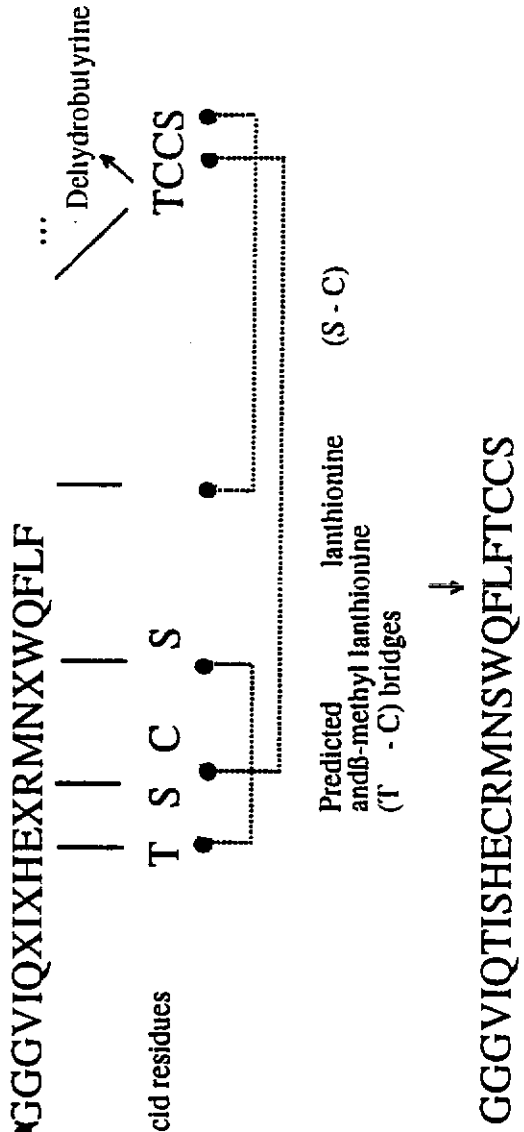
FIGURE 1


End of leader sequence
GGGVIQXIXHEXRMNXWQFLF
Predicted amino acid residues
T S C S
TCCS
Dehydrobutyrine
Predicted lanthionine and ß-methyl lanthionine (T - C) bridges
(S - C)
GGGVIQTISHECRMNSWQFLFTCCS
Deduced amino acid sequence for salB propeptide region

1	TTG	ACT	CTT	GAA	GAA	CTT	GAT	AAC	GTT	CTT	GGT	GCT	GGT	
	Leu	Thr	Leu	Glu	Glu	Leu	Asp	Asn	Val	Leu	Gly	Ala	Gly	13
	-12											-1	↑ +1	
40	GGT	GGA	GTA	ATC	CAA	ACC	ATT	TCA	CAC	GAA	TGT	CGT	ATG	
	Gly	Gly	Val	Ile	Gln	Thr	Ile	Ser	His	Glu	Cys	Arg	Met	26
80	AAC	TCA	TGG	CAG	TTC	TTG	TTT	ACT	TGT	TGC	TCT	TAA		
	Asn	Ser	Trp	Gln	Phe	Leu	Phe	Thr	Cys	Cys	Ser	•		37
											+25			

Figure 2

1	ATG	ATT	GCC	ATG	AAA	AAC	TCA	AAA	GAT	ATT	TTG	AAC	AAT	
	Met	Ile	Ala	Met	Lys	Asn	Ser	Lys	Asp	Ile	Leu	Asn	Asn	13
	-29									-20				
40	GCT	ATC	GAA	GAA	GTT	TCT	GAA	AAA	GAA	CTT	ATG	GAA	GTA	
	Ala	Ile	Glu	Glu	Val	Ser	Glu	Lys	Glu	Leu	Met	Glu	Val	26
							-10							
80	GCT	GGT	GGT	AAA	AGA	GGT	ACA	GGT	TGG	TTT	GCA	ACT	ATT	
	Ala	Gly	Gly	Lys	Arg	Gly	Thr	Gly	Trp	Phe	Ala	Thr	Ile	39
			-1	↑ +1									+10	
120	ACT	GAT	GAC	TGT	CCA	AAC	TCA	GTA	TTC	GTT	TGT	TGT	TAA	
	Thr	Asp	Asp	Cys	Pro	Asn	Ser	Val	Phe	Val	Cys	Cys	•	52
										+20				

Figure 3