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e Serial Number Â .Q: Unable to query MongoDB using RazorJSON I'm trying to query my Mongo database using mongocsharp driver and RazorJSON syntax but I can't manage to get it working. The syntax is `Client.db.Database.GetCollection(MyCollection).AsQueryable();` and the returned value is just 1 document (2 actually but the second document is null). If I use the Code Below I can't get a validation error but the returned doc is null. `string lookup = MyCollection + "." + MyField; Client.db.Database.GetCollection(lookup).AsQueryable();` A: If you are not using the overload of `GetCollection` that uses the `AsQueryable()`, you need to use the overload that uses the `Request` as a parameter. So, you'd need to do something like: `string lookup = MyCollection + "." + MyField; Client.db.Database.Request.GetCollection(lookup).AsQueryable();` Also, you may need to use the `FullSynchronized()` method on the Database, since your query is running asynchronously, rather than synchronously. [Analysis of the EEG spectrum on moderate

hypothermia]. The normal awake and sleep EEGs of 17 patients with moderate hypothermia were analyzed, in an attempt to find out EEG characteristics which can identify these cases. Generalized slowing was more remarkable in the EEGs in which the temperature was under 32 degrees C. The amplitudes of components of the 1-5, 6-9 and 10-20 Hz bands were also significantly reduced. The characteristic decrease in the amplitude of 7-9

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subunit of ribosomes is one of the basic proteins
which, in turn, are constituent components of the
ribosome. The large subunits of ribosomes from
eukaryotes and prokaryotes are distinguished on
the basis of structural differences, differences in
amino acid sequence, their mode of synthesis and
the subsequent processing events which occur
during the assembly process. It has been shown
that in vitro protein synthesis systems, based upon
coupled initiation and termination, can be

reconstituted from individual ribosomal proteins. However, there is a continuing interest in studying the large subunits of ribosomes from different organisms and from different species of a single organism, in order to compare the nature of these two kinds of ribosomes. As a result, there is a need for a model system for the study of large ribosomal subunits, but such a model must be one which is simple, reproducible, and inexpensive. The proposed research is to determine the structural and biochemical characteristics of large ribosomal subunits from a thermophilic organism, *Bacillus stearothermophilus*, which synthesizes a thermophilic ribosome that is indistinguishable from the eubacterial ribosome in its functional properties. The ribosomes of this organism are also easily inactivated by the selective antibiotics, tylosine and tritylosine. The ribosomes will be isolated from a crude ribosomal fraction by the high-salt procedure which has been developed in the laboratory, to be studied in the ribosomal salt wash.

The large subunits will be separated from the small subunits by size exclusion and ion exchange chromatographies. The subunits will be purified by chromatography on heparin-sepharose, phosphocellulose, hydroxylapatite, and phosphocellulose, followed by rechromatography on hydroxylapatite. The large subunits will be characterized with respect to their amino acid composition and their ability to bind substrate during in vitro protein synthesis. The invention relates to a fastening device e79caf774b

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These cookies do not store any personal information. A cyclopropenylchromium(V) precursor to the antiviral agent dianhydride acyclovir. Oligo(phenylenevinylene)s bearing cyclopropenylchromium(V) substituents (PVCs) show excellent

sensitisation efficiencies, and can be efficiently cross-linked to give polymers with much higher T(g)'s and photostabilities than those resulting from non-conjugated photochromic dienes. One such ligand, 3,9-diphenyl-9-propenoic acid-4-(6-methoxy-2-methylphenyl)-6-methylcyclohexa-1,3