

## **SARS-CoV-2 RBD-Tetanus toxoid conjugate vaccine induces a strong neutralizing immunity in preclinical studies**

**Yury Valdes-Balbin,<sup>1\*</sup> Darielys Santana-Mederos,<sup>1\*</sup> Lauren Quintero,<sup>1</sup> Sonsire Fernández,<sup>1\*</sup> Laura Rodriguez,<sup>1</sup> Belinda Sanchez Ramirez,<sup>2\*</sup> Rocmira Perez,<sup>1</sup> Claudia Acosta,<sup>1</sup> Yanira Méndez,<sup>3</sup> Manuel G. Ricardo,<sup>3</sup> Tays Hernandez<sup>2</sup>, Gretchen Bergado,<sup>2</sup> Franciscary Pi,<sup>2</sup> Annet Valdes,<sup>2</sup> Tania Carmentate,<sup>2</sup> Ubel Ramirez,<sup>1</sup> Reinaldo Oliva,<sup>1</sup> Jean-Pierre Soubal,<sup>1</sup> Raine Garrido,<sup>1</sup> Felix Cardoso,<sup>1</sup> Mario Landys,<sup>1</sup> Humberto Gonzalez,<sup>1</sup> Mildrey Farinas,<sup>1</sup> Juliet Enriquez,<sup>4</sup> Enrique Noa,<sup>4</sup> Anamary Suarez,<sup>4</sup> Cheng Fang,<sup>5</sup> Luis A. Espinosa,<sup>6</sup> Yassel Ramos,<sup>6</sup> Luis Javier González,<sup>6</sup> Yanet Climent,<sup>1</sup> Gertrudis Rojas,<sup>2</sup> Ernesto Relova-Hernández,<sup>2</sup> Yanelys Cabrera Infante,<sup>2</sup> Sum Lai Losada,<sup>2</sup> Tammy Boggiano,<sup>2</sup> Eduardo Ojito,<sup>2</sup> Kalet Leon Monzon,<sup>2</sup> Fabrizio Chiodo,<sup>1,7</sup> Françoise Paquet,<sup>8</sup> Guang-Wu Chen,<sup>9\*</sup> Daniel G. Rivera,<sup>3\*</sup> Dagmar Garcia-Rivera,<sup>1\*</sup> Vicente Verez-Bencomo<sup>1\*</sup>**

<sup>1</sup>*Finlay Vaccine Institute, 200 and 21 Street, Havana 11600, Cuba.*

<sup>2</sup>*Center of Molecular Immunology, P.O. Box 16040, 216 St. Havana, Cuba*

<sup>3</sup>*Laboratory of Synthetic and Biomolecular Chemistry, Faculty of Chemistry, University of Havana, Cuba*

<sup>4</sup>*National Civil Defense Research Laboratory, Cuba*

<sup>5</sup> *Shanghai Fenglin Glycodrug Promotion Center, Shanghai 200032, China*

<sup>6</sup>*Center for Genetic Engineering and Biotechnology, Ave 31 e/ 158 y 190, Playa, Havana, Cuba*

<sup>7</sup>*Department of Molecular Cell Biology and Immunology, Amsterdam UMC, Vrije Universiteit Amsterdam, Amsterdam, The Netherlands and Institute of Biomolecular Chemistry, National Research Council (CNR), Pozzuoli, Napoli, Italy*

<sup>8</sup>*Centre de Biophysique Moléculaire, CNRS UPR 4301, rue Charles Sadron, F-45071, Orléans, Cedex 2, France*

<sup>9</sup>*Chengdu Olisynn Biotech. Co. Ltd., and State Key Laboratory of Biotherapy and Cancer Center, West China Hospital, Sichuan University, Chengdu 610041, People's Republic of China.*

Correspondence: [yvbalbin@finlay.edu.cu](mailto:yvbalbin@finlay.edu.cu), [dagarcia@finlay.edu.cu](mailto:dagarcia@finlay.edu.cu), [vicente.verez@finlay.edu.cu](mailto:vicente.verez@finlay.edu.cu)

\*These authors contributed equally.

## **Abstract**

Controlling the global COVID-19 pandemic depends, among other measures, on developing preventive vaccines at an unprecedented pace. Vaccines approved for use and those in development intend to use neutralizing antibodies to block viral sites binding to the host's cellular receptors. Virus infection is mediated by the spike glycoprotein trimer on the virion surface via its receptor binding domain (RBD). Antibody response to this domain is an important outcome of the immunization and correlates well with viral neutralization. Here we show that macromolecular constructs with recombinant RBD conjugated to tetanus toxoid induce a potent immune response in laboratory animals. Some advantages of the immunization with the viral antigen coupled to tetanus toxoid have become evident such as predominant IgG immune response due to affinity maturation and long-term specific B-memory cells. This paper demonstrates that subunit conjugate vaccines can be an alternative for COVID-19, paving the

way for other viral conjugate vaccines based on the use of small viral proteins involved in the infection process.

