question	answer	faithfulness	answer_relevancy	context_recall	answer_correctness	Model
What is the total number of features detected in sample	The total number of featur	1.0	0.99999999999994	1.0	0.961183362732859	GPT-4o-mini
Which genus had the highe abundance in sample 2?	abundance in sample 2 is aeromonas, with an abund	0.5	0.9856356310560584	1.0	0.7158631791192785	GPT-4o-mini
What phylum is dominant i sample 3	The dominant phylum in Sa is proteobacteria.	0.0	0.9732905371968088	1.0	0.9704485128211572	GPT-4o-mini
Is feature 27 present in sample 4?	I don't know.	0.0	0.0	0.0	0.2012993433660251	GPT-4o-mini
What is the NCBI taxonomy	Tderfeatoowthat contribute	s 0.0	0.0	0.0	0.1827492467611333	GPT-4o-mini
most to the total reads in	the most to the total reads yes mine gentle with the with sample 5 is featured with ct highest abundance is 3 50	1.0	0.9999228835912102	1.0	0.8349571958895542	GPT-4o-mini
ls the lg €ਜੰ us with the highe abundance also the domin	1 Agramanac which halange	0.66666666666666	0.8753580487354645	1.0	0.8258739210127118	GPT-4o-mini
Whitehreature among the t in sample 6 has the lowest	alassia alassa a successi al blanchesia	1.0	0.9824290601497666	0.0	0.1943042875886866	GPT-4o-mini
What is the percentage contribution of Feature 1 to	Feature1 to the total reads	1.0	0.9999991340172504	1.0	0.9521885496425868	GPT-4o-mini
Doetsample'8 not have an abundant features with ge if the experiment focuses of		le 3 0.0	0.0	0.0	0.2010285312370062	GPT-4o-mini
rare genera, which top fea frome Sahypdensidealigemeas	Footure 24 which has an	0.66666666666666	0.9457905256967986	1.0	0.5007970052410644	GPT-4o-mini
with known genus names, many top features remain	Ifdomo ttekorboa worteria is exclude	d, 0.0	0.0	0.0	0.1942647963298603	GPT-4o-mini
Which phylum has the high	the phylum with the higher abundance in Sample8 is of firmicutes, with an abunda	1.0	0.9406648787119528	1.0	0.7076278727471649	GPT-4o-mini
in one row and 23 in anoth should it be considered	I don't know.	0.0	0.0	0.0	0.2012993433660251	GPT-4o-mini
which genus would become	The dominant phylum in Sa 7 is proteobacteria, which	imple	0.0	1.0	0.559651310546536	GPT-4o-mini
What can you tell me about dominant phylum in Sample	includes features such as Feature1 (genus: aeromon	0.6666666666666666666666666666666666666	0.9418650869615308	0.0	0.5578445314524447	GPT-4o-mini
I am curious, which feature was most abundant in San	Sample 8 is Feature1, with	1.0	0.9497298284995188	1.0	0.1825834947875367	GPT-4o-mini
Sample 5 that appears mo	The genus from ፕሬተክρle 5 i appears more than once in	1.0	0.9804130186128348	1.0	0.5710130808937879	GPT-4o-mini
Do you know how many re Sample 8 has in total?	Sample 2 has 467 features, Sample 50 has a foral sample of the special reads in sample	0.5% 2 1.0	0.9728658037081152	1.0	0.1779354848071201	GPT-4o-mini
27 is relevant in Sample 2	Adatutakitabisishin thesperbusi	dan0t.0	0.0	0.0	0.1869719401333895	GPT-4o-mini
What is the total number of features detected in sample	for identification to the form of the form	0.0	0.9344171653707646	1.0	0.1932638749836377	Llama-3.2 1B
Which genus had the highe abundance in sample 2?	theogeraidinasiনীয়ে জুখাল্য theothighedaleb)urfdenitye is (patkolibiseteraetele) সুভাগট	0.2	0.9005757895496971	1.0	0.1859976164102122	Llama-3.2 1B
What phylum is dominant is sample 3	gantunbacteraceae), gene gantunbanter 5 Orpeatesessa lowest known taxon	n), 0.0	0.7908968993938629	1.0	0.1755624189918952	Llama-3.2 1B
Is feature 27 present in sample 4?	(madtedinacter/genus) and	0.0	0.0	0.0	0.1938015127306353	Llama-3.2 1B
What is the NCBI taxonomy	Material Track College	es, 0.0	1.0	0.0	0.2037845844592909	Llama-3.2 1B
most to the total reads in	denus with the highest genus with the highest genus order bacteroidales abundance is aeromonas.	vhich ¹²⁵	0.906027299074262	1.0	0.3542149711936261	Llama-3.2 1B
