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APRIL

Date	04/01/2020-04/11/2020
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Aim	Determine the research topic
Participant	All
Content	 The topics discussed in the meeting focused on The Guidelines for the Application of Key Projects of "Synthetic Biology" in 2020 (draft for comments) and The Guidelines for the Application of Key Projects of "Synthetic Biology" in 2019 (draft for comments); Divided into four groups for discussion, group one has no results, group two, group three, and group four considered the related content of protein engineering, modification of E. coli metabolic pathway, environmental heavy metal detection and other topics of interest.
Result	We unanimously believed that the environment is a hot topic at the moment, and it is also a topic of great significance related to human survival, and we planned to explore interesting and meaningful topics in the direction of the environment.

Date	04/12/2020-04/18/2020
Aim	Determine the research topic
Participant	All
Content	Since last week's topic discussion was inconclusive, the instructor suggested a
	possible direction: PET and PETase. We checked relevant literature and news on
	the Internet and learned about PET and PETase.
	The superiority of PETase. Compared with its homologous enzymes, PETase not
	only has the ability to degrade PET with low crystallinity, but more importantly,
	this enzyme also has a strong ability to degrade PET with high crystallinity, which
	is far superior to other homologous enzymes.
Result	By consulting the information, we learned the following:
	The severity and urgency of the plastic pollution problem. A large amount of PET
	is manufactured and used every year, but only a small part of it can be recycled.
	There are a large number of PET products remaining in the environment, which
	poses a huge threat to the ecological environment and will eventually be fed back
	to human beings.
	In contrast, biological enzymatic hydrolysis has outstanding advantages. Among
	them, PETase has shown great advantages. Compared with its homologous
	enzymes, PETase not only has the ability to degrade PET with low crystallinity,
	but more importantly, this enzyme also has a strong ability to degrade PET with
	high crystallinity.
	We are aware of the research potential and significance of this subject, and we
	are determined to carry out research on PETase.

Date	04/19/2020-04/25/2020
Aim	Choose specific transformation directions
Participant	All

Content	1. Reading the literature Enzymatic Degradation of Plant Biomass and Synthetic
Content	
	Polymers etc., combined with daily life, it is found that there are great
	problems in the recycling and degradation of plastic bottles (PET bottles). A
	large number of them are manufactured and used, and only a small amount
	can be recycled. Moreover, the recovery means are physical and chemical
	recovery, which wastes energy, causes pollution and is inefficient, so it is
	difficult to achieve the purpose of recycling;
	2. It is mentioned in the literature that the degradation efficiency of PETase is
	very low and cannot meet the needs of actual production and life, so we
	decided to modify it to improve its degradation efficiency.
Result	The problems in the plastic recycling process, physical and chemical recycling,
	high cost and low recovery rate. We have also put forward many ideas: transform
	the substrate binding pocket, stabilize the catalytic triad, improve the stability of
	PETase and substrate binding, and inhibit contact products and so on. But no
	specific transformation direction was finalized.

Date	04/26/2020-05/02/2020
Aim	Explore the direction of transformation
Participant	All
Content	1. Read the literature An Engineered PET Depolymerase to Break Down and
	Recycle Plastic Bottles, study its subject background, transformation ideas,
	technical details, etc., to prepare for our own research of PETase and its
	substrate PET;
	2. Focused on the research on the modification methods of LCC in the
	literature, the calcium ion binding site in the enzyme was mutated to
	introduce disulfide bonds, and multiple mutations were combined at the
	same time;
	3. Discussed and exchanged about this document, and started the process of
	document research.
Result	1. We have gained a lot of background knowledge related to PET;
	2. Understand the research status of PET degrading enzymes such as LCC and
	PETase, as well as some of the advantages of these enzymes, such as the high
	thermal stability of LCC, and the degradation ability of PETase for high
	crystallinity PET, etc

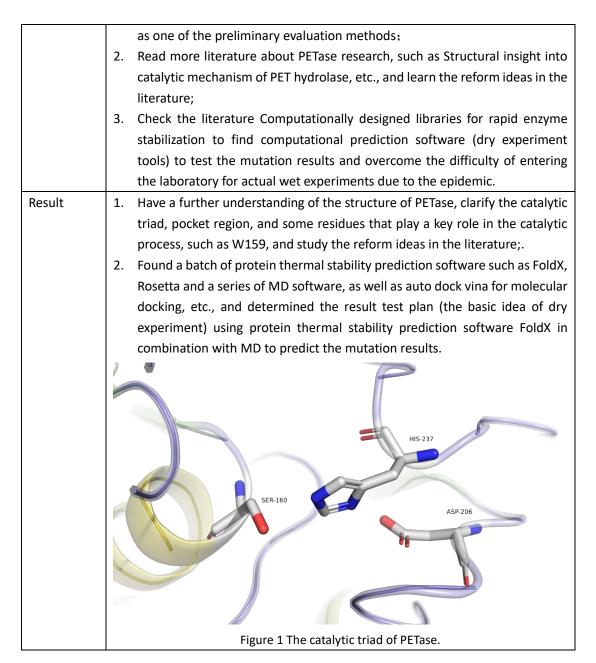
MAY

Date	05/03/2020-05/09/2020
Aim	Determine the transformation direction
Participant	All
Content	Deepen the understanding of the literature last week, especially the
	predecessors' understanding of the modification of the thermal stability of LCC

	and the limitation of PETase that only maintains activity at room temperature
	and degrades PET with low crystallinity.
Result	LCC is limited by its own thermal stability to the degradation efficiency of PET.
	Improving the thermal stability of LCC will be of great significance to the
	improvement of the degradation efficiency of low crystallinity PET. At the same
	time, PETase also has similar problems, so we analogize PETase to improve the
	thermal stability of PETase, so that it can remain stable in the high temperature
	environment in vitro. If the Tm value can be increased above the glass transition
	temperature of PET, the degradation activity of PETase will produce a qualitative
	leap. Therefore, the PETase transformation plan was finally finalized: by
	improving the thermal stability of PETase, improving its catalytic degradation
	ability of high crystallinity PET.

Date	05/10/2020-05/16/2020
Aim	Seek ways to improve enzyme thermal stability
Participant	All
Content	 Learn about a molecular visualization software pymol from our instructor, and start to organize everyone to learn this software; Read the literature Rational Protein Engineering of Thermo-Stable PETase from Ideonella sakaiensis for Highly Efficient PET Degradation, and understand the previous research ideas and transformation methods of PETase; For the need to modify the protein, we have learned the physical and chemical properties of amino acids, such as hydrophobicity, polarity, charging status, etc., in order to select the best alternative amino acid when
Result	performing site mutations. It is known from the literature that the stability of a protein is mainly determined by the internal forces of the protein, such as hydrogen bonds, disulfide bonds, hydrophobic forces, etc. The predecessors started with these forces to improve thermal stability. So we decided on the direction of transformation from different molecular forces, and initially divided the teams into hydrogen bond group, salt bridge group, disulfide bond group, glycosylation group, proline group, hydrophobic force group, etc. The group, each is mainly responsible for the research of one kind of force, different people choose different forces as the transformation direction, and start the initial transformation.

Date	05/17/2020-05/23/2020
Aim	Research from different transformation directions
Participant	Shuying Wu, Yanjiao Lu, Yuan Wang, Guosheng Gao, Yuhan Li, Xujia Liu, Yufei
	Zhou, Dewu Liu, Xinyi Zhang, Tao Fu, Haoyang Shi, He Zhang, Duo Zhang
Content	1. In order to evaluate the local thermal stability better, we introduced B factor



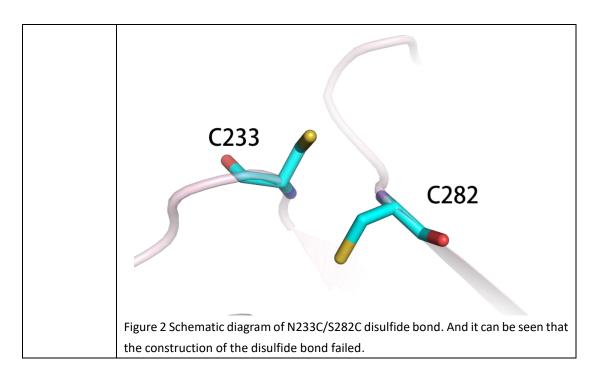
Date	05/24/2020-05/30/2020
Aim	Research from different transformation directions: hydrogen bond, pocket
	hydrophobicity
Participant	Guosheng Gao, Yuhan Li, Xujia Liu, Dewu Liu, Xinyi Zhang, Duo Zhang
Content	1. Hydrogen bond: Reading the literature Enhanced catalytic site thermal
	stability of cold-adapted esterase EstK by a W208Y mutation etc., get ideas;
	2. Pocket hydrophobicity: The substrate PET itself is hydrophobic. While
	enhancing the hydrophobicity of the PETase substrate binding pocket, it can
	not only increase the enzyme activity by promoting the binding of the
	substrate and the enzyme, but also enhance the hydrophobic interaction
	and improve the thermal stability of the enzyme.
Result	1. Knowing that the lipase mutation W208Y would produce increased

- hydrogen bonds, we searched for similar sites in PETase and found the S238D+W159Y mutant. After PyMOL mutation, the result was similar to W208Y, and both produced Additional hydrogen bonds;
- 2. The transformation of the hydrophobic direction has great potential and needs to be further explored.

JUNE

Date	06/01/2020-06/06/2020
Aim	Research from different directions: π - π interaction, glycosylation
Participant	Guosheng Gao, Yuhan Li, Xujia Liu, Xinyi Zhang
Content	1. Glycosylation: After reading Stabilizing leaf and branch compost cutinase (LCC) with glycosylation: Mechanism and effect on PET hydrolysis, it is concluded that LCC has a high aggregation tendency, which is the main reason for its thermal inactivation. Glycosylation can prevent aggregation by improving protein conformational stability or by imposing steric constraints that inhibit protein-protein interactions.
	2. π - π interaction: Reading the literature A conserved and buried edge-to-face aromatic interaction in small ubiquitin-like modifier (SUMO) has a role in SUMO stability and function and Carbohydrate — Protein aromatic ring interactions beyond CH/ π interactions: A Protein Data Bank survey and quantum chemical calculations etc., learn and understand π - π interaction and find ideas.
Result	1. Inspiration from the literature: study whether PETase also has thermal aggregation phenomenon, introduce glycosylation at natural sites or artificially introduce sites, and detect PETase activity at high temperature. Since this is a protein modification work, and there is no related literature that reports PETase thermal aggregation, we temporarily abandoned this research direction.
	2. Understand the essence of π - π interaction, which is a kind of weak hydrogen bonding. The reason for its formation is the mutual attraction of electron clouds between aromatic systems. There are three main types of aromatic interactions, namely F-type accumulation, edge-to-face accumulation and T-type accumulation. The F-type accumulation will adversely affect the system due to the strong repulsion of the electron cloud and reduce the stability. There is no positive repulsion between the latter two electron clouds, and the electron clouds with opposite electrical properties attract each other, making the system more stable. Understand the geometrical requirements for the formation of π - π interaction, and proceed to carry out related mutations in PETase. Finally, the π - π interaction was selected for in-depth research, and the research direction of glycosylation was abandoned.

Date	06/07/2020-06/13/2020
Aim	Research from different transformation directions: proline, disulfide bond
Participant	Shuying Wu, Yuan Wang, He Zhang
Content	1.Proline: Looking at the literature <i>Review: Engineering of thermostable enzymes for industrial applications</i> , it is found that proline substitution can improve the thermal stability of the enzyme: Pro (proline) is more special, containing an imino group, a carboxyl group and a pyrrolidine ring. The unique cyclic structure of the side chain gives Pro a special conformational rigidity relative to other amino acids, and has a special role in protein thermal stability. Use the protein stability prediction website http://kazlab.umn.edu/ to find a suitable proline substitution site, use the modeling website https://web.expasy.org/protparam/ to model the mutant, use The thermal stability prediction website https://servicesn.mbi.ucla.edu/SAVES/ evaluates the stability of the mutant, and uses the software PyMOL to conduct in-depth research on the structure of the mutant. 2.Disulfide bond: We found that the introduction of disulfide bonds into the calcium binding sites in the homologous LCC and Tfcut2 has achieved good results. For Tm, Tfcut2 (D238C/S283C) increased by about 24.9°C (69.8 to 94.7), and LCC (D204C/E253C) increased by about 9.5° C (84.7 to 94.2). We analyzed the structure of calcium ion binding sites in LCC and Tfcut2.
Result	 A suitable proline substitution site (E246P) was found, and it was found that the proline mutation site predicted by the website could not meet the requirements for improving stability. On the one hand, the b-factor value of the site selected by the website was low. It belongs to the region with relatively stable structure; on the other hand, the stability of the protein after mutation decreases, which is contrary to our original intention. Need to continue research. The introduction of disulfide bonds into calcium ion binding sites in LCC and Tfcut2 was successful. It is found that these positions do meet part of the geometric criteria of the disulfide bond structure (mainly the distance between Cα-Cα and Cβ-Cβ). These results indicate that it is feasible to introduce disulfide bonds at the calcium ion binding site. A homology comparison was performed to speculate the calcium binding site of PETase (E175-N204-S253), and the group of mutations N233C/S282C was finally determined from a geometric perspective.



Date	06/14/2020-06/20/2020
Aim	Research from different transformation directions: salt bridge, hydrophobic core
Participant	Guosheng Gao, Yuhan Li, Xujia Liu, Tao Fu, Duo Zhang
Content	 Salt bridge: Check information to learn about salt bridge. Learn about basic mutation methods by reading the literature <i>Evolutionary Stability of Salt Bridges Hints Its Contribution to Stability of Proteins</i>; hydrophobic core: Read the literature <i>Enthalpic and entropic contributions to hydrophobicity</i>, etc. to learn related knowledge, explore and analyze ideas.
Result	 Understand the basic mutation method: single point mutation of endogenous amino acid, select an endogenous amino acid, find the mutation point within 5 amino acids nearby, and try our idea through PyMOL; It is understood that the amino acids in the main hydrophobic binding sites of the protein have larger hydrophobic side chains, forming a tighter hydrophobic core, which can enhance the thermal stability of the enzyme; the amino acids with larger side chains on the β-sheet may reduce the thermal stability, but it can increase enzyme activity and lower the optimal pH (glucose isomerase); the non-polar amino acid Leu on the α-helix may enhance thermal stability (glucose isomerase). We believe that the stability improvement of the hydrophobic core region has great potential and will continue to be studied in the future.

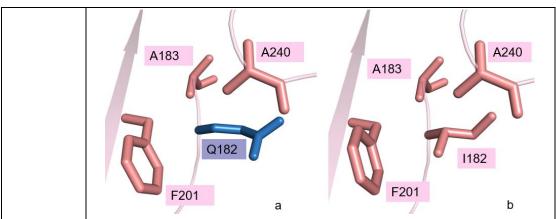
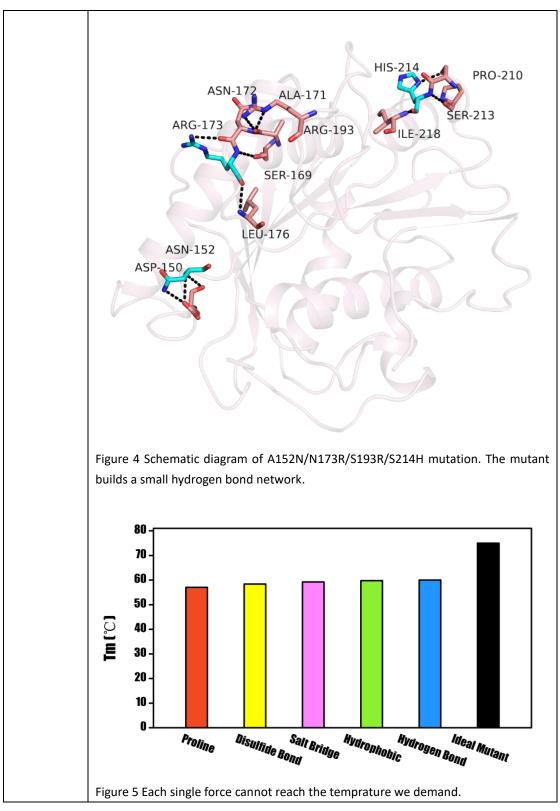


Figure 3 One of the hydrophobic cores we structured successfully. The Polar residue Q182 is showed in blue(a), which is mutated to the hydrophobic residue IIe(b).

Date	06/21/2020-06/30/2020
Aim	Combine mutations within a single force and set the target DDG≤-10kcal/mol
Participant	Shuying Wu, Yanjiao Lu, Guosheng Gao, Yuhan Li, Xujia Liu, Xinyi Zhang, Tao Fu,
	Haoyang Shi, He Zhang, Duo Zhang
Content	1. Hydrogen bond: Reading the literature 分子动力学模拟研究 CGTase 酶活
	性区域的热稳定性 and Extended protein/water Hbond networks in
	photosynthetic water oxidation, etc. gave great inspiration for the
	construction of our hydrogen bond network. We mutated the
	A152N/N173R/S193R/S214H site in an attempt to construct a hydrogen
	bond network;
	2. Salt bridge: Use homologous enzyme sequence alignment to observe the
	salt bridges that exist in the enzymes with high thermostability such as LCC,
	CUT190, TFCUT2 but not in PETase, and introduce mutation points in
	PETase;
Result	1. Hydrogen bond: Mutation was made at the A152N/N173R/S193R/S214H
	sites, but calculated by FoldX, its DDG was -5.43591kJ/mol, which was far
	lower than expected. For this reason, we conducted visual screening and
	found that many new hydrogen bonds in single point mutations did not
	appear in combined mutations. In this regard, we believe that hydrogen
	bonds are greatly affected by conformational changes, and other site
	mutations will have a greater impact on the formation of hydrogen bonds,
	so the expected hydrogen bond network is not formed when combining
	mutations. In the future, we will continue to analyze the hydrogen bond
	network in the combined mutants in order to make a greater contribution to
	the improvement of thermal stability.
	Have a clear understanding of the level of thermal stability of single-point
	mutations and single-force combination mutations, and hope to find more
	excellent mutation sites to achieve the goal of DDG.
	2. Salt bridge: Single point mutations such as A40D, T116D/R, and D220E have
	been made, and new salt bridges have been successfully introduced.