## **Introduction**

Control of SARS-CoV-2 infection focuses on development of preventive vaccines.<sup>1</sup> Viral particles' initial binding is mediated by the receptor binding domain (RBD) of the spike (S)-glycoprotein trimer to the host's cell surface receptor, the angiotensin-converting enzyme 2 (ACE2).<sup>2-5</sup> Most of the 200 COVID-19 vaccines in development<sup>6</sup> aim to block this process.<sup>1</sup> By focusing on the whole S-protein or its RBD as antigen, the primary goal is induction of anti-RBD antibodies that interfere with RBD-ACE2 interaction, blocking the first step of infection. Virus neutralization is mainly associated with antibodies against the receptor binding motif (RBM), a specific RBD region directly interacting with ACE2.<sup>7</sup> This type of antibodies are not involved in antibody dependent enhancement (ADE).<sup>8</sup>

Key advantages of the well-known recombinant subunit vaccine platforms are their safety, stability at 2-8 °C and facility to scale-up the production.<sup>9</sup> While we should expect weak immunogenicity for such a small recombinant RBD protein (30 kDa), requiring repeated vaccination,<sup>10</sup> it was found that recombinant RBD in alum is sufficient to induce a neutralizing immune response in laboratory animals,<sup>11</sup> while its simplicity prompted subsequent evaluation in humans.<sup>12</sup> However, most recombinant vaccines relied on large, RBD-containing macromolecular constructs as a way to increase the immunogenicity and on the use of potent adjuvants.<sup>13</sup> In addition to the lower immunogenicity, small recombinant RBD exposes to the immune system not only the critical RBM surface but also RBD regions that are well-camouflaged at the virus surface. Antibodies directed to camouflaged RBD regions are not neutralizing, and therefore, an

ideal construction should maximize the exposure of the RBM and not of such regions. We hypothesized that the orientation of RBD when conjugated to tetanus toxoid, exposes better the RBM surface increasing the level of neutralizing antibodies.<sup>14-17</sup>

The SARS-CoV-2 RBD comprises 193 amino acid residues from Thr333 to Pro527, including RBM 438-506 that interacts directly with ACE2. It contains eight cysteines forming four disulfide bridges, three of these stabilizing the RBD core and one within the RBM.<sup>6</sup> Our recombinant RBD 319-541 was obtained in CHO-cells with intentionally extended sequence adding S-glycoprotein residues 527 through 541, in order to include an additional Cys538. This cysteine is usually connected to Cys590 in the S-glycoprotein. The extended sequence includes two N-glycosylation sites at residues Asn331 and Asn343 and two O-glycosylation sites at Thr323 and Ser325. The selected sequence results in an unpaired Cys538 intended to be used for chemical conjugation to the highly immunogenic carrier tetanus toxoid (TT). Here we report a promising vaccine candidate based on this high molecular weight conjugate with several copies of recombinant RBD per molecular unit. To our knowledge, chemically conjugated constructs and the immunogenic effect of conjugating viral proteins such as RBD to a protein carrier have not been assessed for SARS-CoV-2 or other coronaviruses. Here we demonstrate that the RBD-TT conjugate induces a potent immune response in laboratory animals, paving the way for their evaluation in human phase I and II clinical trials.<sup>18</sup>

### **Construction of RBD-TT conjugates**

Our design is based on the hypothesis that by conjugating several copies of the extended RBD to a large carrier protein, we can obtain a macromolecular construct displaying multivalent RBD. At

the same time, the RBM will be well exposed (Fig. 1, represented in red) and more available for immune recognition.

Inclusion of an additional free Cys538 in our extended RBD, while potentially useful for conjugation, could jeopardize extended-RBD folding, due to potential S-S rearrangement with the other 8 cysteines (scrambling). Nevertheless, we found that during fermentation and purification, Cys538 is spontaneously protected through an S-S Cys adduct with free cysteine present in the culture media. ESI-MS showed presence of the four native S-S bonds, indicating a correctly folded extended RBD (Fig. S2). Cysteinylation of Cys538 was selectively reduced to free thiol with tris-(2-carboxyethyl)phosphine (TCEP)<sup>19</sup> without affecting ACE2 recognition, while—for example—dithiothreitol (DTT) led to complete loss of its ACE2 binding capacity, suggesting loss of the antigen's 3D structure (Fig. 1B).

To our knowledge, the immunogenic effect of TT as a carrier has not been assessed previously for SARS-CoV-2 or any other coronavirus. We have successfully used TT as a carrier protein for antibacterial carbohydrate-protein conjugate vaccines.<sup>20,21</sup> The presence of multiple T- and B-cell epitopes of this highly immunogenic carrier<sup>22</sup> might potentiate cellular immunity when compared to use of RBD alone. In addition, multimeric RBD-TT can simultaneously activate several B-cell receptors, thus enhancing B cell response.<sup>23</sup>

TT was activated with an average of 20–30 maleimide groups per mol of TT by reaction with *N*-succinimidyl 3-maleimidopropionate (SMP) followed by reaction with 2.5 or, alternatively, 10 equivalents of TCEP-reduced extended RBD, to produce conjugates bearing 2 or 6 mol, respectively, of RBD per mol of TT (Fig. 1c). The RBD<sub>2</sub>-TT and RBD<sub>6</sub>-TT conjugates were produced under good manufacturing practices (GMP) in 72% and 64% yield, respectively, and characterized by SE-HPLC and MS. Both conjugates recognize ACE2 slightly better than the

original RBD, (Fig. 1E), confirming preservation of their structure and, probably, a better exposition of RBM. As convalescent serum usually contains TT antibodies, we prepared a RBD-bovine serum albumin (BSA) conjugate incorporating 6 RBD units per mol of BSA (RBD<sub>6</sub>-BSA), which was recognized well by various convalescent sera, proving conservation of the RBD antigenic properties after conjugation (Fig. 1D).

### **Animal Immunogenicity**

Immunization of BALB/c mice with the four different immunogens (Fig. 2) induces a strong IgG RBD-specific immune response as proven by ELISA. RBD<sub>6</sub>-TT/alum and RBD<sub>2</sub>-TT/alum were compared with RBD alone and with RBD<sub>2</sub>-TT without alum. After the first dose (T7 and T14, Fig. 2C) RBD<sub>6</sub>-TT/alum induces the highest level of anti-RBD antibodies. After the second dose all immunogens adsorbed in alum elicit better anti-RBD IgG levels than without alum (T21 and 28, Fig. 2C). The high and homogeneous early response for RBD<sub>6</sub>-TT/alum could be an important attribute for a vaccine in pandemic times. We explored early response to different dosage of RBD<sub>6</sub>-TT/alum, finding a dose-dependent response at day 7. At day 14, before the second dose, the response was very high even for the lowest dosage (T14, Fig. 2C).

