Description of seqenv's main internal data structures

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Γ		seq_to_gis		One to many		
			[343206452,		199999999]	
ı	C1	->	[234523451,	343204354]		
ı	C2	->	[]			

gi_to_text (For all NCBI GIs) One to one
343206452 -> "the deepest sea mud of the Mariana Trench"
343204354 -> "isolated from terrestorial samples near ..."
199999999 -> ""
343204350 -> "the deepest sea mud of the Mariana Trench"
... etc etc

text_to_matches (Only for sources found)

"the deepest sea mud of the Mariana Trench" -> [ENV0:123123, ENV0:123777, ENV0:123777]

"isolated from terrestorial samples near ..." -> [ENV0:23452345]

text_to_countsOne to one to many"the deepest sea mud of the Mariana Trench" -> (ENV0:00002007 -> 1, ENV0:00002041 -> 2)"isolated from terrestorial samples near ..." -> (ENV0:00001993 -> 1, ENV0:00003452 -> 1)

 seq_to_counts
 One to one to many

 C0 -> (ENV0:00002007 -> 0.667, ENV0:00002041 -> 0.333)

 C1 -> (ENV0:00002007 -> 0.123, ENV0:00002041 -> 0.821)

 C2 -> ()

serial_to_concept lorem

child_to_parents
lorem

concept_to_name
lorem

df_abundances lorem

All numbers are fake
This document just describes a plan for processing data.