

ACID HYDROLYSIS PRODUCTS OF SULFOMYCIN I

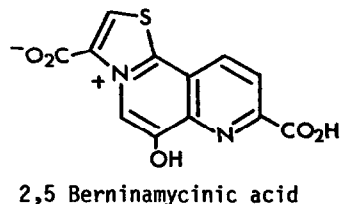
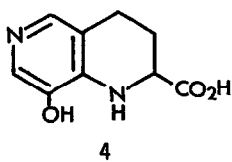
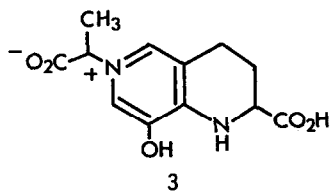
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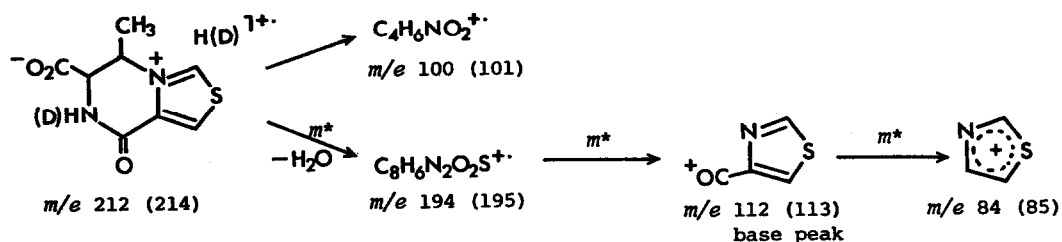
Sulfomycin I (1)<sup>1</sup>, C<sub>55-57</sub>H<sub>56-64</sub>N<sub>15-17</sub>O<sub>20-22</sub>S<sub>2</sub>, a sulfur-containing peptide antibiotic, was hydrolyzed for 6 hours at 110° in conc.-HCl in a sealed tube. Yellow needles which deposited out after cooling were recrystallized from conc.-HCl to give an amphoteric compound (2) as hydrochloride: C<sub>12</sub>H<sub>6</sub>N<sub>2</sub>O<sub>5</sub>·HCl·H<sub>2</sub>O; m.p. >220° (dec.); uv (MeOH): 232 (log ε 4.27) and 277 nm (4.29); ir (KBr): 3400, 3100, 3000-2400 and 1720 cm<sup>-1</sup>. Hydrogenolysis of 2 with Raney nickel catalyst in alkaline aqueous solution gave two novel products assigned 3 and 4. The compounds 3 and 4 were characterized as a methyl and ethyl ester, respectively. Dimethyl ester of 3: C<sub>14</sub>H<sub>18</sub>N<sub>2</sub>O<sub>5</sub>; m.p. 166-167° (dec.); uv (MeOH): 240 (sh., log ε 4.19), 245 (4.21) and 326 nm (4.40); ir (KBr): 1740 cm<sup>-1</sup>; nmr (CDCl<sub>3</sub>): 1.76 (3H,d,J=7), 2.18 (2H,q,J=6), 2.72 (2H,t,J=6), 3.78 (6H,s), 4.32 (1H,t,J=6), 4.72 (1H,q,J=7) and 7.05 (2H,br.s); ms: m/e 294 (M<sup>+</sup>), 236 and 177. Ethyl ester of 4: C<sub>11</sub>H<sub>14</sub>N<sub>2</sub>O<sub>3</sub>; uv (MeOH): 229 and 293 nm; ir (KBr): 1735 cm<sup>-1</sup>; ms: m/e 222 (M<sup>+</sup>) and 149. Thanks to Prof. R.L. Rinehart, Jr., University of Illinois, 2 was identified in his laboratory as berninamycinic acid (5)<sup>2</sup>, a degradation product of berninamycins. Derivation of 3 and 4 from 2 is reasonable to expect.



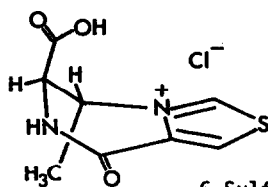
From the filtrate of the above reaction, pyruvic acid, threonine and aminoacetone were recovered. In addition, another new degradation product named sulfomycinine (6) was isolated as hydrochloride: colorless needles; C<sub>8</sub>H<sub>8</sub>N<sub>2</sub>O<sub>3</sub>·HCl; m.p. 205-207° (dec.); uv (MeOH): 230 nm (log ε 3.85), uv (alkaline MeOH): 230 (3.97) and 326 nm (4.20); ir (KBr): 3100, 2900-2500, 1740 and 1695 cm<sup>-1</sup>. The compound 6 was esterified with HCl/EtOH and gave a crystalline ethylester chloride, C<sub>10</sub>H<sub>13</sub>N<sub>2</sub>O<sub>3</sub>·Cl; m.p. 178-180° (dec.); ir (KBr): 3090, 1740 and 1690 cm<sup>-1</sup>; ms: m/e 240 (M<sup>+</sup>-1), 128, 112 (base peak) and 84. In the nmr spectrum of 6 in freshly prepared D<sub>2</sub>O solution, the signals corresponding to CH<sub>3</sub>-CH-CH- system [1.83 (3H,d,J=7), 5.73 (1H,dq,J=2 and 7) and 4.72 (1H,d,J=2)] and aromatic ring protons [8.71 (1H,d,J=2) and 10.25 (1H,d,J=2)] were observed. The latest signal which disappeared on standing in D<sub>2</sub>O solution would be assigned to a proton at C<sub>2</sub>-position of a thiazolium ring. From nmr spectra and high resol-

ution mass spectra of 6 and its deuterized derivative (scheme I), 6 was assumed to be 6-carboxy-5-methyl-8-oxo-5,6,7,8-tetrahydro-thiazolo[3,4-a]pyrazinium chloride. The proposed structure was supported by X-ray crystallography, which further showed that 6 is in DL-form and the conformation is as depicted.

Scheme I



The compound 6 was finally confirmed by synthesis starting from the condensation of 4-thiazolecarboxylic acid with DL-threonine ethylester followed by bromination, cyclization and de-esterification<sup>3)</sup>.



Similarly to thiostrepton<sup>4)</sup>, siomycin<sup>5)</sup> and thiopeptin<sup>6)</sup>, presence of dehydroalanine residue(s) in 1 was suggested by 1) liberation of pyruvic acid, 2) reduction of 1 with sodium borohydride followed by acid hydrolysis to give alanine and 3) reaction of 1 with thioglycolic acid followed by acid hydrolysis to give S-carboxymethylcysteine<sup>7)</sup>. However, 4-( $\alpha$ -hydroxyethyl)-8-hydroxyquinaldic acid and thiostreptine found in alike antibiotics were not detected in the acid hydrolyzate of 1.

## References

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