To evaluate possible immunological advantages of the RBD-TT conjugate, we studied affinity maturation, of antibodies elicited by RBD<sub>6</sub>-TT compared with the rest of immunogens. There was an increase in the avidity index (AI).<sup>24</sup> The highest value of 81% for antibodies induced by RBD<sub>6</sub>-TT is consistent with a more pronounced affinity maturation. (Fig. 2D). The Th1/Th2 balance can be modulated by vaccination and was also evaluated (Fig. 2E). A biased Th2 immune response was observed for RBD<sub>2</sub>-TT/alum (IgG2a/IgG1 ratio 0.54) and RBD/alum (IgG2a/IgG1

ratio 0.40), while RBD<sub>6</sub>-TT/alum displayed more balanced Th1/Th2 immunity (IgG2a/IgG1 ratio 0.81).

Fig. 3 shows the induction of memory antigen-specific B and T-cells, an important property of conjugate vaccines. Mice immunized with conjugate RBD<sub>6</sub>-TT/alum were compared to mice receiving RBD/alum. Both groups developed a primary immunity as shown previously (Fig. 2B). After two doses at T28, splenocytes purified from both groups were intravenously transferred to naïve mice that were then boosted by a single dose of 3 µg RBD/alum (Fig. 3B). Mice receiving splenocytes from RBD<sub>6</sub>-TT/alum responded with a strong secondary RBD-specific IgG response ( titer 10<sup>3</sup>-10<sup>4</sup>), while those receiving splenocytes from RBD/alum did not (results not shown). This finding demonstrated the presence of RBD-specific memory B cells in transferred splenocytes, which were able to be activated in the presence of RBD/alum (alternative SARS-CoV-2 virus) stimuli.

Specific CD8<sup>+</sup> T cells also play an important role in protection, as recently demonstrated.<sup>25</sup> To evaluate the specific T-cell response, we compared RBD<sub>6</sub>-TT/alum and RBD/alum immunized mice. After *in vitro* RBD stimuli, splenocytes from mice immunized with RBD<sub>6</sub>-TT secreted higher levels of IFN $\gamma$  compared to those immunized with RBD/alum (Fig. 3D), suggesting a Th1 pattern, while IL-4 (characteristic of Th2 pattern) and IL-17A (characteristic of Th17 pattern) were not detected. Frequency of CD8<sup>+</sup>CD44<sup>high</sup> memory T-lymphocytes producing IFN- $\gamma$ , TNF- $\alpha$  and Granzyme B increased significantly in RBD<sub>6</sub>-TT immunized mice with respect to control mice (Fig. 3E) as shown by flow cytometry, indicating the relevant activation of cytotoxic T immune memory.

### **Antibody functionality**

We evaluated antibodies' ability to block interaction between the virus and its receptor,<sup>10</sup> using the molecular Virus Neutralization Test (mVNT50)<sup>22</sup> and the conventional Virus Neutralization Test (cVNT50).<sup>23</sup> mVNT50 evaluates inhibition of interaction between recombinant RBD and ACE2 at the molecular level; at the cellular level, cVNT50 evaluates inhibition of interaction between the live virus and Vero E6 cells bearing ACE2 receptors. Antibodies resulting from immunization of Balb/c mice with two doses of RBD<sub>2</sub>-TT/alum and RBD<sub>6</sub>-TT/alum were compared to antibodies elicited after immunization with RBD/alum. mVNT50 showed a high level of inhibition for all sera (Fig. 4a), indicating that all tested antibodies displayed a similar efficacy in interfering with RBD-ACE2 interaction at the molecular level. cVNT50 (Fig. 4B) showed sharp differences between sera from animals immunized with RBD/alum and those with both conjugates. For RBD/alum, the neutralization titer was 232; for both conjugates, there was a higher level of virus neutralization: 1303 for RBD<sub>2</sub>-TT and 2568 for RBD<sub>6</sub>-TT. The mVNT50/cVNT50 ratio was 0.143, 0.732 and 1.08 for RBD, RBD<sub>2</sub>-TT and RBD<sub>6</sub>-TT respectively. While antibodies neutralizing the virus are mainly directed at the RBM,<sup>7</sup> there are antibodies recognizing soluble RBD not only by the RBM but also on a different region, as shown in Fig. 4C. This type of "lateral" antibodies could interfere in mVNT50 with soluble RBD, but probably will not recognize this RBD region that is camouflaged at the virus surface.

Based on the results presented here, GMP batches of the conjugates RBD<sub>2</sub>-TT and RBD<sub>6</sub>-TT were obtained and absorbed on alum as final vaccine candidates. A phase I clinical trial<sup>24</sup> was initiated in October 2020, and after preliminary results confirming a better performance of a vaccine based on RBD<sub>6</sub>-TT/alum, this advanced on December 2020 to a phase II clinical trial with 910 subjects.<sup>25</sup> The encouraging interim results of all those trial envisage moving forward to a phase III trial in March 2021. The resulting vaccine, the first one developed and produced in a Latin-American country, has important advantages to be considered in the future: a) it induces a



strong IgG neutralizing antibody as well as specific T-cell response, b) the well-known safety record of this vaccine platform is further confirmed during clinical trial, c) the storage and distribution temperature of 2-8 °C is the conventional for other vaccines and suitable for a rapid distribution worldwide, and finally, d) it can be adapted to existing vaccine production capacities available at several countries. We hope this vaccine can contribute to win the battle against the SARS-CoV-2 pandemic.