Is the genus with the higher abundance also the domin White Meature among the telegraphic states and the second se		0.25	0.8585873194386816	1.0	0.7540366191794874	Llama-3.2 1B
in sample 6 has the lowest	 	0.0	0.8585822701996867	0.0	0.1942974826641244	Llama-3.2 1B
What-is-the²percentage contribution of Feature1 to ₱vet-sample18 inot have a?	Sample7 has 19 features	0.0	0.8615101288709909	1.0	0.1941272479574317	Llama-3.2 1B
abundant features with ge		1.0	0.0	0.0	0.9312725112391828	Llama-3.2 1B
rare genera, which top fea from Sahyplensidealifereas	Contoyt	0.0	0.0	1.0	0.193156999533462	Llama-3.2 1B
with known genus names, many top features remain if proteobacteria is exclude	Sample 3 has 50 features	0.0	0.8843563762720575	0.0	0.1911313230393792	Llama-3.2 1B
which phylum has the high	Answer: bacteroidota Answer: Feature1 (17383 r	0.0	0.8516292570223564	1.0	0.2109600573688958	Llama-3.2 1B
in one row and 23 in anoth should it be considered	\$a3%pleNylasn5@fedatologste	riaβ.0	0.7960750371605195	0.0	0.1866138193990987	Llama-3.2 1B
which genus would become	genus: aeromonas; Feature (Ætitēxteads, 0.5%, phylum fisæri@ukestjærerWshat can ye	: 0.5	0.0	1.0	0.4906939115532915	Llama-3.2 1B
What can you tell me about dominant phylum in Sample	telols torried izoloobeest; tiFreea toloomea 11:00a n	0.0	0.0	0.0	0.1891826596013451	Llama-3.2 1B
I am curious, which feature was most abundant in Sam can you name a genus fro	(6395 rciads; 0.5%, 78 hylium proteobacteria genus; Samples has 31 clearurgs.	0.7	0.7886683500682995	1.0	0.2118835629998424	Llama-3.2 1B
Sample 5 that appears mo	there are 31 unique feature	0.333333333333333	0.869529132942584	1.0	0.2047301505570253	Llama-3.2 1B
Do you know how many re Sample 8 has in total?	Sample8 has 9595 reads	0.0	0.9670589944730598	0.0	0.1789357399308756	Llama-3.2 1B
27 is relevant in Sample 2	'	0.0	0.8680461067575878	0.0	0.1831106059443738	Llama-3.2 1B
all?	I bacteroraes,					

Yes, it is present in sample
4. Document 3 (table):
sample_id is Sample4,
feature_id is Feature26,
abundance is 0, id is X_0265,
KhmgdumstipbaisteMbjqbhfdaturis
proteophtmetrip_Sample 6
gantheplowtesbabterda,nower is
Booklimbatatis_Mashfabts_mode6.

	bashtha pootestbabterda, noed Docktroketer ia keepta in Samp	26				
question	Brackware/LBadewesgermoswe_	ax fa ithfulness	answer_relevancy	context_recall	answer_correctness	Model
What is the total number of features detected in sample	ictalieatures detected s in blownerika ovamie xon semules i Ecatural E	s 0.0	0.9430634294448604	1.0	0.2015251223213842	OLMo 1B
Which genus had the highed abundance in sample 2?	Handey Have Fire can list of the control of the con	0.5	0.7824618977175825	1.0	0.2021893772413695	OLMo 1B
What phylum is dominant i sample 3	(dan b) de de la composition della composition d	le 4,0.0	0.0	1.0	0.1896007345084115	OLMo 1B
Is feature 27 present in sample 4?	abies danc avision rest violation	nor0g0	0.923141289500946	0.0	0.9361362836770828	OLMo 1B
What is the NCBI taxonomy feature 2? Which feature contributes the	STATES BELT HERENGS SEED OF THE WOOD AND THE	nen 0. 4	0.0	0.0	0.1804138531058456	OLMo 1B
most to the total reads in	baludibacteraceae, genus: Baludibacter-Species; nan Baludibacter-species; nan	1.0	0.8054521248074019	1.0	0.3215544049947998	OLMo 1B
abundance also the domin Whitehfeature among the to	hd Drestimbent da (tablis) : • \$aatatost Contintal S enus i	0.0	0.9326264092511772	1.0	0.2044671960983485	OLMo 1B
In sample 6 has the lowest		ost 0.2857142857142857	0.0	0.0	0.260076091908525	OLMo 1B
What is the percentage contribution of Feature1 to Does sample 8 not have af abundant features with ge If the experiment ocuses or	contributed denoted is clostridic to the common get a common get a clostridium. The fourth man	nu�i9	0.0	1.0	0.2060126263655618	OLMo 1B
abundant features with ge	The third most common get the strict of the	as indine wise	0.0	0.0	0.2039265172947579	OLMo 1B
frome Sahyptensidealigemeas	Beample as a sample ce 30 c	7 1.0 0 Th	0.0	1.0	0.4415639771072537	OLMo 1B
