Properties of Formate Dehydrogenase in *Methanobacterium formicicum*

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Volume 150, no. 1, p. 1, abstract, lines 8 and 9: "(K_m = 13 μM)" should read "(K_m = 44 μM)," and "(K_m = 25 μM)" should read "(K_m = 98 μM)."

Page 2, column 2, lines 24 and 25 from the bottom: "FMN, ε_{450} = 41 mM^{-1} cm^{-1}" should read "FMN, ε_{450} = 12.2 mM^{-1} cm^{-1}," and "FAD, ε_{450} = 44.3 mM^{-1} cm^{-1}" should read "FAD, ε_{450} = 11.3 mM^{-1} cm^{-1}."

Page 4, column 1, line 12: "were 2-, 4-, and 10-fold" should read "were 7-, 16-, and 10-fold."

Page 6: Table 3 should read as follows:

TABLE 3. Kinetic constants of electron acceptors for purified FDH from *M. formicicum*[a]

Acceptor	Sp act (μmol min^{-1} mg^{-1})[b]	K_m (μM)[c]	V_{max} (μmol min^{-1} mg^{-1})[c]
F_{420}	8.2	6	17.4
FO	1.6	63	18.8
FAD		98	26.6
FMN		44	19.5

[a] Assays were performed as described in the text. Reaction mixtures contained 4 μg of protein. The initial rate of reduction with each acceptor was linear, and double-reciprocal plots of initial velocity versus acceptor concentration were also linear, with correlation coefficients of 0.98 or greater.

[b] The concentration of each acceptor was 5 μM.

[c] The ranges of concentrations used were: F_{420}, 5 to 20 μM; FO, 5 to 50 μM; FAD, 20 to 200 μM; and FMN, 20 to 100 μM.