Table 1. Tabulated m/z values of predicted and observed peptide fragments and  $\beta$ -elimination products

Open Cys9-Cys60		
	Predicted	Observed
Fragment	mass, Da	mass, Da
Ala <sub>1</sub> -Cys <sub>18</sub>	772.88	N.D.
itz <sub>9</sub> -Lys <sub>59</sub>	5,086.94	5,080.695*
itz <sub>60</sub> -Ala <sub>82</sub>	2,411.74	2,413.79
Ala <sub>1</sub> -(βCys <sub>9</sub> )Lys <sub>59</sub>	5,764.72	5,770.79
itz <sub>9</sub> -(βCys <sub>60</sub> )Ala <sub>82</sub>	7,421.57	7,447.43 <sup>†</sup>
Open Cys18-Cys54	-1	
Ala <sub>1</sub> -Tyr <sub>17</sub>	1,857.1	1,857.04
itz <sub>18</sub> -Gln <sub>53</sub>	3,454.1	3,436.10*‡
itz <sub>54</sub> -Ala <sub>82</sub>	2,957.4	2,956.85
Ala <sub>1</sub> -( $\beta$ Cys <sub>18</sub> )Gln <sub>53</sub>	5,237.05	5,238.07*
$itz_{18}$ -( $\beta Cys_{54}$ )Ala <sub>82</sub>	6,336.36	N.D.
Open Cys19-Cys45		
Ala <sub>1</sub> -Cys <sub>18</sub>	1,961.23	N.D.
itz <sub>19</sub> -Gly <sub>44</sub>	2,508.92	2,509.55*
itz <sub>45</sub> -Ala <sub>82</sub>	3,800.42	3,796.94*
Ala <sub>1</sub> -(βCys <sub>19</sub> )Gly <sub>44</sub>	4,394.03	4,409.147 <sup>§</sup>
itz <sub>19</sub> -(βCys <sub>45</sub> )Ala <sub>82</sub>	6,233.22	N.D.
Open Cys61-Cys80		
Ala <sub>1</sub> -Cys <sub>60</sub>	5,919.94	5,902.98‡
itz <sub>61</sub> -Asn <sub>79</sub>	2,035.24	2,036.25
itz <sub>80</sub> -Ala <sub>82</sub>	3,16.38	N.D.
Ala <sub>1</sub> -(βCys <sub>61</sub> )Asn <sub>79</sub>	7,878.07	7,894.07 <sup>§</sup>
itz <sub>61</sub> -( $\beta$ Cys <sub>80</sub> )Ala <sub>82</sub>	2,274.51	N.D.

Fragments highlighted in gray are unique to a particular fingerprint. N.D., not detected; itz, iminothiazolidinyl carboxyl residue.  $Cys_x$ - $Cys_y$  designates the disulfide bond that is reduced in the singly reduced and cyanylated species.  $Ala_1$ – $(X_{x-1})$ , itz<sub>x</sub>– $X_{y-1}$ , and itz<sub>y</sub>– $Ala_{82}$ , designates the *S*-cyanyl cleavage products.  $Ala_1$ – $(\beta Cys_x)$ – $X_{y-1}$  and itz<sub>x</sub>– $(\beta Cys_y)$ – $Ala_{82}$  designate the  $\beta$ -elimination products.  $\beta Cys_x$  and  $\beta Cys_y$  refer to the position of the Cys modified by the  $\beta$ -elimination reaction. X = any amino acid residue; subscript symbols refer to the position of designated singly reduced and cyanylated Cys residues.

<sup>\*</sup>Denotes that both the oxidized and non-oxidized M22 are detected.

<sup>&</sup>lt;sup>†</sup>The ion detected is as an adduct with Na<sup>+</sup>.

 $<sup>^{\</sup>ddagger}$ The ion detected has lost  $H_2O$ .

<sup>§</sup>Only the oxidized M22 is detected.