

Supplementary methods: nucleotide sequences of the constructs

1. $\Delta 11S$

```
1 ATGggccatc atcatcatca tcatcatcat ATGgttttta cacttgaaga ttttgtgggt
61 gattgggaac aaactgctgc atataattta gatcaagttt tagaacaggg tggagtgagt
121 tctcttttac aaaatcttgc agtctcagtt acaccaatac aaagaatagt tagaagtgga
181 gaaaatgcat taaagatcga tatacatgta ataattcctt atgagggact atcagcagac
241 caaatggcac aaattgagga agtattcaaa gttgtatatac cagtagacga tcatcacttt
301 aaagtaatat taccttatgg aacttttagta attgatgggtg taacaccaa taatgttaaat
361 tattttggta gaccatacga agggattgca gtttttgatg gaaagaaaat aaccgtaaca
421 ggcactttgt ggaatggaaa taaaataata gatgaaaggt taattacacc tgatTAA
```

2. $\beta 9-G$

```
1 ATGGGCTCCA TGCTGTTCCG AGTAACCATC AACAGTggag gtggatccGG TGGTGGAGGG
61 AGCacctaca aacttGtcAT Taacggtaaa accctgaaag gtgaaaccac caccgaagct
121 gttgacgctg ctaccgcgga aaaagttttc aaacagtacg ctaacgacaa cgggtgttgac
181 ggtgaatgga cctacgacga cgctaccaa acctTcaccg taacggaaGG TGGCGGTAGC
241 catcaccacc atcaccacTG A
```

3. G- $\beta 9$

```
1 ATGggccatc atcatcatca tcatGGTGGGA GGGAGCacct acaaacttGt cATTaacgggt
61 aaaaccctga aaggtgaaac caccaccgaa gctgttgacg ctgctaccgc ggaaaaagtt
121 ttcaaacagt acgctaacga caacgggtgtt gacgggtgaat ggacctacga cgacgctacc
181 aaaacctTca ccgtaacgga aGGTGGCGGT AGCggtggcg gaggctctGG CTCCATGCTG
241 TTCCGAGTAA CCATCAACAG Ttaa
```

4. $\beta 9-G-\beta 9$

```
1 ATGGGCTCCA TGCTGTTCCG AGTAACCATC AACAGTggag gtggatccGG TGGTGGAGGG
61 AGCacctaca aacttGtcAT Taacggtaaa accctgaaag gtgaaaccac caccgaagct
121 gttgacgctg ctaccgcgga aaaagttttc aaacagtacg ctaacgacaa cgggtgttgac
181 ggtgaatgga cctacgacga cgctaccaa acctTcaccg taacggaaGG AGGAGGATCT
241 GGAGGTGGAT CCGGCTCCAT GCTGTTCCGA GTAACCATCA ACAGTGGCGG AAGCCATCAT
301 CACCACCATC ATCACCATTG A
```

5. $\beta 10-RBD$

```
1 ATGCGGCTTC CGGGTGCGAT GCCAGCTCTG GCCCTCAAAG GCGAGCTGCT GTTGCTGTCT
61 CTCCTGTTAC TTCTGGAACC ACAGATCTCT CAGGGCGGTG GaGTGAGCGG CTGGCGGCTG
121 TTCAAGAAGA TTAGCGGTGG CGGAGGATCT GGAGGTGGAT CCaggggtgca gccaacggag
181 tctatcgtgc gctttcctaa taccacaaac ctgtgccat ttggcgaggt gttcaacgca
241 acccgcttcg ccagcgtgta cgctggaat aggaagcgga tcagcaactg cgtggccgac
301 tatagcgtgc tgtacaactc cgctctttc agcaccttta agtgctatgg cgtgtcccc
361 acaaagctga atgacctgtg ctttaccac gtctacgccg attctttcgt gatcaggggc
421 gacgaggtgc gccagatcgc ccccgccag acaggcaaga tcgcagacta caattataag
481 ctgccagacg atttcaccgg ctgcgtgatc gcctggaaca gcaacaatct ggattccaaa
541 gtgggcgga actacaatta tctgtaccgg ctgtttagaa agagcaatct gaagcccttc
601 gagagggaca tctctacaga aatctaccag gccggcagca ccccttgcaa tggcgtggag
661 ggctttaact gttatttccc actccagtc tacggcttc agcccacaaa cggcgtgggc
721 tatcagcctt accgctgggt ggtgctgagc tttgagctgc tgcacgccc agcaacagtg
781 tgcggcccca agaagtccac caatctggtg aagaacaagt gcgtgaactt cGGCGGAAGC
841 CATCATCACC ACCATCATCA CCATTGA
```

