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DISTRIBUTION OF β -LACTAM AND β -LACTONE PRODUCING BACTERIA IN NATURE

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Over one million bacteria were isolated from a large variety of soil, plant and water samples collected from different environments and examined in an extremely sensitive and highly specific screen for β -lactam production.

A group of seven related monocyclic β -lactams (monobactams) were isolated from strains representing four genera–*Agrobacterium*, *Chromobacterium*, *Gluconobacter* and *Pseudomonas*. Monobactam-producing strains of *Agrobacterium* and *Pseudomonas* were isolated only rarely. Producing strains of *Chromobacterium* were isolated from a relatively limited number of habitats while the *Gluconobacter* strains appeared to be widespread in nature. In addition, three closely related β -lactone-containing molecules were isolated from strains representing three genera–*Arthrobacter*, *Bacillus* and *Pseudomonas*. The *Bacillus* and *Pseudomonas* strains were isolated infrequently but from a variety of samples. The producing strain of *Arthrobacter* was isolated only once.

Screening of microorganisms for the production of antibiotics has provided the cornerstone of antibiotic research programs for the past thirty years. The great majority of such studies have been carried out with fungi and the actinomycetes which are capable of producing natural products with widely divergent chemical structures.

Until 1970, only two classes of naturally occurring β -lactam antibiotics, the penicillins and cephalosporins, were known. However, with the advent of new screening and isolation techniques a variety of β -lactam-containing molecules were identified, as evidenced by the discovery of the cephamycins¹⁾, clavulanic acid²⁾, nocardicins³⁾, and the carbapenems^{4~7)}. Moreover, all the novel β -lactam antibiotics identified over the last ten years are products of the actinomycetes.

In an attempt to discover novel β -lactam-containing molecules from nature, we developed a highly sensitive screen, capable of handling large numbers of organisms and highly specific for β -lactam-containing molecules. Employing this technology we screened large numbers of fungi and actinomycetes only to find many of the known β -lactams. It was at this stage, we turned our attention to the bacteria. This paper reports on the isolation from nature of β -lactam- and β -lactone-producing bacteria⁸). With the exception of tabtoxin produced by species of *Pseudomonas*⁹, bacteria have only recently been reported to produce β -lactam-containing molecules^{10,11}.

Methods

Isolation of Bacteria

Soil and plant material (Table 1) were collected in sterile plastic bags (Whirl-pak, American Scientific Products) and brought to the laboratory for immediate examination or stored at 5° C and examined within $24 \sim 48$ hours. Water samples (Table 1), collected in sterile plastic bottles, were treated similarly.

| Collection area | Types of samples | | | |
|----------------------------|---|--|--|--|
| Agricultural crop lands | Soils & plants | | | |
| Agricultural pasture lands | Soils, plants, mushrooms, water from ponds & streams | | | |
| Gardens | Soils, plants, mushrooms & composts | | | |
| Forests | Soils, plants, mushrooms, leaf litter & tree bark | | | |
| Mountains | Soils, plants & waters | | | |
| Pine barren | Soils, plants & waters | | | |
| Desert | Soils | | | |
| Swamps | Muds & plants | | | |
| Salt marshes | Soils, muds, waters & plants | | | |
| Beaches | Sand & waters | | | |
| Garbage dumps | Soils, plants & decaying matter | | | |
| Sewage plants | Waters | | | |
| Industrial wastes | Waters | | | |
| Rivers | Waters, muds & plants | | | |
| Lakes | Waters, muds & plants | | | |
| Canals | Waters, muds & plants | | | |

Table 1. Collection areas for soil, plant and water samples.

Suspensions of bacteria present in soil and plant samples were prepared by adding five grams of soil or one gram of cut up plant material to 99 ml of diluent, e.g. phosphate-buffered water or saline. The samples were mixed by shaking on a rotary shaker for 20 minutes at 150 rpm. Bacteria present in water samples were concentrated on a Millipore filter (0.45μ) and washed off the filter pad with distilled water. Serial dilutions were prepared of all the processed samples and 0.1-ml aliquots of three dilutions were surfaceplated onto at least four isolation agar plates. A standard dehydrated culture medium, e.g. Brilliant green bile, MACCONKEY, ENDO, Eosin methylene blue (BBL), selective for Gram-negative bacteria was always included as one of the four isolation media. The composition of the other media was dependent on the type of samples being examined; pH and salinity were often adjusted to simulate the environmental conditions from which the samples were collected. In various media, soil and plant extracts were substituted for water. All the isolation media contained actidione (50 μ g/ml) to retard fungal growth. Plates were incubated at room temperature for $2 \sim 3$

days. Water samples were plated onto six isolation media and incubated overnight at room temperature and then at 20°C for an additional four days.