## References

1. F. Krammer, SARS-CoV-2 vaccines in development. *Nature* **586**, 516-527 (2020).
2. Y. Chen, Y. Guo, Y. Pan, Z. J. Zhao, Structure analysis of the receptor binding of 2019-nCoV. *Biochem. Biophys. Res. Commun.* **525**, 135-140 (2020).
3. J. Shang, G. Ye, K. Shi, Y. Wan, C. Luo, H. Aihara, Q. Geng, A. Auerbach, F. Li, Structural basis of receptor recognition by SARS-CoV-2. *Nature* **581**, 221-224 (2020).
4. A. C. Walls, Y. J. Park, M. A. Tortorici, A. Wall, A. T. McGuire, D. Velesler, Structure, Function, and Antigenicity of the SARS-CoV-2 Spike Glycoprotein. *Cell* **181**, 281-292 e286 (2020).
5. Y. Wan, J. Shang, R. Graham, R. S. Baric, F. Li, Receptor Recognition by the Novel Coronavirus from Wuhan: an Analysis Based on Decade-Long Structural Studies of SARS Coronavirus. *J. Virol.* **94**, e00127-00120 (2020).
6. J. Lan, J. Ge, J. Yu, S. Shan, H. Zhou, S. Fan, Q. Zhang, X. Shi, Q. Wang, L. Zhang, X. Wang, Structure of the SARS-CoV-2 spike receptor-binding domain bound to the ACE2 receptor. *Nature* **581**, 215-220 (2020).
7. L. Piccoli, Y. J. Park, M. A. Tortorici, N. Czudnochowski, A. C. Walls, M. Beltramello, C. Silacci-Fregni, D. Pinto, L. E. Rosen, J. E. Bowen, O. J. Acton, S. Jaconi, B. Guarino, A. Minola,

- F. Zatta, N. Sprugasci, J. Bassi, A. Peter, A. De Marco, J. C. Nix, F. Mele, S. Jovic, B. F. Rodriguez, S. V. Gupta, F. Jin, G. Piumatti, G. Lo Presti, A. F. Pellanda, M. Biggiogero, M. Tarkowski, M. S. Pizzuto, E. Cameroni, C. Havenar-Daughton, M. Smithey, D. Hong, V. Lepori, E. Albanese, A. Ceschi, E. Bernasconi, L. Elzi, P. Ferrari, C. Garzoni, A. Riva, G. Snell, F. Sallusto, K. Fink, H. W. Virgin, A. Lanzavecchia, D. Corti, D. Veessler, Mapping Neutralizing and Immunodominant Sites on the SARS-CoV-2 Spike Receptor-Binding Domain by Structure-Guided High-Resolution Serology. *Cell* **183**, 1024-1042 e1021 (2020).
8. J. Zang, C. Gu, B. Zhou, C. Zhang, Y. Yang, S. Xu, L. Bai, R. Zhang, Q. Deng, Z. Yuan, H. Tang, D. Qu, D. Lavillette, Y. Xie, Z. Huang, Immunization with the receptor-binding domain of SARS-CoV-2 elicits antibodies cross-neutralizing SARS-CoV-2 and SARS-CoV without antibody-dependent enhancement. *Cell Discov.* **6**, 61 (2020).
9. M. Muhuri, G. Gao, Is smaller better? Vaccine targeting recombinant receptor-binding domain might hold the key for mass production of effective prophylactics to fight the COVID-19 pandemic. *Signal Transduct. Target. Ther.* **5**, 222 (2020).
10. M. Jeyanathan, S. Afkhami, F. Smaill, M. S. Miller, B. D. Lichty, Z. Xing, Immunological considerations for COVID-19 vaccine strategies. *Nat. Rev. Immunol.* **20**, 615-632 (2020).
11. J. Yang, W. Wang, Z. Chen, S. Lu, F. Yang, Z. Bi, L. Bao, F. Mo, X. Li, Y. Huang, W. Hong, Y. Yang, Y. Zhao, F. Ye, S. Lin, W. Deng, H. Chen, H. Lei, Z. Zhang, M. Luo, H. Gao, Y. Zheng, Y. Gong, X. Jiang, Y. Xu, Q. Lv, D. Li, M. Wang, F. Li, S. Wang, G. Wang, P. Yu, Y. Qu, L. Yang, H. Deng, A. Tong, J. Li, Z. Wang, J. Yang, G. Shen, Z. Zhao, Y. Li, J. Luo, H. Liu, W. Yu, M. Yang, J. Xu, J. Wang, H. Li, H. Wang, D. Kuang, P. Lin, Z. Hu, W. Guo, W. Cheng, Y. He, X. Song, C. Chen, Z. Xue, S. Yao, L. Chen, X. Ma, S. Chen, M. Gou, W. Huang, Y. Wang, C. Fan, Z. Tian, M. Shi, F.-S. Wang, L. Dai, M. Wu, G. Li, G. Wang, Y. Peng, Z. Qian, C. Huang, J. Y.-N. Lau, Z. Yang, Y. Wei, X. Cen, X. Peng, C. Qin, K. Zhang, G. Lu, X. Wei, A