JULY

Date	07/01/2020-07/11/2020
Aim	1. Further study single points and provide more single points;

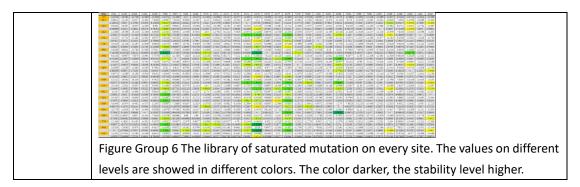
	2. Further analysis of mutants by homology comparison.
Participant	Shuying Wu, Yanjiao Lu, Yuan Wang, Guosheng Gao, Yuhan Li, Xujia Liu, Dewu
	Liu, Xinyi Zhang, Tao Fu, Haoyang Shi, He Zhang, Duo Zhang
Content	Observe and study the unstable regions in PETase, that is, the sites with high B
	factor values. Check the corresponding structure on the PDB website, and hope
	to add molecular forces of proteins in these regions, such as mutations adding
	hydrogen bonds, salt bridges, hydrophobicity, etc.
	1. Hydrogen Bond: Find single-point mutations based on homology alignment
	and website predictions, and read the literature on mutating hydrogen
	bonds to improve PETase thermal stability: Rational Protein Engineering of
	Thermo-Stable PETase from Ideonella sakaiensis for Highly Efficient; at the
	same time, use homologous enzyme sequence comparisons, using b-factor
	as a measure, introduce stable sites in Tfcut2, LCC, Cut190 into PETase;
	2. Proline: Read the literature Increasing protein stability by improving Beta-
	turn Continue to study proline substitution;
	3. Hydrophobic: Through the comparison of homologous enzyme sequences,
	observe the enzymes with high thermostability such as LCC, CUT190,
	TFCUT2, expand the hydrophobic site mutation library.
Result	1. Successfully constructed new hydrogen bonds at S46D, A47R, T56R, T72R,
	R90D, S92D, T116R, S136N, G139R, Y146R, K148R, A152N/S, N173R, S188N,
	I208Q, D220N, S278D and other sites, especially K148R stabilizes two loops,
	Y146R increases hydrophilicity and stabilizes the connection between loop
	and beta sheet;
	2. It is understood that there is a proline residue preference at position i+1 in
	the structure of β-turn, and mutations such as S58P, S141P, S207P, and S223P
	have been determined;
	3. The stable hydrophobic interaction contribution sites in LCC, CUT190,
	TFCUT2 and other enzymes were introduced into PETase, and effective sites
	with reasonable structure, data and empirical explanation were determined,
	such as V68I, A80V, S124A, Q127L, V134L, V149I, S166A, S169I, A179V,
	Q182L, T198V, N246I, K252L, N275I, etc., to further expand and determine
	the hydrophobic site mutation library.

Date	07/12/2020-07/18/2020
Aim	Improve the level of PyMOL mastery and unify the drawing standards;
	2. Determine calculation software and basic plan.
Participant	All
Content	1. 1. Learn to use PyMOL and PyMOL to draw;
	3d the substrate to facilitate docking. Use autodock software for molecular docking;
	3. A protein visualization software YASARA based on FoldX is installed to realize the visualization of FoldX operation and improve the computational efficiency.

Result	1.	The PyMOL drawing standard is determined, and details such as color and
		angle when drawing are uniformly specified;
	2.	In the process of setting flexible residues, if the number of settings exceeds
		three, an error will occur. After consulting the data, it is found that this is a
		bug of the software. Since it is difficult to repair in a short time, I decided to
		change to another backup software auto dock vina. We started to try to
		connect the substrate to the pocket and observe on PyMOL, but the effect is
		not ideal;
	3.	A new experimental protocol was determined: using FoldX to perform
		saturation mutations at all sites of PETase, and adopt a reverse analysis
		method that uses data to guide visual screening to screen out all mutations
		that significantly improve stability, and perform visual screening on them.
		Outstanding mutations under double standards.

Date	07/19/2020-07/25/2020
Aim	Determine the molecular docking experiment protocol;
	2. Use FoldX to make a full-point saturation mutation library.
Participant	Shuying Wu, Yanjiao Lu, Dewu Liu, Haoyang Shi, Kegen Chen, Yansong Bai
Content	1. Looking for software that can construct molecular docking model, tried
	various software such as discovery studio, sybyl, autodock, etc.
	2. After continuous attempts, we began to use FoldX to initially evaluate the
	thermal stability of the mutants designed by the experiment. After that, we
	think it is better to carry out large-scale predictions to screen out the single
	mutations with the most significant improvement in thermal stability, and to
	guide the site selection together with visual structure screening.
Result	1. Finally, autodock was determined as the modeling software; but when testing
	the stability of autodock modeling, it was found that the setting of flexible
	residues would cause the software to report an error docking instance has no
	attribute 'ligMol'. In the end, I chose autodock vina with more complete
	software optimization to do the docking work, and at the same time learned
	the molecular docking tutorial and docking site knowledge. Try to dock PETase
	and BETH small molecules, the docking site is in the active pocket, the effect
	is very good;
	2. We directly performed saturation mutations on 260 sites of PETase,
	constructed a full-site saturation single mutation stability database, and
	finally obtained a total of 4940 single-point mutations, and screened out 514
	DDG drops within the error range (Stable mutations other than 0.46kcal/mol).

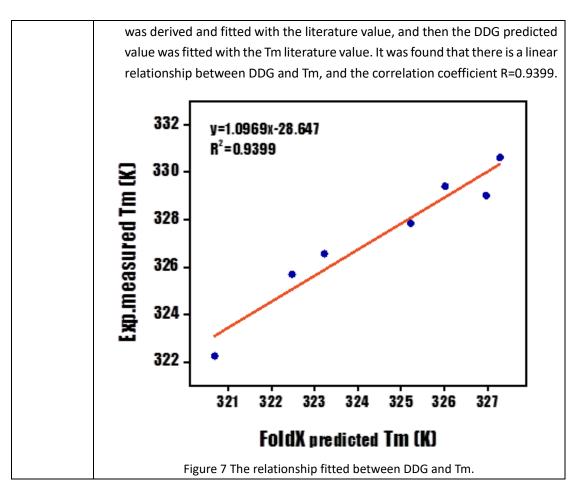




Date	07/26/2020-07/31/2020
Aim	Promote the mastery and experimental progress of Molecular Docking and
	Molecular Dynamics
Participant	Shuying Wu, Yanjiao Lu, Kegen Chen, Yansong Bai
Content	1. Learn MD theory process, install and learn to use MD software amber and
	GROMACS;
	2. Learned the design points of gridbox parameters and literature such as
	Structural insight into molecular mechanism of poly (ethylene terephthalate)
	degradation to determine the setting of flexible residues, and read papers
	on the binding scores output by autodock vina such as Can we use docking
	and scoring for hit-to-lead optimization? and Rescoring ligand docking poses
	to further improve the accuracy of molecular docking.
Result	1. We found that there was a problem with amber, a software used for MD. the
	command pdb4amber could not be used. After asking for help from many
	parties and collecting a lot of data from the network, we still could not solve
	the problem, so we gave up amber and successfully installed and used
	GROMACS;
	2. In the next few molecular dockings, the accuracy of the theoretical docking
	results that have been learned has been improved significantly.

AUGUST

Date	08/01/2020-08/08/2020	
Aim	Calculate the relationship between DDG and Tm	
Participant	Yanjiao Lu, Dewu Liu, He Zhang, Yansong Bai	
Content	1. In order to predict and process data, learn how to use Matlab to perform	
	numerical fitting calculations;	
	2. Through reading literature and mathematical derivation, the PETase and	
	homologous enzyme mutant DDG measured by FoldX are fitted to their Tm	
	literature values to solve the relationship between the two.	
Result	1. Use Matlab to successfully solve the problem of data prediction, and make	
	related charts and PPT;	
	2. According to $\Delta G = -RTInK$, the Tm value corresponding to FoldX predicted DDG	



Date	08/09/2020-08/15/2020
Aim	Combine FoldX and multiple other forecasting websites to conduct in-depth
	analysis of their respective visual screening results
Participant	Shuying Wu, Yanjiao Lu, Yuan Wang, Guosheng Gao, Yuhan Li, Xujia Liu, Dewu Liu,
	Xinyi Zhang, Tao Fu, Haoyang Shi, He Zhang, Duo Zhang
Content	Introduce iMutant, Fireprot and other web pages to assist in the design of salt
	bridges and hydrogen bonds. For disulfide bonds, we searched on PudMed for the
	methods and current research status of artificial disulfide bonds in the past 15
	years. Study the literature Disulfide by Design 2.0: a web-based tool for disulfide
	engineering in proteins, Multi-Template Approach to Modeling Engineered
	Disulfide Bonds, etc Use Biosynthesis software to perform geometric evaluation
	on all sites of the whole enzyme, and select the sites that may form bonds; study
	and research Biosynthesis software such as DbD2, SSBOUND, MODIP, Bridge-D;
	conduct homology modeling of possible bond sites, observe bond formation, and
	measure each bond with PyMOL; FoldX energy evaluation. Study and analyze the
	thermodynamic explanation of the hydrophobic mechanism, the mechanism of
	the hydrophobic interaction between aromatic rings on the atomic surface, the
	analysis of the hydrophobic force between aliphatic chains at the atomic level, and
	the integration of different hydrophobic single point mutations.
Result	Design salt bridges and hydrogen bond networks.

2. The introduction of disulfide bonds in the homology region failed, and the mutant predicted by FoldX did not form disulfide bonds. There are 7 intersections between 10 MODIP Grade A sites and 40 DbD prediction sites: L43C-K252C, Y69C-A102C, V156C-S166C, A170C-V196C, P197-F261C, A202-E231C, A256C-F271C. Visually screen out two unreasonable mutations located in the protein, Y69C-A102C, V156C-S166C. FoldX screened out 3 Y69C-A102C, A170C-V196C, P197-F261C. Finally, successfully 2 L43C-K252C, A202-E231C.

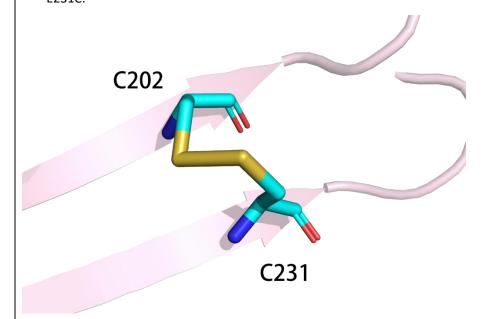


Figure 8 A202-E231C disulfide bond. The disulfide bond formed is showed in yellow.

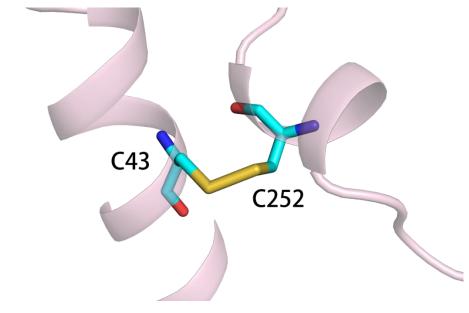
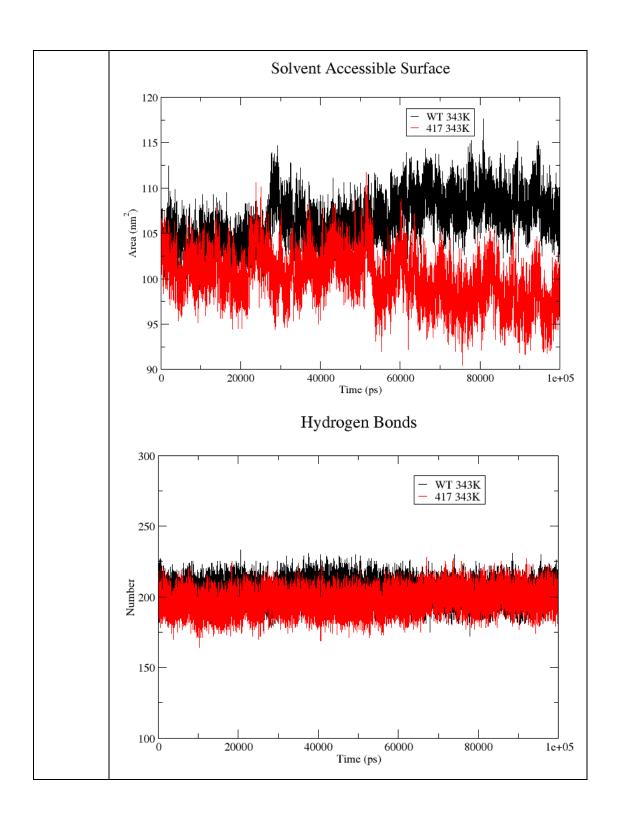
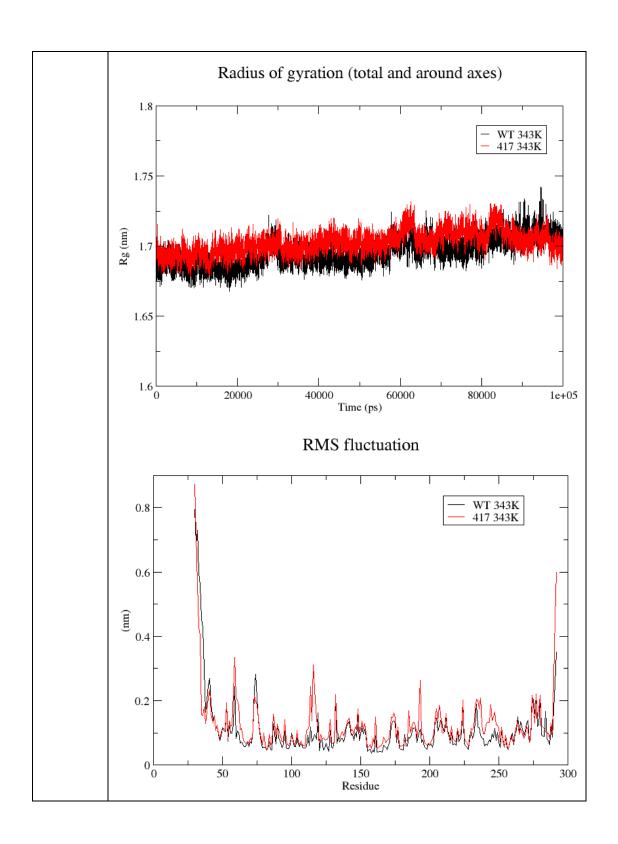


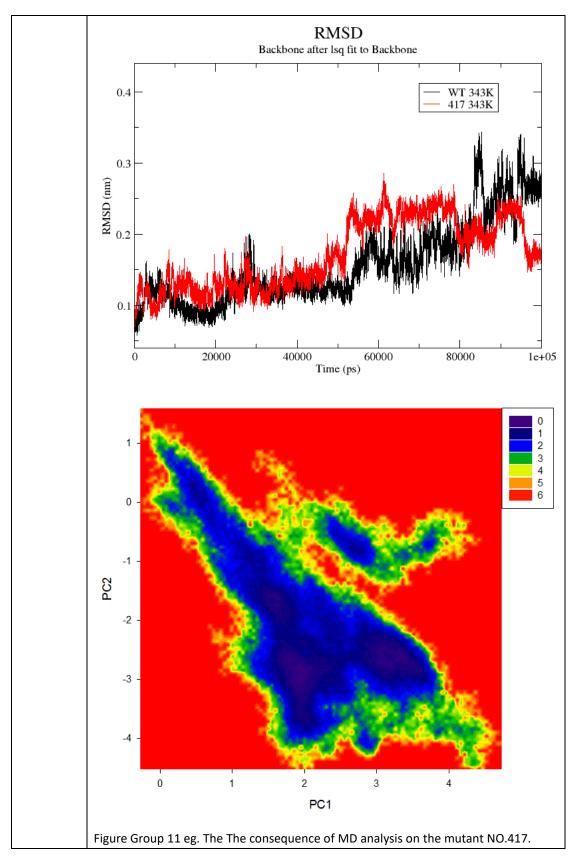
Figure 9 L43C-K252C disulfide bond. The disulfide bond formed is showed in yellow.

Date	08/16/2020-08/22/2020
Aim	1. Speed up MD operation;

2 Determine MD parameters and analysis methods and nut MD into practice
2. Determine MD parameters and analysis methods, and put MD into practice.
Yanjiao Lu, Kegen Chen, Yansong Bai
1. Try to run Gromacs on a high-configuration computer to increase computing
power;
2. Read the literature on the use of MD to study the thermostability of enzymes,
and make preliminary attempts on software and analytical methods.
Using software such as VMD xmgrace, we successfully analyzed the RMSD, RMSF,
gyration radius, solvent accessible surface area, hydrogen bond, salt bridge, free
energy topography, motion trajectory animation and other related content of small
molecule proteins at different temperatures. The MD simulation process is very
smooth. First, the protein structure energy is minimized, followed by protein
position restriction simulation (NVT and NPT), and finally simulation.
Molecular docking has been mastered and matured and can be directly applied.
Figure 10 Connection diagram of substrate binding pocket and PET

