ulation J (τ = 538 fs) simultaneously with K (τ = 656 fs) (Scheme II). Relatively slow decay of J back to bR results in the apparent generation of a more blue-shifted intermediate ($\Delta t = 10 \text{ ps}$, Figure 1g). Time resolved resonance Raman spectroscopy has the potential of detecting J and testing the validity of our model. Furthermore, the transition from $J(S_1)$ to the ground state is potentially observable by using time resolved fluorescence spectroscopy. We predict this transition to be very broad ($\Delta \tilde{\nu} \approx \sim 6000$ cm⁻¹) with a Franck-Condon maximum at $\sim 1.6 \mu$.

The Biosynthesis of LL-C10037 α from the Shikimate Pathway

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LL-C10037 α , an antitumor metabolite of Streptomyces LL-C10037 first assigned structure 1,² has recently been shown to have the corrected structure 2.3 Several related structures have been isolated from a wide variety of microorganisms. For ex-



ample, sarubicin A, 3, contains a quinone moiety which we have shown to be derived from the shikimate metabolite 6-hydroxyanthranilic acid, 4.4 In addition, the hydroxyazaquinone 5^5 is derived from shikimic acid 6 via p-aminobenzoic acid (PABA),^{5,6} while chaloxone 7,7 isolated with its cometabolite methyl anhy-

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Cultures of S. LL-C10037 were grown in shake flasks (200 mL of broth in a 1-L Erlenmeyer) at 28 °C as previously described.² After incubation for 122 h, the broth was obtained free of the mycelia, adjusted to pH 4.7 (KH_2PO_4), and saturated with $(NH_4)_2SO_4$. Extraction with ethyl acetate, concentration in vacuo, and column chromatography (20 g silica gel 60, eluted with 20% hexanes in ethyl acetate) typically gave 30 mg of crystalline LL-C10037 α which was then recrystallized from methanol.

For feeding experiments cultures were usually grown for 96 h before potential precursors were added in a sterile manner through a Millipore filter unit, and the broth was worked up after a total of 122 h. A mixture of sodium $[1^{-14}C]$ acetate (9 μ Ci), 11a, sodium acetate (23.6 mg), 11, and sodium [2-13C]acetate



(68.3 mg), 11b,¹² was fed first, and workup yielded 60 mg of 2a (1.3% incorporation of ¹⁴C). The 100.6-MHz ¹³C NMR spectrum¹³ of **2a** revealed enrichment (6.4%) only in the acetamide methyl group, revealing that the carbocycle is not polyketide in origin.

Feeding¹⁴ a mixture of $[1-1^4C]$ -D-glucose, **12a**, (5.5 μ Ci) and [1-13C]-D-glucose (0.996 g), 12b, 15 yielded 35 mg of 2b. The 13C NMR spectrum of enriched 2b showed labeling at C-2 (5.94%) and C-4 (7.02%), clearly indicative of a shikimate-type pathway;¹⁶ the acetamide methyl was also enriched (4.69%), due to the incorporation of glycolysis-derived acetylCoA. The ring labeling pattern is inconsistent with the intermediacy of PABA (C-2 and C-6 would have been labeled in this case).¹⁷

To determine the correct orientation of the apparent shikimic

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- (13) Spectrum taken in 0.4 mL MeOH- d_4 with a Bruker AM 400 spectrometer at 100.6 MHz (sweep width = 2500 Hz, 64K data points, pulse width = 32.5°, acquisition time = 1.311 s, no. scans = 2839).
- (14) In this experiment the production broth was harvested 193 h after inoculation. (15) [1-¹³C]-D-Glucose (99 atom % C-13) was obtained from Omicron
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acid precursor relative to 2, a mixture of $[1-^{14}C]$ -p-ervthrose (5.5 μ Ci), 13a, and [1-¹³C]-D-erythrose (336 mg), 13b,¹⁸ was administered and afforded 8 mg of 2c. From the ¹³C NMR spectrum it was clear that 2c was only labeled at C-1 (5.13%), rather than at C-5. Thus, the nitrogen substituent of 2 had been introduced at C-6 of the shikimate intermediate, rather than at C-2. This is consistent with the involvement of either 4, 10,¹⁹ or anthranilic acid, 14.20

