CAST(computer aided synbio tool) combines the design automation, innovative models and algorithms, and wetlab validation, making complex biological systems accessible to researchers who want rapid, intuitive, and high-quality accomplish their designs.

Get started in CAST

I. Download CAST from https://github.com/igemsoftware/SYSU-Software2013



II. Open CAST

1. Open 'project' folder in 'SYSU-Software2013-master'. Then click on one of sever_.bat (choose x64 if your system is 64-bit; otherwise, 32bit is your best choice)



2. It will occur something like this. Just minimize the window.



3. Open your browser(such as chrome) and input 127.0.0.1:5000 in the address bar.



4. Sucess!



- III. Register & Sign in
- 1. Register if you don't have an account.



2. Fill in the form and finish signing up. Attention! Gender must be 0(male) or 1(female).



3. Sign in with your username and password.

welcome
king
Remember me Sign up for CAST?

IV. CAST at a glance



Biobrick Panel: You can choose any biobricks from iGEM registry in 'Protein' or your own biobricks in 'Coding. Apart, regulation factors are available in 'Regulation Factors'.

Information Panel: You can view your account information in 'Menu' as well as details of the biobrick chosen by you.

Function Panel: Select one of the functions and go to next step.

Working Panel: It's a white paper on which you can draw a regulation network.

Plan Your Experiment

Choose one folder and select the biobrick you need and click on 🖸 to add. (For instance, choose 'DNA-modification' -> 'BBa_I11020')



II.Work on your biobricks.

On the working panel, you are able to design regulation networks and manage them. When clicking on one of the biobricks on the working panel,



configuration of this biobrick can be review on the Information Panel.(Right-hand side)

la constal	Menu 🌱
lasmid	Configuration 👻
	Component Config
	Part_id
	5248
	Part_name
	BBa_I11020
	Part_short_name
	I11020
	Part_short_desc
<	integrase from E. coli phage lai
	Part_type
	Coding
	Part_status
	Available
	Part_results
	None
	Part_nickname

III. Generate gene circuit.

Click on 'gene circuit' after building regulation networks.



'Gene circuit' interface at a glance.

1. Dashboard: On the dashboard, each biobrick in the regulation network has a control



panel, which allows users to adjust its configurations, such as PoPS, RiPS and so on. After setting, the rate calculated is showed on the upper panel.



biobricks in the circuit(promoter, RBS, etc.) automatically, according to the value you set.



2. **Simulation graph**: According to the configuration, CAST calculates the expression efficiency of each biobrick and show it as curves.



2400 3000

View more

00

If you need more information or more complex simulation, you can click to enter 'Simulation' interface.



3. **Regulation network editor**: You can return to 'Regulation Network' interface for further edition.



Click and go back to 'Regulation Network'

4. **Gene circuits manager**: The information of Gene circuits on the plasmid is displayed on this panel.

You can add one more plasmid if you want

1003	BBa_B0013	cis		New plasm	nd) Ad	d a new	plasmic	1		
You ca	n simply	move c	circuits b	y clicking	< >					
BBa_1712074	BBa_J61104	BBa_C0060	BBa_B0013	cis BBa_J64951	BBa_J61104	BBa_C0061	BBa_J61104	BBa_K518003	BBa_B0013	<> cis
				V						<>
BBa_J64951	BBa_J61104	BBa_C0061	BBa_J61104	BBa_K518003	BBa_B0013 Cis	BBa_I712074	BBa_J61104	BBa_C0060	BBa_B0013	cis

At last, if you think you have done something wrong, you can click



III. View the plasmid

To see check the plasmid, just click 'Plasmid' on the Function panel, or 'plasmid' button in the Gene circuit manager.





interface at the glance

In the plasmid viewer, users can drag their mouses and rotate the plasmid, in order to check each part of it. In the meanwhile, the sequence of the part your are viewing is displayed on the upper side. You can also see the whole sequence of this plasmid.

On the console, you can choose different types of plasmid

Assembly Standard	-
Assembly Standard RFC 10	
RFC 20	
RFC 21	
RFC 23	
RFC 10 RFC 20 RFC 21 RFC 23	

You can download the sequence or save the plasmid graph if you like.

