Supplementary Files: Figure S1

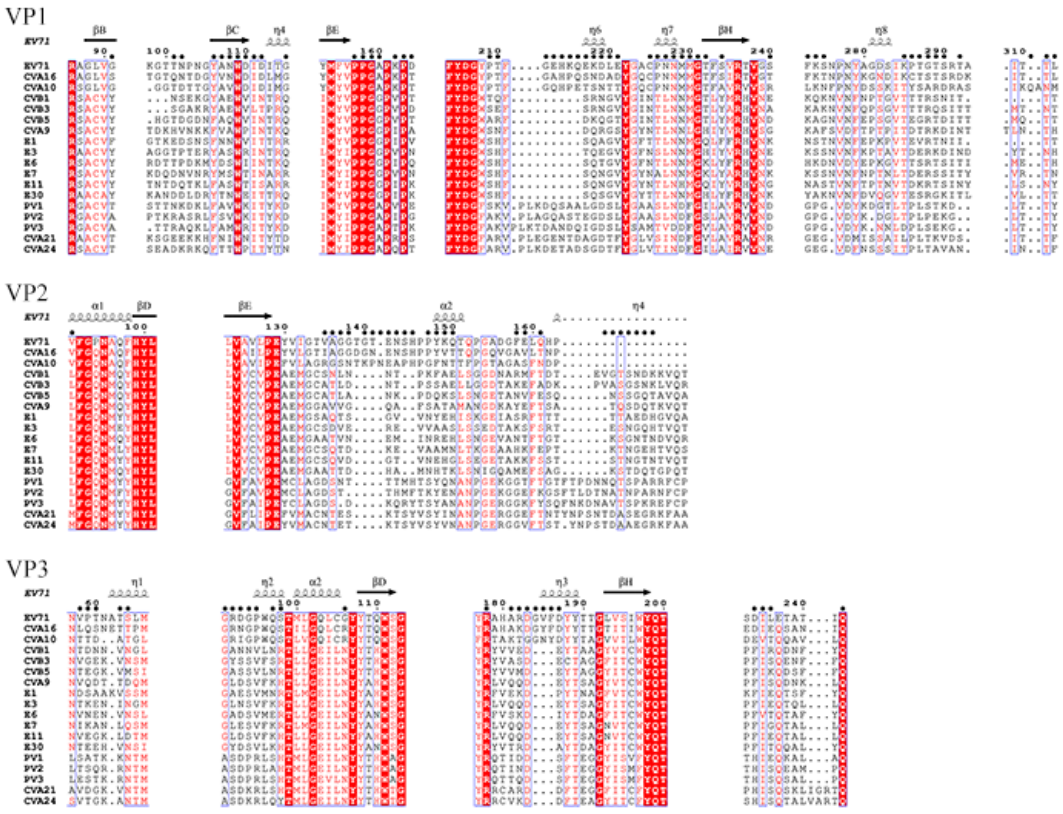

Figure S1. Sequence alignment results of binding sites for uncoating receptors of enteroviruses. The secondary structural regions of the amino acid sequences of enterovirus VP1-VP3 were intercepted. These regions contained 104 receptor-binding sites, which were marked with solid black dots.