with known genus names, many top features remain ir proteobacteria is exclude	Sevento most common ger	us 0\$0	0.8507419175097953	0.0	0.1835042359400822	OLMo 1B
which phylum has the high	Makare mental and a company of the c	<u>۔۔</u>	0.0	1.0	0.1948591213536165	OLMo 1B
in one row and 23 in anoth		t ₂ 0.63636363636364	0.8391676902352762	0.0	0.4041639711111374	OLMo 1B
which genus would become	Sample: Features: Sample: Figures: Joseph Sample: Samp	nie)is)	0.0	1.0	0.1964562319898224	OLMo 1B
What can you tell me abou dominant phylum in Sampl		இருந்த 333333333333333333333333333333333333	0.8560159929138916	0.0	0.1810617123316813	OLMo 1B
I am curious, which feature was most abundant in Sam Can you name a genus from		100 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	0.0	1.0	0.1853447133521872	OLMo 1B
Sample 5 that appears mo	estusas para la la la comenta a vial del Marco a para la la comenta de l	0.0 logs	0.8474356219174872	1.0	0.8969381408198872	OLMo 1B
Do you know how many re Sample 8 has in total?		lino.o siniss:	0.0	0.0	0.1835726832289773	OLMo 1B
27 is relevant in Sample 2		de a01.d777777777777778	0.9787986145513408	0.0	0.1871818451185229	OLMo 1B
What is the total number o features detected in sampl	nete ne mi Wirds nese da easinetas Postaranin (22 gradas meesie xietaja, yeds	an 0.0	0.8985502857275599	1.0	0.2762664231385769	Pythia 1.4B
Which genus had the higher abundance in sample 2?		pe to 25	0.0	1.0	0.3891047933692881	Pythia 1.4B
What phylum is dominant i sample 3	propress First off T vou do	off 0.0	0.0	1.0	0.1646665353028396	Pythia 1.4B
Is feature 27 present in sample 4?		ano.o	0.0	0.0	0.2990636998184648	Pythia 1.4B
What is the NCBI taxonomy feature 2? Which feature contributes !		or 0.0 Geografia	0.0	0.0	0.1834382899481278	Pythia 1.4B
I most to the total reads in I		arie 0.333333333333333333333333333333333333	0.0	1.0	0.1990002775290602	Pythia 1.4B
sthe genus with the higher abundance also the domin Whith feature among the to		\$\$₹ _{0.0} Qr	0.7187931654560217	1.0	0.1787431488201952	Pythia 1.4B
in sample 6 has the lowest		1002 1002 1004	0.8059465504760045	0.0	0.1885404787436742	Pythia 1.4B
What is the percentage contribution of Feature 1 to Does sample 8 not have at	orajakaidak augeaja ja ja ja kasakaja akkenika a ja	Geš. gat.	0.8118611219201525	1.0	0.1868237594002829	Pythia 1.4B
abundant features with ge	ाकाराम् अस्ति । स्वति । स्वति । स्वति । स्वति	Maile (1254	0.7885609407422981	0.0	0.4878707046011177	Pythia 1.4B
rare genera, which top fea ffowe தெந்நம் நிழ்வர்க்கை	្ត្រាប់ ទៅលើ ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុ ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង ខេត្តក្រុង	ofictivities)	0.7555528322253077	1.0	0.1880218122748171	Pythia 1.4B
with known genus names, many top features remain if proteobacteria is exclude		eveet 0	0.0	0.0	0.186884567322476	Pythia 1.4B
which phylum has the high		g 0.0 g,	0.0	1.0	0.1862151284282007	Pythia 1.4B
in one row and 23 in anoth	obe to to an inventive and a set in roman see it easiles of Not south as consider when	his0.125	0.0	0.0	0.4233493861320965	Pythia 1.4B
which genus would become		gs 0.0 Wead	0.0	1.0	0.6844700318351403	Pythia 1.4B
What can you tell me abou dominant phylum in Sampl		it 0.0 ame.	0.0	0.0	0.1752113736627249	Pythia 1.4B
I am curious, which feature was most abundant in Sam	Salet A Chrow Viewbrastork Cerned Heover 100 et net Las ose puewine die saitses ovi	ed 0.0	0.0	1.0	0.1705435819236642	Pythia 1.4B
Sample 5 that appears mo			0.0	1.0	0.2625740183401832	Pythia 1.4B
Do you know how many re Sample 8 has in total?	in e tinsothradswick e ayro'ta tredidifa tideaealbarue soerene anfiythotraan	ⁿ 0.0	0.0	1.0	0.1672344130074508	Pythia 1.4B
27 is relevant in Sample 2	features but were unable t get those ideas into your o	0.0	0.0	0.0	0.1734122688965698	Pythia 1.4B
JIII		leyn erawe				

get those ideas into your of the law we were to remove duplicate values from consideration.