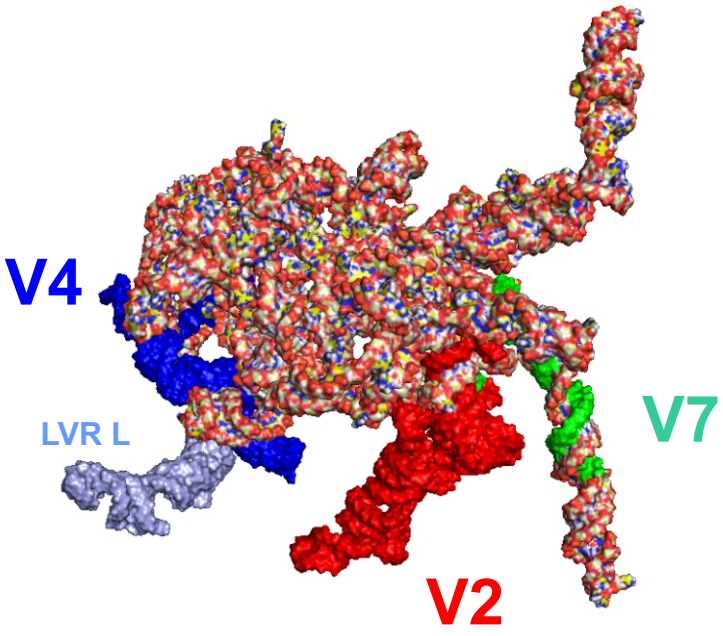
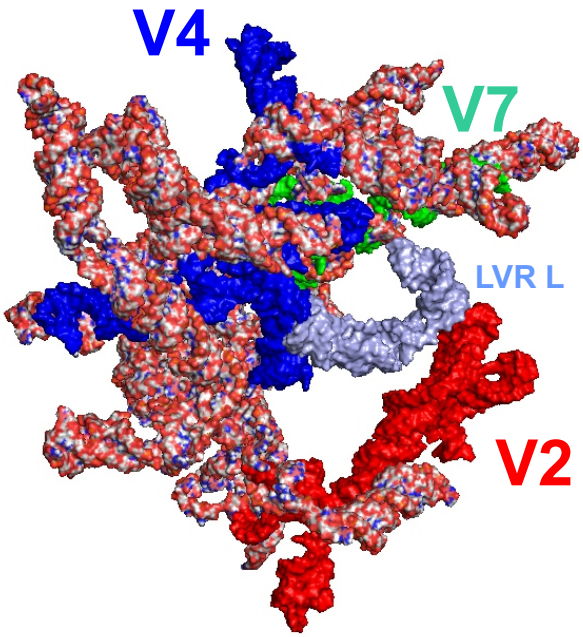
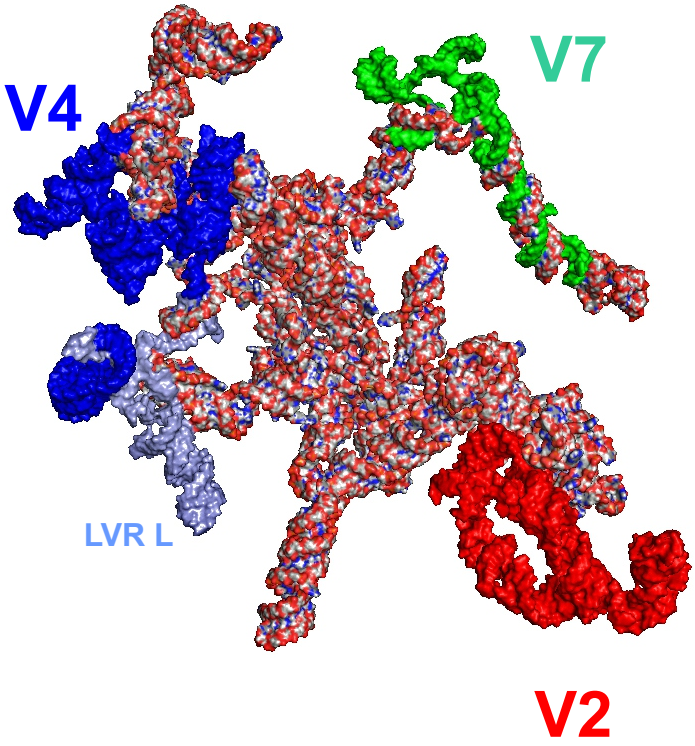
File S6. The LVR L position (marked in light blue) within the hypervariable region V4 in the tertiary structure models of the 18S rRNA gene. The hypervariable regions are marked in red (V2), dark blue (V4), and green (V7). The sequences were not aligned, and each is displayed in such a position that the LVR L can best be seen.



