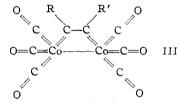
II (R = R' = CH₂OH), light-orange-red needles from ethanol-water, m.p. *ca.* 135° with decomposition. Calcd. for $C_{10}H_{\theta}O_{8}Co_{2}$: C, 32.28; H, 1.63; Co, 31.69. Found: C, 32.24; H, 1.67; Co, 31.67.

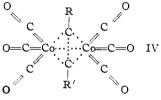
II (R = R' = H), a dark-red oil at room temperature; m.p. $13.0-13.6^{\circ}$; b.p. $64-66^{\circ}$ (3.5-4mm.) Calcd. for C₈H₂O₆Co₂: C, 30.80; H, 0.65. Found: C, 30.76; H, 0.62.

The infrared spectra of these compounds contain a characteristic group of three sharp bands at 2090, 2050 and 2025 cm.⁻¹, which are similar to the bands of the terminal carbonyl groups in dicobalt octacarbonyl. The 1859 cm.⁻¹ band corresponding to the bridge carbonyl group in I as well as the bands of the $-C \equiv C-$ group, are absent. The spectra of II (R = R' = H), II (R = CH₃(CH₂)₄, R' = H) and II (R = CH₂OH, R' = H) contain a band at 3096 cm.⁻¹, characteristic of an ethylenic (or aromatic) carbon-hydrogen bond.

It is reasonable to assume that the C–C bond is either parallel to the Co–Co bond, such as in III



or perpendicular to the Co-Co bond, such as in IV.



The analytical, spectroscopic, magnetic and dipole moment data, in conjunction with the high volatility and the solubility in organic solvents of II, are compatible with a structure such as III, where cobalt possesses a noble-gas configuration.

In IV the C-C Co-Co bonds may be either in the same plane (IVa) or in different planes (IVb). Structure IVa is very unlikely³ in view of the dipole moment of II ($R = R' = C_{6}H_{5}$). Neither the structure of IVa nor that of IVb can be represented readily in terms of localized bonds.

(3) Compound II ($\mathbf{R} = \mathbf{R}' = \mathbf{C}_{6}\mathbf{H}_{5}$) has a molar polarization of 204 ± 4 at 30° and a calculated molecular fraction for the D sodium line of 121. For II ($\mathbf{R} = \mathbf{R}' = \mathbf{C}_{6}\mathbf{H}_{5}$) to be symmetrical it would have to have an extremely high atomic polarization (cf. I. E. Coop and L. E. Sutton, J. Chem. Soc., 1269 (1938)).

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A FORMYLATION COFACTOR¹

Sir:

This paper reports the biosynthesis of a formylation cofactor and the transfer of its one-carbon

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group to a purine precursor. Formylation of IRMP- $5^{2,3,4}$ in pigeon liver extract may occur as

$$RH_4X + HCOOH + ATP \longrightarrow FRH_4X + ? (1)$$

 $FRH_4X + IRMP-5 \longrightarrow RH_4X + IMP-5$ (2) The over-all reaction exhibits almost an absolute requirement for IRMP-5, HCOOH, ATP, Mg^{++} and a cofactor. FRH₄X is a heat-stable cofactor obtained enzymatically from FRH4 or from $RH_{4^{5}}$ and formate by ATP-dependent reactions. The following three experiments provide evidence for the above mechanism: (1) C^{14} -(formyl)-FRH₄X transformylated directly to IRMP-5 (Reaction 2) in a large pool of unlabeled formate to yield IMP-5 with a specific activity more than 10 times that of the formate. The C^{14} -activity of the FRH₄X was displaced by the unlabeled formate in the presence of ATP (Reaction 1) while without acceptor IRMP-5, FRH₄X lost no activity. (2) Purified FRH4X formylated IRMP-5 in the absence of ATP, but FRH_4 alone was inactive. (3) An excess of FRH₄X obtained from FRH₄ plus ATP converted IRMP-5 completely to IMP-5 in the absence of formate. The resulting RH₄X accepted added C14-formate to yield C14-FRH4X (paper chromatography) equivalent in quantity to 80% of the initial IRMP-5 (Reaction 1). Exchange between HC¹⁴OOH and IMP-5⁶ was insignificant. Loss of X during transformylation (Reaction 2) is not excluded.

Table I shows that the synthesis of FRH₄X, which is measured by its cofactor effect on IMP-5 synthesis, depends on the presence of both ATP and FRH₄ during preincubation.

TABLE I

INTERACTION OF LEUCOVORIN AND ATP

Preincubation additions: 20 mg. lyophilized extract (acetone powder extracted with 0.05 M KHCO₃ and treated with Dowex-1 chloride), 2.5 μ M·ATP, 7 μ M, phosphoglycerate, 3 μ M·MgCl₂, 0.5 mg. lyophilized muscle extract fraction,⁷ 7.5 μ M·KHCO₃, 0.2 μ M·FRH₄, 8 μ H·DL-homocysteine; vol., 0.6 ml.; time, 10 min.; temp., 38°; in air. After preincubation 0.36 μ M·IRMP-5, 2 μ M. Cl⁴-formate and the omitted compounds were added and the reaction continued 5 min.; total vol. 1.0 ml.

Omitted in prein-

cubation	None	FRH_4	ATP	FRH_4^a	ATP^a
Δ Amine, μ M	0.262	0.084	0.057	0.024	0.002
C ¹⁴ fixed, µM	.294	.116	.093	. 010	.0

^a Also omitted after preincubation.

IMP-5 synthesis is greatest when FRH₄X is preformed. One mole of diazotizable amine⁸ disappears per mole of C^{14} formate fixed into IMP-5. On paper chromatograms FRH₄X appears as a blue-fluorescing compound.

