or 5 (SEQ ID NO: 29)) amino acid sequence at its N-terminus,

iv) a fourth polypeptide with sortase activity whereby the polypeptide is derived from Staphylococcus aureus sortase A, and

v) a fifth polypeptide with sortase activity whereby the polypeptide is derived from Streptococcus pyogenes sortase A,

or

c) i) a first polypeptide comprising the amino acid sequence LPXTA (SEQ ID NO: 31, wherein X can be any amino acid residue),

ii) a second polypeptide that has an oligo-alanine A_m (m = 2 (SEQ ID NO: 26), or 3 (SEQ ID NO: 27), or 4 (SEQ ID NO: 28), or 5 (SEQ ID NO: 29)) amino acid sequence at its N-terminus and that comprises the amino acid sequence LPXTG (SEQ ID NO: 20, wherein X can be any amino acid residue),

iii) a third polypeptide that has an oligo-glycine G_m (m = 2 (SEQ ID NO: 22), or 3 (SEQ ID NO: 23), or 4 (SEQ ID NO: 24), or 5 (SEQ ID NO: 25)) amino acid sequence at its N-terminus,

iv) a fourth polypeptide with sortase activity whereby the polypeptide is derived from Staphylococcus aureus sortase A, and

v) a fifth polypeptide with sortase activity whereby the polypeptide is derived from Streptococcus pyogenes sortase A,

and

- recovering the conjugate from the reaction mixture and thereby producing the enzymatic conjugation product of three polypeptides.

2. The method according to claim 1, wherein the first, second and third polypeptide are present at equimolar concentration.

3. The method according to any one of claims 1 to 2, wherein the oligo-alanine is a di-alanine (SEQ ID NO: 26) or a tri-alanine (SEQ ID NO: 27).