc \leftrightarrow F26BPc + ADPc	K _{i592}	0.7	DHAPc	-1×10 ⁻⁵	Markham and Kruger (2002)
59	F6Pc + ATPc \leftrightarrow F26BPc + ADPc	K _{E59}	590		-1×10 ⁻⁷	Cornish-Bowden (1997)
60	F6Pc + ATPc \leftrightarrow F26BPc + ADPc	K _{m601}	0.042	ADPc	NA	Kimura and Shimada (1988)
60	F6Pc + ATPc \leftrightarrow F26BPc + ADPc	K _{m602}	1.66	ATPc	NA	Kimura and Shimada (1988)
60	F6Pc + ATPc \leftrightarrow F26BPc + ADPc	K _{m603}	0.28	UDPc	NA	Jong and Ma (1991)
60	F6Pc + ATPc \leftrightarrow F26BPc + ADPc	K _{m604}	16	UTPc	NA	Fukuchi et al (1994)
60	F6Pc + ATPc \leftrightarrow F26BPc + ADPc	K _{E60}	16	Equili.	NA	Lynn and Guynn (1978)
61	SUCPc \leftrightarrow SUCc + Pic	K _{E61}	1.2*10 ⁷	Equili.	-4×10 ⁻³	Flodgaard and Fleron (1974)
62	SUCc \leftrightarrow Sink	K _{m621}	5	Sucrose	-5×10 ⁻⁷	Weschke et al. (2000)

1 ^{a b c d e} The description is same as in the Table C1.

1 **Table E2**

2 The maximum rate of each enzyme (V_m) as given in Table 1 for enzymes in the metabolic
 3 pathway leading to sucrose synthesis, but here normalized on the maximum Rubisco
 4 carboxylation activity (V_1).
 5

Maximum Velocity	Reaction	V_m/V_1	Reference
V_{51}	DHAPc + PGAc \leftrightarrow FBPc	0.037	Chen et al (2005), Strand et al. (2000), Strand et al (1999), Chen et al. (2005)
V_{52}	FBPc \leftrightarrow F6Pc + Pic	0.022	Same as above
V_{55}	G1Pc + UTPc \leftrightarrow GDPc + UDPGc	0.040	Same as above
V_{56}	UDPGc + F6Pc \leftrightarrow SUCPc + UDPc	0.019	Same as above
V_{57}	SUCPc \leftrightarrow Pic + SUCC	0.19	Same as above
V_{58}	F26BPc \leftrightarrow F6Pc + Pic	0.007	Same as above
V_{59}	F6Pc + ATPc \leftrightarrow F26BPc + ADPc	0.002	Villadsen and Nielsen (2001)
V_{60}	ATPc + UDPc \leftrightarrow UTPc + ADPc	1	Villadsen and Nielsen (2001)

6 ^a see Table C2

1 **Table E3**

2 The concentrations of metabolites in the metabolic pathway leading to sucrose synthesis in the
3 cytosol.

4

Metabolite	Locat- ion ^a	Concentration (mmol l ⁻¹)	Reference
TPc	Cyt	2.3	Stitt et al. (1980), Stitt et al. (1985), Gerhardt et al. (1987), Laisk et al. (1989)
FBPc	Cyt	2	As above
F26BPc	Cyt	7×10 ⁻⁶	As above
UTc	Cyt	1	As above
HexPc	Cyt	6	As above
UDPG	Cyt	0.6	As above
PTc	Cyt	15	As above
ATc	Cyt	1	As above