In addition to direct examinations, many soil and plant samples were enriched by the addition of colloidal chitin, cellulose, pectin or various amino acids to enhance the numbers of specific types of bacteria. After a $7 \sim 10$ -day incubation period at room temperature, dilutions were prepared and the samples were plated onto the isolation plates as previously described.

Antibiotic Characterization

Preliminary characterization of antibiotics was based on thin-layer chromatography (F1440 cellulose Schleicher and Schuell, Keine, NJ, and polysilicic acid gel impregnated glass fiber sheets, Gelman Instrument Co., Ann Arbor, MI) employing acetonitrile - water (7: 3, 4: 1 or 5: 1) and high voltage electrophoresis at 2,000 volts for 30 minutes at pH 2, 7 and 9. Antibiotics were visualized by bioassay using *B. licheniformis* SC 9262. In addition, all compounds were tested for stability to a range of β lactamases as described by SYKES, *et al.*¹¹⁾.

Characterization of Producing Organisms

All producing cultures were examined initially for their Gram reaction and by phase contrast microscopy for morphology and motility. Additional standard taxonomic characterization was carried out based on BERGEY'S Manual of Determinative Bacteriology (8th edition). Antibiograms were performed against a battery of 24 antibiotics using a disc diffusion assay (Table 2).

Screening

The screen developed to identify novel β -lactam-containing molecules employed *Bacillus licheni*formis SC 9262 as the assay organism¹¹). Over a million bacteria were tested leading to the discovery of seven related β -lactam and three related β -lactone-containing molecules.

Results

β -Lactam-Producing Bacteria

Monobactams (monocyclic derivatives of 3-amino-2-oxoazetidine-1-sulfonic acid)¹¹⁾ have been

| Compound | Sensi-Disc (µg/disc) | Chromo- bacterium violaceum | Agro- bacterium radiobacter | Glucono- bacter sp. | Pseu- domonas sp. | Bacillus sp. | Arthro- bacter sp. |
|---------------------|-------------------------|-----------------------------------|-----------------------------------|---------------------------|-------------------------|-----------------|--------------------------|
| Amikacin | 10 | 18.4ª | 20.3 | 14.2 | 18.9 | 26.5 | 19.7 |
| Ampicillin | 10 | b | | | | 13.6 | 16.2 |
| Bacitracin | 10 | 9.5* | — | — | | 18.7 | 34.4 |
| Carbenicillin | 50 | | 22.2 | | | 12.4 | 13.8 |
| Cephalothin | 30 | | | | | 12.7 | 27.6 |
| Chloramphenicol | 30 | 17.2 | 9.3* | _ | | 30.6 | 32.9 |
| Colistin | 10 | | 15.4 | | 13.5 | | 14.8 |
| Erythromycin | 15 | 21.7 | 12.9 | <u></u> | - | 31.2 | 35.2 |
| Gentamicin | 10 | 16.2 | 19.6 | 13.2 | 17.0 | 27.6 | 19.2 |
| Kanamycin | 30 | 23.7 | 20.5 | 18.8 | 23.5 | 23.8 | 21.4 |
| Nafcillin | 1 | | | | | | 10.8 |
| Nalidixic acid | 30 | 57.4 | | 21.9 | 15.2 | 26.9 | 16.8 |
| Neomycin | 30 | 17.6 | 19.4 | 13.2 | 16.9 | 24.8 | 19.6 |
| Nitrofurantoin | 300 | 36.0 | 11.7* | | | 21.9 | - |
| Novobiocin | 30 | 43.6 | 27.7 | 18.5 | - | 25.9 | 36.5 |
| Penicillin G | 10 units | | | | | 10.2 | 11.9 |
| Polymyxin B | 300 units | | 17.9 | | 15.5 | 10.9 | 17.6 |
| Lincomycin | 2 | | | | | 13.3 | |
| Rifampin | 5 | 47.5 | 26.4 | 15.7 | 16.4 | 21.0 | 41.8 |
| Streptomycin | 10 | 23.0 | 16.9 | 9.5* | 18.0 | 23.9 | 21.8 |
| Sulfachloropyrazine | 250 | 15.4* | | | | 28.3 | 28.9 |
| Tetracycline | 30 | 39.0 | 39.0 | 23.8 | 29.4 | 25.2 | 36.6 |
| Tobramycin | 10 | 18.2 | 17.1 | 12.2 | 17.0 | 19.9 | 14.6 |
| Vancomycin | 30 | 19.6* | 14.6 | | | 19.9 | 31.0 |

