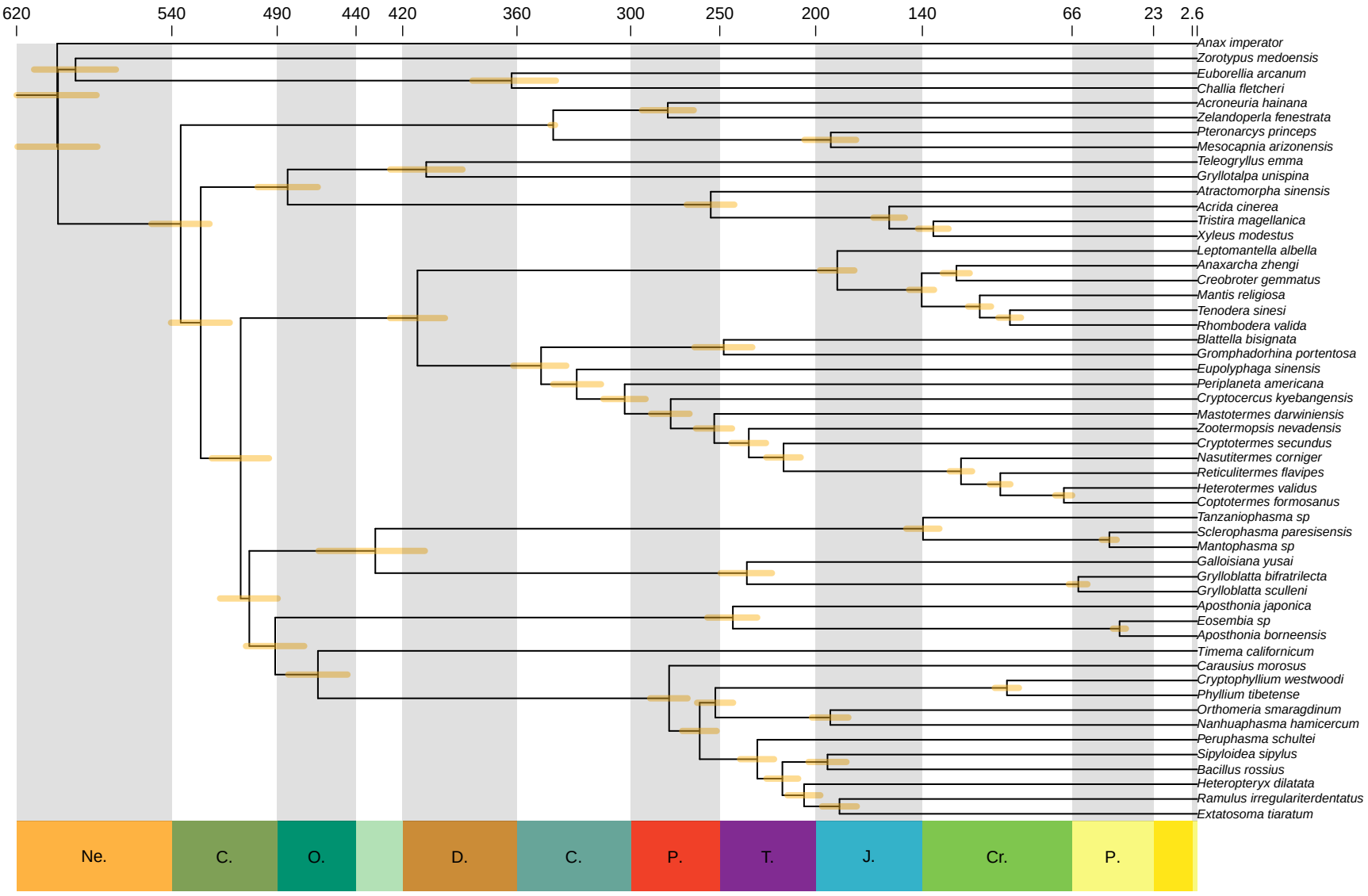
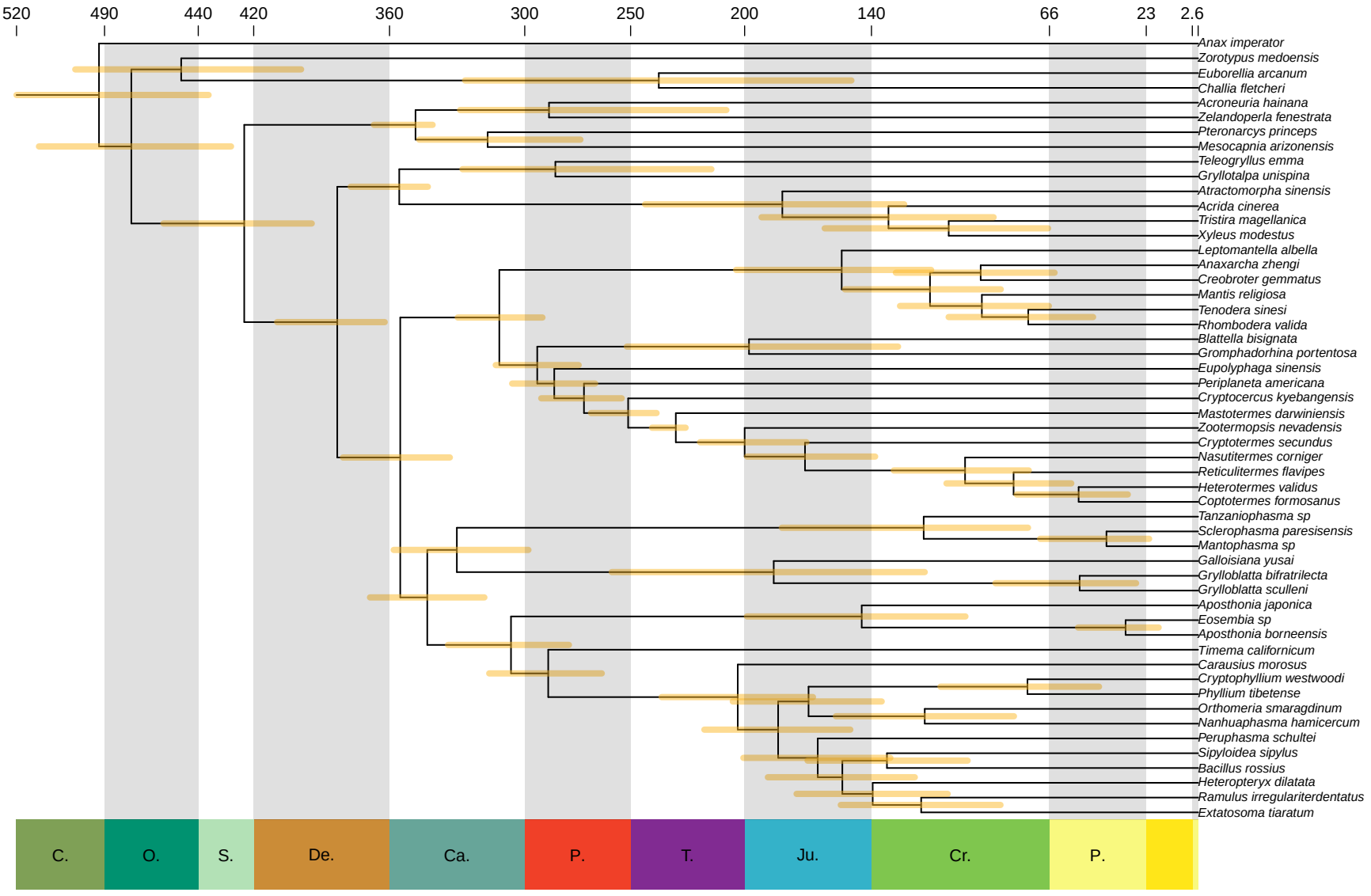