6. RBD- $\beta 10$

```
1 atgttcgtct tcttggctct gctgcctctg gtctcctcac agaggggtgca gccaacggag
61 tctatcgtgc gctttcctaa taccacaaac ctgtgccat ttggcgaggt gttcaacgca
121 acccgcttcg ccagcgtgta cgctggaat aggaagcgga tcagcaactg cgtggccgac
181 tatagcgtgc tgtacaactc cgctctttc agcaccttta agtgctatgg cgtgtcccc
241 acaaagctga atgacctgtg ctttaccac gtctacgccg attctttcgt gatcaggggc
```

301 gacgaggtgc gccagatcgc ccccggccag acaggcaaga tgcagacta caattataag
 361 ctgccagacg atttcaccgg ctgctgtatc gcctggaaca gcaacaatct ggattccaaa
 421 gtggggcggca actacaatta tctgtaccgg ctgttttagaa agagcaatct gaagcccttc
 481 gagagggaca tctctacaga aatctaccag gccggcagca ccccttgcaa tggcgtggag
 541 ggctttaact gttatttccc actccagtcc tacggcttcc agcccacaaa cggcgtgggc
 601 tatcagcctt accgcgtggt ggtgctgagc tttgagctgc tgcacgcccc agcaacagtg
 661 tgcggcccca agaagtccac caatctggtg aagaacaagt gcgtgaactt cGGCGGAGGA
 721 GGATCTGGAG GTGGATCCGT GAGCGGCTGG CGGCTGTTCA AGAAGATTAG CGGCGGAAGC
 781 CATCATCACC ACCATCATCA CCATTGA

7. β 10-RBD- β 10

1 ATGCGGCTTC CGGGTGCAT GCCAGCTCTG GCCCTCAAAG GCGAGCTGCT GTTGCTGTCT
 61 CTCCTGTTAC TTCTGGAACC ACAGATCTCT CAGGGCGGTG GaGTGAGCGG CTGGCGGCTG
 121 TTCAAGAAGA TTAGCGGTGG CGGAGGATCT GGAGGTGGAT CCagggtgca gccaacccag
 181 tctatcgtgc gctttcctaa tatcacaac ctgtgcccac ttggcgaggt gttcaacgca
 241 accgccttcg ccagcgtgta cgcttgaat aggaagcggg tcagcaactg cgtggccgac
 301 tatagcgtgc tgtacaactc cgctctttc agcaccttta agtgctatgg cgtgtcccc
 361 acaaagctga atgacctgtg ctttaccac gtctacgccc attctttcgt gatcaggggc
 421 gacgaggtgc gccagatcgc ccccggccag acaggcaaga tgcagacta caattataag
 481 ctgccagacg atttcaccgg ctgctgtatc gcctggaaca gcaacaatct ggattccaaa
 541 gtggggcggca actacaatta tctgtaccgg ctgttttagaa agagcaatct gaagcccttc
 601 gagagggaca tctctacaga aatctaccag gccggcagca ccccttgcaa tggcgtggag
 661 ggctttaact gttatttccc actccagtcc tacggcttcc agcccacaaa cggcgtgggc
 721 tatcagcctt accgcgtggt ggtgctgagc tttgagctgc tgcacgcccc agcaacagtg
 781 tgcggcccca agaagtccac caatctggtg aagaacaagt gcgtgaactt cGGCGGAGGA
 841 GGATCTGGAG GTGGATCCGT GAGCGGCTGG CGGCTGTTCA AGAAGATTAG CGGCGGAAGC
 901 CATCATCACC ACCATCATCA CCATTGA

8. β 10-S

1 ATGCGGCTTC CGGGTGCAT GCCAGCTCTG GCCCTCAAAG GCGAGCTGCT GTTGCTGTCT
 61 CTCCTGTTAC TTCTGGAACC ACAGATCTCT CAGGGCGGTG GaGTGAGCGG CTGGCGGCTG
 121 TTCAAGAAGA TTAGCGGTGG CGGAGGATCT GGAGGTGGAT CCtgcgtcaa tctgacaact
 181 cggactcagc tgccacctgc ttataactaat agcttcacca gaggcgtgta ctatcctgac
 241 aagggtgttta gaagctccgt gctgcactct acacaggatc tgtttctgcc attcttttagc
 301 aacgtgacct ggttccacgc catccacgtg agcggcacca atggcacaaa gcggttcgac
 361 aatcccgtgc tgctttttaa cgatggcgtg tacttgcct ctaccgagaa gagcaacatc
 421 atcagaggct ggatctttgg caccacactg gactccaaga cacagtctct gctgatcgtg
 481 aacaatgcca ccaacgtggt catcaagggtg tgcgagttcc agttttgtaa tgatcccttc
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3421	cactggtttg	tgacacagcg	caatttctac	gagccccaga	tcataccacc	agataacacc
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3781	CCCGAGGCC	CtCGCGAtGG	aCAGGctTAC	GTGCGCAAGG	ACGGCGAGTG	GGTGTCTGCTG
3841	AGCACCTTCC	TGGGCGGAAG	CCATCATCAC	CACCATCATC	ACCATTGA	

