

Predicted epitopes:

| No. | Start Position | End Position | Peptide | Peptide Length |
|------------|---------------------------|-------------------------|---|---------------------------|
| 1 | 18 | 54 | WWTPPAWAIGPPVVDAAAQPPSGDPGPVAPMEQRGAC | 37 |
| 2 | 59 | 73 | VIPGTDPGVPTPSQT | 15 |
| 3 | 82 | 86 | QFSRG | 5 |
| 4 | 97 | 126 | GVQPGPRLPNVDAGGDFVESTDGLTDCDGH | 30 |
| 5 | 135 | 148 | AGQPGNDGFSGVAP | 14 |
| 6 | 161 | 175 | KFSPRTSGGDPQLAQ | 15 |
| 7 | 211 | 212 | MV | 2 |
| 8 | 226 | 226 | V | 1 |
| 9 | 236 | 269 | AGNTGASGSVSASCDNPLTDLSRPDDPRNWAGV | 34 |
| 10 | 272 | 272 | V | 1 |
| 11 | 275 | 277 | PSW | 3 |
| 12 | 289 | 331 | TSAGQPSKFSMPGPWVGIAAPGENIASVSNSGDGALANGLPDA | 43 |
| 13 | 341 | 346 | TSYAAG | 6 |
| 14 | 359 | 363 | YPGLN | 5 |
| 15 | 376 | 386 | HRGARESSNIV | 11 |
| 16 | 401 | 431 | LPAEPGGGAAPAKPVADPPVPAPKDTTPRNV | 31 |