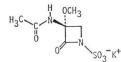
Table 2. Antibiograms of bacteria producing β -lactams and β -lactones.

^a Diameter (mm) of zone of inhibition. ^b No zone of inhibition.

* Indicates variation among strains tested.

detected from strains of *Chromobacterium violaceum*, *Gluconobacter* sp., *Pseudomonas* sp., and *Agrobacterium radiobacter*. The simplest molecule SQ 26,180 (Fig. 1) is produced by strains of *Chromobacterium violaceum*¹²⁾. Although these organisms were isolated frequently from nature, monobactam-producing strains were isolated from a relatively limited number of habitats¹³⁾ (Table 3). In addition, a large number of pigmentless *C. violaceum* strains producing SQ 26,180 were isolated from samples collected in the New Jersey pine barrens. These isolates possessed the same key biochemical characters (Table 4) and antibiotic susceptibility pattern (Table 5) as *C. violaceum* SC 11,378 and represent natural variants of this species. *C. violaceum* SC 11,378, producing SQ 26,180, has been deposited with the American Type Culture Collection (ATCC) under the accession number of ATCC 31,532. This organism was isolated from a soil sample collected in a cedar forest in the New Jersey pine barrens.

The most commonly encountered monobactam was SQ 26,445 (Fig. 2). This compound was produced by strains of bacteria identified as *Gluconobacter* sp.¹¹⁾ and strains of a fluorescent *Pseudomonas* sp. SQ 26,445¹⁴⁾ is identical with sulfazecin, a monocyclic β -lactam produced by strains of *Pseudomonas acidophila*^{10,15)}. Samples of *Gluconobacter* sp. producing SQ 26,445 were deposited with the ATCC under the accession number of ATCC 31,581. *Gluconobacter* sp. ATCC 31,581 was initially isolated from a sample of ground moss collected at New Hope, Pennsylvania. Additional SQ 26,445-producing strains Fig. 1. Structure of SQ 26,180.





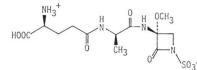
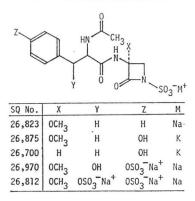


Fig. 3. Structures of monobactams isolated from *Agrobacterium radiobacter* fermentations.



| Sample | Collection site | | |
|--------------------------------------|--|--|--|
| Cedar forest soil | New Jersey pine barrens, New Jersey | | |
| Swamp water | New Jersey pine barrens, New Jersey | | |
| Lake water | New Jersey pine barrens, New Jersey | | |
| Creek water | New Jersey pine barrens, New Jersey | | |
| Water lily (leaves and stem) | New Jersey pine barrens, New Jersey | | |
| Cranberry bog | New Jersey pine barrens, New Jersey | | |
| Blue Spruce needle compost pile | Waterbury, Connecticut | | |
| Small unidentified plant | Barnegat Bay, New Jersey | | |
| Sulfur-iron bog sediment | Great Swamp Refuge, New Jersey | | |
| Decaying root mass of swamp plant | Mercer County Park, New Jersey | | |
| Forest soil | West Windsor, New Jersey | | |
| Oak-leaf litter | Hacklebarney State Park, New Jersey | | |

Table 3. Collection sites of *Chromobacterium violaceum* strains producing SO 26,180.

