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The images depict models of rLHβα with different linkers using the I-TASSER server. The resulting models for (left) 3x(Gly-Ser); (center left) 5x(Gly-Ser); (center right) 2x(Gly4Ser); (right) 3x(Gly4Ser); the common α-subunit is marked in purple, the β-subunit in light blue, the His-tag in red and the linker in orange. Binding sites in the α-subunit (residues 192, 199, 218) are marked in dark purple and in the β-subunit (14, 26, 60–64, 75–79) in dark blue. From Aizen et al. 232 391–402.

Credit: J Aizen & B Levavi-Sivan (The Hebrew University of Jerusalem, Rehovot, Israel)