5

6 Comment: ^a Chl: chloroplast stroma; Mit: Mitochondrion; Cyt: Cytosol;

1

2 Appendix F

3 Molecular weight and catalytic number of the enzymes in photosynthetic carbon metabolism

4

Enzyme Name	EC	Molecular Weight (D)	Catalytic number ¹ (s ⁻¹)	Reference
Rubisco	4.1.1.39	588000	2*	Spreitzer and Salvucci (2002)
PGA Kinase	2.7.2.3	45000	540	Fifis and Scopes (1978), Bentahir et al (2000)
GAP Dehydrogenase	1.2.1.12	180000	50	Speranza and Ferri (1982)
Aldolase	4.1.2.13	143000	65	Krueger and Schnarrenberger (1983), Moorhead and Pplaxton (1990)
FBPase	3.1.3.11	160000	22.9	Tang et al. (2000), Reichert et al.(2000)
Transketolase	2.2.1.1	160000	69	Nilsson et al (1998)
SBPase	3.1.3.37	66000	81	Teige et al. (1989)
PRK	2.7.1.19	90000	615	Cadet et al. (1988), Cadet and Meunier (1987)
ADPG Pyrophosphorylase	2.7.7.27	210000	546	Surek et al. (1985), Porter et al. (1986)
Phosphoglycolate phosphatase	3.1.3.18	100000	292	Kleczkowski et al. (1991), Li and Preiss(1992)
Glycerate Kinase	2.7.1.31	47000	200	Kim et al. (2004), Kerr and Gear (1974)
Glycolalate oxidase	1.1.1.79	125000	437	Kleczkowski et al. (1985), Kleczkowski and Randall (1988)
Serine Glyoxylate aminotransferase	2.6.1.45	85000	97	Kleczkowski et al. (1986), Zelitch(1955)
Glycerate dehydrogenase	1.1.1.29	90000	1629	Ireland and Joy (1983; Paszkowski and Niedzielska, 1990)
Glutamate Glyoxylate aminotransferase	2.6.1.44	70800	54	Julliard and Breton-Gilet (1997), Izumi et al.(1990)
Glycine decarboxylase	1.4.4.2	270000	18	Paszkowski and Niedzielska (1989)
6-phosphofructo-2-kinase	2.7.1.105	390000	9300	Hiraga and Kikuchi (1980), Kochi and Kikuchi (1974)
fructose-2,6-bisphosphate 2-phosphatase	3.1.3.46	390000	1550	Villadsen and Nielsen (2001), Baez et al. (2003)
UDP Glucose pyrophosphorylase	2.7.7.9	53000	400	Pilkis et al. (1987), Villadsen and Nielsen (2001), Gustafson and Gander (1972), Sowokinos et al (1993)
Sucrose phosphate synthase	2.4.1.14	480000	640	Sonnewald et al. (1993)
Sucrose phosphatase	3.1.3.24	120000	2500	Echeverria and Salerno (1994), Lunn et al. (2000)

