Name: GRADER ID:______

Consider the following characteristic matrix of two sets: S1 and S2.

и/
H2_
2
2 3 4
4
5
6
)
1

[6 points] What are the two minhash values of S1 and S2 based on the permutation using h1(x) = (x + 1) mod 7 (3pts) and h2(x) = (x + 2) mod 7 (3pts). You need to calculate the minhash values with only one scan on the entire table.

	one scan on the entire table.	15, 52	131
S1 S2	$\frac{h_1 + \frac{S_1 + S_2}{1 + \infty}}{h_1 + \frac{S_1 + S_2}{1 + \infty}}$	$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	$h_1 \mid 1 \mid 2$ $h_2 \mid 2 \mid 3$ [3points]
12 00 00	Rawo h2 21	* O *	S1 , S1 (3 PLU)
	S1 82 h,	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	- hi 1 0 >
→ h1 -> -	$\begin{array}{c c} 1 & 2 \\ \hline 2 & 2 \\ \hline \end{array}$	2 3 Rows h2 0 3	Rowb h2 0 1
Row 3 h2	1213 Rows 42		L>[3points]

 [4 points] Construct a signature for S1 and S2 based on the minhash values obtained from h1(x) and h2(x) above. Estimate the Jaccard similarity of S1 and S2 using the signature. What is the actual Jaccard similarity of S1 and S2 (2pt)? Is the estimate close to the actual Jaccard similarity? If not, suggest a way to improve the estimate (2pts).

Actual Faccard similarity of si and si = 3/6 = 1/2

[I point]

[I point]

[Sestimated]

NO, the estimate is not close to the actual Jaccard Similarity. To improve, increasing the number of pamentations or hash frenctions will help

[2 points]