Polymorphism III HR1 Predicted linear epitope LII RBD Glycosite Accessibility 27 AYTNSFTRGVYYPDKVFRSSVLHSTQDLFLPFFSNVTWFHAIHVSGTNGTKRFDNPVLPFNDGVYFASTE 96 307 T V E K G I Y Q T S N F R V Q P T E S I V R F P N I T N L C P F G E V F N A T R F A S V Y A W N R K R I S N C V A D Y S V L Y N S A S F S T 376 . N D L C F T N V Y A D S F V I <mark>R G D E V R Q I A P G Q T G K I A D Y</mark> N Y K L P D D F T G C V I A W <mark>N S N N L</mark> D S <mark>K</mark> NYLYRL FRKSNLKPFERDISTEIY QAGSTPCNGVEGFNCYFPLQSYGFQPTNGVGYQPYRVVVLSFE 516 APATVCG<mark>PKKSTNLVKNK</mark>CVNFNFNGLTGTGVL<mark>TESNKKFLP</mark>FQQFGRDIADTTDAVRDP<mark>QTLE</mark>ILD 586 657 NNSYECDIPIGAGICASYOTOTNSPRRARSVASOSIIAYTMSLGAENSVAYSNNSIAIPTNFTI TKTSVDCTMYICGDS<mark>TE</mark>CSNLLLQYGSFCTQLNRALTGIAVEQDKNTQEVFAQV<mark>KQ</mark>IYKTPP K V T L A D A G F I K Q Y G D C L G D I A 867 DEMIAQYTSALLAGTITSGWTFGAGAALQIPFAMQMAYRFNGIGVTQNVLYENQKLIANQFNSAIGKIQD 936 937<mark>S L S S T A S A L G K L Q D V V N Q N A Q A L N T L V K Q L S S N F G A I S S V L N D I L S R L D K V E A E V Q I D R L I T G R L Q S L Q T 1006</mark> 1077 TAPAICHDGKAHFPREGVFVSNGTHWFVTQRNFYEPQIITTDNTFVSGNCDVVIGIVNNTVYDPLQ 1142