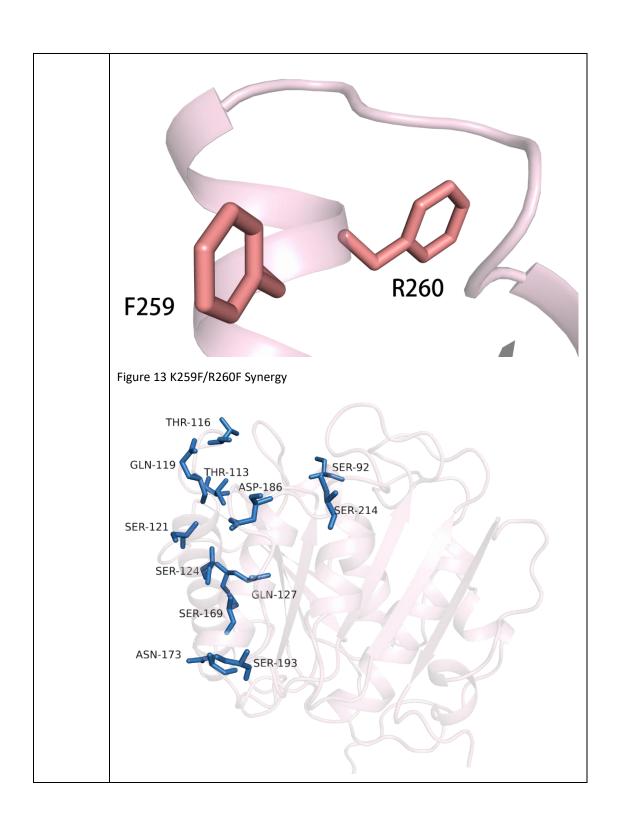


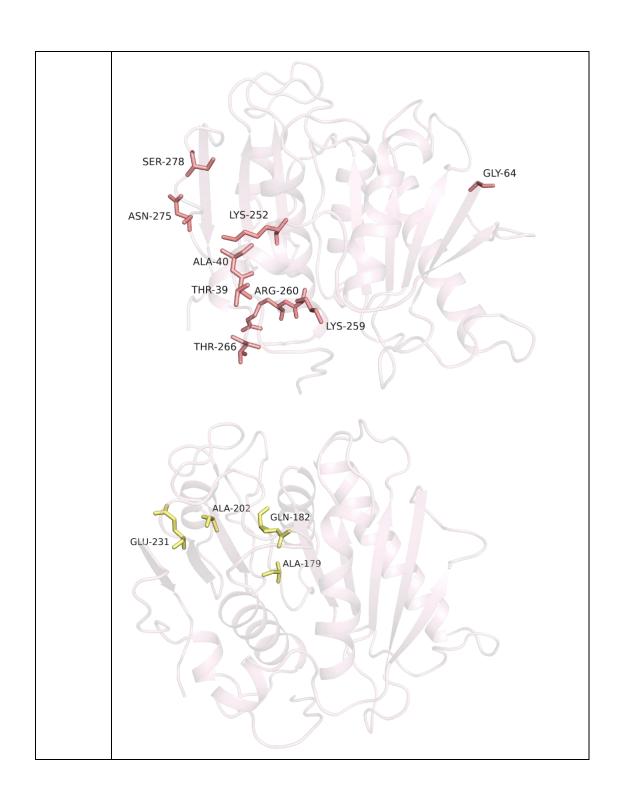


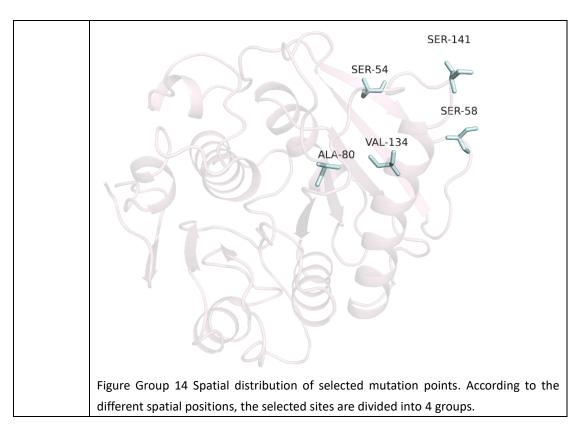


Date	08/23/2020-08/31/2020
Aim	1. Explore the feasibility of reducing DDG to a lower level by combining different

		force mutation points
	,	force mutation points.;
Doubleins	2.	Prepare the preliminary model for modeling.
Participant	All	
Content	2.	A combination of mutations with a single force cannot achieve the expected stabilization effect, try a combination of mutations with multiple forces. Salt bridges, hydrogen bonds, disulfide bonds, proline, and hydrophobic parts begin to integrate their mutations, and use FoldX to predict their DDG; Investigate all Best Models in the IGEM team in the past four years. Considering that our team's project is not suitable for biochemical reaction modeling, the modeling team collected relevant data on molecular docking, FoldX, Rosetta, and MD, and supplemented the prediction of TM based on FoldX.
Result	2.	Hundreds of multi-point combination mutants containing multiple forces were made. When multiple forces were combined on the same mutant, most of them had no effect on each other, but some combinations showed strong synergistic effect; Statistical summary of all the point mutations that significantly improved the structure of the team and significantly reduced the free energy of folding under the FOLDX prediction, and identified more than 20 point mutations that were finally used for combined mutations: T29D, A40P, S54W, S58P, A80V, S92P, T113P, T116R, Q119D, S121E/P, Q127L, S141P, Q182I, D186F, S193R, A202C, S214F, E231C, K252L, R260F, T266P, N275P/F. The basic principle of our combined mutation strategy is to average the single mutation in the spatial structure distribution and force type, and take into account the disulfide bond constructed by S121E / D186F, A202C/ E231C, S58P / S141P and other pairs of point mutations, or the synergistic effect of DDG value on improving thermal stability, a total of 543 twelve combination mutations were obtained, and energy prediction was started.
	Figu	ure 12 S121E/D186F Synergy
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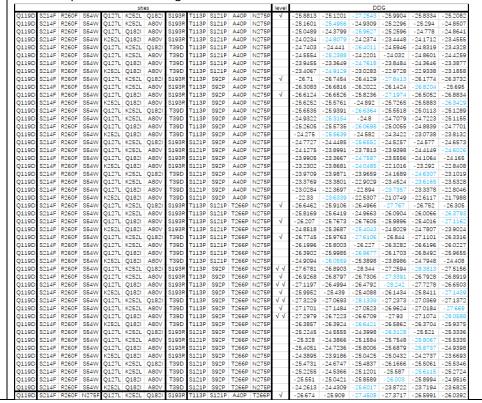
SEPTEMBER

WEEK1-2

09/01/2020-09/12/2020
1. Calculate 543 combination mutations of twelve-point mutations and screen
the results;
2. Analysis of combined mutant structure
Yuan Wang, Yuhan Li, Xujia Liu, Yufei Zhou, Dewu Liu, Haoyang Shi, He Zhang
Calculate the calculation of 543 combined mutations. For the obtained 543 combination mutants with superior DDG values, the two mutant combinations satisfying DDG<-28, -28 <ddg<-27 analysis,="" analyze="" analyzed,="" analyzing="" and="" assigned="" bonds="" bonds,="" bridges,="" changes="" charge="" different="" disulfide="" fields.="" hydrogen="" hydrophobicity.<="" in="" integrated="" jmol="" mutant,="" mutants="" number="" of="" performs="" person="" proline="" protein="" pymol,="" respectively,="" salt="" selected="" structural="" structure="" td="" the="" then="" these="" to="" using="" viewing="" visually="" were="" with=""></ddg<-27>
Completed the calculation of 543 combined mutations. 40 mutants satisfying DDG<-28 and nearly a hundred mutants satisfying -28 <ddg<-27 12="" 75f="" a202c="" best="" combination="" e231c="" initially="" mutants="" n114r="" n2="" n275<="" n37r="" n73r="" out.="" q119d="" q127l="" r260f="" s214f="" s54w="" screened="" screening,="" selected:="" t266p="" t270r="" t39d="" t72e="" t77e="" td="" the="" through="" visual="" were=""></ddg<-27>

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for subsequent MD screening.