S. LL-C10037 was next fermented (2×200 mL broths) in the presence of ¹⁸O molecular oxygen,²¹ yielding 10 mg of **2d**. The ¹³C NMR spectrum exhibited ¹⁸O-induced isotope shifts²² for C-4 (2.0 Hz), C-5 (3.5 Hz), and C-6 (3.0 Hz), revealing that the secondary alcohol and epoxide oxygens had been introduced by oxidative processes rather than having been retained from the organic precursor. The origin of the C-4 hydroxyl from molecular oxygen ruled out 4 as an intermediate.

[5-2H]-3-Hydroxyanthranilic acid 15²³ was then fed (200 mg to 200 mL of broth) yielding²⁴ 5.0 mg of 2e that was analyzed by ²H NMR.²⁵ A 67% enrichment of the signal at δ 3.8 (H-5) was observed, representing a 2.2% incorporation of 15 (50 mg was recovered).

The biosynthesis of 2 represents a novel aromatic amino acid metabolism. In view of our work, metabolites such as paulomycin A,²⁶ the enaminomycins,²⁷ MM14201,²⁸ and G7063-2²⁹ are most likely also derived from the shikimate pathway. We are currently investigating the biosynthesis of the latter two.

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(18) [1-13C]-D-Erythrose (99 atom % C-13) was obtained from Omicron Chemicals, Ithaca, NY

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Soc., Chem. Commun. 1980, 183 (23) Obtained by exchange (86%) with deuteriotrifluoroacetic acid at 80 °C in a sealed tube for 48 h. In the ¹H NMR, the dd at δ 7.44 was nearly gone and the signals at δ 7.33 and 7.82, previously doublets, were now singlets.

(24) The fermentation was harvested 135 h after innoculation.

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Mechanism-Based Isocoumarin Inhibitors for **Trypsin-like Serine Proteases Involved in Blood** Coagulation[†]

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Human plasma contains a number of proteins (zymogens) which are precursors of serine proteases with trypsin-like specificity. The

[†] Dedicated to Professor George Büchi on the occasion of his 65th birthday.



Figure 1. Proposed mechanism of inactivation of serine proteases by substituted isocoumarins.

interaction of activated serine proteases with these zymogens and with natural plasma protease inhibitors in a cascade of enzymatic reactions forms the basis of the blood coagulation pathway. Although intravascular clotting is a major health problem in the United States, almost no new anticoagulant drugs have been developed in recent years. A number of heterocyclic compounds have previously been shown to be suicide substrates of serine proteases.¹⁻⁶ Substituted isocoumarins have been reported to be mechanism-based inhibitors for elastases and a variety of chymotrypsin-like enzymes.^{4,5} Here we report the synthesis of three new isocourmarins with basic functional groups (guanidino or isothiureidoethoxy) attached to the isocoumarin ring system. These compounds are potent inhibitors of coagulation enzymes and are the first effective mechanism-based inhibitors of clotting in human plasma.

Incubation of 4-chloro-3-(2-isothiureidoethoxy)isocoumarin (1), 7-guanidino-3-methoxyisocoumarin (2), and 4-chloro-7guanidino-3-methoxyisocoumarin $(3)^7$ with several coagulation enzymes and trypsin results in time-dependent loss of enzymatic activity (Table I). Human protein D, human leukocyte elastase, human leukocyte cathepsin G, porcine pancreatic elastase, chymotrypsin, and human plasma plasmin were also inactivated by compound 3 with k_{obsd} [I] values of 120, 6400, 11000, 860, 7200, and $3600 \text{ M}^{-1}\text{s}^{-1}$, respectively. Compound 3 was quite selective among serine proteases and has its most potent inhibitory activity toward trypsin, thrombin, and human plasma kallikrein with k_{obsd} /[I] values of 2-3 × 10⁵ M⁻¹ s⁻¹. It was an order of magnitude

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