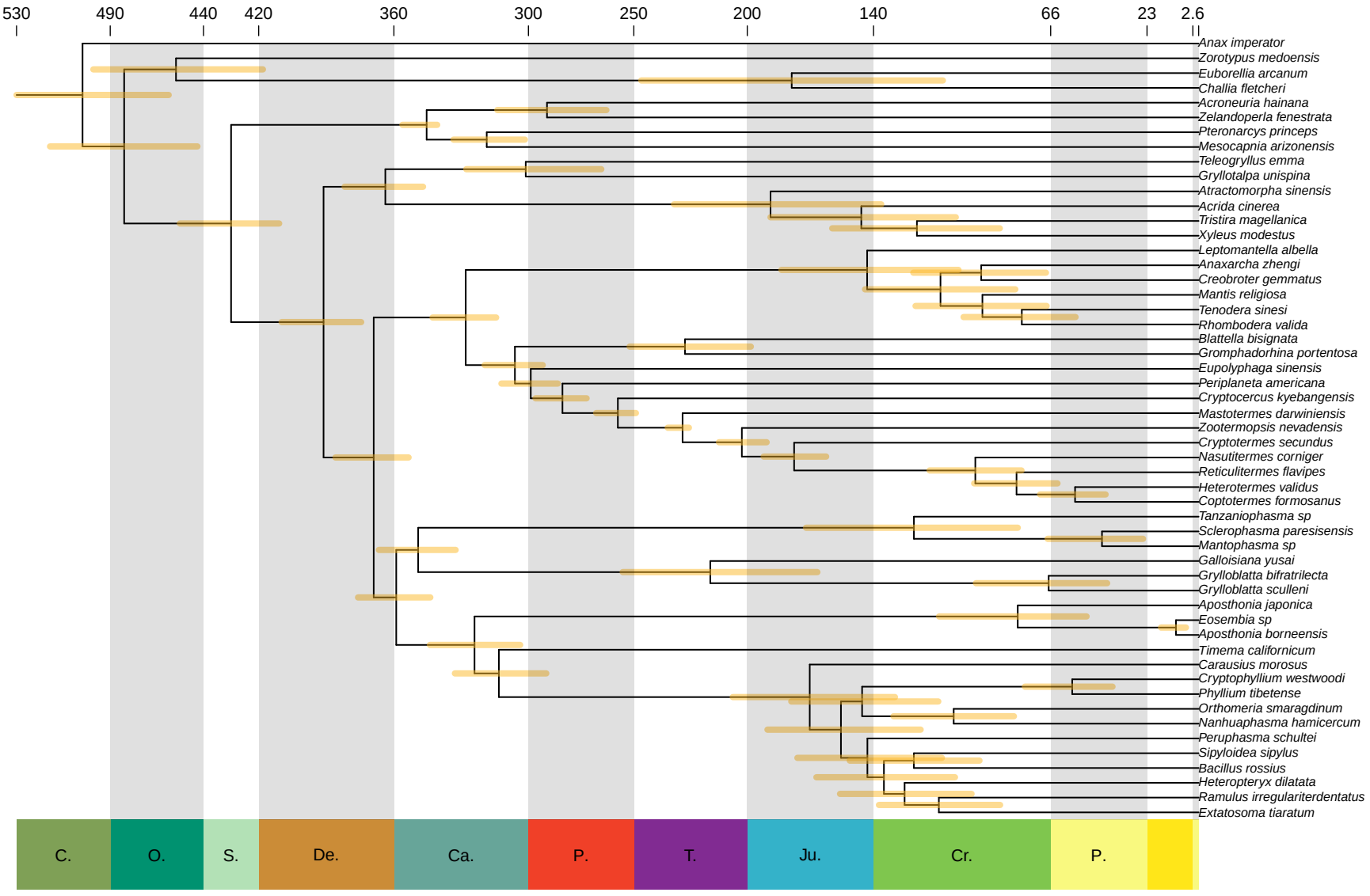


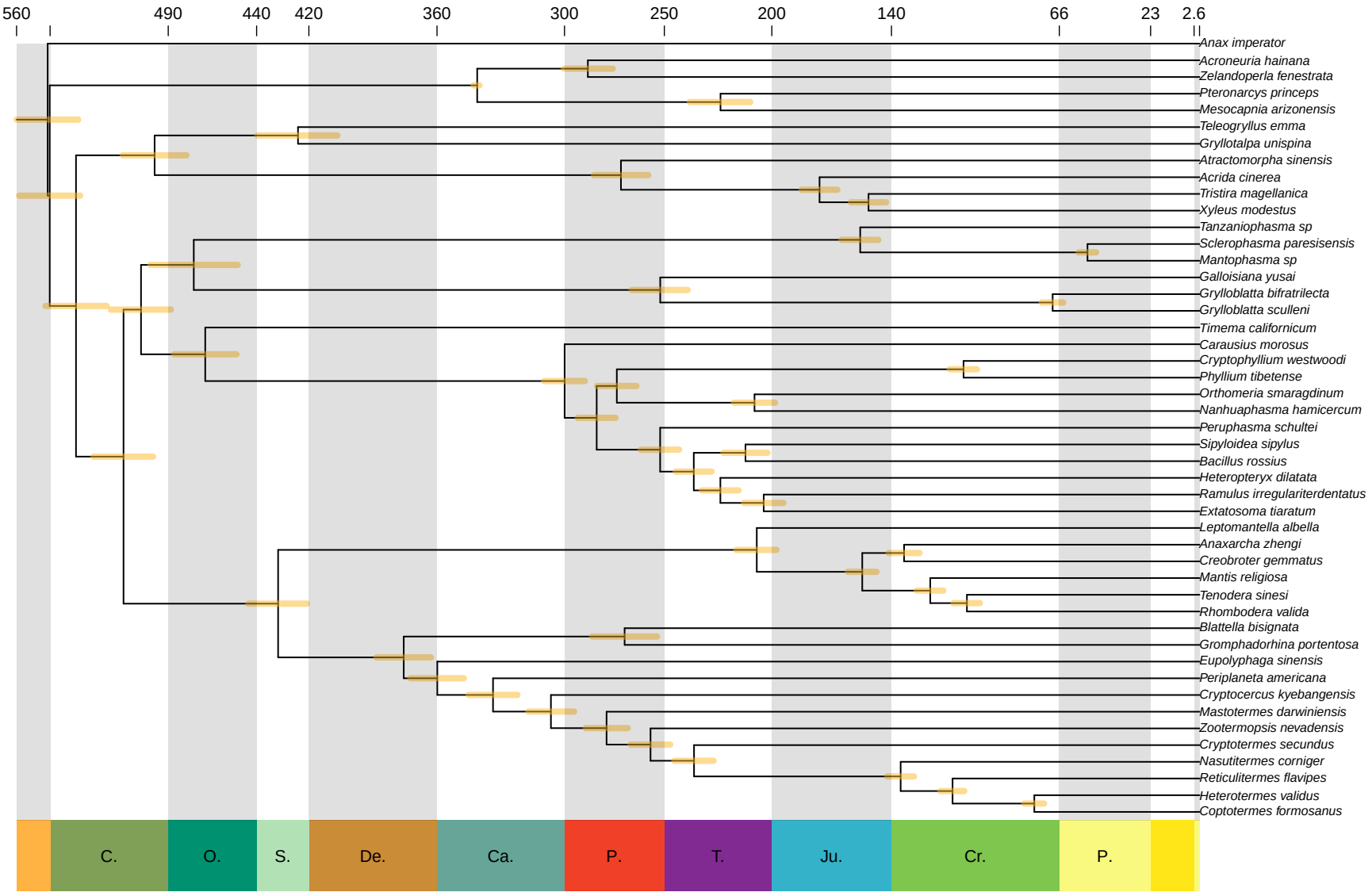
Mitochondrial phylogeny generated using MCMCtree with a strict clock model and the nucleotide sequences including Embioptera and Zoraptera.



