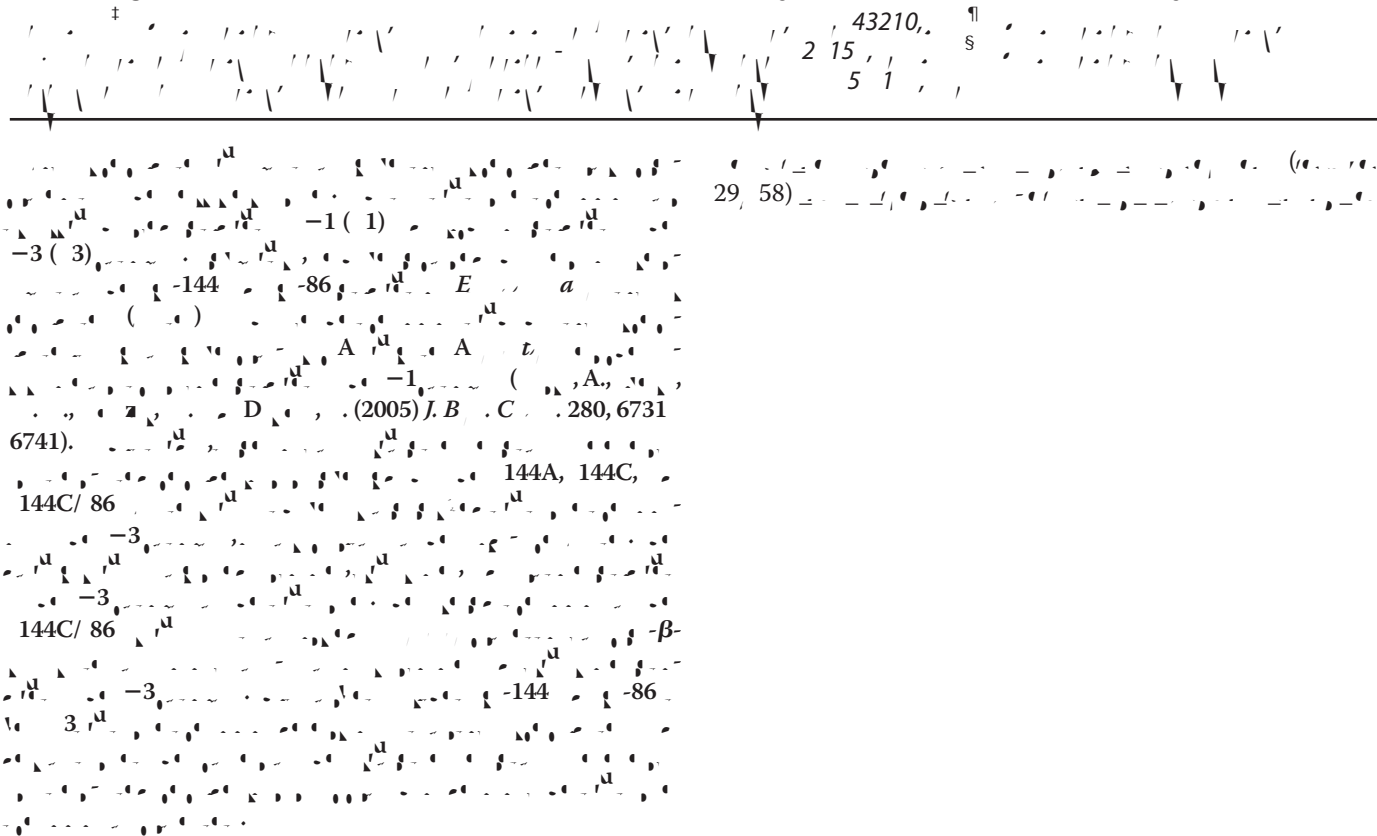


Altered –3 Substrate Specificity of Signal Peptidase 1 Mutants as Revealed by Screening a Combinatorial Peptide Library*

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Pr. ... (1). T ... (2) ... D ... (SP ... I)² ... 37 D (3). T ...

... α -...-4-... (10
... 500 μ ... 0.1% ... 500 μ ... CH₃CN).
T... MALDI-TOF ... 1 μ
... " ...

..... T.....
.....
Peptide Sequencing P..... E..... / ..

5:1
N-
(F₁, 2B). T.
MALDI-MS
I PED/MS
T. NH₂

F
(F₁, 1B). A.
SP N-
SP T. (B)
(a, F₁, 1B). T. F
E (F₁, 1B). T.
N- I MALDI-
TOF NH₂

SP₂ (F₁.3D).
 P2F β₂ (F₁.3D).
 I144C β₂
 -3 G. β₂
 (F₁.3E),
 A 207(6)

(28,908 D). T₁ (232.4 D). H₁-Pr₁ (234.3 D). NH₂-H₁-Pr₁ β₂ A -3 A₁ P₁ -1 M13 (39).

The SPase I144C/I86T Mutant Can Cleave Pre-β-lactamase Mutants with Non-canonical -3 Residues B₁ +2 β₂ Pr₁-2 A F₁.3A, SP₂ -3 A₁ P2F β₂ I144C/I86T SP₂ -3 A₁ P2F β₂ S I144C/I86T SP₂ -3 G. β₂ (F₁.3B), -3 T β₂ (F₁.3C). T₁ SP₂ -3 G. β₂ (F₁.3B). I

I₁-101
I₂-132
T₁
T₂-3
T₃

DISCUSSION

I₁-86 I₂-144
E. coli SP₁-3 U₁ FRET I₁-144

PED/MS
20 30 1 \$1/
(
I
H

C

()