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Q1190 Q1190	RESOF RESO	N275F R260F	SSAW SSAW SSAW SSAW SSAW SSAW SSAW SSAW	Q1271 Q127	K252L	Q1821 A80V A80V A80V A80V A80V A80V A80V A80V	\$193R \$193R	T113P S121P	992P 992P	A400P	72669 72669	\frac{1}{\sqrt{1}}	27.4007 -26.3686 -26.2708 -27.3812 -26.5094 -26.6399 -27.6374 -26.7693 -21.8184 -26.928 -25.973 -24.8184 -26.909 -25.9026 -25.2476 -26.909 -24.3621 -26.902 -24.9258 -24.9258 -24.4869 -26.5869 -2	-26.8755 -26.4855 -26.4855 -26.4856 -26.8344 -26.4679 -27.1276 -26.3896 -27.3819 -26.5132 -26.5132 -26.5132 -26.5132 -27.1085 -26.2752 -27.1085 -26.2752 -24.4847 -25.5844 -24.9776 -26.6645 -23.8938 -24.3518 -24.3518 -24.3518 -24.3518 -24.3518 -24.3518 -25.38037 -24.3518	-28.0379 -28.5707 -28.5707 -25.54511 -26.8321 -24.999 -26.3432 -26.7558 -27.459 -28.8669 -27.64054 -25.1286 -27.622 -25.892 -26.6187 -26.5728 -26.834 -24.1318 -24.9866 -25.327 -26.6187	28.2084 26.2077 26.9429 28.217 27.0215 27.70215 27.7619 22.4086 22.53694 24.5168 27.0741 25.1542 24.0887 25.932 26.8376 24.4263 26.4578 26.3032 22.2032 22.2032 23.3007 24.2087 25.3032 24.2087 25.3032 26.	-27.187 -26.9014 -26.9014 -27.8763 -27.8763 -26.7019 -26.4536 -27.5686 -27.5686 -27.5686 -27.5686 -27.5686 -27.5882 -25.533 -26.2217 -25.7479 -25.9659 -24.1925 -26.1116 -26.032 -26.0	-26.703 -26.7048 -26.2138 -27.5124 -26.6669 -26.5667 -26.5667 -26.315 -24.2035 -26.3665 -24.315 -25.9076 -24.315 -25.9076 -24.9459 -26.1624 -24.825 -24.8002 -24.8002 -24.8002 -24.908 -24.708 -25.9076 -24.8556 -25.9076 -25.9076 -25.9076	
Q1190 Q119	RESOF RESO	N275F	SSAW N276F	91271. 91271.	K252L	Q1821 A80V A80V A80V A80V A80V A80V A80V A80V	S199R S19PR S19PR	T113P S121P	992P 992P 992P 992P 992P 992P 992P 992P	A40P A40P A40P A40P A40P A40P A40P A40P	72669 T2669	\frac{1}{\sqrt{1}}	27.4007 -26.3686 -26.2708 -26.2708 -26.5034 -26.6928 -26.6339 -27.3812 -26.6339 -27.3814 -26.7693 -24.8184 -26.985 -	-26.8755 -26.4455 -26.4679 -26.8344 -26.4679 -27.1276 -26.3896 -27.3819 -26.5132 -25.75.48 -26.271 -26.512 -27.1085 -26.2762 -24.8447 -24.8447 -24.8762 -24.8011 -24.4762 -24.8011 -24.4762 -23.8938 -24.3518 -24.8037 -23.8938 -24.3518 -24.8037 -23.8938 -24.3518 -24.8037 -23.8938 -24.3518 -24.8037 -23.8938 -24.3518 -25.51305 -2	-28.0379 -28.5707 -28.5707 -28.5707 -28.9321 -24.999 -26.3432 -26.7658 -27.459 -28.266 -28.266 -28.266 -28.5092 -26.6187 -26.5726 -28.5374 -28.3374 -28.3374 -28.3374 -28.3374 -28.3374 -28.5226 -28.6384 -28.428877 -28.6684 -28.2684 -28.2684 -28.27 -28.26884 -28.217 -28.2884 -28.217 -28.2884 -28.217 -28.2884 -28.217 -28.2884 -28.217 -28.6884 -28.217 -28.6884 -28.217 -28.6884 -28.217 -28.6884 -28.217 -28.6884 -28.217 -28.6884 -28.217 -28.6884	28.2084 26.2027 -26.9429 -28.217 -27.0215 277.838 27.8411 -27.6413 -24.5168 -27.0741 -25.1542 -24.0887 -25.3657 -26.7619 -25.3657 -24.4083 -24.518 -26.502 -24.2518 -26.502 -24.2518 -26.503	-27.1787 -26.9014 -25.9014 -25.9014 -25.9014 -25.9014 -25.9014 -25.9014 -26.436 -27.8763 -26.436 -27.8666 -27.8666 -27.8666 -27.8666 -27.8669 -24.925 -26.1166 -27.8669 -24.925 -26.1166 -26.032 -26.0488 -26.8688 -24.8688	-26.703 -26.703 -26.703 -26.703 -26.703 -26.2138 -27.5124 -26.6669 -26.5637 -26.5637 -26.3615 -24.2035 -26.3615 -24.2035 -26.3615 -24.2035 -24.315 -25.4947 -25.1305 -24.9459 -24.8002 -23.5634 -24.8002 -23.5634 -24.8002 -24.2708 -24.4877 -25.9076 -24.4877 -25.9076 -24.8002 -24.909 -24.4177 -25.9076 -24.4877 -25.9076 -24.4877 -25.9076 -24.4877 -25.9076 -24.4877 -25.9076 -24.4877 -25.9076 -24.4877 -25.9076 -24.48856 -24.9076 -24.88856 -24.9076 -24.88856 -24.89076 -24.88856 -24.89076 -24.88856 -24.89076	
Q1190 Q1190	ESSOP ESSO	N275F R260F R260F R275F R260F R275F R260F R275F R260F R275F R260F R275F R260F	SSAW SSAW SSAW SSAW SSAW SSAW SSAW SSAW	91271. 91271.	K2561. K2581. K2581. K2581. K2581. Q1821 Q1821 Q1821 Q1821 Q1821 Q1821 Q1821 K2581.	Q1821 A80V A80V A80V A80V A80V A80V A80V A80V	Signar S	THISP SICIP	992P 992P 992P 992P 992P 992P 992P 992P	A40P A40P A40P A40P A40P A40P A40P A40P	72669 72669	\frac{1}{\sqrt{1}}	27.4007 -26.3686 -26.3686 -27.3812 -26.5094 -26.6399 -27.6374 -26.7693 -24.8184 -25.973 -24.8184 -25.973 -24.8184 -25.973 -24.8184 -26.6399 -25.3026 -26.505 -26.505 -26.505 -24.3621 -26.6902 -24.9258 -24.4789 -25.5006 -25.5006 -25.5006 -26	-26.8755 -26.4455 -26.4456 -26.4456 -26.8344 -26.4679 -27.1276 -26.3696 -27.3819 -26.5132 -25.7548 -25.2748 -25.2844 -27.1085 -26.275 -26.844 -25.944 -25.944 -25.944 -25.944 -24.951 -24.48011 -24.4705 -26.645 -23.9515 -25.844 -24.951 -25.844 -23.9515 -25.844 -23.9515 -25.844 -23.9515 -25.844 -23.9515 -25.844 -23.9515 -25.844 -23.9515 -25.844 -23.9515 -25.85844 -23.9515 -25.85844 -24.951 -25.85844 -24.951 -25.85844 -24.951 -25.85844 -24.951 -25.85844 -24.951 -25.85844 -24.951 -25.85844 -24.951 -25.85844 -24.951 -26.6545 -23.9515 -25.85844 -24.951 -24.951 -24.951	-28.0379 -28.5707 -28.5707 -25.5517 -28.9321 -24.999 -26.3432 -26.7558 -27.459 -25.8668 -27.459 -25.8668 -25.992 -25.75692 -25.75692 -25.6384 -24.1318 -24.8877 -25.0824 -25.3374 -24.1318 -24.76622 -25.728 -26.728 -26.728	28.2084 -26.2207 -26.9207 -28.217 -27.0215 -27.6213 -27.6413 -27.6713 -27.6713 -23.694 -24.5168 -27.0741 -25.1542 -24.0887 -26.7619 -25.3653 -26.4678 -26.4678 -26.5023 -26.5023 -26.5023 -26.5023 -26.5025 -26.50	-27.187 -26.9014 -26.9014 -26.9014 -27.8763 -27.8763 -27.8763 -26.102 -28.1166 -27.5686 -27.5686 -27.5686 -27.5686 -27.5686 -25.5686 -25.5681 -26.032	-26.703 -26.7049 -26.7049 -26.2138 -27.5124 -26.6669 -26.5637 -26.5122 -27.584 -26.4085 -24.2035 -26.3615 -24.2035 -24.2035 -24.215 -25.1305 -27.5941 -24.9459 -24.8856 -24.9459 -24.9459 -24.4477 -25.9424 -24.525 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9459 -24.9526 -24.9856	
Q1190 Q119	ENORGE E	N275F R260F	SSAW SSAW SSAW N275F SSAW SSAW SSAW N275F SSAW SSAW N275F SSAW SSAW SSAW SSAW SSAW SSAW SSAW SSA	91271, 912711, 912711, 912711, 912711, 912711, 912711, 912711, 912711, 9	K252L	Q1821 A80V A80V A80V A80V A80V A80V A80V A80V	11938 51938	T113P S121P	992P 592P	A40P A40P A40P A40P A40P A40P A40P A40P	72669 72669	\frac{1}{\sqrt{1}}	27.4007 26.3686 26.2708 27.3812 26.6928 26.6339 27.6374 26.7693 22.937 24.8184 26.937 24.8184 26.938 25.937 24.8184 26.938 25.2476 26.938 27.2482 28.2476 26.938 27.2482 28.2476 28.2476 24.9258 25.5036 24.174 24.3051 25.5036 24.174 24.3051 25.2476 26.505 27.2482 27.2	-26.8755 -26.4555 -26.4565 -26.8344 -26.4679 -27.1276 -26.3696 -27.3819 -26.5132 -25.7548 -25.2943 -25.2943 -25.2943 -24.4847 -25.59424 -24.4847 -25.59424 -24.4847 -25.59424 -24.4847 -25.59424 -24.4847 -25.59424 -24.4705 -25.59424 -24.4705 -25.59424 -24.4705 -25.59424 -24.4705 -25.59424 -24.4705 -25.5942 -24.4705 -25.5942 -24.4705 -25.5942 -24.5945 -25.5942 -24.5945 -25.5942 -24.5945 -25.5942 -24.5945 -25.5942 -24.5945 -25.5942 -24.5945 -25.5942 -24.5945 -25.5942 -24.5945 -25.5942 -24.5945 -25.5942 -24.5945 -25.5942 -24.5945 -25.5942 -24.5945 -25.5942 -24.5945 -25.5944 -24.5945 -25.5944 -24.5945 -25.5944 -24.5945 -25.5944 -24.5945 -25.5944 -24.5945 -25.5944 -25.5945 -25.5944 -25.5945 -25.5944 -25.5945 -25.5944	-28.0379 -28.5707 -28.5707 -28.5707 -24.999 -26.3432 -26.7558 -27.459 -28.6688 -28.1286 -27.0622 -25.892 -25.5992 -25.66187 -26.5728 -26.6187 -26.6187 -26.6187 -26.6187 -26.6187 -26.6187 -26.6187 -26.6187 -26.6187	28.2084 26.2027 -26.9429 -28.217 -27.0215 277.8418 -27.6418 -24.6168 -24.5168 -24.5168 -24.5169 -24.5169 -24.0887 -25.3657 -26.7619 -26.7619 -26.7619 -26.3063 -26.5032 -24.818 -26.5032 -24.818 -25.3039 -25.3039 -26.7619 -	-27.1787 -26.9014 -25.9014 -25.9014 -25.9014 -25.9014 -25.9014 -25.9016 -25.9701 -26.4536 -25.9701 -24.9469 -25.523 -25.523 -25.523 -26.9701 -26.9696 -25.9701 -26.9696 -26.9701 -26.9696 -26.9701 -26.97	-26.703 -26.703 -26.704 -26.2138 -27.5124 -26.6669 -26.5637 -26.1512 -27.584 -26.4085 -26.3615 -24.2035 -26.1956 -24.315 -24.2035 -26.1956 -24.315 -24.9459 -26.1624 -24.9459 -24.856 -24.826 -24.826 -24.826 -24.9444 -27.5941 -24.4847 -27.5941 -24.4847 -27.5941 -24.4847 -24.4856 -24.8866	
Q1190 Q1190	ECOP	N275F R250F N275F R260F N275F R260F N275F R260F N275F R260F N275F R260F	SSAW SSAW N275F SSAW SSAW SSAW SSAW SSAW SSAW SSAW SSA	91271, 912711, 912711, 912711, 912711, 912711, 912711, 912711, 912711, 9	KCSSLL KCSLL KCSSLL KCSLL KCSSLL KCSLL KCSLL KCSL KCS	Q1821 A80V A80V A80V A80V A80V A80V A80V A80V	Signar S	THISP SICIP	992P 992P 992P 992P 992P 992P 992P 992P	A40P A40P A40P A40P A40P A40P A40P A40P	12669 12669	\frac{1}{\sqrt{1}}	27.4007 26.3686 26.2708 27.3812 26.6092 26.6339 27.6374 26.7692 25.937 24.8184 26.937 25.937 24.8184 26.938 25.2476 26.938 25.2476 26.938 25.2476 26.938 27.2482 24.1376 25.24489 24.1376 25.2476 26.2482 27.2482	-26.8755 -26.4555 -26.4565 -26.8344 -26.4679 -27.1276 -26.3896 -27.3819 -26.5132 -25.75.48 -25.942 -24.8477 -25.9424 -24.9776 -26.6545 -23.9515 -24.750 -26.024 -23.8938 -24.711 -24.366545 -23.8938 -24.712 -24.366545 -25.5646 -23.8938 -24.713 -24.366545 -25.5646	-28 0379 -28 5707 -28 5707 -28 5707 -28 5821 -24 599 -26 3432 -25 5982 -27 459 -28 6688 -26 4054 -25 1286 -25 5992 -25 5992 -25 5992 -26 5872 -26 5	28.2084 26.2027 -26.9429 -28.217 -27.0215 -27.7036 -27.6411 -26.713 -24.6168 -24.6168 -24.6168 -24.6169 -25.1642 -24.0887 -25.3657 -24.0887 -25.0987 -24.0887 -24.0887 -24.0887 -24.0887 -25.0987 -24.0887 -24.0887 -24.0887 -25.0987 -24.0887 -24.0887 -25.0987 -24.0887 -24.0887 -25.0987 -24.0887	-27.1787 -26.9014 -25.9014 -25.9014 -25.9014 -25.9014 -25.9016 -26.1409 -26.1409 -26.1409 -27.5686 -25.9741 -24.9489 -25.533 -25.533 -25.533 -26.0419 -26.032 -26.0488 -25.533 -24.5742 -25.56869 -25.5032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.032 -26.032 -26.032 -26.032 -26.032 -26.032 -26.032 -26.032 -26.032 -26.032 -26.032 -26.032 -26.032 -26.038	-26.703 -26.7049 -26.7049 -26.2049 -26.2138 -27.5124 -26.6669 -26.5637 -26.5122 -27.5844 -26.4085 -24.2035 -24.2035 -24.2035 -24.315 -25.49459 -26.15624 -24.4525 -24.525 -24.2085	
Q1190 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q1	RESOF RESOFT RESOF RES	N275F R260F	SSAW SSAW N275F SSAW SSAW SSAW SSAW SSAW SSAW SSAW SSA	91271, 91	K2551. K2521. K2	Q1821 A80V A80V A80V A80V A80V A80V Q1821 Q1821 A80V A80V A80V A80V A80V A80V A80V A80V	11938 11	THISP SICIP	992P 992P 992P 992P 992P 992P 992P 992P	A40P A40P A40P A40P A40P A40P A40P A40P	72669 72669	\frac{1}{\sqrt{1}}	27.4007 -26.3686 -26.2708 -26.2708 -26.5034 -26.6928 -26.6339 -26.6339 -27.3812 -26.6339 -26.6339 -21.8184 -26.928 -24.8184 -26.928 -25.2476 -26.909 -24.3621 -26.909 -24.3621 -26.909 -24.4925 -24.4925 -24.4925 -24.4925 -26.902 -24.4826 -26.903 -24.4826 -26.903 -24.4826 -26.903 -24.4826 -26.903 -24.4826 -26.903 -24.4826 -26.903 -24.4826 -26.903 -24.4826 -26.903 -24.4826 -26.903 -24.4826 -26.903 -24.3621 -26.903 -24.4826 -26.903 -26.	-26.8755 -26.4855 -26.4856 -26.8344 -26.4679 -27.1276 -26.3896 -27.3819 -26.5132 -26.5132 -27.1285 -26.2752 -27.1285 -26.2752 -24.4847 -25.5844 -24.4847 -25.5844 -24.48611 -24.476 -26.624 -23.8938 -24.3518 -26.024 -24.88037 -25.5844 -24.88037 -25.5844	-28.0379 -28.5707 -28.5707 -28.5707 -28.9321 -24.999 -26.3432 -26.7558 -27.459 -28.26668 -27.622 -25.892 -26.6187 -26.5728 -26.6384 -24.1318 -24.8877 -26.024 -25.3274 -26.6384 -24.1318 -24.8877 -26.672	28.2084 -26.2077 -26.9479 -28.217 -27.0215 -27.70215 -27.6713 -27.4395 -24.5168 -27.0741 -25.1542 -24.0887 -25.3657 -26.7619 -25.932 -24.818 -26.4576 -26.3630 -24.2544 -25.132 -26.3630 -24.2544 -25.132 -26.3630 -24.2544 -25.132 -26.3630 -24.2544 -26.3630 -24.2544 -26.3630 -24.2544 -26.3630 -24.2544 -26.3630 -24.2544 -26.3630 -24.2544 -26.3630 -24.2544 -26.3630 -24.2544 -26.3630 -24.2544 -26.3630 -2	-27.1787 -26.9014 -26.9014 -27.8763 -27.8763 -27.8763 -26.1409 -26.14536 -27.8686 -27.8686 -27.8686 -27.8686 -27.8686 -28.7228 -26.533 -26.21925 -26.1116 -24.9489 -25.5971 -26.4799 -26.9659 -24.1925 -26.1116 -26.0488 -26.86888 -26.8688 -26.868888 -26.868888 -26.86888 -26.868888 -26.868888 -26.868888 -26.868888 -26.868888 -26.868888 -26.868888 -26.868888 -26.8688888 -26.8688888 -26.8688888 -26.86888888 -26.868888888888888888888888888888888888	-26.703 -26.7048 -26.2138 -27.5124 -26.6669 -26.5667 -26.5667 -26.315 -24.2035 -26.3666 -24.315 -25.9076 -24.9459 -26.1624 -24.8052 -24.8052 -24.2088	
Q1190 Q119	RESOF RESOFT RESOF RESOFT RESOF RESOFT RESOTT	N275F R260F	SSAW N275F	01271. 01271.	K2551. K2521. K2	Q1821 A80V A80V A80V A80V A80V A80V A80V A80V	11938 51	THISP SHIP SHIP SHIP SHIP SHIP SHIP SHIP	992P 992P 992P 992P 992P 992P 992P 992P	A40P A40P A40P A40P A40P A40P A40P A40P	1266p	\frac{1}{\sqrt{1}}	27.4007 26.3686 26.2708 27.3812 26.6928 26.6928 26.6329 27.6374 26.7692 27.25937 24.8184 25.973 24.8184 25.973 24.8184 25.973 24.8184 26.909 25.2476 26.6505 25.1764 26.0099 24.3621 25.902 24.3621 25.902 24.1376 25.5497 24.4899 24.1408 25.1764 26.009 24.1376 25.1764 26.009 24.1376 25.2476 26.25036 26.1376 26.27174 24.001 25.006 25.1764 26.009 24.1376 25.1764 26.009 24.1376 25.1764 26.009 24.1376 25.1764 26.009 24.1376 25.1764 26.009 24.1376 25.1764 26.009 24.1376 25.1764 26.009 24.1376 25.1764 26.009 24.1376 25.1764 26.009 24.1376 25.1764 26.009 24.1376 25.1764 26.009 24.1376 25.1764 26.009 24.1376 25.1764 26.009 24.1476 24.14776 24.1476 24.1476 24.1476 24.1476 24.1476 24.1476 24.14776 24.1476 24.1476 24.1476 24.1476 24.1476 24.1476 24.14776 24.1476 24.1476 24.1476 24.1476 24.1476 24.1476 24.147776 24.14776 24.14776 24.14776 24.14776 24.14776 24.14776 24.14776 24.14776 24.14776 24.14776 24.14776 24.14776 24.14776 24.14776 24.14776 24.14776 24.14776 24.14776 24.147776 24.147776 24.14776 24	-26.8755 -26.4555 -26.8344 -26.4679 -27.1276 -26.3696 -27.3819 -26.5122 -26.7548 -26.277,1025 -26.275,202 -26.275,	-28 0379 -28 5707 -28 5707 -28 5707 -24 999 -26 3432 -26 7598 -27 459 -28 5698 -27 459 -28 5698 -27 459 -28 5698 -26 4054 -25 126 -25 5992 -26 5892 -26 5892 -26 5892 -26 5892 -26 5892 -26 5892 -26 5892 -26 5892 -26 5892 -26 5892 -26 5892 -26 5892 -26 5892 -24 1365 -24 1365 -24 1365 -24 1365 -24 126 572 -25 5931 -26 572 -26 572 -26 572 -26 572 -27 479 -26 5872 -	28.2084 26.2027 -26.9429 -28.217 -27.0215 -27.7038 -27.6411 -27.6713 -24.6169 -24.5169 -24.5169 -24.0887	-27.1787 -26.9014 -25.9118 -27.8763 -26.7019 -26.4536 -26.1409 -28.1166 -27.5686 -25.9741 -24.9489 -27.2882 -25.533 -25.5217 -25.7479 -26.1166 -25.9619 -24.1925 -24.1925 -24.1925 -24.1925 -24.1925 -25.9659 -24.1925 -25.9659 -24.1925 -25.9659 -24.1925 -25.9659 -24.1925 -26.116 -25.9659 -24.1925 -26.116 -25.9659 -24.1925 -26.116 -25.9659 -24.1925 -26.116 -26.0032 -26.0488 -25.9659 -24.1925 -26.116 -26.0032 -26.0488 -26.032 -26.0488 -28.5931 -26.0583 -26.0583 -26.0583 -26.0583 -26.0584 -26.0583 -26.0584 -26.0583 -26.0584 -26.0584 -26.0585 -	-26.703 -26.703 -26.704 -26.2138 -27.5124 -26.6669 -26.5637 -26.5637 -26.4085 -26.3615 -24.2035 -26.1582 -26.3615 -24.315 -25.4947 -25.1305 -24.315 -25.4947 -25.1305 -24.315 -24.49459 -24.4526 -24.826	
Q1190 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q1	ECOP	N275F R260F R260F R275F R260F R250F	SSAW N275F	91271. 91271.	K2551. K2521. K2	Q1821 A80V A80V A80V A80V A80V A80V Q1821 Q1821 A80V A80V A80V A80V A80V A80V A80V A80V	11938 51938	THISP SICIP	992P 992P 992P 992P 992P 992P 992P 992P	A40P A40P A40P A40P A40P A40P A40P A40P	72669 72669	\frac{1}{\sqrt{1}}	27.4007 -26.3686 -26.2708 -26.2708 -26.5024 -26.5024 -26.5024 -26.5024 -26.7693 -27.3812 -26.5034 -26.7693 -24.8184 -26.9857 -24.8399 -25.3026 -25.2476 -26.9857 -24.3621 -25.5026 -24.3621 -24.9258 -25.5036 -24.4309 -24.4809 -24.4809 -24.4809 -24.2524 -26.505 -26	-26.8755 -26.4455 -26.4679 -26.8344 -26.4679 -27.1276 -26.3896 -27.3819 -26.5132 -25.75.48 -26.271 -26.2806 -27.3819 -26.5132 -26	-28 0379 -28.5707 -28.5707 -28.5821 -24.999 -26.3432 -26.7658 -27.459 -28.3660 -28.2660 -28.5992 -26.6187 -26.374 -28.384 -28.	28.2084 26.2027 -26.9429 -28.217 -27.0215 277.8411 -27.6413 -27.4985 -24.5168 -27.0741 -25.1542 -24.0887 -25.3657 -26.7619 -25.3657 -26.7619 -24.4813 -26.4578 -26.3063	-27.1787 -26.9014 -26.9014 -27.8763 -27.8763 -27.8763 -26.7019 -26.4336 -27.8664 -27.8666 -27.8666 -27.8666 -27.8666 -27.86669 -26.102 -26.102 -26.032 -26.0488 -26.8680	-26.703 -26.7048 -26.2138 -27.5124 -26.6669 -26.5637 -26.5637 -26.5124 -26.3615 -24.2035 -26.3615 -24.2035 -26.3615 -24.2035 -24.315 -25.4947 -25.1305 -24.9459 -24.152 -24.9459 -24.952	
Q1190 Q119	ECOP	N275F R260F	SSAW N275F	91271. 91271.	K2551, K2	Q1821 A80V A80V A80V A80V A80V A80V A80V Q1821 A80V A80V A80V A80V A80V A80V A80V A80V	11938 51	THISP SHIP SHIP SHIP SHIP SHIP SHIP SHIP	992P 992P 992P 992P 992P 992P 992P 992P	A40P A40P A40P A40P A40P A40P A40P A40P	7266 72	\frac{1}{\sqrt{1}}	27.4007 26.3686 26.2708 27.3812 26.6928 26.6339 27.677 26.7673 22.873 22.873 23.873 24.8184 26.987 25.4339 25.207 26.8309 27.20987 26.8309 27.20987 28.809 27.20987 28.8090 24.8090 24.9258 25.5036 24.9258 25.5049 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 24.9258 25.5049 24.9258 25.5049 24.9258 25.5049 24.9258 25.9258 26.9258	-26.8755 -26.4855 -26.4854 -26.4679 -27.1276 -26.3696 -27.3819 -26.5132 -25.7548 -25.2943 -25.2943 -25.2943 -24.4847 -25.2943 -24.4847 -25.2943 -24.4847 -25.2943 -24.4847 -25.2943 -24.4705 -25.29542 -24.4847 -25.29545 -25.2954	-28 0379 -28.5707 -28.5707 -28.5821 -24.999 -26.3432 -26.7558 -27.459 -28.6688 -28.1286 -27.0622 -25.892 -25.5992 -26.6187 -26.5728 -26.6187 -26.5728 -26.6187 -26.5728 -26.6187 -26.5728 -26.6187 -26.5728 -26.57	28.2084 26.2027 -26.9429 -28.217 -27.0215 277.8418 -27.6418 -24.5168 -24.5168 -24.5168 -24.5169 -24.5169 -24.0887 -25.3657 -26.7619 -26.7619 -26.3063 -26.5002 -24.8181 -25.3063 -26.5002 -24.2087 -24.2087 -25.3063 -26.5002 -24.20887 -24.2088 -26.3063	-27.1787 -26.9014 -25.9014 -25.9014 -25.9014 -25.9014 -25.9014 -25.9016 -26.7019 -26.4536 -25.9741 -24.9469 -25.523 -25.523 -25.523 -26.924 -26.923 -26.924 -2	-26.703 -26.703 -26.7048 -26.2138 -27.5124 -26.6669 -26.5637 -26.1512 -27.584 -26.4085 -26.3615 -24.2035 -26.1956 -24.315 -24.2035 -26.1956 -24.315 -24.2035 -24.9459 -26.1624 -24.9459 -26.1624 -24.952 -26.2036 -24.8566 -24.8002 -23.9687 -24.2708 -24.7919 -24.44847 -27.5941 -24.9459 -24.44847 -27.5941 -24.9459 -24.4705 -26.238 -24.2708 -24.4856 -24.8002 -24.4705 -26.238 -24.27108 -24.24.8866 -24.8002 -24.4705 -26.238 -24.2711 -26.238 -24.2711 -26.238 -24.38037	
Q1190 Q119	ESOP	N275F R260F	SSAW SSAW N275F SSAW N	91271. 91271.	K2521, K2	Q1821 A80V A80V A80V A80V A80V Q1821 Q1821 A80V A80V A80V A80V A80V A80V A80V A80V	1032 1032 1032 1032 1032 1032 1032 1032	T113P S121P S12P S12	992P 992P 992P 992P 992P 992P 992P 992P	A40P A40P A40P A40P A40P A40P A40P A40P	1266p 126p 12	\frac{1}{\sqrt{1}}	27.4007 -26.3686 -26.2708 -27.3812 -26.5034 -26.6928 -26.6339 -27.6374 -26.7693 -24.8184 -25.5036 -25.277 -24.8184 -26.000 -25.277 -24.8184 -26.000 -25.277 -24.8184 -26.000 -25.277 -24.3621 -26.000 -24.3621 -26.902 -24.9258 -24.1376 -26.503 -26.503 -26.	-26.8755 -26.4855 -26.4856 -26.48679 -27.1276 -26.3896 -27.3819 -26.5132 -26.5132 -26.5132 -27.1208 -26.5132 -27.1208 -26.5132 -27.1208 -26.5132 -24.4847 -25.5844 -24.4847 -25.5844 -24.9776 -26.6545 -23.9515 -25.1589 -24.3818 -24.4705 -26.6545 -23.9515 -25.52844 -24.8037 -23.8938 -24.3818 -24.8037 -23.8938 -24.3818 -24.58694 -23.8938 -24.3818 -24.58694 -23.8938 -24.3818 -24.977 -25.1305 -25.51899 -24.3818 -24.977 -25.1305 -25.51899 -24.3818 -24.977 -25.1305 -25.51899 -24.3818 -24.977 -25.1305 -25.51899 -24.3818 -24.977 -25.1305 -25.51899 -24.3818 -24.977 -25.2891 -24.977 -25.2891 -25.24708 -24.4777 -25.2891 -25.247919 -24.44777 -25.2891 -25.247919 -25.247919	-28.0379 -28.5707 -28.5707 -28.5707 -28.9321 -24.999 -26.3432 -26.7558 -27.459 -28.9669 -27.0622 -25.892 -25.7656 -28.5092 -26.6187 -24.1318 -24.8877 -25.0824 -25.3374 -24.9806 -25.57266 -25.6726 -25.5726 -26.6187 -26.0824 -26.1383 -24.0684 -26.217 -26.0824 -26.217 -26.0824 -26.217 -26.0824 -26.217 -26.0824 -26.217 -26.0824 -26.217 -26.0824 -26.217 -26.0824 -26.218 -26.217 -26.0824 -26.218	28.2084 26.2027 26.2027 26.2027 27.0215 27.70316 27.6713 27.6713 27.6713 27.6713 22.4895 24.5168 27.0741 25.1542 24.0827 25.932 25.8376 24.4813 26.4578 25.3032 24.2544 25.132 25.3032 24.2544 25.132 26.2532 27.254	-27.1787 -26.9014 -26.9014 -26.9014 -27.8763 -27.8763 -27.8763 -26.102 -26.102 -26.102 -26.102 -26.102 -26.102 -26.102 -26.102 -26.102 -26.202 -26.203	-26.703 -26.7048 -26.7048 -26.2138 -27.5124 -26.6669 -26.5637 -26.5612 -27.584 -26.4085 -26.315 -24.2035 -26.3615 -24.2035 -24.315 -25.49459 -26.1562 -24.315 -24.9459 -26.1624 -24.952 -24.95	
Q1190 Q119	RESOF RESOFT RESOF RESOFT RESOF RESOFT RESOF	N275F R260F	SSAW SSAW N275F SSAW N275F SSAW N275F SSAW SSAW N275F SSAW SSAW N275F SSAW SSAW N275F SSAW SSAW N275F SSAW SSAW N275F SSAW SSAW N275F SSAW SSAW SSAW SSAW SSAW SSAW SSAW SSA	01571 01	K2521, K2	Q1821 A80V A80V A80V A80V A80V A80V A80V A80V	11938 51	THISP SHIP SHIP SHIP SHIP SHIP SHIP SHIP	992P 992P 992P 992P 992P 992P 992P 992P	A40P A40P A40P A40P A40P A40P A40P A40P	7.266p	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	27.4007 -26.3686 -26.2708 -27.3812 -26.5034 -26.6928 -26.6339 -27.6374 -26.7693 -24.8184 -25.5036 -25.277 -24.8184 -26.000 -25.277 -24.8184 -26.000 -25.277 -24.8184 -26.000 -25.277 -24.3621 -26.000 -24.3621 -26.902 -24.9258 -24.1376 -26.503 -26.503 -26.	-26.8755 -26.4855 -26.4854 -26.4679 -27.1276 -26.3896 -27.3819 -26.5132 -25.75.48 -25.2943 -27.1095 -26.295 -27.1095 -26.295 -26.295 -27.1095 -26.295 -26.295 -27.1095 -26.295	-28 0379 -28 5707 -28 5707 -28 5707 -28 5821 -24 599 -26 3432 -25 5982 -27 459 -28 6688 -26 4054 -25 1286 -25 5992 -25 5992 -26 5872	28.2084 26.2027 -26.9429 -28.217 -27.0215 -27.7036 -27.6416 -25.3694 -24.5168 -24.5169 -24.5169 -24.0887 -25.3657 -25.3657 -24.0887 -24.0887 -24.0887 -24.0887 -24.0887 -24.0887 -24.0887 -24.0887 -24.0887 -24.0887 -24.0887 -24.0887 -24.0887 -24.0887 -24.0887 -25.3063 -25.3063 -25.3063 -26.5069 -24.2648 -26.5069 -24.2648 -26.5069 -24.2648 -26.5069 -24.2648 -26.5069 -24.2648 -26.5069 -24.2644 -25.3069 -24.2644 -25.3069 -24.2644 -25.3069 -24.2644 -25.3069 -24.2644 -25.3069 -24.2644 -25.3069 -24.2644 -25.3069 -24.26889 -24.2	-27.1787 -26.9014 -25.9014 -25.9014 -25.9014 -25.9014 -25.9014 -26.436 -26.1409 -26.1409 -26.1409 -27.5686 -25.9741 -24.9489 -25.533 -25.533 -25.533 -26.0419 -25.0032 -26.0489 -25.533 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.032 -26.0488 -26.0382	-26.703 -26.703 -26.7048 -26.2138 -27.5124 -26.6669 -26.5637 -26.1582 -27.5841 -26.1986 -24.315 -24.2035 -26.1986 -24.315 -24.315 -24.2035 -24.315 -24.315 -24.49459 -26.1624 -24.9459 -26.1624 -24.9459 -26.1624 -24.8002 -23.8938 -24.711 -24.94847 -25.944847 -25.946 -24.8002 -24.4002 -24.4002 -24.4002 -24.4002 -24.4002 -24.4002 -24.4002 -24.4002 -24.4002 -24.4002 -24.4002 -24.4002 -24.5002 -24.2002	