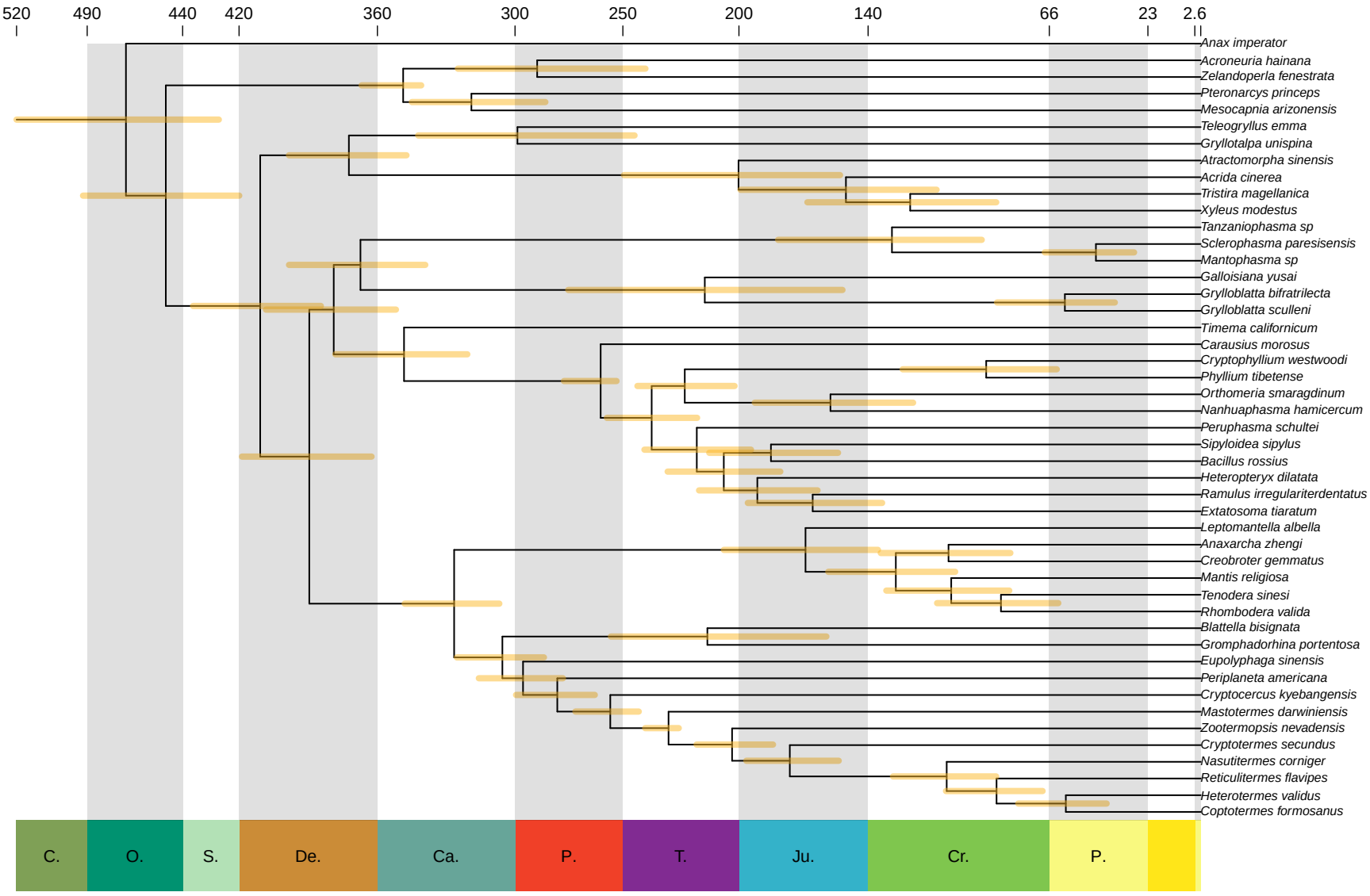
Mitochondrial phylogeny generated using MCMCtree with an indipendent rates clock model and the nucleotide sequences including Embioptera and Zoraptera.



