

TABLE 3 CO<sub>2</sub> produced by various metabolic reactions and refixed by RuBisCO<sup>a</sup>

Substrate	% of substrate converted to CO <sub>2</sub> (relative to amt of substrate consumed)		% of CO <sub>2</sub> refixed by Calvin cycle (relative to amt of substrate converted to CO <sub>2</sub> )		Net CO <sub>2</sub> yield (% relative to amt of substrate consumed)	
	WT	NifA*	WT	NifA*	WT	NifA*
Fumarate <sup>b</sup>	40 ± 4	44 ± 4	21 ± 9	6 ± 1	32 ± 2	42 ± 2
Succinate	37 ± 3	40 ± 2	49 ± 7	30 ± 5	19 ± 2	28 ± 2
Acetate <sup>c</sup>	22 ± 2	23 ± 1	68 ± 11	13 ± 3	6 ± 1	18 ± 1
Butyrate-HCO <sub>3</sub> <sup>-</sup>	16 ± 1	15 ± 3	180 ± 16 <sup>e</sup>	149 ± 36 <sup>e</sup>	-16 ± 1 <sup>f</sup>	-10 ± 3 <sup>f</sup>
Butyrate <sup>d</sup>		23 ± 3		76 ± 17		6 ± 1

<sup>a</sup> Average values with 90% confidence intervals were derived from the fluxes shown in Fig. 1. Minor variations between CO<sub>2</sub> yields in Tables 2 and 3 are due to changes made by the fitting algorithm to find the most likely set of fluxes to explain all of the data.

<sup>b</sup> All values were calculated by grouping malate and fumarate as a single pool. This grouping results in different CO<sub>2</sub> yields between Tables 2 and 3, because the CO<sub>2</sub> yields in Table 2 were normalized to fumarate alone so that the amount of malate produced could also be reported. If fumarate and malate were grouped in Table 2, the CO<sub>2</sub> yields would be the same as those reported in Table 3.

<sup>c</sup> The acetate data were previously published (7).

<sup>d</sup> Wild-type cells do not grow without the NaHCO<sub>3</sub> supplement.

<sup>e</sup> One hundred percent of the butyrate converted to CO<sub>2</sub> was refixed along with CO<sub>2</sub> from the NaHCO<sub>3</sub> supplement.

<sup>f</sup> The negative values indicate that there was a net uptake of CO<sub>2</sub> from the NaHCO<sub>3</sub>.