- vaccine targeting the RBD of the S protein of SARS-CoV-2 induces protective immunity. *Nature* **586**, 572-577 (2020).
12. Clinical trial phase II NCT04718467, Clinical trial phase II RPCEC00000306, Clinical trial phase I NCT04522089.
  13. C. Keech, G. Albert, P. Reed, S. Neal, J. S. Plested, M. Zhu, S. Cloney-Clark, H. Zhou, N. Patel, M. B. Frieman, R. E. Haupt, J. Logue, M. McGrath, S. Weston, P. A. Piedra, I. Cho, A. Robertson, C. Desai, K. Callahan, M. Lewis, P. Price-Abbott, N. Formica, V. Shinde, L. Fries, J. D. Linkliter, P. Griffin, B. Wilkinson, G. Smith, G. M. Glenn, First-in-Human Trial of a SARS CoV 2 Recombinant Spike Protein Nanoparticle Vaccine. *medRxiv*, 2020.2008.2005.20168435 (2020).
  14. P. J. M. Brouwer, T. G. Caniels, K. van der Straten, J. L. Snitselaar, Y. Aldon, S. Bangaru, J. L. Torres, N. M. A. Okba, M. Claireaux, G. Kerster, A. E. H. Bentlage, M. M. van Haaren, D. Guerra, J. A. Burger, E. E. Schermer, K. D. Verheul, N. van der Velde, A. van der Kooi, J. van Schooten, M. J. van Breemen, T. P. L. Bijl, K. Sliepen, A. Aartse, R. Derking, I. Bontjer, N. A. Kootstra, W. J. Wiersinga, G. Vidarsson, B. L. Haagmans, A. B. Ward, G. J. de Bree, R. W. Sanders, M. J. van Gils, Potent neutralizing antibodies from COVID-19 patients define multiple targets of vulnerability. *Science* **369**, 643-650 (2020).
  15. T. F. Rogers, F. Zhao, D. Huang, N. Beutler, A. Burns, W. T. He, O. Limbo, C. Smith, G. Song, J. Woehl, L. Yang, R. K. Abbott, S. Callaghan, E. Garcia, J. Hurtado, M. Parren, L. Peng, S. Ramirez, J. Ricketts, M. J. Ricciardi, S. A. Rawlings, N. C. Wu, M. Yuan, D. M. Smith, D. Nemazee, J. R. Teijaro, J. E. Voss, I. A. Wilson, R. Andrabi, B. Briney, E. Landais, D. Sok, J. G. Jardine, D. R. Burton, Isolation of potent SARS-CoV-2 neutralizing antibodies and protection from disease in a small animal model. *Science* **369**, 956-963 (2020).

16. M. Yuan, H. Liu, N. C. Wu, C. D. Lee, X. Zhu, F. Zhao, D. Huang, W. Yu, Y. Hua, H. Tien, T. F. Rogers, E. Landais, D. Sok, J. G. Jardine, D. R. Burton, I. A. Wilson, Structural basis of a shared antibody response to SARS-CoV-2. *Science* **369**, 1119-1123 (2020).
17. S. J. Zost, P. Gilchuk, J. B. Case, E. Binshtein, R. E. Chen, J. P. Nkolola, A. Schafer, J. X. Reidy, A. Trivette, R. S. Nargi, R. E. Sutton, N. Suryadevara, D. R. Martinez, L. E. Williamson, E. C. Chen, T. Jones, S. Day, L. Myers, A. O. Hassan, N. M. Kafai, E. S. Winkler, J. M. Fox, S. Shrihari, B. K. Mueller, J. Meiler, A. Chandrashekar, N. B. Mercado, J. J. Steinhardt, K. Ren, Y. M. Loo, N. L. Kallewaard, B. T. McCune, S. P. Keeler, M. J. Holtzman, D. H. Barouch, L. E. Gralinski, R. S. Baric, L. B. Thackray, M. S. Diamond, R. H. Carnahan, J. E. Crowe, Jr., Potently neutralizing and protective human antibodies against SARS-CoV-2. *Nature* **584**, 443-449 (2020).
18. IFV/COR/06. <https://rpcec.sld.cu/trials/RPCEC00000347>
19. D. J. Cline, S. E. Redding, S. G. Brohawn, J. N. Psathas, J. P. Schneider, C. Thorpe, New water-soluble phosphines as reductants of peptide and protein disulfide bonds: reactivity and membrane permeability. *Biochemistry* **43**, 15195-15203 (2004).
20. V. Verez-Bencomo, V. Fernandez-Santana, E. Hardy, M. E. Toledo, M. C. Rodriguez, L. Heynngnezz, A. Rodriguez, A. Baly, L. Herrera, M. Izquierdo, A. Villar, Y. Valdes, K. Cosme, M. L. Deler, M. Montane, E. Garcia, A. Ramos, A. Aguilar, E. Medina, G. Torano, I. Sosa, I. Hernandez, R. Martinez, A. Muzachio, A. Carmenates, L. Costa, F. Cardoso, C. Campa, M. Diaz, R. Roy, A synthetic conjugate polysaccharide vaccine against Haemophilus influenzae type b. *Science* **305**, 522-525 (2004).
21. C. P. Dotres, R. Puga, Y. Ricardo, C. R. Brono, B. Paredes, V. Echemendia, S. Rosell, N. Gonzalez, D. Garcia-Rivera, Y. Valdes, D. Goldblatt, V. Verez-Bencomo, G. Laboratory-Pneumococci, G. Havana-Pneumococci, Safety and preliminary immunogenicity of Cuban