5

1 Appendix G: List of Abbreviations and their units

2

Abbreviation ^a	Full Name	Units
A	The photosynthetic CO ₂ uptake rate	μmol m ⁻² s ⁻¹
ADPG	ADP-glucose	mmol l ⁻¹
ADPGPP	ADP glucose pyrophosphorylase	NA
ATc	Total ADP and ATP concentration in cytosol	mmol l ⁻¹
ATPase	ATP synthase	NA ^b
[CO ₂]	CO ₂ concentration	μmol mol ⁻¹ or mmol l ⁻¹
CA	Total adenylate nucleotide in the chloroplast stroma including ATP and ADP	mmol l ⁻¹
CN	Total of NADP ⁺ and NADPH in chloroplast stroma	mmol l ⁻¹
CP	The total concentration of phosphate in chloroplast stroma	mmol l ⁻¹
DHAP	Dihydroxyacetone-phosphate	mmol l ⁻¹
DPGA	1,3-bisphosphoglycerate	mmol l ⁻¹
E4P	Erythrose 4-phosphate	mmol l ⁻¹
E _t	Total Rubisco concentration	mmol l ⁻¹
F6P	Fructose 6-phosphate	mmol l ⁻¹
FBP	Fructose 1,6-bisphosphate	mmol l ⁻¹
F26BP	Fructose 2,6-bisphosphate	mmol l ⁻¹
G1P	Glucose 1-phosphate	mmol l ⁻¹
G6P	Glucose 6-phosphate	mmol l ⁻¹
GAP	Glyceraldehyde 3-phosphate	mmol l ⁻¹
GAPDH	Glyceraldehyde 3-phosphate dehydrogenase	NA
GCA	Glycollate	mmol l ⁻¹
GCEA	Glycerate	mmol l ⁻¹
GDC	Glycine decarboxylase	NA
GLUc	Glutamate	mmol l ⁻¹
GLYc	Glycine in cytosol	mmol l ⁻¹
GOA	Glyoxylate	mmol l ⁻¹
HexP	Hexose phosphate, includes F6P, G6P, and G1P	NA
GGAT	Glycine glyoxylate aminotransferase	NA
GSAT	Glyoxylate serine aminotransferase	NA
HPR	Hydroxypyruvate	mmol l ⁻¹
KGc	α-Ketoglutarate	mmol l ⁻¹
ODE	Ordinary differential equation	NA
[O ₂]	Oxygen concentration in atmosphere	mmol mol ⁻¹
OPOP	Pyrophosphate	mmol l ⁻¹
PCOP	Photosynthetic carbon oxidation pathway	NA
PCRC	Photosynthetic carbon reduction cycle	NA
PenP	Pentose phosphate including Ri5P, Ru5P, Xu5P	mmol l ⁻¹
3-PGA	3-Phosphoglycerate	mmol l ⁻¹
PGCA	3-Phosphoglycolate	mmol l ⁻¹
PRK	Ribulose-5-phosphate kinase	NA
PGCA Pase	Phosphoglycolate phosphatase	NA

PGA Kinase	3-phosphoglycerate kinase	NA
Ri5P	Ribose 5-phosphate	mmol l ⁻¹
PRK	Phosphoribulose kinase	mmol l ⁻¹
PTc	Total phosphate concentration in cytosol	mmol l ⁻¹
Ru5P	Ribulose 5-phosphate	mmol l ⁻¹
Rubisco	Ribulose1,5-bisphosphate Carboxylase/Oxygenase	NA
R _t	Total RuBP concentration in stroma	mmol l ⁻¹
RuBP	Ribulose 1,5-biphosphate	mmol l ⁻¹
S7P	Sedoheptulose 7-phosphate	mmol l ⁻¹
SBP	Sedoheptulose 1,7-bisphosphate	mmol l ⁻¹
SBPase	Sedoheptulosebisphosphatase	NA
SERc	Serine in cytosol	mmol l ⁻¹
SPP	Sucrose phosphate phosphatase	NA
SPS	Sucrose phosphate synthetase	NA
TPU	Triose phosphate utilization	NA
SUCc	Sucrose in cytosol	mmol l ⁻¹
SUCPc	Sucrose phosphate in cytosol	mmol l ⁻¹
T3P	Triose phosphate including DHAP and GAP	mmol l ⁻¹
V ₁₁₁	The rate of RuBP oxygenation	μmol m ⁻² s ⁻¹ or mmol l ⁻¹ s ⁻¹
UDPGc	Uridine Diphosphate Glucose	mmol l ⁻¹
UDPGP	UDP glucose pyrophosphorylase	NA
UT	Total UDP and UTP concentration in cytosol	mmol l ⁻¹
V ₁	The rate of RuBP carboxylation	μmol m ⁻² s ⁻¹ or mmol l ⁻¹ s ⁻¹
V _{atpg}	The rate of ATP formation in cytosol	μmol m ⁻² s ⁻¹ or mmol l ⁻¹ s ⁻¹
V _{pga_use}	The rate of PGA utilization in cytosol	μmol m ⁻² s ⁻¹ or mmol l ⁻¹ s ⁻¹
Xu5P	Xylulose 5-phosphate	mmol l ⁻¹

1 ^aA suffix c was added to the name of metabolites appeared in cytosol if the metabolite also exists in stroma. For example, PGA is the
 2 phosphoglycerate in stroma; while PGAc is the phosphoglycerate in cytosol.

3 ^bNA: Not applicable

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