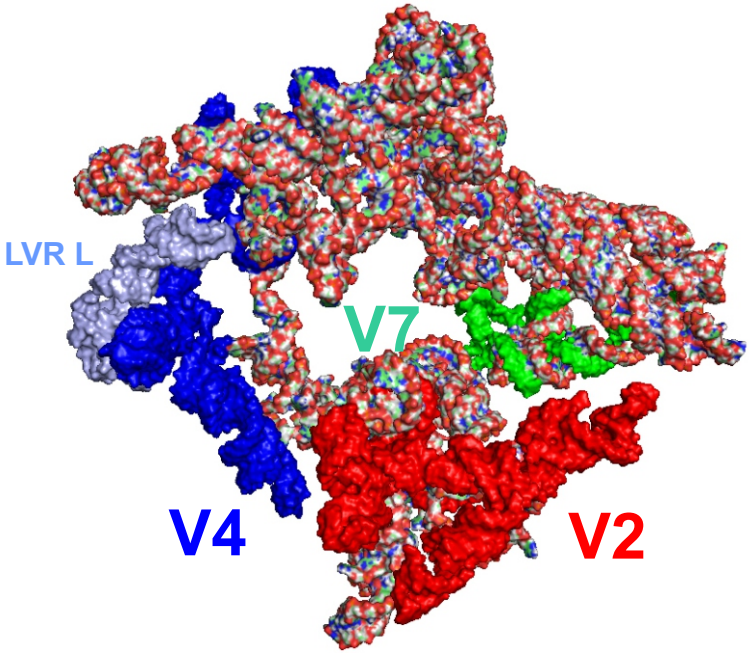
Adomerus biguttatus



Canopus sp.