9. S-β10

1	atgttCGTct	tcctgggtcct	gctgcctctg	gtctcctcac	agtgcgtcaa	tctgacaact
61	cggactcagc	tgccacctgc	ttataactaat	agcttcacca	gaggcgtgta	ctatcctgac
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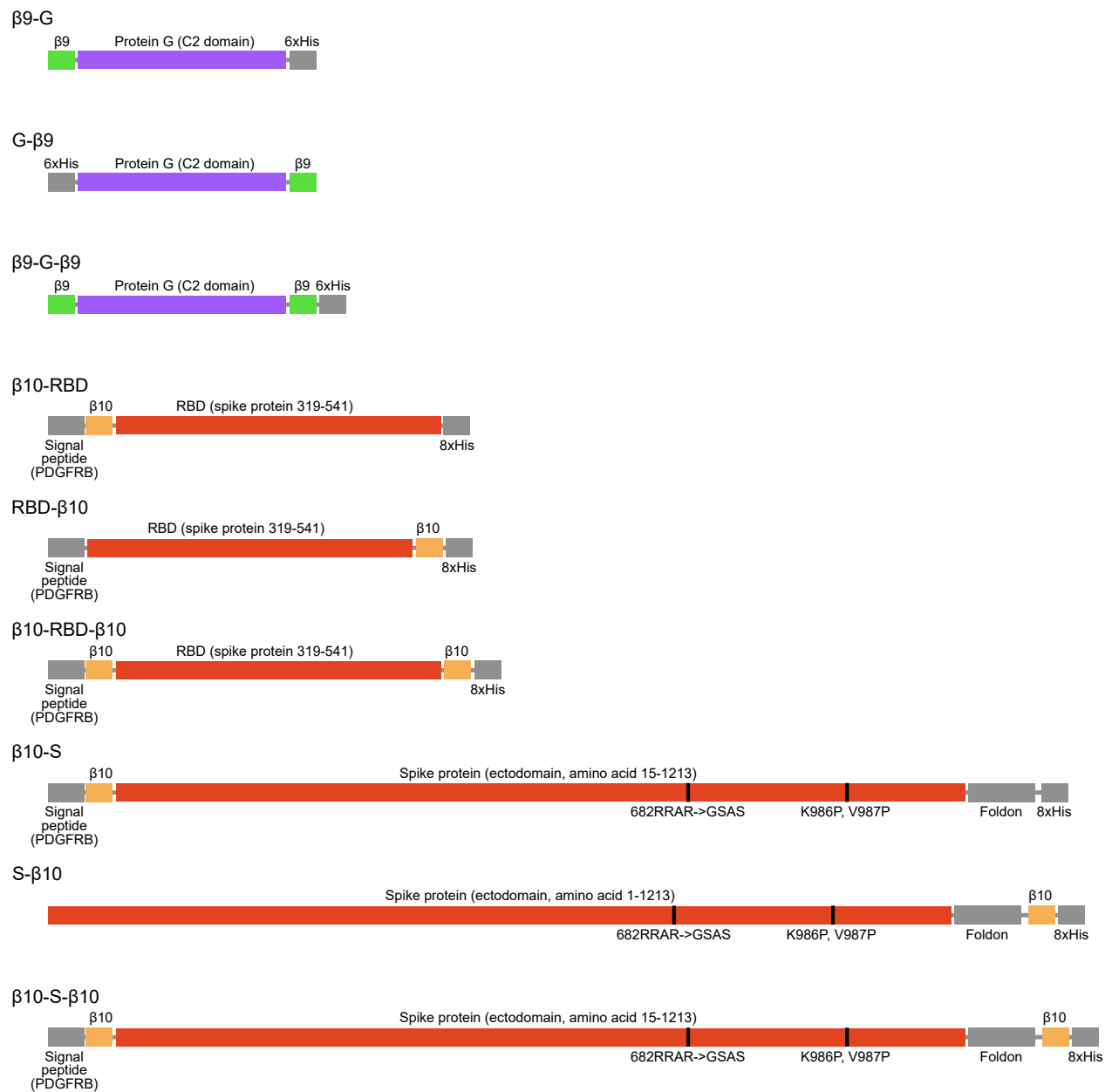
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1741 accctggaga tcctggacat cacaccctgc tctttcggcg gcgtgagcgt gatcacacc
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3601 caggagctgg gcaagtatga gcagtacatc aagtggcccG GaGGCAGCGG aGGCTACATC
3661 CCCGAGGCCC CtCGCGAtGG aCAGGctTAC GTGCGCAAGG ACGGCGAGTG GGTGCTGCTG
3721 AGCACCTTCC TGGGCGGAGG AGGATCTGGA GGTGGATCCG TGAGCGGCTG GCGGCTGTTC
3781 AAGAAGATTA GCGGCGGAAG CCATCATCAC CACCATCATC ACCATTGA