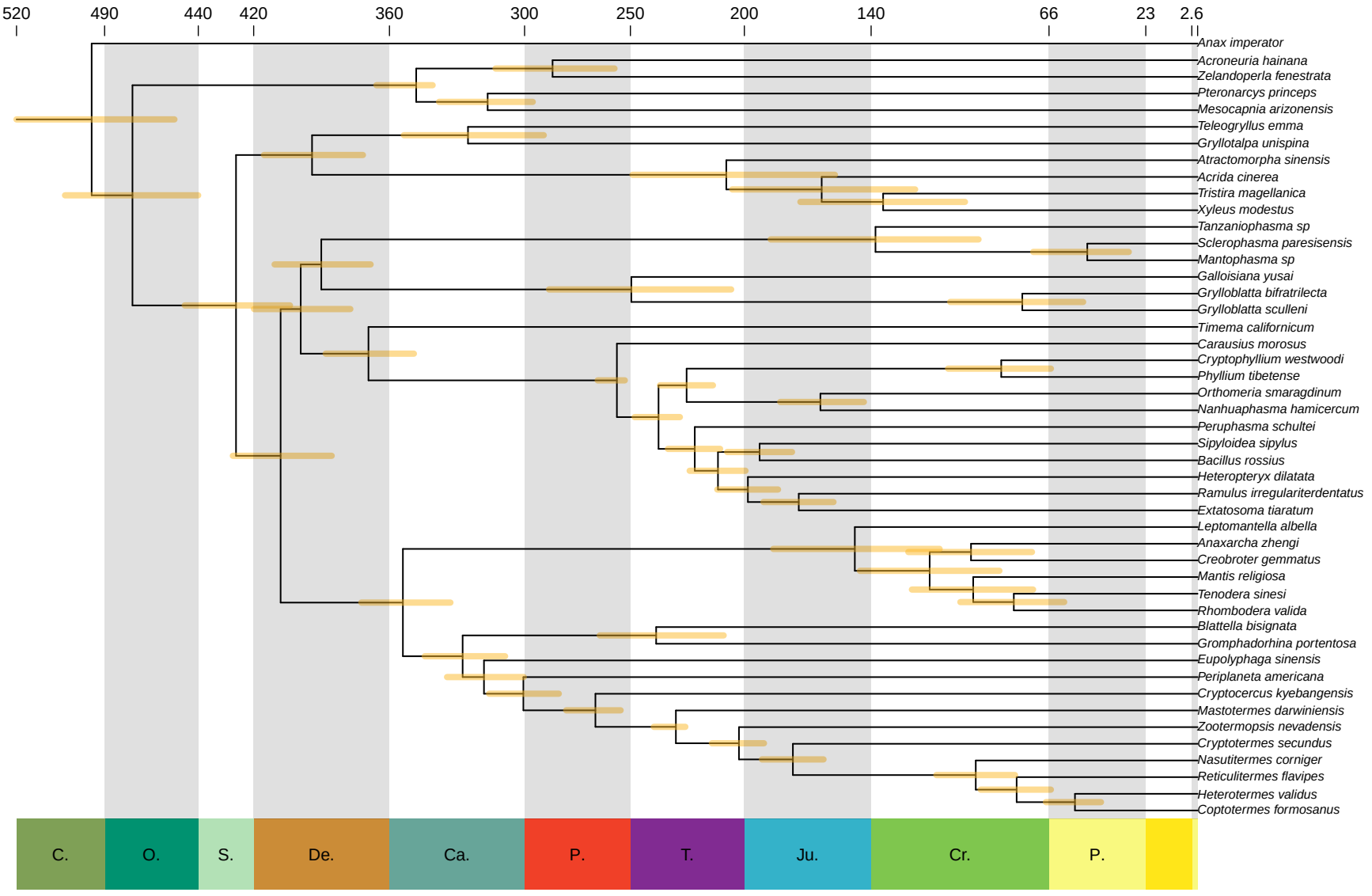
Mitochondrial phylogeny generated using MCMCtree with a correlated rates clock model and the nucleotide sequences including Embioptera and Zoraptera.