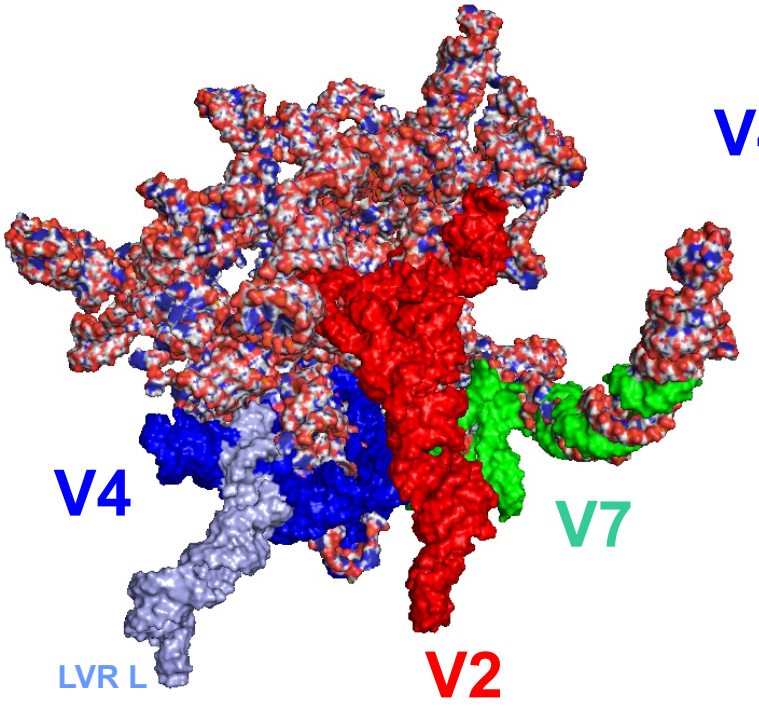


Cantao ocellatus

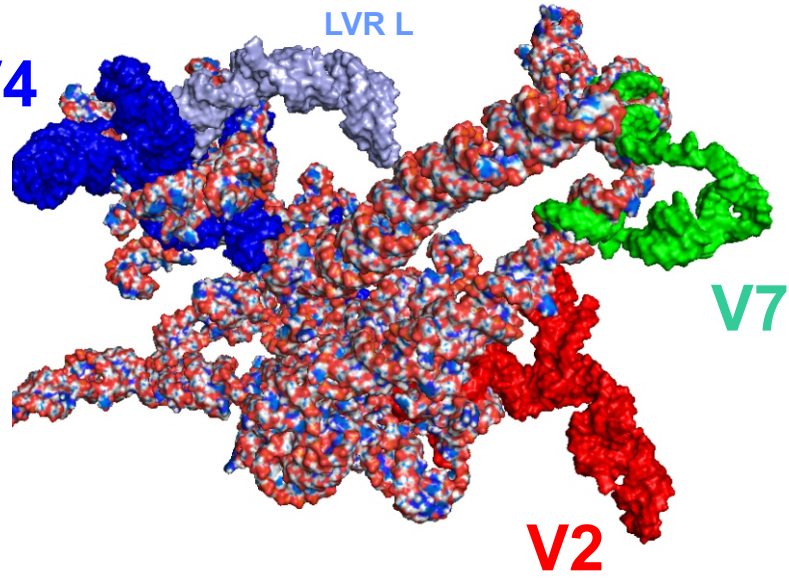


Elasmostethus interstinctus

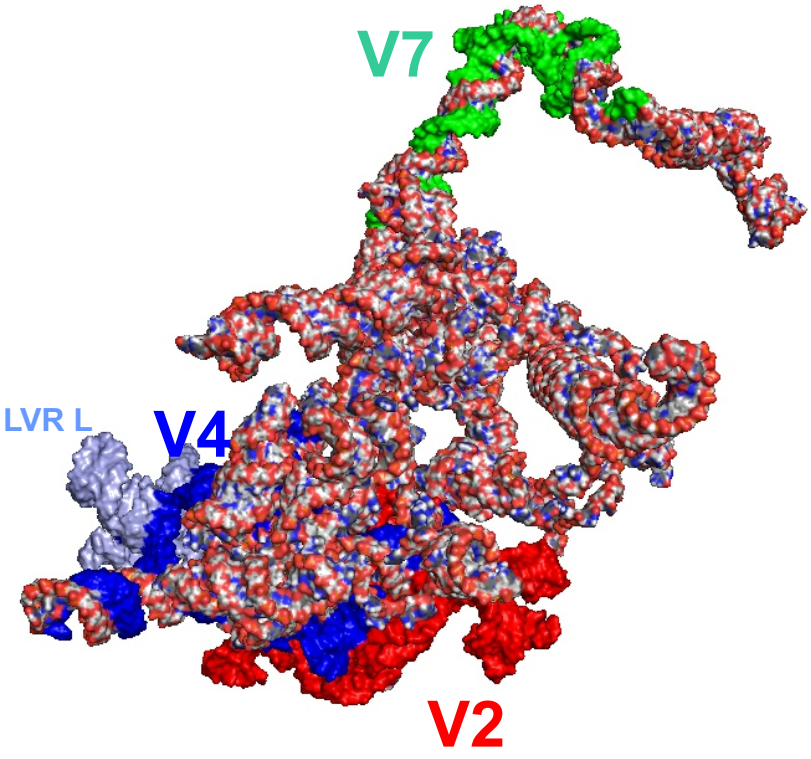
File S6. The LVR L position (marked in light blue) within the hypervariable region V4 in the tertiary structure models of the 18S rRNA gene. The hypervariable regions are marked in red (V2), dark blue (V4), and green (V7). The sequences were not aligned, and each is displayed in such a position that the LVR L can best be seen.



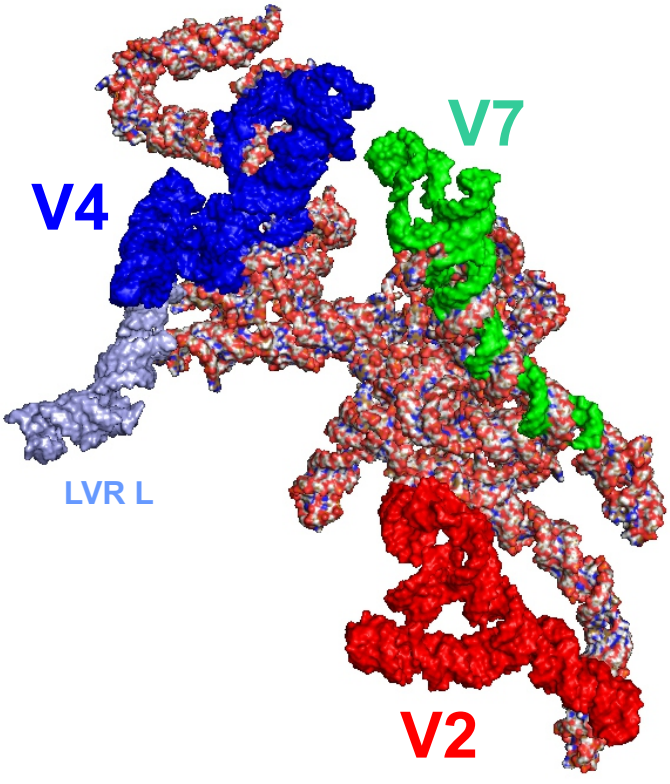
Eurydema maracandica



Fromundus pygmaeus



Megymenum sp.



Thyreocoris scarabaeoides