	Q119D	S214F	R260F	\$54W	Q127L	K252L	A80V	T39D	T113P	S121P	N275P	T116R		-25.8938	-25.7434	-25.8448	-25.5908	-25.8286	-26.4613
Column C		S214F	R260F	S54W	Q127L	Q182I	A80V	T39D	T113P	S121P	N275P	T116R		-25.6838	-25.1699	-26.0729	-25.298	-26.5782	-25.3
Column C	Q119D	S214F	R260F	S54W	Q127L	K252L	Q182I	S193R	T113P	S92P	N275P	T116R	√	-27.0682	-26.1135	-27.099	-27.5565	-27.2723	-27.2998
Column C													-1						
Quick Start Start Start August Start August August	· · · · · · · · · · · · · · · · · · ·	S214F	R260F		Q182I	K252L	A80V	S193R	T113P	S92P	N275P	T116R	•	-25.3054	-25.9222	-25.2792	-24.7835	-24.9591	-25.5831
Quick Claim South South South Claim Claim South Claim Clai	•																		
Quino State State State Quino State				S54W	Q127L	Q182I	A80V	T39D	T113P	S92P	N275P	T116R			-26.1396	-27.4489	-26.2706	-26.6171	-25.8572
Quick Size Size Size Quick Color Anno Size Size													√						
QUILD SLIP PORT SERV QUIL NELL AUV STEP CHIP SERV PORT TARK CARRY	Q119D	S214F	R260F	S54W	Q127L	K252L	A80V	S193R	S121P	S92P	N275P	T116R		-24.7186	-25.2963	-25.2315	-24.7329	-24.5208	-23.8114
CLISP CLIP SEAR SEAR CLIP																			
QUISS DIAP SEAR SEAR QUIT, QUIT, QUIT, QUIT, AND TABLE DIAP TABLE DIAP C. SEAR DIAP DIAP C. SEAR DIAP DIAP DIAP C. DIAP DIAP C. DIAP DIAP C. DIAP DIAP	Q119D	S214F	R260F	\$54W	Q127L	K252L	Q182I	T39D	S121P	S92P	N275P	T116R		-25.1994	-25.3581	-25.8951	-24.8468	-25.4586	-24.4386
QUISSO BLIFF FORE SEAN QUITA CELLS AREN TROP SILVEY SEAN PROTECTION AND ADMINISTRATION								T39D	S121P	S92P	N275P	T116R							
QUISSO DILLEY FORCE SEAN QUITA, CRISTA, ARON SURPH TILES CRISTA	Q119D	S214F	R260F	S54W	Q182I	K252L	A80V	T39D	S121P	S92P	N275P	T116R		-23.6428	-23.1681	-23.0629	-23.4339	-24.5413	-24.008
QUISS DILE* PROF. SEMP QUITY, QUITY, QUITY, AND SEMP SEMP SEMP TIPP TIPP													√						
Q1100 2114 PROF. 5440 Q271, C281, Q381 790 7118 AAD NOTE 7118 26034 2679 24871 24112 25468 24182	Q119D	S214F	R260F	S54W	Q127L	Q182I	A80V	S193R	T113P	A40P	N275P	T116R		-26.1364	-25.9085	-26.0543	-26.9029	-26.4365	-25.3801
Q1100 2114 EXCOR 5540 Q177, Q151, A007 300 7119 A00 A077 71140																			
Quies California Series California	Q119D	S214F	R260F	S54W	Q127L	K252L	A80V	T39D	T113P	A40P	N275P	T116R		-25.0038	-24.3835	-24.7652	-25.5782	-24.6388	-25.6535
Q1906 2114 PEGGS 56440 Q271, NSEL Q081 3189 7118 7168 7178 7108																			
Califor Colley Record Servey (March 1980) Servey No. Servey No.																			
Q1100 D116F RODG SAMP Q110F RODG ASSAN Q100F Q110F Q100F Q110F Q100F Q110F Q100F Q110F Q100F Q110F Q100F Q110F Q11	Q119D	S214F	R260F	\$54W	Q127L	K252L	A80V	S193R	T113P	T266P	N275P	T116R		-26.4418	-26.6974	-27.0528	-26.7541	-25.3889	-26.3157
Q1100 G114F REGG SEAW G171, MSSL, Q182, 1300	Q119D	S214F	R260F	S54W	Q182I	K252L	A80V	S193R	T113P	T266P	N275P	T116R		-25.8863	-26.4121	-26.3069	-25.5212	-25.9046	-25.2867
Q1100 S114 R500 S404 Q171 Q181 A00 T300 T310 T300 T300 T310	Q119D	S214F	R260F	S54W	Q127L	K252L	Q182I	T39D	T113P	T266P	N275P	T116R		-27.3528	-27.2208	-27.089	-27.7903	-27.6867	-26.9772
Q1150 S116 R500 S404 Q121 K511 A80 T300 T310 T300 T310 T500 M178 T116 C500 C5004 C5005 C4004 C5055 C4010 C3115 C505 C3115 C400 K511 C400 C5004 C5055 C4010 C3115 C5055 C5004 C5055								T39D	T113P	T266P	N275P	T116R							
QUISS DILF REGOT Servi QUITT, INSEL, 1807 1908 SUISF ALGO NUTSF TLER 2.0242 23.646 2.0262 2.046 2.026	Q119D	S214F	R260F	S54W	Q182I	K252L	A80V	T39D	T113P	T266P	N275P	T116R		-26.0504	-25.6584	-25.8367	-25.3886	-26.5773	-26.7913
Q1100 S114 REGO S640 Q171 KSC1 A80 S288 S112 A80 N278 T168 2.027 2.077 2.052 2.162 2.052 2.072 2.062 2.072 2.062 2.072 2.062 2.072 2.062 2.072 2.062 2.072 2.062 2.072 2.062 2.072 2.062 2.072 2.062 .														-25.3265 -24.2424	-25.1646	-24.7384 -24.2262	-25.5625 -23.6619	-25.6179 -24.2422	-24.3126 -23.9174
Q1100 0114F R200F SEAVQ Q12T, ISSL. [Q120] YBDD 5111F ADDF WCTSF T1160	Q119D	S214F	R260F	\$54W	Q127L	Q182I	A80V	S193R	S121P	A40P	N275P	T116R		-24.2457	-23.9071	-24.7052	-24.1149	-24.9306	-23.5706
Q1100 S114F R260F SEWQ G1271, ICSSL, ABOV TOD S1112F AGD9 NCTSP T1168														-24.3081	-25.2265	-23.4992	-25.2191	-23.465	-24.1308
Q1100 S114F R650F S84W Q17T K551, Q851 S108F S112F T660F N75F T116F	Q119D	S214F	R260F	S54W	Q127L	K252L	A80V	T39D	S121P	A40P	N275P	T116R		-23.5639	-24.1682	-23.5427	-23.311	-23.2039	-23.5935
Q1100 S114F R650F S84W Q17T K551, Q851 S108F S112F T660F N75F T116F														-23.4814	-23.6476 -23.6062	-24.338 -23.9201	-23.9701	-22.774 -22.578	-22.6774
Q1100 S114F R260F S8AW Q17T, Q1821 ABOV S108P S112F TS6P N37F T116F 25.017 22467 224	Q119D	S214F	R260F	S54W	Q127L	K252L	Q182I	S193R	S121P	T266P	N275P	T116R		-25.8865	-25.2731	-25.9749	-25.2303	-26.7769	-26.1776
Q1150 S114F REOF SEAVY Q12F1 RESTAL ABOV S1398 S112F TESEP N17FF T116F 2.69073 2.3427 2.38077 2.4411 2.36683 2.1368 Q1150 S114F REOF SEAVY Q117L RESTAL ABOV T800 S112F TESEP N17FF T116F 2.4564 2.45077 3.6667 2.6067																			
Q1150 S114 RESOF SEAV Q171 RESIL ABOV T800 S110 TESEP NTEP T1168 7. S5277 S2607 S2					Q182I	K252L	A80V	S193R	S121P	T266P	N275P	T116R		-23.9073	-23.4427	-23.8917	-24.411	-23.6563	-24.1348
Q1100 S114 REOF SEAV Q121 Q121 A507 T300 S112 T569 N1759 T1168 7. 25578 244313 52480 261073 25.669 25.6002 Q1100 S114 REOF SEAV Q171 N551 Q825 S1389 S026 A00 N1759 T1168 7. 25.6382 05.7170 C5.6582 0								T39D	S121P	T266P	N275P	T116R		-26.4391	-26.6606	-26.0517 -25.3627	-26.4924	-26.6272	
Q1100 S114 P360F S84W Q171 K151 Q180 S198 S02P A00P A75F T168 7.5 -56.552 .56.572 .56.575 .56.555 .56.552 .56.575 .5													?						
Q1190 2114 P360F SAW Q171 (1261 A80V 1398 S027 A09 N75F T1168													2						
Q1190 S114F P60F S54W Q12TL KSSLI ABOV S198R S0F AAGP N.75F T116R -24179 -24385 -246982 -24394 -24888 -20373 -55968 Q1190 S114F P60F S54W Q12TL KSSLI ABOV T900 S0F AAGP N.75F T116R -242784 -24384 -24390 -24386 -23766 -25566 Q1190 S114F P60F S54W Q12TL KSSLI ABOV T900 S0F AAGP N.75F T116R -242714 -25189 -24300 -25666 -23300 -25566 Q1190 S114F P60F S54W Q12TL KSSLI ABOV T900 S0F AAGP N.75F T116R -232785 -24867 -23980 -226042 -23496 -23678 Q1190 S114F P60F S54W Q12TL KSSLI ABOV T900 S0F AAGP N.75F T116R -232785 -24867 -23890 -226042 -23496 -23678 Q1190 S114F P60F S54W Q12TL KSSLI ABOV S198R S0F T50F N.75F T116R -25609 -24855 -258915 -266327 -26576 -26182 -268842 Q1190 S114F P60F S54W Q12TL KSSLI ABOV S198R S0F T50F N.75F T116R -26099 -24855 -25950 -72459 -7256 -26182 -268842 Q1190 S114F P60F S54W Q12TL KSSLI ABOV S198R S0F T50F N.75F T116R -261810 -26591 -72478 -26182 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382 -265842 -24382													- 2						
Q1190 S114F PEOPS S44W Q171, KSSL Q181 7900 S0P AGP N758 71188															-24.2609	-25.1667	-25.987	-25.8427	-25.0617
Q1190 S114F Resoft S8AW Q1171 Q181 ABOV T900 S9CP AGOP N.759 F1146F															-24.7702	-24.6082	-24.6838	-24.0973	-25.2966
Q1190 S214F REGOF S44W Q17T K52L A80V T39D S9CP A60P N278P T116F -2.82765 -2.88967 -2.28890 -2.2642 -2.34862 -2.30732 Q1190 S214F REGOF S44W Q17T K52L A80V S198F S9CP T56F N278P T116F -2.60369 -2.4206 -2.88915 -2.6337 -2.86768 -2.6919 Q1190 S214F REGOF S44W Q17T K52L A80V S198F S9CP T56F N278P T116F -2.60369 -2.4206 -2.88915 -2.6337 -2.86768 -2.6919 Q1190 S214F REGOF S44W Q17T K52L A80V S198F S9CP T56F N278P T116F -2.48115 -2.5160 -2.58944 -2.4506 -2.48915 -2.5769 -2.48915 -2.5769 -2.48915 -2.5769 Q1190 S214F REGOF S44W Q17T K52L A80V T39O S9CP T56F N278P T116F -2.48915 -2.5769 -2.4011 -2.49604 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.44010 -2.49009 -2.44865 -2.49009 -2.49009 -2.44865 -2.49009 -2.44865 -2.49009 -2.49009 -2.44865 -2.49009 -2.49009 -2.44865 -2.49009 -2.49009 -2.49009 -2.49009 -2.49009 -2.49009 -2.49009 -2.49009														-24.2381	-24.2381	-23.3706	-24.4206	-23.7969	-25.2641
Olifo Silf REGOF SAW QLTI KISL QNES SADS SODE TEGO NITSP TILGE V - 26.587 -28.814 274.14 26.337 25.688 26.198 26.198 26.842 20.118 20.11															-24.3057	-24.6564	-24.8032	-24.3103	-28.0674
Q1100 S114F R600F SSAW Q127 C1521 A80V S198 S92F T266P N375F T116R	Q119D	S214F	R260F	S54W	Q127L	K252L	Q182I	S193R	S92P	T266P	N275P	T116R	√		-25.8514	-27.4214	-26.3436	-26.1928	-26.8842
Q1190 S114F R250F SSAW Q17L K25L A80V S198 S9CF T256P NATSP T116R -248115 -25.2494 -725070 -25.0986 -23.7575 Q1190 S114F R250F SSAW Q17L K25L A80V T39O S9CF T256P NATSP T116R -248988 -25.2475 -24.0014 -24.0004 -24.7596 -35.921 -37.7596 Q1190 S114F R250F SSAW Q17L (A12L) A80V T39O S9CF T256P NATSP T116R -24.0988 -25.2475 -24.0014 -24.0004 -24.0906 -25.0215 -24.0014 -24.0004 -24.0906 -24.0906 -25.014 -24.0004 -24.0906													<i>!</i> √						
Q119D S214F R260F S54W Q12TL (K251L ABOV T39D S92F T266F) N275F T116F				S54W	Q182I	K252L	A80V	S193R	S92P	T266P	N275P	T116R							
O1190 S214F R260F SSAW Q127L Q182L A80V T39D S92F T266F N275F T116F C.													٧						
Q1190 S214F R256F S4W Q17L K252L Q182L S198F T113F S121P A40P T116F V - 2.6 207 -2.5 84 -2.5 744 -5.6 207 -7.5 4477 -7.5 2447 -7.5 2549 -7.5 544 -7.5 2549 -7.5 544 -	Q119D	S214F	R260F	S54W	Q127L	Q182I	A80V	T39D	S92P	T266P	N275P	T116R		-24.9098	-24.4856	-23.4400	-26.5502	-24.9290	-25.1440
Q1190 S214F R250F S54W Q127L K252L Q182L S198F T113F S121P A40P T116F C5516 C55147 C52140 C56065 C5655F C565F C5	4												√						
Q1190 R260F N275F S4W Q127L K25L A80V S198F T113F S121P A40P T116F -252472 -253036 -253267 -25326 -251252 -25594 C119D S124F R260F S42W Q127L K25L A80V S198F T113F S121P A40P T116F -24274 -247011 -24333 -243435 -243482 -243482 C119D S124F R260F S42W Q127L K25L A80V S198F T113F S121P A40P T116F -25516 -2528F -260487 -253356 -252121 -257658 C119D S124F R260F S42W Q127L K25L A80V S198F T113F S121P A40P T116F -25516 -2528F -260487 -253356 -252112 -257658 C119D S124F R260F S42W Q127L Q182L A80V S198F T113F S121P A40P T116F -248892 -243481 -243422 -25769 -25366 -252765 -252	Q119D	S214F	R260F	S54W	Q127L	K252L	Q182I	S193R	T113P	S121P	A40P	T116R		-25.364	-24.5744	-26.0237	-25.4477	-25.2449	-25.5291
Q1190 S214F R260F S24W Q17L K252L A80V S198F T113F S212F A40P T116F -2.4274 -2.42701 -2.4233 -2.5386 -2.5125 -2.51382 -2.52688 Q119D S214F R276F S54W Q127L K252L A80V S198F T113F S212F A40P T116F -2.5166 -2.528F -2.6487 -2.5383 -2.52112 -2.57688 Q119D S214F R260F R276F S24W Q127L K252L A80V S198F T113F S212F A40P T116F -2.5144 -2.5365 -2.528F -2.5285 -2.52																			
Q1190 S214F N275F S54W Q127L K252L A80V S198F T118F S121P A40P T116F -25.516 -25.288 -26.0487 -25.3335 -25.2112 -25.7588 Q119D S214F R260F N275F S54W Q127L Q182L A80V S198F T118F S121P A40P T116F -24.7569 -24.8569 -24.7569 -25.3755 -24.8484 -24.9342 -24.7569 -25.3755 -24.8484 -24.9342 -24.7569 -25.3755 -24.8484 -24.9342 -24.7569 -25.3855 -24.712 -25.0765 -24.7233 -24.7233 -24.7269 -25.3855 -24.712 -25.0765 -24.7233 -24.7269 -25.3855 -24.712 -25.0765 -24.7233 -24.7269 -25.3855 -24.8484 -24.7233 -24.7269 -25.3855 -25.2858 -26.5658 -25.0785 -25.0858 -26.0858 -26.5688 -25.0835 -25.0858 -26.085	Q119D	S214F	R260F	N275F	Q127L	K252L	A80V	S193R	T113P	S121P	A40P	T116R		-25.2472	-25.0247	-25.3356	-25.1225	-25.1382	-25.5944
Q119D R260F N275F S44W Q127L C25L A80V S198R T113P S121P A40P T116R -2.5141 2.50.045 -2.56.496 -2.5.755 -2.4.848 -2.4.9342 Q119D S114F R260F N275F Q127L Q1821 A80V S198R T113P S121P A40P T116R -2.47269 -2.38314 -2.43422 -2.5.6833 -2.50.076 -2.47233 Q119D S114F R275F S44W Q127L Q1821 A80V S198R T113P S121P A40P T116R -2.58744 -2.4.335 -2.50.156 -2.5.6833 -2.50.076 -2.4.7233 Q119D S114F R260F N275F S44W Q127L Q1821 A80V S198R T113P S121P A40P T116R -2.58744 -2.4.335 -2.4.0156 -2.5.0356 -2.5.0353 -2.2.433 -2.2.4335 -2.4.0156 -2.5.0356																			
Q1190 S214F R756F S54W Q127L Q1821 A80V S198F T113F S121P A40P T116F -247569 -258744 -254335 -250566 -256233 -25795 Q1190 R216F R757F S54W Q127L Q1821 A80V S198F T113F S121P A40P T116F -24622 -244335 -24798 -24247 -242432 -258393 Q1190 S214F R756F S54W Q127L Q1821 A80V S198F T113F S121P A40P T116F -24672 -239314 -243675 -23675 -24071 -23951 Q1190 S214F R756F S54W K252L Q1821 A80V S198F T113F S121P A40P T116F -24873 -24853 -250856 -250515 -242571 Q1190 R256F R75F S54W K252L Q1821 A80V S198F T113F S121P A40P T116F -248573 -246534 -24739 -256651 -24954 -255574 Q1190 R256F R75F S54W K252L Q1821 A80V S198F T113F S121P A40P T116F -248573 -246562 -24676 -24739 -250651 -24954 -24564 -24567 -24566 -24739 -250651 -24954 -24564 -24567	Q119D	R260F	N275F	S54W	Q127L	K252L	A80V	S193R	T113P	S121P	A40P	T116R		-25.1441	-25.0345	-25.6495	-25.2765	-24.8484	-24.9342
Q1190 S214F R275F S54W Q127L Q182L A80V S198F T113F S121P A40P T116F -2462 -24385 -250156 -258083 -262433 -258905 Q1190 S214F R260F R275F S54W C25L Q182L A80V S198F T113F S121P A40P T116F -24672 -23931 -250356 -250568 -25051 -24237 -259905 Q1190 S214F R275F S54W K25L Q182L A80V S198F T113F S121P A40P T116F -24672 -23893 -250356 -250518 -24237 -24237 -22937 Q1190 S214F R275F S54W K25L Q182L A80V S198F T113F S121P A40P T116F -24857 -246533 -22333 -250356 -250518 -24237 -23951 -24237 -24237																			
Q119D S214F R260F N37FF K2SL Q1821 A80V S193R T113P S121P A40P T116R	Q119D	S214F	N275F	S54W	Q127L	Q182I	A80V	S193R	T113P	S121P	A40P	T116R		-25.8744	-25.4335	-25.0156	-25.8083	-26.2433	-26.7993
Q1190 S114F R75FF S44W K25L Q182 A80V S198F T113F S121P A40P T116F -2.88F 24.8673 2-23.035 -23.6775 -24.0971 -2.9951 Q119D R260F N275F S54W K25L Q182 A80V S198F T113F S121P A40P T116F -2.4873 2-26553 -24.7893 -2.65631 -2.2735 -2.65631 -2.2735 -2.2565 -2.256531 -2.2735 -2.25653 -2.256531 -2.2735 -2.25653 -2.256531 -2.2735 -2.25653	Q119D	R260F S214F	N275F R260F	S54W N275F															
Q1190 R260F N275F S4W K25L Q182 A80V S198 T113P S121P A40P T116R -2.45462 -2.48768 -2.47613 -2.49656 -2.38836 -2.42321 Q1190 S214F R260F S45W Q127 K25L Q182 T390 T113P S121P A40P T116R -2.5832 -2.56808 -2.56007 -2.6139 -2.5242 -2.27608 Q1190 S214F N275F S45W Q127 K25L Q182 T390 T113P S121P A40P T116R -2.5832 -2.56808 -2.56108 -2.54766 -2.5476 -2.58066 -2.5476 -2.58066 -2.5476 -2.58066 -2.5476 -2.58066 -2.5476 -2.58066 -2.5476 -2.58066 -2.5476 -2.58066 -2.5476 -2.58066 -2.5476 -2.58066 -2.5476 -2.58066 -2.5476 -2.58066 -2.5476 -2.58066 -2.5676 -2.56068 -2.56108 -2.56118 -2.56108	Q119D	S214F	R260F	S54W	K252L	Q182I	A80V	S193R	T113P	S121P	A40P	T116R		-23.89	-24.6313	-23.1035	-23.6775	-24.0971	-23.9351
Q1190 S214F R260F N275F G127L K252L G182L T390 T113P S212P A40P T116R -58.828 -25.8002 -26.6907 -26.013P -24.7808 Q119D S214F R260F S44W Q127L K252L Q182L T39D T113P S212P A40P T116R -25.8035 -24.7908 -24.6908 -25.6108																			
Q119D S214F R260F N275F S54W Q127L K252L A80V T39D T113P S121P A40P T116R	Q119D	S214F	R260F	N275F	Q127L	K252L	Q182I	T39D	T113P	S121P	A40P	T116R		-25.8285	-25.5602	-26.6907	-26.0139	-26.1324	-24.7808
Q119D R260F N275F S44W Q127L K252L A80V T39D T113P S121P A40P T116R -245326 -24524 -299668 -240988 -2511 -249912 -249912 -249																			
Q1190 S214F R260F S54W Q127L K252L A80V T39D T113P S121P A40P T116R -23.9246 -24.6823 -23.2113 -24.656 -23.7586 -23.8585 Q119D R260F R275F S54W Q127L K252L A80V T39D T113P S121P A40P T116R -24.777 -24.9706 -25.2908 -24.138 -24.138 -24.6822 -23.813 -24.6828 -24.138 -25.8873 Q119D S214F R260F S54W Q127L Q182l A80V T39D T113P S121P A40P T116R -24.0767 -24.0755 -24.6922 -23.7817 -24.955 -24.892 -23.7817 -24.955 -24.992 -23.7817 -24.955 -24.992 -23.7817 -24.955 -24.992 -23.7817 -24.955 -24.992 -23.7817 -24.955 -24.992 -23.7817 -24.955 -24.992 -23.7817 -24.955 -24.992 -23.7817 -24.955 -24.992 -23.7817 -24.955 -24.992 -23.891 -24.955 -24.992 -23.993 -24.492 -24.955 -24.9	Q119D	R260F	N275F	S54W	Q127L	K252L	Q182I	T39D	T113P	S121P	A40P	T116R		-24.5336	-24.524	-23.9668	-24.0998	-25.11	-24.9912
Q119D S214F N275F S54W Q127L K252L A80V T39D T113P S121P A40P T116R																			
Q119D R360F N275F S4W Q127L K352L A80V T39D T113P S121P A40P T116R	Q119D	S214F	N275F	S54W	Q127L	K252L	A80V	T39D	T113P	S121P	A40P	T116R		-25.0527	-24.7906	-25.2908	-24.138	-25.157	-25.8873
Q1190 S214F R250F S44W Q127L Q1821 A80V T39D T113P S121P A40P T116R								T39D	T113P	S121P	A40P	T116R							
Q119D S214F N275F S54W Q127L Q1821 A80V T39D T113P S121P A40P T116R		S214F	R260F	\$54W	Q127L	Q182I	A80V	T39D	T113P	S121P	A40P	T116R		-23.5257	-24.0927	-23.5834	-24.5466	-22.8195	
Q119D S214F R260F N276F K252L Q1821 A80V T39D T113P S121P A40P T116R -23565 -249403 -233369 -22982 -225406 -240251 Q19D S214F N276F S54W K252L Q1821 A80V T39D T113P S121P A40P T116R -334994 -230434 -234996 -225854 -245346 -225838 Q119D S214F N276F S54W K252L Q1821 A80V T39D T113P S121P A40P T116R -24237 -254361 -241543 -230294 -246644 -238342 Q119D R260F N276F S54W K252L Q1821 A80V T39D T113P S121P A40P T116R -236389 -24882 -241197 -227294 -236733 -233913		S214F	N275F	\$54W	Q127L	Q182I	A80V	T39D	T113P	S121P	A40P	T116R				-23.34	-24.1061	-25.1977	
Q119D S214F R260F S54W K252L Q1821 A80V T39D T113P S121P A40P T116R -234994 -230434 -234996 -225854 -245345 -225338 Q119D S214F N275F S54W K252L Q1821 A80V T39D T113P S121P A40P T116R -242237 -254361 -241543 -230294 -246644 -238342 Q119D R260F N275F S54W K252L Q1821 A80V T39D T113P S121P A40P T116R -236399 -24882 -241197 -227294 -230733 -233913	Q119D	REDUT	19275F	354W	Q127L	Q182I	ABOV	T39D	T113P	S121P	A40P	T116R							
Q119D R260F N275F S54W K252L Q182L A80V T39D T113P S121P A40P T116R -23.6389 -24.882 -24.1197 -22.7284 -23.0733 -23.3913	Q119D Q119D Q119D	S214F	R260F	N275F	K252L														
	Q119D Q119D Q119D Q119D	S214F S214F	R260F	\$54W	K252L	Q182I	A80V												
Q119D 3214F R260F N276F Q127L K252L Q182L 3193R T113P 592P A40P T116R V -264044 -260312 -278646 -275494 -255364 -250406	Q119D Q119D Q119D Q119D Q119D	S214F S214F S214F	R260F N275F	\$54W \$54W	K252L K252L	Q182I Q182I	A80V A80V	T39D	T113P	S121P	A40P	T116R		-24.2237 -23.6389	-25.4361 -24.882	-24.1543 -24.1197	-23.0294 -22.7284	-24.6644 -23.0733	-23.8342 -23.3913