Table 4. Key taxonomic characters demonstrating similarity of pigmentless bacteria with *C. violaceum* SC 11,378.

| | SQ 26,180 producing strains | | | | | | | |
|------------------------|-----------------------------|-----------|--------|--------|---------------------------|--|--|--|
| Key characters | | Pigmented | | | | | | |
| | F-3600 | F-3615 | F-3616 | F-3617 | C. violaceum SC 11,378 | | | |
| Gram reaction | - | | | | - | | | |
| Motility | + | + | + | + | + | | | |
| Oxid./Ferm. | F | F | F | F | F | | | |
| Casein hydrolysis | + | + | + | + | + | | | |
| Acid from trehalose | + | + | + | + | + | | | |
| Aesculin hydrolysis | _ | - | _ | _ | _ | | | |
| Production of HCN | + | +- | + | + | + | | | |
| Arginine decarboxylase | + | + | + | + | + | | | |
| Violacein production | _ | - | _ | | + | | | |

of *Gluconobacter* sp. were isolated from a variety of environments as shown in Table 6. The fluorescent *Pseudomonas* that produces SQ 26,445 was isolated on only two occasions from water samples collected at the same site in Regensburg, Germany. Ten strains of *Gluconobacter* and *Acetobacter* (Table 7) obtained from the ATCC produced SQ 26,445.

A mixture of monobactams was found to be produced by bacterial isolates identified as strains of

| Compound | Sensi-disc (µg/disc) | C. violaceum SC 11,378 | F-3600 | F-3615 | F-3616 | F-3617 |
|---------------------|-------------------------|---------------------------|--------|--------|--------|--------|
| Amikacin | 10 | 14.7ª | 15.0 | 13.7 | 14.5 | 14.5 |
| Ampicillin | 10 | b | | | | |
| Bacitracin | 10 | - | | | | |
| Carbenicillin | 50 | - | | | | |
| Cephalothin | 30 | _ | | | | |
| Chloramphenicol | 30 | 16.5 | 17.8 | 15.7 | 13.3 | 15.0 |
| Colistin | 10 | - | | | | |
| Erythromycin | 15 | 21.1 | 26.4 | 12.9 | 19.4 | 23.9 |
| Gentamicin | 10 | 15.6 | 15.4 | 15.0 | 15.5 | 14.7 |
| Kanamycin | 30 | 21.5 | 21.5 | 21.5 | 20.2 | 20.9 |
| Nafcillin | 1 | | | | | |
| Nalidixic Acid | 30 | 43.4 | 45.5 | 40.6 | 43.8 | 45.3 |
| Neomycin | 30 | 15.5 | 15.1 | 9.9 | 14.7 | 14.7 |
| Nitrofurantoin | 300 | 29.1 | 28.7 | 22.1 | 29.0 | 28.7 |
| Novobiocin | 30 | 37.3 | 41.0 | 39.6 | 39.7 | 37.6 |
| Penicillin G | 10 units | | | | | |
| Polymyxin B | 300 units | | | | | |
| Lincomycin | 1 | | | | | _ |
| Rifampin | 5 | 27.2 | 28.7 | 32.0 | 27.8 | 30.9 |
| Streptomycin | 10 | 18.9 | 17.7 | 17.2 | 18.0 | 19.0 |
| Sulfachloropyrazine | 250 | 18.7 | 24.0 | 21.4 | 17.8 | 17.9 |
| Tetracycline | 30 | 32.0 | 34.6 | 34.0 | 34.3 | 34.3 |
| Tobramycin | 10 | 15.4 | 14.5 | 14.5 | 14.3 | 14.8 |
| Vancomycin | 30 | 9.5 | 10.7 | 11.6 | 12.4 | 10.0 |

Table 5. Susceptibility of *C. violaceum* SC 11,378 (ATCC 31,532) and four pigmentless strains of *C. violaceum* to 24 antibiotics.

^a Diameter (mm) of zones of inhibition.

