|  |
| --- |
|  **S9 Table. Peptide docking results**  |
| **Peptide sequence** | **CPD domain I** |  | **CPD domain II** |
| **Correct poses (%)** | **Experimental ratio** |  | **Correct poses (%)** | **Experimental ratio** |
| TITGQKR | 15.84 |  0.10 \* |  | 11.88 | 0.16 |
| LVTDLTK |  5.94 | 0.83 |  |  7.92 | 0.29 |
| GAAAGQR |  7.00 | 0.56 |  |  5.94 | 0.83 |
| FAATSFR | 12.87 |  0.10 \* |  |  12.87 | 0.24 |
| VDPVNFK |  1.98 | 1.05 |  |  12.87 |  0.10 \* |
| SALGEPKK |  6.93 | 1.04 |  |  1.98 |  0.94 |
| Peptide sequences and experimental ratios were selected from table 5 of good and weak substrates for CPD domains I and II. \*For experimental values lower than 0.10 (< 0,1), we assigned an exact value of 0.1.  |