				Q127L				T113P		A40P	T116R	√					-26.5387	
		N275F N275F		Q127L Q127L				T113P T113P		A40P A40P	T116R T116R	√ -/	-27.0365				-27.3825 -26.7902	
		R260F		Q127L				T113P		A40P	T116R	٧					-26.7902	
		R260F		Q127L				T113P		A40P	T116R						-24.9273	
				Q127L				T113P		A40P	T116R	√	-26.4329				-26.2252	
		N275F R260F		Q127L Q127L				T113P T113P		A40P A40P	T116R T116R						-25.4741 -24.6559	
		R260F		Q127L				T113P		A40P	T116R						-24.7357	
Q119D	S214F	N275F	S54W	Q127L	Q182I	A80V	S193R	T113P	S92P	A40P	T116R	√	-26.4164			-26.221	-25.9292	-27.0121
		N275F		Q127L				T113P		A40P	T116R		-25.6833	-26.491	-25.7968		-25.305	
Q119D Q119D		R260F	N275F S54W	K252L K252L				T113P T113P		A40P A40P	T116R		-25.0658 -24.7502		-25.9842		-24.7828 -24.9116	
	S214F		S54W	K252L				T113P		A40P	T116R	√					-24.422	
	R260F		S54W	K252L	Q182I	A80V		T113P		A40P	T116R		-25.0255	-26.3087	-25.1205	-23.4394	-25.2455	-25.0133
	S214F S214F		N275F S54W	Q127L Q127L				T113P T113P		A40P A40P	T116R T116R		-25.9596 -25.7717				-26.5527	
	S214F		S54W	Q127L				T113P		A40P	T116R	√	-26.6322		-26.2936		-25.8805 -26.113	-24.7794
Q119D				Q127L				T113P		A40P	T116R			-26.0107		-26.1849	-24.6816	-25.4431
Q119D		R260F		Q127L				T113P		A40P	T116R			-24.2933			-24.5625	
Q119D 0119D		R260F N275F	S54W	Q127L Q127L				T113P T113P		A40P A40P	T116R T116R		-24.5926 -25.3628	-24.8676 -26.2475			-24.1479 -25.4323	
Q119D		N275F	S54W	Q127L			T39D	T113P		A40P	T116R		-24.9551				-24.5689	
Q119D			N275F	Q127L			T39D	T113P		A40P	T116R		-24.5397				-24.4924	
			S54W	Q127L				T113P		A40P	T116R		-24.5454				-23.9867	
		N275F N275F		Q127L Q127L				T113P T113P		A40P A40P	T116R T116R		-25.3151 -25.2347				-25.533 -25.0607	
		R260F		K252L				T113P		A40P	T116R		-23.6563	-24.4074	-24.5538	-24.1579	-23.1519	-22.0107
		R260F		K252L				T113P		A40P	T116R		-23.8509				-23.4553	
				K252L K252L				T113P T113P		A40P A40P	T116R T116R	٧	-24.8101 -24.4959				-23.9385 -23.7767	
Q119D	S214F	R260F	N275F	Q127L	K252L	Q182I	S193R	S121P	S92P	A40P	T116R		-24.8006	-24.7469	-25.0723	-24.7919	-24.7575	-24.6345
Q119D	S214F	R260F	S54W	Q127L	K252L	Q182I	S193R	S121P	S92P	A40P	T116R		-23.5794				-23.3274	
				Q127L Q127L				S121P S121P		A40P A40P	T116R T116R		-25.4319 -24.8819				-25.6372 -24.8341	
				Q127L				S121P		A40P	T116R						-24.8341	
				Q127L				S121P		A40P	T116R						-24.0801	
				Q127L				S121P		A40P A40P	T116R T116R						-25.3119 -24.4052	
				Q127L Q127L				S121P S121P		A40P	T116R						-25.6442	
				Q127L				S121P		A40P	T116R						-24.0857	
				Q127L				S121P			T116R						-23.9333	
				Q127L K252L				S121P S121P			T116R T116R						-25.0156 -23.2624	
				K252L				S121P			T116R		-23.0009					
				K252L				S121P			T116R		-24.0588				-23.69	
				K252L Q127L				S121P S121P			T116R T116R						-22.6828 -24.5469	
				Q127L				S121P		A40P	T116R		-24.1442	-23.5436	-24.0621	-24.1763	-23.9695	-24.9694
				Q127L				S121P			T116R		-25.1396				-25.2916	
		N275F R260F		Q127L Q127L				S121P S121P			T116R	-				-23.8616 -24.8305	-24.4967 -25.555	
				Q127L				S121P			T116R						-23.0301	
O110D			CE AWAY		MODEL	AROV	TOOD	S121P	S92P	A40P	T116R		22.0024	-241727	220220	00.0000	0.1.7007	-24.5021
	S214F												-23.9924	-24.2101	-22.9230	-23.6229		
Q119D	R260F	N275F	S54W	Q127L	K252L	A80V	T39D	S121P	S92P	A40P	T116R		-23.4831	-23.1391	-23.5009	-24.7553	-23.2034	-22.8168
Q119D	R260F S214F		S54W N275F	Q127L Q127L	K252L Q182I	A80V A80V	T39D T39D		S92P S92P	A40P A40P A40P	T116R T116R T116R		-23.4831 -23.8026	-23.1391 -23.3242 -21.9929	-23.5009 -23.9325	-24.7553 -23.2842	-23.2034 -25.2608 -23.5818	-22.8168 -23.2111 -22.4654
Q119D Q119D Q119D Q119D	R260F S214F S214F S214F	N275F R260F R260F N275F	S54W N275F S54W S54W	Q127L Q127L Q127L Q127L Q127L	K252L Q182I Q182I Q182I	A80V A80V A80V A80V	T39D T39D T39D T39D	S121P S121P S121P S121P	S92P S92P S92P S92P	A40P A40P A40P	T116R T116R T116R		-23.4831 -23.8026 -22.7046 -23.8619	-23.1391 -23.3242 -21.9929 -23.278	-23.5009 -23.9325 -23.4208 -24.1086	-24.7553 -23.2842 -22.062 -23.2927	-23.2034 -25.2608 -23.5818 -24.8813	-22.8168 -23.2111 -22.4654 -23.749
Q119D Q119D Q119D Q119D Q119D	R260F S214F S214F S214F R260F	N275F R260F R260F N275F N275F	\$54W N275F \$54W \$54W \$54W	Q127L Q127L Q127L Q127L Q127L Q127L	K252L Q182I Q182I Q182I Q182I	A80V A80V A80V A80V A80V	T39D T39D T39D T39D T39D	S121P S121P S121P S121P S121P	S92P S92P S92P S92P S92P	A40P A40P A40P A40P	T116R T116R T116R T116R		-23.4831 -23.8026 -22.7046 -23.8619 -23.0732	-23.1391 -23.3242 -21.9929 -23.278 -22.2145	-23.5009 -23.9325 -23.4208 -24.1086 -23.775	-24.7553 -23.2842 -22.062 -23.2927 -22.9667	-23.2034 -25.2608 -23.5818 -24.8813 -23.6304	-22.8168 -23.2111 -22.4654 -23.749 -22.7795
Q119D Q119D Q119D Q119D	R260F S214F S214F S214F R260F S214F	N275F R260F R260F N275F	S54W N275F S54W S54W S54W N275F	Q127L Q127L Q127L Q127L Q127L Q127L K252L	K252L Q182I Q182I Q182I Q182I	A80V A80V A80V A80V A80V A80V	T39D T39D T39D T39D T39D T39D	S121P S121P S121P S121P	S92P S92P S92P S92P S92P S92P	A40P A40P A40P	T116R T116R T116R		-23.4831 -23.8026 -22.7046 -23.8619	-23.1391 -23.3242 -21.9929 -23.278 -22.2145 -23.8512 -23.0878	-23.5009 -23.9325 -23.4208 -24.1086 -23.775 -22.6108	-24.7553 -23.2842 -22.062 -23.2927	-23.2034 -25.2608 -23.5818 -24.8813 -23.6304 -22.2066	-22.8168 -23.2111 -22.4654 -23.749 -22.7795 -22.3098
Q119D Q119D Q119D Q119D Q119D Q119D Q119D Q119D Q119D	R260F S214F S214F S214F R260F S214F S214F S214F	N275F R260F R260F N275F N275F R260F R260F N275F	S54W N275F S54W S54W S54W N275F S54W S54W	Q127L Q127L Q127L Q127L Q127L K252L K252L K252L	K252L Q182l Q182l Q182l Q182l Q182l Q182l Q182l Q182l	A80V A80V A80V A80V A80V A80V A80V A80V	T39D T39D T39D T39D T39D T39D T39D T39D	S121P S121P S121P S121P S121P S121P S121P S121P S121P	S92P S92P S92P S92P S92P S92P S92P S92P	A40P A40P A40P A40P A40P A40P A40P	T116R T116R T116R T116R T116R T116R T116R		-23.4831 -23.8026 -22.7046 -23.8619 -23.0732 -22.8683 -22.4603 -23.2643	-23.1391 -23.3242 -21.9929 -23.278 -22.2145 -23.8512 -23.0878 -25.25	-23.5009 -23.9325 -23.4208 -24.1086 -23.775 -22.6108 -22.7522 -22.7396	-24.7553 -23.2842 -22.062 -23.2927 -22.9667 -23.3631 -21.4746 -21.5099	-23.2034 -25.2608 -23.5818 -24.8813 -23.6304 -22.2066 -22.1123 -22.9735	-22.8168 -23.2111 -22.4654 -23.749 -22.7795 -22.3098 -22.8746 -23.8487
Q119D Q119D Q119D Q119D Q119D Q119D Q119D Q119D Q119D Q119D	R260F S214F S214F S214F R260F S214F S214F S214F R260F	N275F R260F R260F N275F N275F R260F R260F N275F N275F	S54W N275F S54W S54W S54W N275F S54W S54W S54W	Q127L Q127L Q127L Q127L Q127L K252L K252L K252L K252L	K252L Q182l Q182l Q182l Q182l Q182l Q182l Q182l Q182l Q182l	A80V A80V A80V A80V A80V A80V A80V A80V	T39D T39D T39D T39D T39D T39D T39D T39D	S121P S121P S121P S121P S121P S121P S121P S121P S121P S121P	S92P S92P S92P S92P S92P S92P S92P S92P	A40P A40P A40P A40P A40P A40P A40P A40P	T116R T116R T116R T116R T116R T116R T116R T116R		-23.4831 -23.8026 -22.7046 -23.8619 -23.0732 -22.8683 -22.4603 -23.2643 -22.8598	-23.1391 -23.3242 -21.9929 -23.278 -22.2145 -23.8512 -23.0878 -25.25 -23.6236	-23.5009 -23.9325 -23.4208 -24.1086 -23.775 -22.6108 -22.7522 -22.7396 -23.2659	-24.7553 -23.2842 -22.062 -23.2927 -22.9667 -23.3631 -21.4746 -21.5099 -21.8455	-23.2034 -25.2608 -23.5818 -24.8813 -23.6304 -22.2066 -22.1123 -22.9735 -22.8388	-22.8168 -23.2111 -22.4654 -23.749 -22.7795 -22.3098 -22.8746 -23.8487 -22.7251
Q119D Q119D Q119D Q119D Q119D Q119D Q119D Q119D Q119D Q119D Q119D	R260F S214F S214F S214F R260F S214F S214F S214F R260F S214F	N275F R260F R260F N275F N275F R260F R260F N275F N275F R260F	S54W N275F S54W S54W S54W N275F S54W S54W S54W N275F	Q127L Q127L Q127L Q127L Q127L K252L K252L K252L K252L Q127L	K252L Q182I Q182I Q182I Q182I Q182I Q182I Q182I Q182I K252L	A80V A80V A80V A80V A80V A80V A80V A80V	T39D T39D T39D T39D T39D T39D T39D T39D	S121P S121P S121P S121P S121P S121P S121P S121P S121P S121P T113P	\$92P \$92P \$92P \$92P \$92P \$92P \$92P \$92P	A40P A40P A40P A40P A40P A40P A40P A40P	T116R T116R T116R T116R T116R T116R T116R T116R T116R	√ √	-23.4831 -23.8026 -22.7046 -23.8619 -23.0732 -22.8683 -22.4603 -23.2643 -22.8598	-23.1391 -23.3242 -21.9929 -23.278 -22.2145 -23.0878 -25.25 -23.6236 -26.6761	-23.5009 -23.9325 -23.4208 -24.1086 -23.775 -22.6108 -22.7522 -22.7396 -23.2659 -27.162	-24.7553 -23.2842 -22.062 -23.2927 -22.9667 -23.3631 -21.4746 -21.5099 -21.8455 -27.2489	-23.2034 -25.2608 -23.5818 -24.8813 -23.6304 -22.2066 -22.1123 -22.9735 -22.8388 -26.0644	-22.8168 -23.2111 -22.4654 -23.749 -22.7795 -22.3098 -22.8746 -23.8487 -22.7251 -25.8844
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Q1190 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q1	R260F S214F S214F R260F S214F	N275F R260F N275F R260F R260F R260F R260F R260F R260F R260F R260F R275F R260F R275F R260F R275F R260F R275F R260F R260F R250F R260F R250F	\$54W N.275F \$54W S.54W \$54W N.275F \$54W S.54W \$54W \$54W \$554W \$555W \$554W \$555W \$555	Q1271 Q1271 Q1271 Q1271 Q1271 K2521 K2521 K2521 Q1271 Q1271 Q1271 Q1271 Q1271 Q1271 Q1271 Q1271 K2521 K2521 K2521 K2521 Q1271	K252L Q182l Q182l Q182l Q182l Q182l Q182l Q182l Q182l K252L K252L K252L K252L Q182l Q182l Q182l Q182l Q182l Q182l Q182l Q182l Q182l Q182l X252L K252L X252L	A80V A80V A80V A80V A80V A80V A80V A80V	T39D T39D T39D T39D T39D T39D T39D T39D	S1210 S1210 S1210 S1210 S1210 S1210 S1210 S1210 S1210 S1210 T1130	\$92P \$92P \$92P \$92P \$92P \$92P \$92P \$92P	A40P A40P A40P A40P A40P A40P A40P A40P	T11GR T11GR	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	-23.4831 -23.8026 -22.7046 -23.8619 -23.0732 -22.8683 -22.4603 -23.2643 -22.8598 -26.6071 -25.1313 -26.207 -25.2313 -26.7093 -25.72313 -25.0776 -25.9712 -25.72313 -25.0776 -25.72313 -25.0776 -25.72313 -25.0776 -25.7231	-23.1391 -23.3242 -21.9929 -23.278 -22.2145 -23.88512 -23.0878 -25.25 -23.6236 -26.6761 -25.4302 -26.2717 -26.4579 -26.2717 -26.4579 -25.9753 -25.8055 -25.8055 -25.8055 -25.8055 -25.8055 -25.6866 -25.4784 -26.8086 -26.4127 -26.8086	-23,5009 -23,4208 -24,4086 -23,4208 -24,1086 -23,775 -22,6108 -22,7522 -22,7396 -23,2659 -27,162 -25,5005 -27,814 -27,1661 -26,1527 -24,6045 -26,181 -26,8217 -26,6734 -25,8237 -26,6734 -25,8237 -26,6734 -25,8237 -26,6734 -26,9832 -26,9736 -27,9834 -28,4484 -27,2371 -26,3929	-24.7553 -23.2842 -22.062 -23.2927 -22.9667 -23.3631 -21.4746 -21.5099 -27.4299 -27.429 -27.429 -27.429 -27.429 -23.56866 -26.1961 -26.673 -24.429 -24	-23.2034 -23.5818 -24.8813 -24.8813 -22.966 -22.1123 -22.9735 -22.8388 -26.0644 -25.2214 -26.5676 -26.4978 -25.7309 -25.7309 -25.3264 -24.436 -24.436 -24.436 -24.7229 -25.3838 -24.7229 -25.3838 -24.7229 -25.3838 -24.7229 -25.3838 -24.7229 -25.3838 -24.7229 -25.3838 -24.7229 -25.3838 -24.7229 -25.3838 -24.7229 -25.3838 -24.7229 -25.3838 -26.9146 -26.9167 -26.9167 -26.9167 -26.9167 -26.9167 -26.9167 -26.9167 -26.9167 -26.9167 -26.9167 -26.9167	-228.168 -23.2111 -22.4684 -23.2111 -22.4684 -23.749 -22.7795 -22.3098 -22.8746 -23.8487 -22.7251 -25.8884 -25.9627 -27.6596 -25.7869 -25.7869 -25.8876 -25.28387 -26.0775 -26.0775 -26.0775 -26.0775 -26.0775 -26.4077 -26.4077 -26.4077 -27.2377 -27.2377 -27.2377 -27.2377 -25.4885
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Q1190 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q1	R2507	N275F R260F N275F R260F N275F R260F N275F R260F	\$54W N275F	01271. 01271. 01271. 01271. 01271. 02271. 02271. 01	K2521	ABOV ABOV ABOV ABOV ABOV ABOV ABOV ABOV	T39D T39D T39D T39D T39D T39D T39D T39D	S111P S121P S121P S121P S121P S121P S121P S121P S121P T113P	992P 992P 992P 992P 992P 992P 992P 992P	A40P A40P A40P A40P A40P A40P A40P A40P	T116R T116R	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	-23.4831 -23.8026 -22.7046 -23.8619 -23.0732 -22.8683 -22.4603 -23.2643 -22.8598 -26.6071 -25.913 -26.207 -25.2313 -26.207 -25.2313 -26.207 -25.2313 -26.207 -25.2313 -26.207 -25.2313 -26.207 -25.2313 -26.207 -25.2313 -26.207 -25.2313 -26.7076 -25.7913 -26.4615 -26.4617 -26.7649 -25.8733 -27.4379 -26.4012 -26.485 -25.2625 -26.8863 -25.8563 -25.8563 -25.8563 -25.8563 -25.8569 -25.8569 -25.8569 -25.8569	-23.1391 -23.3242 -21.9929 -23.278 -22.2148 -23.8512 -23.8578 -25.25 -23.6276 -25.25 -25.4302 -26.4579 -25.0763 -25.6362 -25.4302 -25.8055	-23,5009 -23,925 -23,4208 -24,1086 -24,1086 -23,7552 -22,7396 -27,522 -27,326 -27,521 -27,521 -26,527 -24,6045 -26,128 -25,8417 -25,6539 -27,944 -26,128 -26,1	-24.7553 -23.2842 -22.062 -23.2927 -22.9667 -23.3631 -21.4746 -21.5099 -27.44931 -27.12 -22.45498 -27.4991 -27.1 -27.12 -25.4358 -26.8713 -25.9479 -23.56869 -24.429 -25.4488 -26.973 -24.8028 -25.8458 -27.219 -26.4103 -25.6159 -27.216 -27.	-23.2094 -25.2608 -24.8813 -23.8904 -22.2066 -22.1123 -22.9735 -22.9738 -26.0644 -25.2214 -26.5676 -26.4978 -27.2069 -26.0141 -25.3436 -24.436 -25.7803 -26.5578 -26.5578 -27.2069 -26.217 -26.59536 -27.207 -26.59536	-228.68 -232.111 -22.4684 -23.7795 -22.7795 -22.3098 -22.8746 -23.8487 -22.7759 -23.8287 -22.7559 -25.7869 -25.7869 -25.7869 -25.8841 -25.66.0519 -26.0570 -26.0519
Q1190 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q1	R2507	N275F R260F	SS4W SS4W SS4W SS4W SS4W SS4W SS4W SS4W	01271. 01	K2521 (1922) (19	ABOV ABOV ABOV ABOV ABOV ABOV ABOV ABOV	T39D T39D T39D T39D T39D T39D T39D T39D	S121P S1	992P 992P 992P 992P 992P 992P 992P 992P	A40P A40P A40P A40P A40P A40P A40P A40P	T116R T116R	\$\frac{1}{\sqrt{1}}\$	-23.4831 -23.8026 -22.7046 -23.8026 -23.0732 -23.6833 -22.4603 -23.2463 -22.8598 -26.6071 -25.911 -25.911 -26.26523 -26.207 -25.2313 -26.7036 -26.25723 -25.7036 -26.25723 -26.207 -25.2013 -26.207 -25.2013 -26.207 -25.2013 -26.207 -26.2613 -26.207 -26.2613 -26.207 -26.2613 -26.207 -26.2613 -26.207 -26.2613 -26.2613 -26.2613 -26.2613 -26.2613 -26.2613 -26.2613 -26.2613 -26.2613 -26.2613 -26.2613 -26.2613 -26.2613 -26.2613	-23.1391 -23.3242 -21.9929 -23.278 -22.2145 -23.88512 -23.0878 -25.25 -26.6761 -25.4302 -26.7379 -25.0763 -26.4579 -25.0763 -26.4579 -25.0763 -26.4579 -25.0763 -26.4579 -25.0763 -26.4579 -25.0763 -26.4579 -25.0763 -26.4579 -25.0763 -26.4579 -25.0763 -26.4579 -25.0763 -26.4579 -25.0763 -26.4579 -25.0763 -26.516 -26.0763 -26.516 -26.0784 -26.8784 -26.8784 -26.8782 -26.926 -25.7784 -26.8784 -26.8782 -26.926 -26.4784 -26.8782 -26.926 -26.4784 -26.86811 -26.8081 -26.84416 -26.84416 -26.84416 -26.84416 -26.84416 -26.84416	-23,5009 -23,4208 -24,1086 -24,1086 -23,752 -23,2659 -27,162 -23,2659 -27,162 -25,2059 -27,162 -25,2059 -27,165 -26,108 -26,10	-24.7553 -23.2642 -22.062 -23.2927 -22.9667 -23.3631 -21.4746 -21.5099 -21.8455 -27.2489 -27.4403 -27.491 -27.1 -27.12 -25.4558 -26.8713 -25.9753 -25.9479 -23.4564 -26.073 -24.8022 -26.6038 -25.8458 -27.218 -26.61961 -25.6159 -24.7429 -23.4264 -26.073 -24.8022 -26.6038 -25.8458 -27.216	-23.2034 -25.2608 -23.5818 -24.8813 -23.6304 -22.2066 -22.1123 -22.9735 -22.8388 -26.0644 -25.2214 -26.5676 -26.4978 -25.7309 -25.74634 -25.7464 -25.74644 -25	-228.68 -227.4684 -227.49 -227.79 -227.97 -22.308 -22.7749 -22.8746 -22.8746 -22.8746 -22.8746 -22.8746 -22.8746 -23.8844 -25.9627 -25.7869 -25.7869 -25.7869 -25.7869 -25.7869 -25.7878 -26.0775 -26.0775 -26.0775 -26.0775 -26.0775 -26.0775 -26.0775 -26.0775 -26.0775 -26.0775 -26.0775 -26.0775 -26.0776 -26.3887 -26.0775 -26.0775 -26.0776 -26.3887 -26.0776 -26.3887 -26.0776 -26.3887 -26.0776 -26.38887 -26.0776 -26.38887 -26.0776 -26.38887 -26.0776 -26.38887 -26.0776 -26.38887 -26.0776 -26.388887 -26.0776 -26.0776 -26.0774 -26.38881 -26.0788882 -26.078882 -26.07882 -26.078882 -26.078882 -26.078882 -26.078882 -26.078882 -26.0788
Q1190 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q1	R2507	N275F R260F	SS4W SS4W SS4W SS4W SS4W SS4W SS4W SS4W	01271. 01271.	K2521 (1922) (19	ABOV ABOV ABOV ABOV ABOV ABOV ABOV ABOV	T39D T39D T39D T39D T39D T39D T39D T39D	S1210 S1210	992P 992P 992P 992P 992P 992P 992P 992P	A40P A40P A40P A40P A40P A40P A40P A40P	T116R T116R	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	-23.4831 -23.8026 -22.7046 -23.8026 -23.0732 -23.0732 -22.6863 -22.4603 -23.2643 -22.8598 -26.6071 -25.911 -25.911 -26.207 -25.213 -26.207 -25.213 -26.207 -25.213 -26.207 -25.213 -26.207 -25.213 -26.207 -25.213 -26.207 -26.213 -26.207 -26.213 -26.207 -26	-23.1391 -23.2322 -21.929 -23.278 -22.2145 -23.88512 -23.8978 -25.25 -25.66761 -25.4302 -26.6761 -25.4302 -26.6761 -25.4302 -26.6761 -25.4302 -26.5732 -25.056 -25.516 -25.056 -25.516 -25.25 -25.056 -25.25 -25.056 -25.25 -25.056 -25.25 -25.056 -25.25 -25.056 -25.25 -25.056 -25.25 -2	-23,5009 -23,4208 -24,1086 -24,1086 -24,1086 -23,775 -22,6108 -27,512 -22,7396 -27,162 -23,2659 -27,162 -23,2659 -27,165 -26,185 -26,1	-24.7553 -23.2642 -22.062 -23.2927 -22.9667 -23.3631 -21.4746 -21.5099 -27.4403 -27.499 -27.1 -2	-23.2034 -25.2608 -23.5818 -24.8813 -23.5304 -22.2066 -22.1123 -22.9735 -22.8288 -26.0644 -25.2214 -26.5676 -26.4978 -26.5871 -27.2069 -25.871 -27.2069 -25.871 -27.2069 -25.873 -26.1542 -25.7803 -26.1542 -25.7803 -26.1542 -27.807 -25.5833 -26.1542 -27.807 -25.5833 -26.1542 -27.807 -25.5833 -26.1542 -27.807 -25.5833 -26.1542 -27.807 -25.5833 -26.1542 -27.807 -25.5833 -27.2071 -25.5833 -27.2071 -25.5833 -27.2071 -25.5833 -27.2071 -25.5833 -27.2071 -25.5833 -27.2071 -25.5833 -27.3073 -26.5060 -26.608 -27.3073 -25.5808 -27.3073 -26.5008 -27.3073 -26.5008 -27.3073 -26.4038 -27.50608	-228.68 -23.749 -22.779 -22.779 -22.779 -22.3098 -22.779 -22.3098 -22.725 -23.8487 -22.725 -23.8487 -22.725 -25.8481 -25.962 -25.7869 -25.7869 -25.7869 -25.8487 -26.0775 -26.
Q1190 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q10 Q1	R2507	N275F R260F	SS4W SS4W SS4W SS4W SS4W SS4W SS4W SS4W	01271. 01271. 01271. 01271. 01271. 01272. 01	K2521 (1922) (19	ABOV ABOV ABOV ABOV ABOV ABOV ABOV ABOV	T39D T39D T39D T39D T39D T39D T39D T39D	S121p	992P 992P 992P 992P 992P 992P 992P 992P	A40P A40P A40P A40P A40P A40P A40P A40P	T116R T116R	? \\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	-23.4831 -23.8026 -22.7046 -23.8026 -23.0732 -23.0732 -22.6863 -22.4603 -23.2643 -22.8598 -26.6071 -25.911 -25.911 -26.207 -25.213 -26.207 -25.213 -26.207 -25.213 -26.207 -25.213 -26.207 -25.213 -26.207 -25.213 -26.207 -26.213 -26.207 -26.213 -26.207 -26	-23.1391 -23.2322 -21.929 -23.278 -22.2145 -23.88512 -23.8978 -25.25 -25.66761 -25.4302 -26.6761 -25.4302 -26.6761 -25.4302 -26.6761 -25.4302 -26.5732 -25.056 -25.516 -25.056 -25.516 -25.25 -25.056 -25.25 -25.056 -25.25 -25.056 -25.25 -25.056 -25.25 -25.056 -25.25 -25.056 -25.25 -2	-23,5009 -23,4208 -24,1086 -24,1086 -24,1086 -23,775 -22,6108 -27,512 -22,7396 -27,162 -23,2659 -27,162 -23,2659 -27,165 -26,185 -26,1	-24.7553 -23.2642 -22.062 -23.2927 -22.9667 -23.3631 -21.4746 -21.5099 -27.4403 -27.499 -27.1 -2	-23.2094 -23.5818 -24.8813 -23.6304 -22.2066 -22.1123 -22.9735 -22.9388 -26.0644 -25.2144 -25.2144 -25.2144 -25.2144 -25.21496 -26.0141 -25.3468 -25.7803 -26.5528 -25.1831 -24.7229 -25.3838 -25.1841 -27.2069 -25.73893 -26.1542 -27.1807 -25.5368 -25.1841 -27.2069 -25.73893 -26.1542 -27.1807 -25.5368 -25.1841 -27.2069 -25.36838 -25.1841 -27.2069 -25.2069 -25.36838 -25.2069 -25.36838 -25.2069 -25.36838 -25.2069 -25.36838 -25.2069 -25.36838 -25.207 -27.207 -26.5068 -25.6068 -25.6068 -25.6068 -25.6068 -25.607 -27.207 -26.3068 -25.6068 -27.3073 -26.4471	-228.68 -23.749 -22.779 -22.779 -22.779 -22.3098 -22.779 -22.3098 -22.725 -23.8487 -22.725 -23.8487 -22.725 -25.8481 -25.962 -25.7869 -25.7869 -25.7869 -25.8487 -26.0775 -26.