^b No zone of inhibition.

| Table | 6. T | ypes | and | nu | mbers | of | sampl | es co | ntaining |
|-------|---------|-------|------|----|-------|-----|-------|-------|----------|
| SQ | 26,44 | 5-pro | duci | ng | Gluco | nob | acter | and | Pseudo- |
| mon | as stra | ins. | | | | | | | |

| Table 7. | SQ | 26,445-pro | oducing | bacteria | obtained |
|----------|-----|------------|---------|----------|------------|
| from | the | American | Type | Culture | Collection |
| (ATCC | C). | | | | |

| Sample | No. Locations | Culture | ATCC No. |
|--|------------------|---|-----------------|
| <u> </u> | | Acetobacter aceti subsp. aceti | 15973 |
| Ground mosses (Pennsylvania and New Jersey): | 4 | Acetobacter aceti subsp. liquefaciens | 23751 |
| Fungi (New Jersey): <i>Polyporus</i> sp., <i>Amanita</i> sp. and unidentified species of bird's-nest | 11 | Acetobacter pasteurianus subsp. pasteurianus | 6033 |
| fungi, white bracket fungi and a | | Acetobacter peroxydans | 12874 |
| brown gilled fungus | | Acetobacter sp. | 21760 |
| Soils (New Jersey, California, Iowa and Germany): | 13 | Gluconobacter oxydans subsp. oxydans | 15178 and 19357 |
| Plants (New Jersey): | 4 | Gluconobacter oxydans subsp. | |
| Decaying plants (Pennsylvania, New | | suboxydans | 19441 and 23773 |
| Jersey, New York, Maine and Germany): | 13 | Gluconobacter oxydans subsp. industrius | 11894 |
| Water (Germany): | 1 | | 11074 |

Agrobacterium radiobacter^{11,16}). The structures of five of these compounds are shown in Fig. 3¹⁷). SQ 26,700 was the first non-methoxylated monobactam derivative isolated in quantity. Samples of A. radiobacter producing monobactams were deposited at the ATCC under the accession number of ATCC 31,700. Monobactam-producing A. radiobacter strains were isolated on only three occasions, twice from plant samples collected in Germany and once from mud and leaf litter collected in New Jersey.

β-Lactone-Producing Bacteria

During our search for β -lactam-producing bacteria we isolated a number of β -lactone containing molecules^{8,18)}. These compounds contain a monocyclic ring system in which oxygen replaces nitrogen of the β -lactam.

The first β -lactone encountered, SQ 26,517 (Fig. 4), contains the same acyl side chain as the monobactam, SQ 26,180, but has a β -methyl group at position-4. The bacterium producing SQ 26,517 is an aerobic, Gram-positive, spore-forming rod and was therefore assigned to the genus Bacillus. Strains of SQ 26,517-producing bacilli were isolated relatively infrequently but from samples collected worldwide (Table 8). An antimicrobial evaluation of racemic SQ 26,517, synthesized from DL-allothreonine,

Fig. 4. Structure of SQ 26,517.

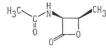


Fig. 5. Structure of SQ 27,012.

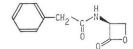


Fig. 6. Basic structure of EM5357 and EM5395.



Table 8. Types and number of samples containing SQ 26,517-producing Bacillus strains.

| Sample | No. Locations |
|--|------------------|
| Soils (Hawaii, Iowa, Louisiana, New Jersey, Argentina, Brazil, Chile and Germany): | 10 |
| Moss (Germany): | 1 |
| Sand (Germany): | 1 |
| Plant debris (Pennsylvania): | 1 |
| Mud (New Jersey): | 1 |
| Water (Pennsylvania): | 1 |
| | |

| Organism | | | $(\mu g/ml)$ |
|------------------------|----|-------|--------------|
| Staph. aureus | SC | 1276 | >100 |
| Staph. aureus | SC | 2399 | >100 |
| Staph. aureus | SC | 2400 | >100 |
| Staph. aureus | SC | 10165 | >100 |
| Strep. faecalis | SC | 9011 | >100 |
| Strep. agalactiae | SC | 9287 | 50 |
| Micro. luteus | SC | 2495 | 100 |
| E. coli | SC | 8294 | >100 |
| E. coli | SC | 10857 | >100 |
| E. coli | SC | 10896 | >100 |
| E. coli | SC | 10909 | >100 |
| K. pneumoniae | SC | 10440 | >100 |
| K. pneumoniae | SC | 9527 | >100 |
| Prot. mirabilis | SC | 3855 | >100 |
| Prot. rettgeri | SC | 8479 | >100 |
| Prot. vulgaris | SC | 9416 | 100 |
| Sal. typhosa | SC | 1195 | >100 |
| Shig. sonnei | SC | 8449 | >100 |
| Ent. cloacae | SC | 8236 | >100 |
| Ent. aerogenes | SC | 10078 | >100 |
| Citro. freundii | SC | 9518 | >100 |
| Ser. marcescens | SC | 9783 | >100 |
| Ps. aeruginosa | SC | 9545 | >100 |
| Ps. aeruginosa | SC | 8329 | >100 |
| Acineto. calcoaceticus | SC | 8333 | >100 |

Minimum inhibitory concentrations were determined by a two fold agar dilution method on DST agar (Oxoid). Final inoculum level was 104 colony-forming units.

