

Band 1

Protein name: Chain A, Crystal Structure Of Efb-C From *Staphylococcus aureus*
Alphalyse number: ALPHA09150
GI-number: gi|145579437
MW: 7184
pI: 10,23
Mascot score: 179
Sequence coverage: 65%

1 IKKEQKLIQA QNLVREFEKT HTVSAHRKAO KAVNLVSFEY KVKKMVLQER
51 IDNVLKQGLV R

Band 2

Protein name: leukocidin LukS component [*Staphylococcus aureus* subsp. *aureus*]
Alphalyse number: ALPHA09148
GI-number: gi|13549150
MW: 32563
pI: 9,09
Mascot score: 273
Sequence coverage: 39%

1 ANDTEDIGKG SDIEIKRTE DKTSNKWGV T QNIOFDFVKD KKYNKDALIL
51 KMOGFISSRT TYNYKKTNH VKAMRWPFQY NIGLKTNDKY VSLINYLPKN
101 KESTNVSQT LGYNIGGNFQ SAPSLGGNGS FNYSKISYT QQNYVSEVEQ
151 QNSKSVLWGV KANSFATESG QKSAFDSDLF VGYKPHSKDP RDYFVPDSEL
201 PPLVQSGFNP SFIATVSHEK GSSDTSEFEI TYGRNMDVTH AIKRSTHYGN
251 SYLDGHRVHN AFVNRNYTVK YEVNWKTHEI KEKGQN

Band 3

Protein name: leukotoxin, LukE, putative [*Staphylococcus aureus* subsp. *aureus* NCTC 8325]
Alphalyse number: ALPHA09148
GI-number: gi|88195648
MW: 33666
pI: 9,33
Mascot score: 294
Sequence coverage: 38%

1 MSVGLIAPLA SPIQESRANT NIENIGDGAE VIKRTEDVSS KKWGVTONVO
51 FDFVKDKKYN KDALIVKMOG FINSRTSFSV VKGSGYELTK RMIWPFQYNI
101 GLTTKDPNVS LINYLPKNKI ETTDVGQTLG YNIGGNFQSA PSIGGNQSFN
151 YSKTISYTK SYVSEVDKQN SKSVKGVKA NEFVTPDGKK SAHDRYLFVQ
201 SPNGPTGSAR EYFAPDNQLP PLVQSGFNPS FITTLSHEKG SSSDTSEFEIS
251 YGRNLDITYA TLPRTGIYA ERKHNAFVNR NFVVR YEVNWKTHEIKVKGH
301 N