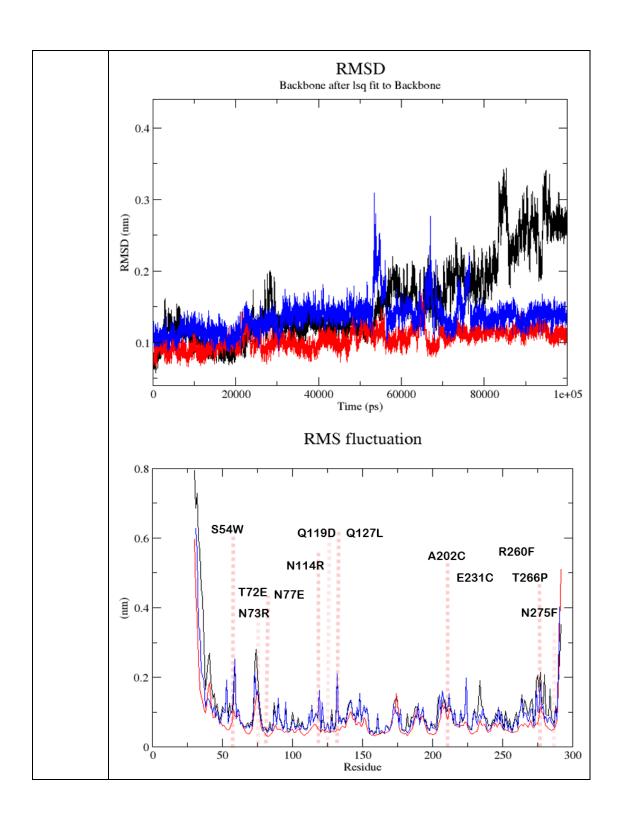
Q119	02145	R260F	CEANA	0127	V2E2I	A 9.0\/	C102D	T113P	coan	TOGER	Titen	√	26 5261	25.2042	-26.0959	27.1022	-26.7378	25 2201	
Q119		N275F		Q127L	K252L	A80V	S193R	T113P	S92P	T266P	T116R	√	-27.6349	-25.455	-27.0236	-27.8567	-28.0344	-27.8048	
Q119 O119		N275F R260F		Q127L Q127L			S193R S193R	T113P		T266P T266P		√ √		-25.9875 -27.0689	-26.6522	-27.6635 -24.008	-26.2854 -24.6431	-26.0625 -27.3851	
Q119		R260F		Q127L			S193R			T266P			-26.212				-25.9098		
		N275F N275F		Q127L Q127L				T113P T113P				√ √		-26.7695	-27.8263	-27.0365 -27.0311	-27.7146 -27.7627		
		R260F		K252L				T113P				٧	-26.0667	-26.7918	-26.0365	-26.9077	-25.1866	-25.411	
		R260F N275F		K252L				T113P T113P				√ √	-25.6582 -27.1627				-25.7667 -26.3827		
		N275F						T113P									-26.4529		
		R260F R260F						T113P T113P				√ √	-26.0163 -27.2276				-25.5241 -27.0757		
		N275F					T39D	T113P	S92P	T266P	T116R	√ √	-28.2017	-28.1346	-28.8366	-27.8974	-27.824	-28.3158	
		N275F R260F						T113P T113P				√ √					-28.4377 -27.537		
Q119	S214F	R260F	S54W	Q127L	K252L	ABOV	T39D	T113P	S92P	T266P	T116R	√	-26.947	-26.6706	-27.0743	-27.0599	-26.1121	-27.3683	
		N275F N275F						T113P T113P				√ √	-27.7115				-27.2934 -26.449		
		R260F					T39D	T113P	S92P	T266P	T116R	√	-26.841	-27.0025	-27.3042	-26.2091	-26.7956	-26.8938	
		R260F N275F						T113P T113P				√ √	-26.3006 -27.4201				-26.8209		
		N275F						T113P				√	-26.583	-26.2101	-27.0914	-25.9764	-26.3056	-27.3314	
		R260F R260F						T113P T113P									-26.9061 -26.2352		
		N275F						T113P				√					-26.7097		
		N275F						T113P				√ √					-27.1142		
		R260F										٧	-25.5226 -24.5884				-24.5887		
		N275F										√	-26.4653						
		N275F R260F					S193R	S121P	S92P	T266P	T116R						-24.7889 -25.9277		
		R260F						S121P					-24.5794	-24.1542	-24.2471	-24.5348	-24.4981	-25.463	
		N275F N275F						S121P S121P					-25.7824	-25.5299	-25.3354	-25.8394 -25.8337	-25.8245 -24.6159	-26.1826	
		R260F						S121P					-25.5451	-25.2107	-25.6375	-26.2406	-25.6403	-24.9967	
		R260F N275F						S121P S121P					-24.424	-24.4573	-25.047	-23.1201 -24.81	-24.3758 -25.7027	-25.6289	
Q119	D R260F	N275F	S54W	Q127L	Q182I	A80V	S193R	S121P	S92P	T266P	T116R		-24.6441	-24.7002	-24.8116	-23.2707	-25.6613	-24.7769	
		R260F R260F		K252L K252L				S121P S121P					-24.4719	-24.4809	-23.9905	-23.916	-23.3536 -23.0599	-24.4219 -24.3239	
Q1191		N275F		K252L	Q1821	A80V	S193R	S121P	S92P	T266P	T116R		-25.0731	-25.0077	-25.9878	-25.0109	-25.0358	-24.3234	
Q119		N275F R260F						S121P S121P				√		-24.9341	-24.7774		-24.4948 -25.1316		
Q1191 Q1191		R260F N275F		Q127L				S121P S121P				J		-24.4861			-24.8945 -27.4242		
		N275F						S121P				√	-25.5283	-24.6909	-25.7408	-24.5341	-25.1582	-27.5175	
Q119		R260F R260F		Q127L Q127L				S121P S121P							-26.4209 -24.8814		-26.5679 -24.7666		
		N275F		Q127L			T39D	S121P	S92P	T266P	T116R		-25.8883	-25.6718	-25.7066	-26.1058	-25.3903	-26.5672	
		N275F R260F		Q127L Q127L				S121P S121P									-24.9568 -26.9462		
Q119	D S214F	R260F	S54W	Q127L	Q182I	A80V	T39D	S121P	S92P	T266P	T116R		-24.9972	-24.8332	-24.9782	-25.0421	-25.3172	-24.8151	
		N275F N275F						S121P S121P					-25.8542 -25.4173				-25.876 -25.9887		
Q119	D S214F	R260F	N275F	K252L	Q182I	A80V	T39D	S121P	S92P	T266P	T116R		-25.1877	-26.1812	-25.9402	-25.2621	-24.0476	-24.5075	
		R260F N275F					T39D	S121P S121P	S92P S92P	T266P	T116R						-24.2067 -24.2731		
Q119	D R260F	N275F	S54W	K252L	Q182I	A80V	T39D	S121P	S92P	T266P	T116R						-23.6709		
		R260F						T113P T113P				√ √	-27.6765 -26.4758				-27.8291		
		N275F N275F						T113P T113P				√ √					-28.5727 -27.3496		
		R260F						T113P				√ -√		-26.3319			-26.9418		
		R260F N275F						T113P				./					-25.8834 -27.2131		
Q119	D R260F	N275F	S54W	Q127L	K252L	A80V	S193R	T113P	A40P	T266P	T116R	•	-26.596	-26.8334	-26.3256	-26.6339	-26.5864	-26.6009	
		R260F										√ √		-25.5604 -25.2464			-26.7526 -25.6023		
Q119	D S214F	N275F	S54W	Q127L	Q182I	A80V	S193R	T113P	A40P	T266P	T116R	√ √	-27.0762	-26.1573	-27.7335	-26.6311	-28.0564	-26.8026	
		N275F R260F										√					-26.9014 -25.6435		
Q119	D S214F	R260F	S54W	K252L	Q182I	A80V	S193R	T113P	A40P	T266P	T116R		-25.5384	-25.8305	-25.942	-25.0113	-25.0672	-25.8408	
		N275F N275F		K252L K252L			S193R S193R	T113P	A40P A40P	T266P	T116R		-25.8366 -25.5983		-25.8814 -25.1254		-26.0702 -26.2941	-25.244 -24.9315	
		R260F		Q127L	K252L	Q182I		T113P				√ √	-26.9182	-26.5112	-26.4811	-28.0433			
		R260F N275F						T113P T113P				√					-25.6729 -26.3249		
		N275F					T39D	T113P T113P	A40P	T266P	T116R		-25.9178	-26.5143	-26.3666	-26.7393	-25.1733 -25.5755	-24.7958	
		R260F						T113P									-25.0175		
		N275F N275F						T113P T113P				√	-26.559 -25.9625	-26.1471		-26.8995	-25.9395 -25.1759	-27.5074	
Q119	D S214F	R260F	N275F	Q127L	Q182I	A80V	T39D	T113P	A40P	T266P	T116R		-25.8812	-24.767	-24 7784	-27146	-26 2645	-26.4501	
		R260F N275F					T39D	T113P T113P	A40P A40P	T266P	T116R		-25.0134 -26.3955	-24.3599 -25.779	-24.2591 -26.6166	-26.195 -25.9615 -26.651	-24.9977 -26.8038	-25.2552 -26.8164	
Q119	D R260F	N275F	S54W	Q127L	Q182I	A80V	T39D	T113P	A40P	T266P	T116R		-25.4528	-24.6878	-24.309	-26.651	-25.8912	-25.8149	
		R260F		K252L K252L				T113P T113P									-24.5798 -24.119		
Q119	D S214F	N275F	S54W	K252L	Q182I	A80V	T39D	T113P	A40P	T266P	T116R		-25.6232	-26.0709	-25.4589	-25.5434	-24.9847	-26.0581	
Q119	D S214F	N275F R260F	N275F	Q127L	K252L	Q182I	139D S193R	T113P S121P	A40P A40P	1266P T266P	T116R		-25.9297	-25.0316	-25.7927	-26.9244	-25.2179 -25.9818		
Q119	D S214F	R260F N275F	S54W	Q127L	K252L	Q182I	S193R	S121P S121P	A40P	T266P	T116R	.,	-24.721	-24.6156	-24.3173	-25.213	-23.6289	-25.83	
Q119	D R260F	N275F	S54W	Q127L	K252L	Q182I	S193R	S121P	A40P	T266P	T116R	٧	-25.4031	-24.7054	-25.8109	-26.4651	-25.7732 -24.9227	-25.1115	
Q119	D S214F	R260F R260F	N275F	Q127L	K252L	A80V	S193R	S121P S121P	A40P	T266P	T116R						-26.6604 -24.8698		
Q119	D S214F	N275F	S54W	Q127L	K252L	A80V	S193R	S121P	A40P	T266P	T116R		-25.3349	-25.3165	-24.5447	-26.288	-25.0308	-25.4945	
Q119	D R260F	N275F R260F	S54W	Q127L	K252L	A80V	S193R	S121P	A40P	T266P	T116R						-25.0816 -25.9595		
ÓIIa	0214	NZ60F	NEISE	Y121L	4101	AUUV	orack	DIZIP	A+0P	12002	11104		-20.0435	-20.0421	-20.05/1	-20./3/6	-20.9595	-20.921	

