Erratum to: *In silico* study of binding motifs in squalene synthase enzyme of secondary metabolic pathway of solanaceae

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Following section of the original version of the article unfortunately contains errors. The corrected version is given below.

1. The manuscript title, “*In silico* study of binding motifs in squalene synthase enzyme of secondary metabolic pathway solanaceae family” may be read as “*In silico* study of binding motifs in squalene synthase enzyme of secondary metabolic pathway **of** solanaceae”.
2. Under result section, subheading motif analysis, lines 2, 3, 4, 5, 6, 7, 8 and 9 “(Fig. 2). All protein sequence of SQS reported two conserved motifs viz., motif 1

“YCHYVAGLVGLGLSKL” which was 16 residue long and motif 2 “MGLFLQKTNIIRDYLEDI-NEIPKSRMFWP” which was 29 residue long. In positions of motifs lied between 170 and 186 for motif 1 and 205 and 234 for motif 2. Only in the case of *N. tabacum* the length of motif 2 varied. These motif sequences were previously reported in other plants with several residues substitution [22, 23].” may be treated as deleted.

3. Under subheading “Molecular docking” the line, “FPP was found to bind with two residues LYS²²³ and VAL^{221 263 261} of the motif 2 (Fig. 6a).” may be read as “FPP was found to bind with two residues LYS²²³ and VAL²²¹ of the motif 2 (Fig. 6a).”

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