

Status: G/U	11G, 15U, 5?	U	U	G	G	U	U	G	G	U	U	G	G	G	U	U	G	U	U	U	U	U	?	?	?	?	?	.		
Credit?	26Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	?	?	?	?	?	.		
Combining?	21Y	Y	Y	Y	Y	N	Y	Y	Y	Y	N	A	Y	Y	N	N	Y	Y	Y	Y	Y	Y	?	?	?	?	?	.		
[A]pple [B]anana [C]herry [S]trawber	11S, 7B, 6A, 2C	A	S	B	B	B	S	A	S	S	B	S	C	A	S	A	S	A	A	C	B	S	B	S	B	S	.			
Can you program in perl?	1Y	N	N	N	N	Y	N	N	N	N	N	N	N	N	N	N	N	N	N	N	N	N	N	N	N	N	N	.		
Can you program in C?	5Y	Y	N	N	N	Y	N	N	N	N	Y	N	N	N	N	N	N	Y	Y	N	N	N	N	N	N	N	N	.		
Single-variable calculus?	25Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	N	A	Y	Y	Y	Y	Y	Y	Y	Y	.			
Do you have a Pantheon Account?	26Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	.			
Read and create web pages? YES	10Y	Y	N	Y	N	Y	Y	N	N	N	Y	N	N	Y	Y	N	N	A	Y	N	N	N	N	Y	Y	N	.			
DNA, RNA	2.8	3	3	3	3	3	3	3	3	3	2	3	3	3	3	3	2	3	3	3	3	2	2	2	2	2	.			
Cell nucleus	2.8	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3	2	3	2	2	2	2	.			
Genetic code	2.7	3	3	3	3	3	3	3	3	3	3	3	3	3	3	2	3	2	1	2	2	3	1			.				
alpha-helix	2.7	3	3	3	3	3	3	3	3	3	2	3	3	3	3	3	2	3	2	2	2	3	2	1		.				
ATP, NAD	2.7	3	3	3	3	3	3	3	3	3	2	2	3	2	3	3	2	2	3	2	3	2	3	3	2	1	.			
Proteins are tightly packed	2.3	3	3	2	3	2	3	3	2	3	3	2	3	2	3	3	2	2	3	1	2	1	2	1	1	1	.			
Bell-shaped Distribution	2.3	3	3	3	2	2	3	3	2	3	2	2	3	1	1	2	2	2	2	2	2	3	2	1	2	.				
Calculation of Standard Deviation	2.2	3	3	2	2	3	3	2	3	2	2	2	3	1	2	2	2	2	2	2	3	1	2	0	3	.				
E. coli is gram negative	2.2	1	3	3	3	2	2	3	3	1	3	1	2	3	3		3	3	3	1	3	3	1	1	1	1	.			
Protein families	1.8	2	1	3	3	3	2	2	2	2	2	2	2	3	1	3	2	2	1	2	1	1	1	1	1	0	.			
A worm is a metazoa	1.8	3	3	3	3	1	1		3	1	3	1	3	2	1	3	1	2	0	2	2	3	0	2	1	0	.			
What chemokines are	1.7	0	2	3	3	2	3	2	3	1		1	3	2	2		3	3	3	1	1	2	0	1	0	0	.			
BLAST search	1.5	0	2	3	3	1	1	2	2	2	3	1	2	1	1	1	2	2	2	2	1	1	0	1	0	1	1	.		
Force as the Derivative	1.3	3	1	3	2	1	1	3	0	2	3	2	1	0	2	1	3	0	1	1	0	0	1	0	0	2	0	.		
3D rotations, translations	1.2	3	3	1	2	1	2	2	2	1	1	0	1	1	2	2	1	0	0	2	1		1	0	0	0	0	.		
A P-value of .01	1.1	3	2	1	1	1	3	1	1	2	0	1	2	2	0	2	0	1	2	0	0	2	1	0	0	0	0	.		
Artificial Intelligence	0.8	3	2	1	1	3	0	0	1	1	0	2	1	1	0	1	1	1	1	1	0	0	0	0	1	0	0	.		
Protein alignment algorithms	0.8	1	2	1	0	1	1	2	2	1	0	1	1	1	1	1	1	0	1	1	0	1	0	1	0	0	0	0	.	
What GroEL does	0.8	0	0	3	3	1	0	2	0	2	3	3	0	0	2	0	0	0	1	0	0	0	0	0	0	0	1	0	.	
Poisson-Boltzman Equation	0.7	2	2	1	1	1	0	2	0	1	1	1	0	0	1	1	0	1	0	1	0	0	1	0	0	0	0	0	.	
Decision trees	0.6	2	0	0	1	2	2	1	0	1	0	1	0	1	0	0	0	0	0	1	2	0	0	1	0	0	0	0	.	
Joining together 2 tables	0.5	2	2	1	0	1		0	0	1	0	1		2	0		0	0	1		0	0	0	0	0	0	0	0	.	
Robotics	0.5	1	2	1	1	1	0	0	1	1	0	0	1	1	0	1	0	0	0	2	0	0	0	0	0	0	0	0	.	
An Extreme Value Distribution	0.5	3	0	2	0	0	2	0	0	1	0	1	0	1	0	1	0	0	0	0	0	1	1	0	0	0	0	0	.	
Genetic algorithms	0.5	3	1	1	1	0	0	0	1	1	0	1	0	1	0	1	1	0	0	0	0	0	0	0	0	0	0	0	.	
Simulated annealing	0.5	1	0	0	1	1	2	0	2	1	0	0	2	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	.	
Constraint Satisfaction	0.4	0	0	0	1	2	0	0	0	1	2	1	0	0	1	2	0	0	0	0	0	0	1	0	0	0	0	0	.	
A Hashing Function	0.4	3	2	0	0	2	1	0	0	0	0	1	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	.	
Neural nets	0.3	2	0	0	1	0	0	1	1	0	0	1	0	1	0	1	0	0	0	1	0	0	0	0	0	0	0	0	.	
Dynamic Programming	0.3	1	1	1	0	2	1	0	0	1	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	.	
Sequence homology twilight zone	0.3	0	0	0	1	1	1	1	1	1	0	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	.	
Bayesian probability	0.3	3	0	0	0	0	0	0	0	1	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	1	.	
A Recursive Descent Parser	0.0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	.	
Belief nets	0.0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	.	
Average		1.9	1.6	1.6	1.6	1.5	1.5	1.5	1.4	1.4	1.4	1.4	1.4	1.4	1.3	1.3	1.3	1.3	1.2	1.1	1.1	1.0	1.0	0.9	0.8	0.7	0.7	0.6	0.5	.