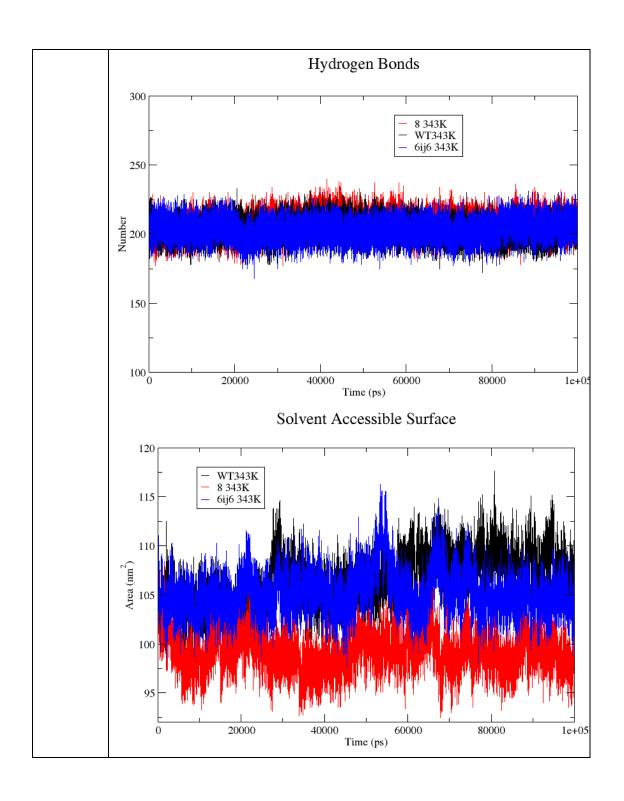
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25.9567 - 24.5388 - 25.9857 -26.4866 -24.9118 S141P S141P 26.337 -25.964 -26.4535 -26.610 S58F S141P 26.5055 -25.4607 26.7969 27.7647 -26.4532 -26.4631 -27.1586 S58P S141P 27 9109 28 2138 -27.961 25.2037 25.5557 -26.4941 -26.3949 26.8939 26.3856 26.6418 -26.684 -26.2827 -26.186 -25.9783 -26.6261 27.2271 26.6295 26.0554 26.5127 25.1734 25.8018 26.035 S141P 26.0207 -25.658 25.6275 25.8573 -25.141 27.1602 26.7399 25.4719 27.6739 -25.3012 25.4192 25.5538 -27.981 -26.186 -25.6091 -26.5302 27.9732 26.4385 26.3753 26.4899 26.7195 26.7266 27.2976 26.9831 27.091

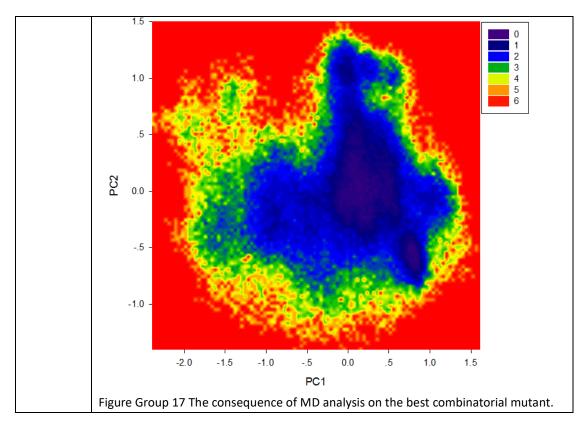
Figure Group 15 All of the DDG value of each Combinatorial mutant calculated. The first few columns are the mutated sites combined in each mutant, which are followed by the stability level that more check marks mean higher stability. The DDG values are showed at the last few columns. The first column is the average of the five values, and the lowest value among the five is showed in blue.

Date	09/13/2020-09/19/2020
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Aim	Try to use ROSETTA for stability analysis similar to FoldX;
	2. Use MD to screen mutants with the best thermal stability.
Participant	Shuying Wu, Dewu Liu, Haoyang Shi, Kegen Chen, Yansong Bai
Content	Learn to install and use ROSETTA to predict mutant thermal stability;
	2. MD and molecular docking analysis were performed on the combined
	mutation screening results.
Result	1. There were many problems with ROSETTA compilation, such as virtual machine
	downtime and installation failure, so we decided to use PYROSETTA with similar
	functions to work.
	2. Determine the best mutant:
	S54W/T72E/N73R/T77E/N114R/Q119D/Q127L/A202C/E231C/R260F/T266P/N275
	F
	PHE-275
	CYS-231 PRO-266
	PHE-260
	CYS-202
	V Spr
	GLU-77
	ARG-73
	ASP-119 GLU-72
	TRP-54
	ARG-114
	Figure 16 The best combinatorial mutant. It contain the mutated sites
	S54W/T72E/N73R/T77E/N114R/Q119D/Q127L/A202C/E231C/R260F/T266P/N275F.







Date	09/20/2020-09/30/2020
Aim	Try to replace Rosetta with pyrosetta
Participant	Dewu Liu, Kegen Chen
Content	Learn to install and use PyROSETTA to predict the thermal stability of mutants.
Result	The virtual machine configuration cannot meet the requirements of compiling
	pyrosetta, and finally the attempt to use ROSETTA and pyrosetta can only be
	abandoned.