Table 9. Antibacterial activity of racemic SQ 26,517.

MIC

showed weak activity (Table 9). SQ 27,012, a synthetic analog of SQ 26,517 (Fig. 5), produced no zone of inhibition when tested against a limited number of microorganisms. Interaction of SQ 27,012 with β -lactamases is shown in Table 10. With the exception of the K-l enzyme from *Klebsiella pneumoniae*, the compound was stable to β -lactamase attack.

 β -Lactones EM5395 and EM5357 shown in Fig. 6 were produced by strains of bacteria identified as *Pseudomonas* sp. and a soil coryneform, respectively. The bacterium producing EM5395 is a Gram-

| Table | 10. | Action | of | β -lactamases | on | SQ | 27,012.* |
|-------|-----|--------|----|---------------------|----|----|----------|
|-------|-----|--------|----|---------------------|----|----|----------|

| Compound | Relative rate of hydrolysis with β -lactamase type | | | | | |
|---------------|--|-----|------|--|--|--|
| compound | TEM-2 | K-1 | P-99 | | | |
| Cephaloridine | 100 | 100 | 100 | | | |
| SQ 27,012 | <0.01 | 0.6 | 0.04 | | | |

 Studies were performed using spectrophotometric assays at 25°C in 0.1 м phosphate buffer, pH 7.0.

negative polar flagellated rod. Growth did not occur at pH 4.5 nor at temperatures of 41°C. The culture was cytochrome-oxidase and argininedihydrolase positive. Fluorescent pigment was produced on KING's medium B¹⁰⁾. On the basis of these characteristics, the organism was assigned to the genus *Pseudomonas*. EM5395-producing strains of *Pseudomonas* were isolated most frequently from a variety of mushrooms (Table 11). The bacterium producing EM5357 is a Grampositive, pleomorphic non-motile rod. Based on

| Sample | Site |
|--|---|
| Hygrophorus cantharellus mushroom | Washington Crossing State Park, New Jersey |
| Yellow Hygrophorus sp. | Washington Crossing State Park, New Jersey |
| Tan-grey, small, umbonated mushroom on dead tree stump | Washington Crossing State Park, New Jersey |
| Decaying Armillariella sp. mushroom | Princeton, New Jersey |
| Amanita chlorinosma mushroom | Princeton, New Jersey |
| Tan gilled mushroom from dead tree | Princeton, New Jersey |
| Decaying unidentified mushroom | Princeton, New Jersey |
| Unidentified mushroom on rotting log | Hacklebarney State Park, New Jersey |
| Leaf litter and soil | Hacklebarney State Park, New Jersey |
| Water and leaves from small spring | Jamesburg, New Jersey |
| Algae and beach sand | Bahamas |

Table 11. Collection sites of Pseudomonas strains

producing the β -lactone EM5395.

morphology and analysis of cell wall hydrolysates, the culture was assigned to the soil coryneform group and tentatively placed in the genus *Arthrobacter*. The culture was isolated only once from a sample of decaying leaf litter collected at a site in New Jersey.

Discussion

The ability to produce monocyclic β -lactams and β -lactones is possessed by strains of bacteria representing a number of genera. Although some of the strains appeared in a limited number of samples, others were isolated from a wide variety of natural environments.

The discovery of these compounds has opened up a new era of antibiotic research. One exciting aspect of this research has been the total chemical synthesis of these compounds. Furthermore, as with the penicillins and cephalosporins, chemical modification of the monobactams has led to compounds with superior properties over the naturally occurring molecules²⁰. One of these compounds, SQ 26,776 (Azthreonam) a highly active β -lactamase-stable compound has been developed for clinical evaluation²¹.

Acknowledgements

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820

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