10. β 10-S- β 10

1 ATGCGGCTTC CGGGTGCAT GCCAGCTCTG GCCCTCAAAG GCGAGCTGCT GTTGCTGTCT
61 CTCCTGTTAC TTCTGGAACC ACAGATCTCT CAGGGCGGTG GaGTGAGCGG CTGGCGGCTG
121 TTCAAGAAGA TTAGtGGTGG CGGAGGATCT GGAGGTGGAT Cctgctcaa tctgacaact
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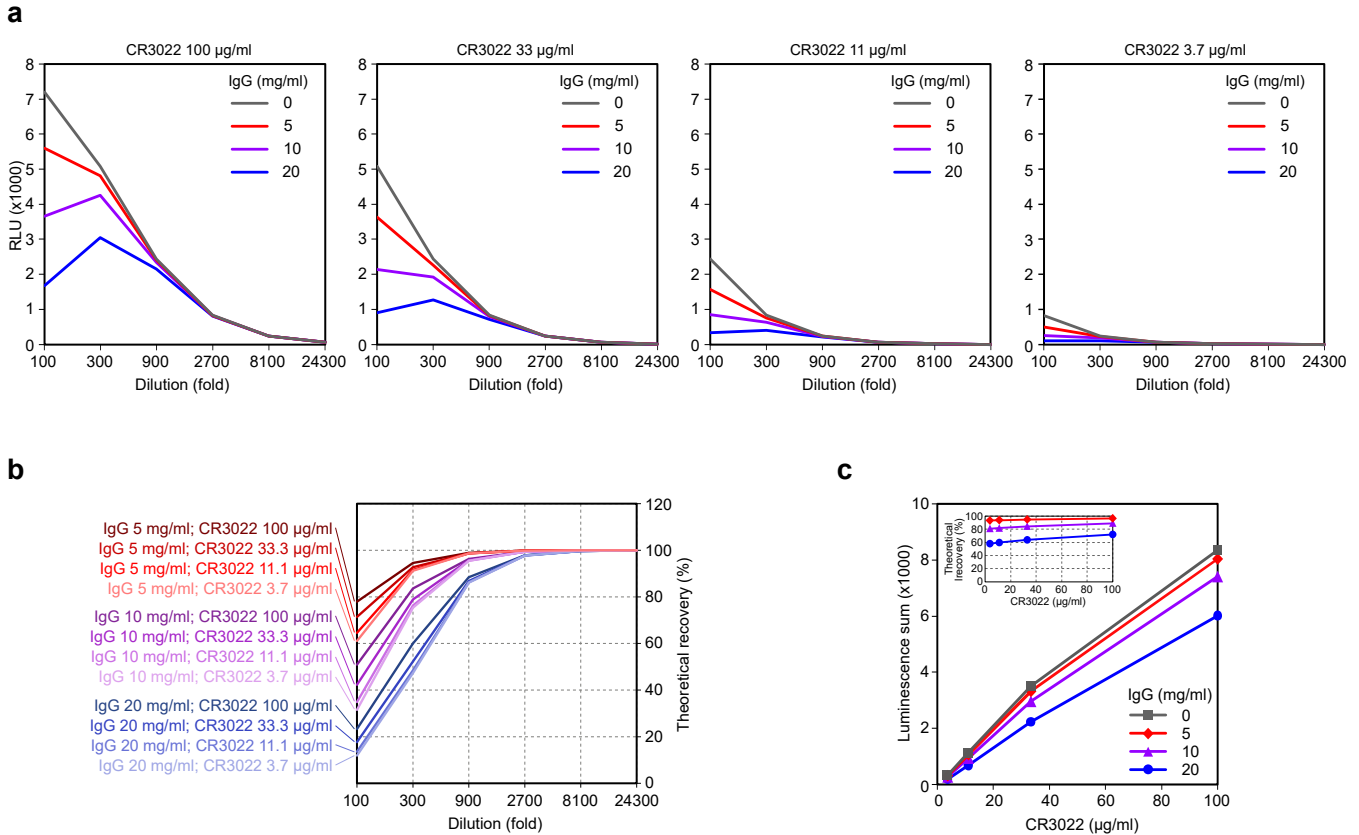
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Supplementary Fig. 1