- pneumococcal conjugate vaccine candidate in healthy children: a randomized phase I clinical trial. *Vaccine* **32**, 5266-5270 (2014).
22. M. E. Pichichero, Protein carriers of conjugate vaccines: characteristics, development, and clinical trials. *Hum Vaccin Immunother* **9**, 2505-2523 (2013).
  23. A. C. Walls, B. Fiala, A. Schafer, S. Wrenn, M. N. Pham, M. Murphy, L. V. Tse, L. Shehata, M. A. O'Connor, C. Chen, M. J. Navarro, M. C. Miranda, D. Pettie, R. Ravichandran, J. C. Kraft, C. Ogohara, A. Palser, S. Chalk, E. C. Lee, K. Guerriero, E. Kepl, C. M. Chow, C. Sydeman, E. A. Hodge, B. Brown, J. T. Fuller, K. H. Dinnon, 3rd, L. E. Gralinski, S. R. Leist, K. L. Gully, T. B. Lewis, M. Guttman, H. Y. Chu, K. K. Lee, D. H. Fuller, R. S. Baric, P. Kellam, L. Carter, M. Pepper, T. P. Sheahan, D. Veessler, N. P. King, Elicitation of Potent Neutralizing Antibody Responses by Designed Protein Nanoparticle Vaccines for SARS-CoV-2. *Cell* **183**, 1367-1382 e1317 (2020).
  24. D. Goldblatt, A. R. Vaz, E. Miller, Antibody avidity as a surrogate marker of successful priming by Haemophilus influenzae type b conjugate vaccines following infant immunization. *J. Infect. Dis.* **177**, 1112-1115 (1998).
  25. K. McMahan, J. Yu, N. B. Mercado, C. Loos, L. H. Tostanoski, A. Chandrashekar, J. Liu, L. Peter, C. Atyeo, A. Zhu, E. A. Bondzie, G. Dagotto, M. S. Gebre, C. Jacob-Dolan, Z. Li, F. Nampanya, S. Patel, L. Pessaint, A. Van Ry, K. Blade, J. Yalley-Ogunro, M. Cabus, R. Brown, A. Cook, E. Teow, H. Andersen, M. G. Lewis, D. A. Lauffenburger, G. Alter, D. H. Barouch, Correlates of protection against SARS-CoV-2 in rhesus macaques. *Nature*, (2020).
  26. C. W. Tan, W. N. Chia, X. Qin, P. Liu, M. I. C. Chen, C. Tiu, Z. Hu, V. C.-W. Chen, B. E. Young, W. R. Sia, Y.-J. Tan, R. Foo, Y. Yi, D. C. Lye, D. E. Anderson, L.-F. Wang, A SARS-CoV-2 surrogate virus neutralization test based on antibody-mediated blockage of ACE2–spike protein–protein interaction. *Nat. Biotechnol.* **38**, 1073-1078 (2020).

27. A. Manenti, M. Maggetti, E. Casa, D. Martinuzzi, A. Torelli, C. M. Trombetta, S. Marchi, E. Montomoli, Evaluation of SARS-CoV-2 neutralizing antibodies using a CPE-based colorimetric live virus micro-neutralization assay in human serum samples. *J. Med. Virol.*, (2020).
28. IFV/COR/06 (phase-1-40). <https://rpcec.sld.cu/trials/RPCEC00000340-En>.
29. IFV/COR/08 (phase-2-910). <https://rpcec.sld.cu/trials/RPCEC00000347>.

### **Acknowledgments**

We thank Rolando Pérez, Luis Herrera, Agustin Lage and Eduardo Martinez (BioCubaFarma), for advice and support to the project, Lila Castellanos and Gail Reed for editing English version; as well as the Fondo de Ciencia e Innovacion (FONCI) for financial support (Project-2020-20). We are grateful to the Leibniz Institute of Plant Biochemistry, Germany, for support in structural characterization.

### **Author contributions**

Y.V.B., D.G.R., and V.V.B. designed and lead the study. S.F. is the manager of the project. D.S.M. and D.G.R. designed conjugation protocols and L.Q., U.R., J.P.S., Y.M., H.G., and M.G.R. performed chemical conjugation. L.R., B.S.R., R.P., C.A., T.H., G.B., F.Pi., A.V., performed the immunologic assays, M.F. and R.O. led the clinical care of the animals, F. C., R.G., M.L. led the analytical chemistry of the conjugates and vaccines J.E., N.G., and A.S. performed the virologic assays L.A.E., Y.R., and L.J.G. performed mass-spectra studies of the conjugates G.R., E.R-H., Y.C.I., S-L.L., T.B., E.O., K.L.M., C.F, and G.W.C. led the CHO-cells RBD preparation, Y.C., F.C. and F.P. supported and contributed to the study design and analysis. All authors revised the manuscript and approved submission.

## **Competing interests**

The authors declare no financial conflicts of interest. Y.V.B., D.S.M, S.F., M.R., L.R., U.R., D.G.R., T.B., E.O., D.G.R., D.G.R., and V.V.B. are co-inventors on provisional SARS-CoV-2 vaccine patents (Cu 2020-69).

## **Supplementary material**

### **Material and Methods**

#### **Table I-II**

#### **Fig. S1-S9**

## Figures caption.

**Fig. 1. Synthesis of RBD-TT conjugates.** (A) Reduction of RBD Cys538 using TCEP. (B) Recognition of RBD and reduced-RBDs by convalescent sera. (C) Conjugation of RBD with TT and representation of RBD<sub>2</sub>-TT and RBD<sub>6</sub>-TT. (D) Recognition of RBD-BSA conjugates by convalescent sera (CS), n=1-4. (E). Binding to ACE2 of RBD<sub>2</sub>-TT and RBD<sub>6</sub>-TT.

**Fig. 2.** Immunization of BALB/c mice with RBD<sub>2</sub>-TT/ alum and RBD<sub>6</sub>-TT/alum compared to RBD and RBD<sub>2</sub>-TT. The serum of individual mice is represented by RBD/alum ▼, RBD<sub>2</sub>-TT/alum ▲, RBD<sub>2</sub>-TT ◆, RBD<sub>6</sub>-TT/alum ● (A) Immunization protocol. (B) anti-RBD-specific IgG at days 7, 14, 21, and 28. (C) Dose response to RBD<sub>6</sub>-TT/alum at days 7 and 14. (D) Avidity index of antibodies elicited at T28. (E) RBD-specific IgG1 and IgG2a.