Save Plasmid	Graph
Sequence	downloadSeq

IV. Design the experiments

1. Click 'Protocol' button to enter next interface.

lasmid	Protocol	Simulation

2. On 'Protocol' interface, you can find standard experiment methods that give you guidance in wetlab.

Step 1	BBa_I712074 BBa_J81104	BBa_C0060 BBa_B	013 BB	.R0062 BB.J61104	BBa_C0081	BBa_J01104	 BBa_K518
Step 2	BBa_1712074 BBa_J61104 BBa_C000	0 BBa_B0013	BBs_R0062 BBs	_J61104 BBa_C0061 BB	a_J61104	BBa_K518003 BBa_B0013	3
Step 3	BBa_1712074 BBa_J01104 BBa_C000	0 BBa_B0013 BBa_R0	062 BBa_J61104 BBa	BBa_J61104	BBa_K518003	BBa_B0013	
Step 4							
	BBs_1712074 BBs_J61104 BBs_C006	30 BBa_B0013 BBa_R(062 BBa_J61104 BBa	a_C0061 BBa_J61104 BBs	_K518003 BBa_B0013		
4							
	Step 1 Step 2 Step 3 Step 4	Step 1 Step 2 Step 3 Step 4 $BBa_1712074 BBa_161104 BBa_0000$ $BBa_1712074 BBa_161104 BBa_0000$ Step 4 $BBa_1712074 BBa_161104 BBa_0000$	Step 1 Step 2 Step 3 Step 4 4 4 5 5 5 5 5 5 5 5 5 5 5 5 5	Step 1 Step 2 Step 3 Step 4 Step 4	Step 1 Step 2 Step 3 Step 4	Step 1 Step 2 BBs_1712074 BBs_161104 BBs_C0060 BBs_B0013 BBs_R0062 BBs_161104 BBs_C0061 BBs_161104 BBs_C0060 BBs_160113 BBs_160062 BBs_161104 BBs_C0061 BBs_161104 BBs_C0061 BBs_161104 BBs_C0061 BBs_161104 BBs_C0061 BBs_161104 BBs_C0061 BBs_161104 BBs_160061 BBs_180113 BBs_161104 BBs_16000000000000000000000000000000000000	Step 1 Step 1 Be00001 Be00104 Be00000 BeB0013 BeR0002 Be00104 Be00001 Be00104 Be00001 Be00104 Be00001 Be00000 Be00013 BB00002 Be00101 Be00001 Be00000 Be00001 Be00000 Be00001 Be00001 Be00000 Be00001 Be00000 Be0

V. Manage your files

1. Return to 'Gene circuit' or 'Plasmid'. Open information panel and click 'Save' after naming this file.



2. You can find your file by clicking on 'My Files'.



Bobby	Your File L	.ist
	Filename	Туре
Logout	234234	genecircuit
My Files	Bobby	plasmid

3. You can manage your files. (Share, for example)

Your File List

Filename	Туре	operation			
234234	genecircuit	opent	delete 💼	shared your file	
Bobby	plasmid	open	delete 💼	shared your file	
abcd	plasmid	open	delete 💼	shared your file	

Your file extract code

72d0a666a3fa0e063a218763dd4628d12feb2c27

You can acquire others' shared files by inputing extract code. It's the same to others.

Shared File List

K

×



VI. Add your own biobricks

1. Open information panel on 'Regulation Network' interface. Click on 'Create New Part' to add your own biobricks.



2. If you want to add a part based on registry, you can choose one biobrick from biobrick panel.



3. If you want to add a part from sequence, you can input the sequence in this blank.

Add a part from sequence

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••	•••	•••	~	٠.		-	•••

Add a part from sequence	tttttgggagagtt
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4. Click 'Accept' after finishing. Then provide the entire information of the new biobrick.

				_		
	A	Accept				
		Ļ				
BASIC INFORMATIC	N	•				
Туре:	Regulatory	▼ Short description:	An inducer			
Name:	BBa_70020	Nick name:	IPTG			
Short name:	Indu	Author:	Bobby			
MODELING PARAMETERS						
Promoter						
Name:	LuxR	Number:	BBa_I1051			
PoPS:	0.2	MPPromoter:	12			
LeakageRate:	0.3	Туре:	Regulatory			
K1:	0	Activator:	0			
Repressor:	0	Source:	Nature			
		Accept				

Click 'Accept' to finish.



5. You can find you own parts enter 'Your Part List' from 'My Files' and manage them.



VII. Finish & Log out

When you finish all your work, just log out and exit for relaxation. You files will be saved well.

Click 'Log out', then close the window and exit CAST.



VIII. Other functions

1. Change your account information Click your name or photo to enter your account profile, where you can change your information.



Bobby

profile	>			
password	>	Username: Bobby		
	*			
		Gender:		
		● Male ○ Female Email:		
		101@qq.com123		
		Submit		