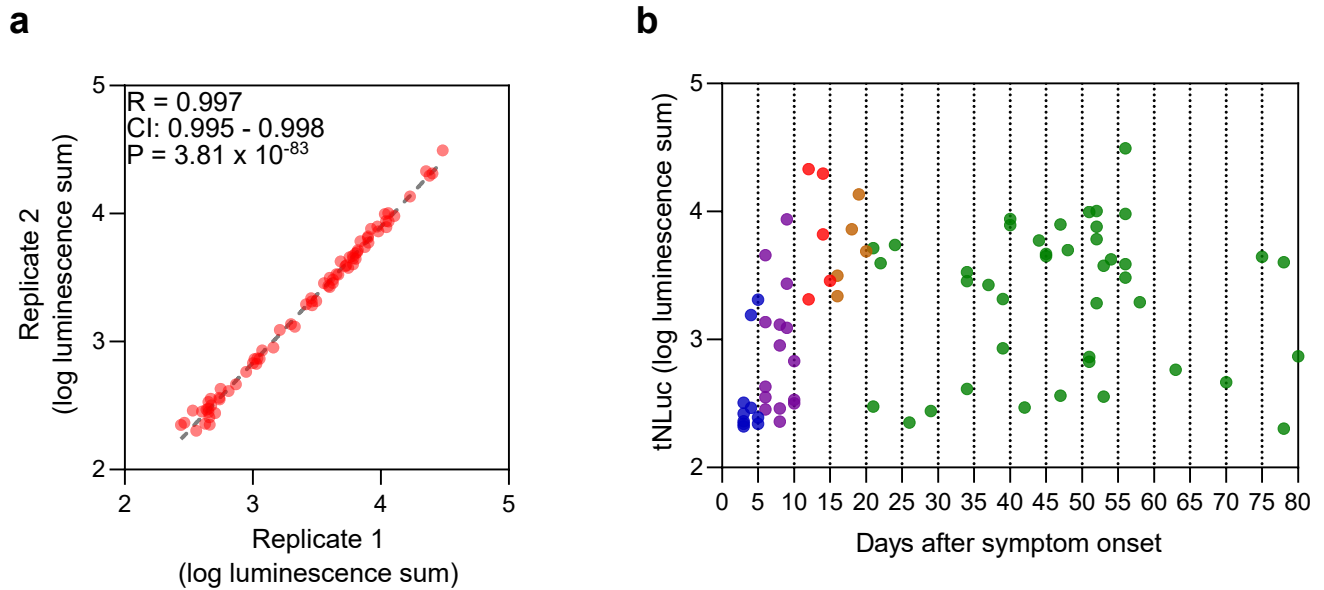
Supplementary Figure 1. Schematic representation of SATiN probes.

Supplementary Fig. 2



Supplementary Figure 2. Computational simulation of inhibitory effects of IgG on the SATIN assay. **A.** Virtual samples containing CR3022 (3.7, 11, 33 and 100 $\mu\text{g}/\text{mL}$) in the presence of IgG at different levels (5, 10, and 20 mg/mL) were serially diluted as indicated. Their tNLuc signals at each dilution endpoint were calculated using the chemical kinetics model derived from the experimental data shown in **Fig. 2B**. **B.** Theoretical recovery rates of samples in **A** at different dilutions were calculated by dividing the calculated reading of a virtual sample in **A** by the corresponding virtual sample not containing IgG. **C.** The overall tNLuc signal of each virtual sample was calculated by luminescence summation of the calculated signals at the 1:300, 1:900 and 1:2700 dilutions obtained in **A**. Overall recovery rates are presented inset.

Supplementary Figure 3



Supplementary Figure 3. Detection of α -SARS CoV-2 antibody in serum samples with the SATiN assay. **A.** 70 serum samples collected from COVID-19 patients or convalescents were analyzed in two independent replicates using the SATiN assay. The two sets of data are presented as a scatter plot. R and P values were obtained from two tailed Pearson analysis. **B.** SATiN luminescence signal of 80 samples plotted against the time of sample collection.