**Fig. 3. Memory B and T cells induced by RBD<sub>6</sub>-TT.** (A) Primary immune response to RBD<sub>6</sub>-TT/alum (green arrows). (B) Classical passive transfer of splenocytes from RBD<sub>6</sub>-TT/alum BALB/c and stimulated with RBD/alum (strong secondary response after day 7). (C) T-cell stimulation with RBD. (D) Cytokine secretion after *in vitro* RBD stimulation (E). % RBD-specific memory T CD<sub>8</sub><sup>+</sup>CD<sub>44</sub><sup>high</sup>IL-4<sup>+</sup>; % RBD-specific memory T CD<sub>8</sub><sup>+</sup>CD<sub>44</sub><sup>high</sup>Granzyme<sup>+</sup>; % RBD-specific memory T CD<sub>8</sub><sup>+</sup>CD<sub>44</sub><sup>high</sup>IFN $\gamma$ <sup>+</sup>; % RBD-specific memory T CD<sub>8</sub><sup>+</sup>CD<sub>44</sub><sup>high</sup>TNF $\alpha$ <sup>+</sup>

**Fig. 4. Virus neutralization by RBD antibodies induced by conjugates.** (A) mVNT<sub>50</sub> representing the serum dilution giving 50% inhibition ACE2-RBD interaction. (B) cVNT<sub>50</sub> measured as serum dilution giving 50% of virus neutralization (C) mVNT50/cVNT50 ratio and a possible schematic representation of the differences found.



