FAQ (2022.11.28)

Q1	What resources do we need to use the CLOUD platform (STB CLOUD)?
A1	You need a device to access internet. The next step is to register on the STB CLOUD. The STB CLOUD is an automated platform that provides a list of compounds potentially binding to a target protein (Al-Hit). To use this platform, users will only need to select the target protein of interest.
Q2	What are the benefits of CLOUD computing instead of using software?
A2	STB CLOUD computes automatically using the compound library and protein structure data through the supercomputer system. Using STB CLOUD can reduce maintenance, labors, and software and hardware costs. In addition, users can easily access to the STB CLOUD on any device, anytime anywhere.
Q3	Do we need to be trained for the CLOUD platform? Do you provide any CLOUD platform training?
А3	STB CLOUD is an automated, complete platform that requires no special training. In case, we'll provide instructional manuals and video materials.
Q4	What is the process of creating a CLOUD account?
A4	You can create an account as Guest temporarily by entering the user's email, password and name. When the STB CLOUD use agreement is signed, your account will be fully activated to start your project.
Q5	During the process, can we stop the project and alter the target?
A5	Within two hours after submitting a project, you can cancel the project upon your request once. Please refer to the contract for use.
Q6	Can we fully own the results? And not share with your company?
A6	The ownership of outcomes of a project, according to the STB CLOUD service contract, belongs to the user. Please refer to Paragraph 2, Article 20 (Ownership of Rights) in STB CLOUD Use Agreement.
Q7	I think DeepMatcher® is a drug screening platform, so if we tell you the name of one target protein we want, are you going to screen active substance candidates?
A7	Yes. DeepMatcher® is a solution that finds active substance candidates for a target protein desired by customers.
Q8	How is input and output provided when computing with DeepMatcher®?

A8	Users input a target protein name of their interest and a selection of chemical library data is provided in the STB CLOUD. For output, the result reports (PDF, Excel) including the ranked Al-Hit candidate list are provided. In addition, you can check binding interactions between the candidates and the target protein in 3D.
Q9	What is the cost of using DeepMatcher® for one protein?
А9	Promotion: We guarantee the results until March 2023. Please see DeepMatcher® Use Agreement. Hit1 -100,000\$ (Derivation of AI hit candidates) Hit2 - 150,000\$ (Derivation of AI Hit candidates + in vitro assay by Syntekabio)
Q10	Is it possible to own and register an IP after validations under only user's rights or sharing it with Syntekabio?
A10	The intellectual property rights essentially belong to the user but depends on the agreement.
Q11	After results are provided, how can we proceed if we want to collaborate with you further?
A11	For follow-up collaboration inquiries, please contact at admin.usa@syntekabio.com. BD@syntekabio.com
Q12	What is DeepMatcher®-Hit?
A12	DeepMatcher® is Syntekabio's AI small molecule drug discovery platform that provides active substance candidates by predicting interactions between protein (target protein) and compounds through AI deep learning physics calculations.
Q13	What are the advantages of DeepMatcher®?
A13	DeepMatcher® is an automatic Al-Hit discovery solution that derives effective substance candidates potentially binding to a target in three weeks through physics theory and Al deep learning. Facilitating supercomputing system and CLOUD platform can provide quick and accurate discovery of effective substance candidates and compound optimization. Physics theory and deep learning are used to perform 3D-based docking, generate 24,000 conformers per compound, and find the best poses. MD simulation based on the best poses is performed to screen out the compounds binding to the target with wrong poses.
Q14	What is the range of molecular weights that can be calculated?
A14	DeepMatcher® is designed for small molecular drug discovery. Typically, M.W. of small molecular drugs ranges from 200 to about 600 Dalton.

Q15	Where is the source of your database?
A15	The Zinc database is being applied to DeepMatcher®. Users can have access to a library of 1 billion commercially available compounds.
Q16	How long will it take to provide AI-Hit candidates using DeepMatcher®?
A16	You can receive the list of Al-Hit candidates in 3 weeks after submitting a Hit 1 project on the STB CLOUD. You can receive the validated Hits in 14 weeks after submitting a Hit 2 project (Hit discovery + in vitro assay by STB)
Q17	When substance candidates are screened in silico, how many libraries will be used and how many candidates will be provided?
A17	We have access to and utilize 1.2 billion commercially available compound library, and 100 to 200 Al-Hit candidates will be provided based on the Grades.
Q18	Drug-protein interaction analysis in 3-D structures can differ among other algorithms. Why are there different results in terms of accuracy?
A18	Each algorithm uses different physical and chemical potentials. Different potentials give different binding energies between target and the ligands and governs the accuracy of the hit candidates.
Q19	Does the calculation result remain on the server when using STB CLOUD?
A19	No. The calculation results are not remained on the server. In addition, the meta-data generated in the process is deleted when the contract is terminated. For further information on data storage and related security policies, please see the Security System Introduction.
Q20	Where does your server physically reside? Are there any legal ramifications regarding our data privacy?
A20	The server is located at KT-IDC Center in Guro, Seoul, South Korea. (KT-IDC has international standard certification systems such as ISO14001, Green Data Center, ISO27001, and ISMS, and strictly protects and controls facilities, security management and cyber security management.)
Q21	Who has access to our data in the CLOUD? What is your company policy for ensuring only authorized employees can access our data?

A21	access. We conduct security training for system operators to control data access. In addition to controlling data access to physical facilities, security settings for storage access are applied for only authorized personnel to allow to use it.
Q22	What security measures do you take?
A22	Services and data are protected by physical security settings, blocking cyber attacks to security equipment and session encryptions.
Q23	Which specific data transmissions do you encrypt?
A23	From the moment the user connects to the CLOUD system, all the data transmitted and received through the session (login account information, input and receive various project related data, etc.) are encrypted with the SSL method to prevent malicious extortion and data decryption by third parties.
Q24	What level of technical support is included in your standard SLA?
A24	Failure and problem management (failure propagation, failure handling rate management, etc.), security measures (security measures, preventive inspections, security incident management, etc.). Technical support for data backup and performance management (maintenance of operation rate) are included.
Q25	What are your procedures for suspected security violations on our data or your data?
A25	Please figure out the extent of damages to your data and proceed with blocking/recovery measures for the data. Legal follow-up and damage compensation procedures will be carried out in accordance with the internal regulations and the contents of the contract.

Data access is limited to system-specific operation personnel who are granted secure