(2) Abbreviations: IRMP-5, 5-amino-4-imidazolecarboxamide-5'-phosphoribotide; ATP, adenosine triphosphate; IMP-5, inosine-5'-phosphate; RH, 7,8-dihydrofolic acid; RH, 5,6,7,8-tetrahydrofolic acid; FRH, N-5-formyl-5,6,7,8-tetrahydrofolic acid (leucovorin); X, product of the interaction of a folic acid compound and ATP DPNH, reduced diphosphopyridine nucleotide.

(3) G. R. Greenberg, Federation Proc., 12, 211 (1953).

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(6) J. M. Buchanan and M. P. Schulman, J. Biol. Chem., 202, 241 (1953).

(7) S. Ratner and A. Pappas, *ibid.*, **179**, 1183 (1949).

(8) J. M. Ravel, et al., *ibid.*, **172**, 67 (1948). Diazotization was preceded by acetic anhydride treatment.

The conversion of RH₄ to FRH₄X occurs under conditions such as shown in Table I at a rate comparable to IMP-5 synthesis. Preincubation studies without formate suggest that RH4 and ATP first react to form RH_4X . RH_2^{5} and DPNH replace RH4 in FRH4X synthesis. Folic acid did not react under these conditions. These experiments implicate RH₂ and RH₄ as intermediates in the synthesis of FRH₄ from folic acid.

C¹⁴FRH₄X is converted to N-10-C¹⁴-formylfolic acid during isolation and by dilute acid.9

Leucovorin is known to catalyze the exchange between HC14OOH and IMP-5.6 RH4,10 ATP and DPN¹¹ have been involved in FRH₄ synthesis. Rauen and Jaenicke¹² have reported a cofactor derived from folic acid derivatives. The expected interrelationship of FRH4X with other 1-carbon acceptor systems¹³ has been discussed previously.¹⁴

(9) Compare M. Silverman and J. C. Keresztesy, Federation Proc. 12, 268 (1953). The author gratefully acknowledges the aid of Drs. Silverman and Keresztesy in identifying N-10-formylfolic acid.

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(14) Dr. E. L. R. Stokstad, Lederle Laboratories, kindly supplied the calcium leucovorin and part of the dihydrofolic acid employed.

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LOSS OF THE α -AMINO GROUP IN LYSINE METAB-OLISM TO FORM PIPECOLIC ACID

Sir:

The early steps of the metabolism of lysine have been the subject of speculation for many years. It has recently been found in this laboratory that pipecolic acid is a metabolite of L-lysine- ϵ -C¹⁴ in the rat.¹ Pipecolic acid must be formed at an early stage of lysine metabolism since it still contains six carbon atoms but has only one amino group. This fact, along with the in vivo "metabolic overloading" technique devised to isolate specific metabolites of isotopic precursors^{1,2} has afforded a means of ascertaining which amino group of lysine is removed first. This was determined by injecting intraperitoneally into a 24-hour fasted rat a solution containing 74 mg. of DL-lysine-e-N¹⁵·HCl $(25.3 \ atom \ \% \ excess \ N^{15})$ and 360 mg. of non-isotopic L-pipecolic acid.³ The effective dose of lysine- ϵ - N^{15} -HCl is 37 mg. since D-lysine, under similar conditions, does not contribute significantly

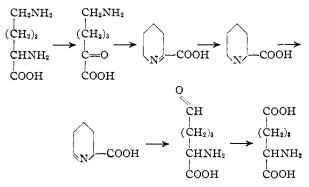
(1) M. Rothstein and L. L. Miller, THIS JOURNAL, 75, 4371 (1953).

M. Rothstein and L. L. Miller, J. Biol. Chem., January, 1954.
 The authors wish to thank Dr. F. C. Steward of Cornell Uni-

versity for supplying the L-pipecolic acid used in this research.

to the formation of pipecolic acid.⁴ After isolation from the urine as previously reported¹, 140 mg. of pure pipecolic acid was isolated and found to contain 2.0 atom % excess N^{15.5} This large enrichment of N¹⁵ in the pipecolic acid is remarkable in view of the large dilution of biologically formed material with non-enriched pipecolic acid. This is in accord with the concept established with C¹⁴, namely, that the formation of pipecolic acid is both a major and a primary step in lysine metabolism. Furthermore, this is conclusive evidence that the ϵ -amino group of lysine remains in large measure intact until after the loss of the α -amino group, and lends support to the hypothesis that lysine forms an α -keto analog. Work with Neurospora crassa is in accord with this.6 The possibility of concomitant formation of pipecolic acid by loss of the ϵ -amino group of lysine and subsequent cyclization is at present being investigated with lysine- α -N¹⁵.

If, as seems probable, the conversion of lysine to pipecolic acid is part of the pathway between lysine and α -aminoadipic acid, the most likely mechanism is



This is a pathway whereby the ϵ -amino group could be oxidized by what amounts to an intramolecular transamination reaction. It is of interest to note that α -aminoadipic acid- ϵ -C¹⁴ does not lead to radioactive pipecolic acid under conditions where lysine- ϵ -C¹⁴ with a radioactive count of similar magnitude leads to pipecolic acid containing 1.2 \times 10⁵ disintegrations/min./mmole, indicating the irreversibility of the pathway.

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(4) M. Rothstein, C. G. Bly and L. L. Miller, Arch. Biochem. and Biophys., in press

(5) We are indebted to Glenn Happ of the Department of Analytical Chemistry, Eastman Kodak Co., Rochester, N. Y., for performing the N¹⁵ assay

(6) P. H. Lowy, J. T. Holden and R. S. Schweet, Abstracts, Atlantic City Meeting, A. C. S., 1952, p. 44c.