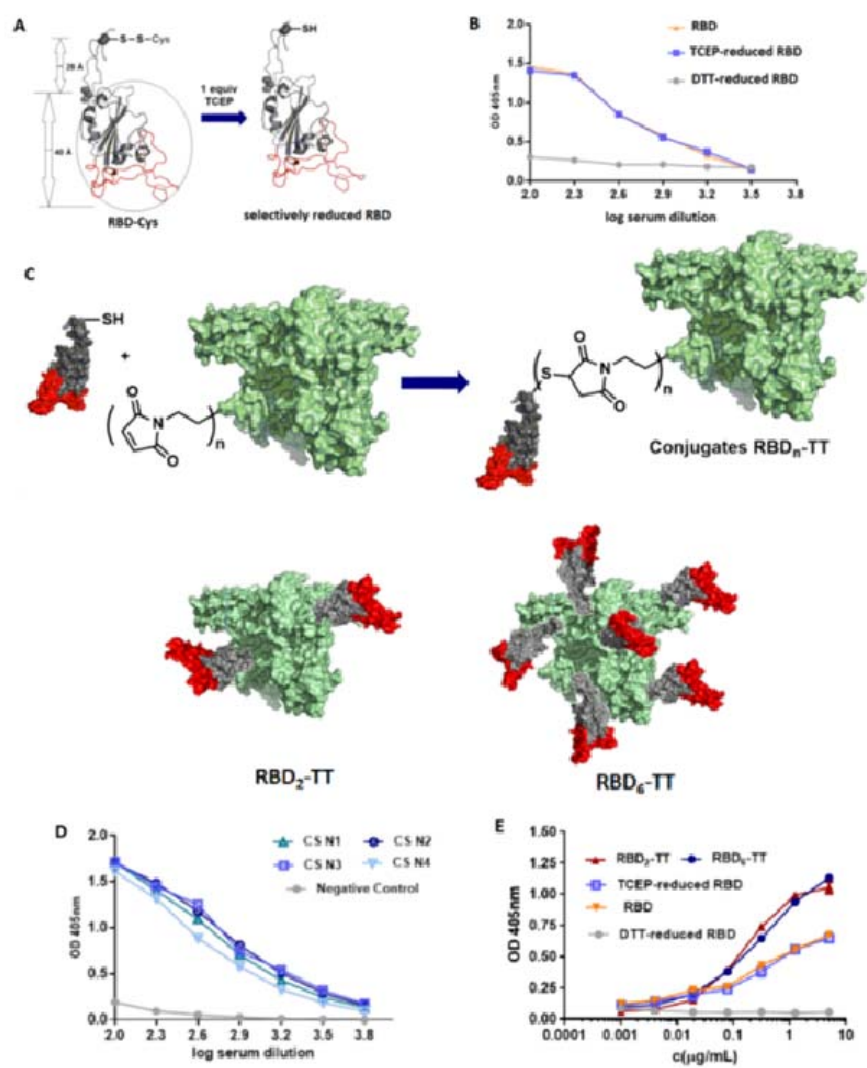


Fig.1

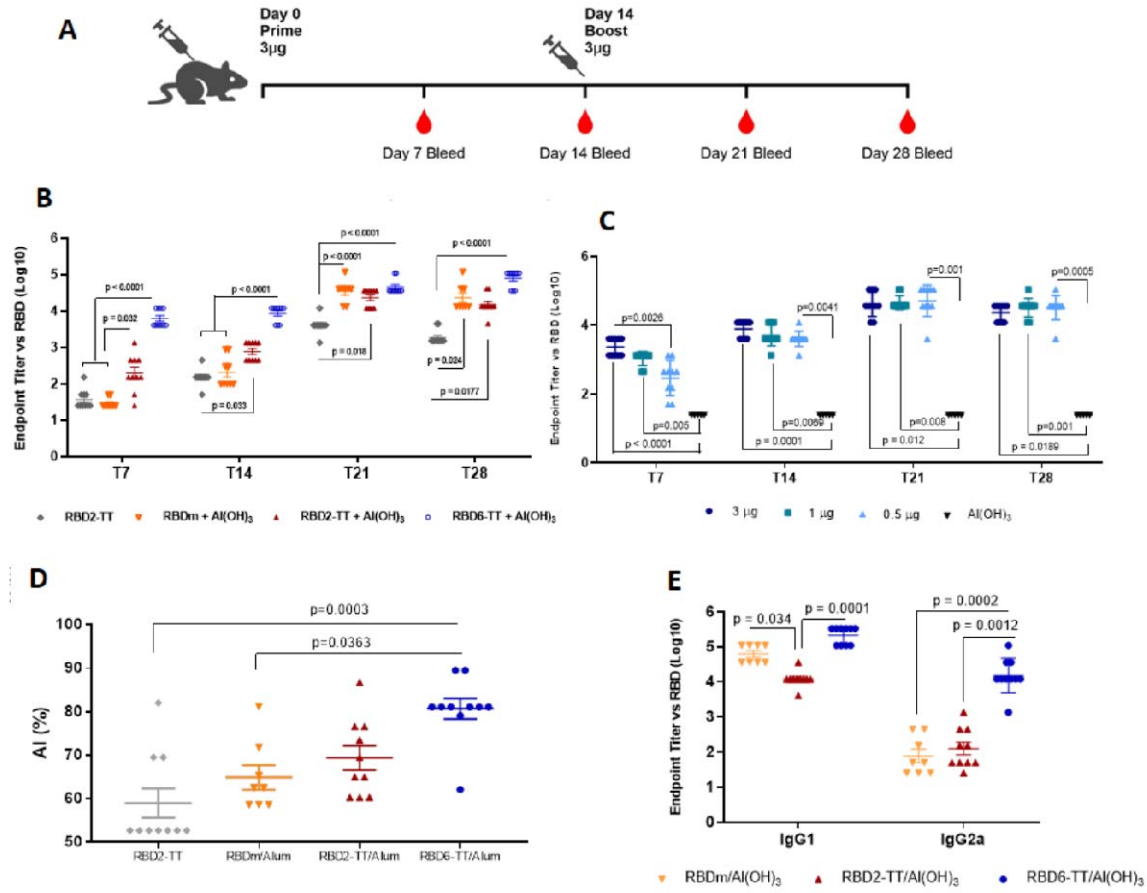


Fig.2

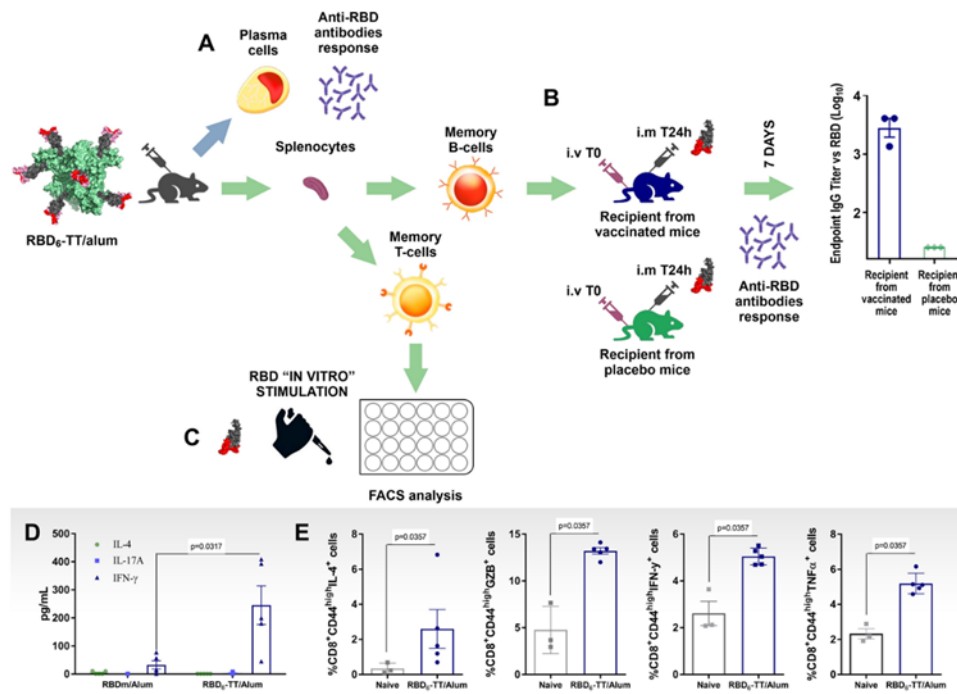


Fig.3.

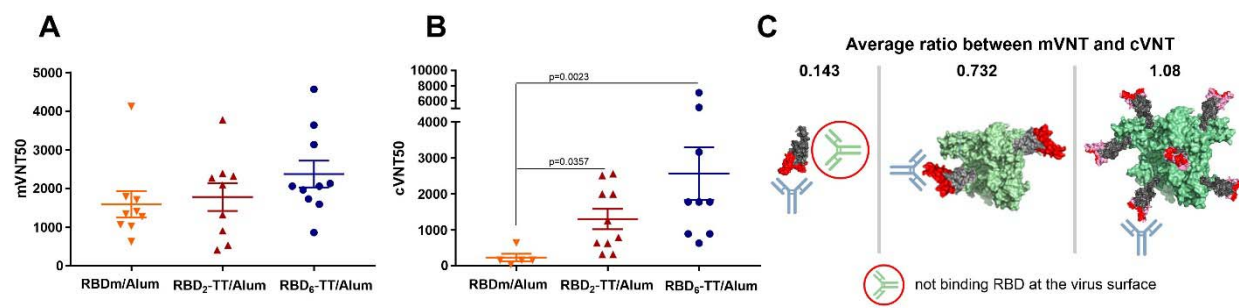


Fig.4.