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| Type/subtype | Position | length | sequences | vaxijen | Bepi pred median |
| .A/H1 | 229 | 17 | **ARPQVNGQSGRIDFHWTI** | **1.9766** | 0.501556 |
| A/H1 | 23 | 37 | **GYHANNSTDTVDTVLEKNVTVTHSVNLLEDSHNGKLC** | **0.5868** | 0.548333 |
| A/H1 | 231 | 24 | **QTAAQKAMMDQVRESRNPGNAEFE** | **0.5114** | 0.507 |
| A/H3 | 387 | 25 | **AIRTRSGGNTNQQRASAGQISIQPT** | **1.6861** | 0.504 |
| A/H3 | 78 | 17 | **RFVQNALNGNGDPNNMD** | **0.7434** | 0.504444 |
| A/H3 | 7 | 19 | **KRSYEQMETDGERQNATE** | **1.4941** | 0.575 |
| A/H3 | 159 | 29 | **MDPRMCSLMQGSTLPRRSGAAGAAVKGVG** | **0.8779** | 0.522333 |
| A/H3 | 425 | 31 | **IMAAFNGNTEGRTSDMRTEIIRMMESARPED** | **0.9847** | 0.504556 |
| A/H3 | 362 | 16 | **GVQIASNENMETMESS** | **0.8436** | 0.545889 |
| A/H3 | 78 | 17 | **RFVQNALNGNGDPNNMD** | **0.7434** | 0.504444 |
| A/H3 | 46 | 38 | **TNDQIEVTNATELVQSSSTGRICDSPHRILDGKNCTLI** | **0.6291** | 0.579889 |
| A/H3 | 211 | 11 | **GRKTRSAYERM** | **1.1406** | 0.498444 |
| A/H3 | 56 | 9 | LIQNSLTIE | **0.7807** | 0.592222 |
| A/H3 | 264 | 12 | LILRGSVAHKSC | **0.4368** | 0.503222 |
| A/H3 | 199 | 36 | HHPGTNNDQISLYTQASGRITVSTKRSQQTVIQNIGS | **0.6939** | 0.505778 |
| A/H3 | 5 | 10 | **TEVETYVLSI** | **1.1699** | 0.506778 |
| B | 5 | 16 | **DIDGINTGTIDKTPEEITSGTSGTTR** | **1.4912** | 0.521444 |
| B | 341 | 10 | **KTPLKLANGT** | **0.6755** | 0.504333 |
| B | 173 | 17 | **WAVPKDNNKNATNPLTV** | **0.4123** | 0.502778 |
| B | 5 | 40 | IVLLMVVTSNADRICTGITSSNSPHVVKTATQGEVNVTGV | **0.7561** | 0.526889 |
| B | 102 | 12 | **RFGETKGNSAPL** | **1.1730** | 0.502778 |
| B | 41 | 20 | **PSNKRTRNPSPERATTSSED** | **1.1352** | 0.510889 |
| B | 404 | 10 | **TGTEFKPRSA** | **1.4082** | 0.503333 |
| B | 68 | 15 | **KKQTPTEIKKSVYNM** | **0.3777** | 0.49 |
| B | 522 | 11 | **MFQISDKNKTN** | **0.7705** | 0.501556 |
| B | 247 | 32 | **PNQAEDGGLPQSGRIVVDYMVQKSGKTGTITY** | **0.6222** | 0.503 |
| B | 373 | 16 | **RVGDDAKDKSQLFFM** | **0.7021** | 0.500667 |

Conserved B\_cell epitope located o the surface of HA, predicted using vaxi

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