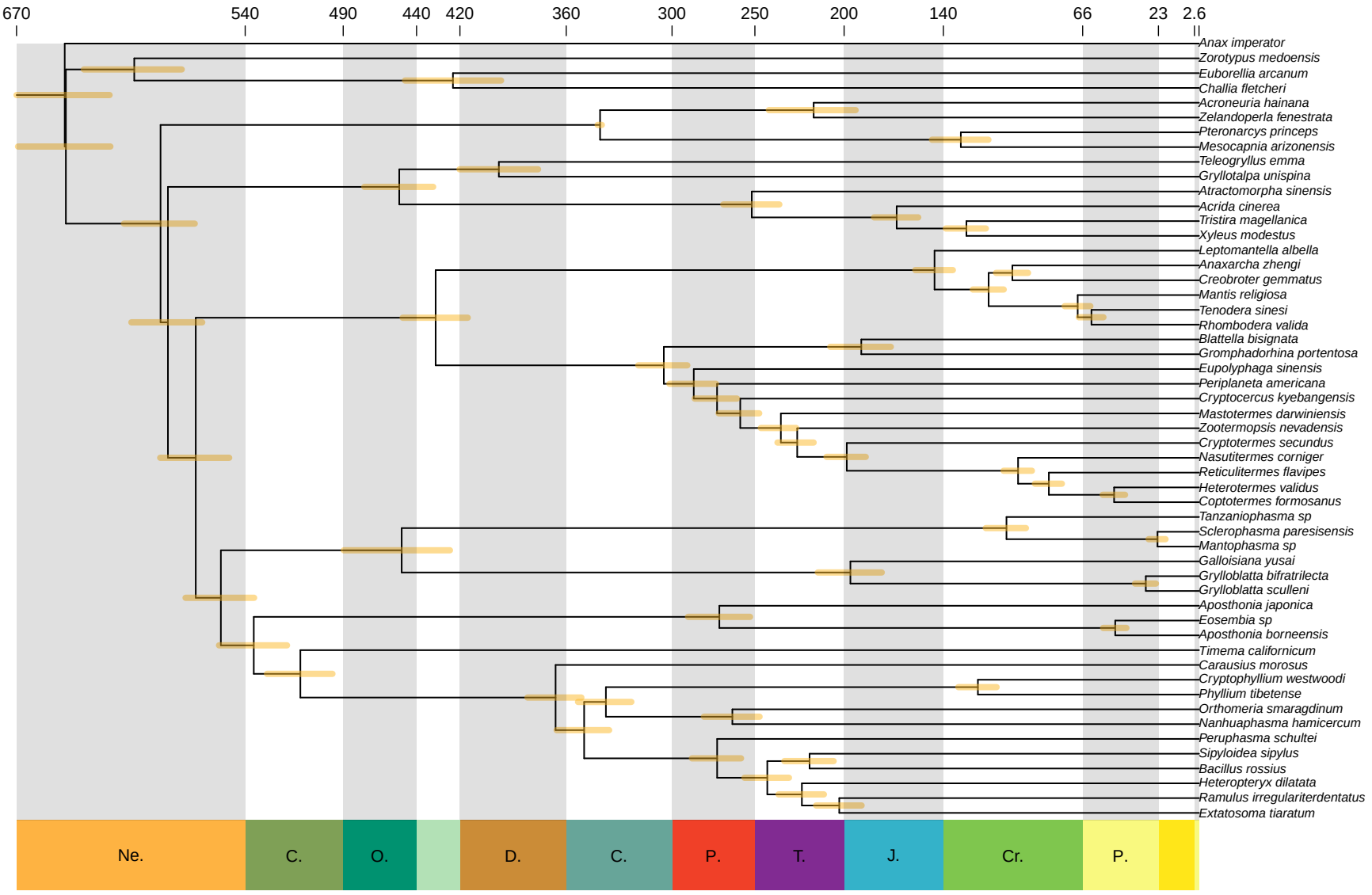
Mitochondrial phylogeny generated using MCMCtree with an strict clock model and the nucleotide sequences without Embioptera and Zoraptera.



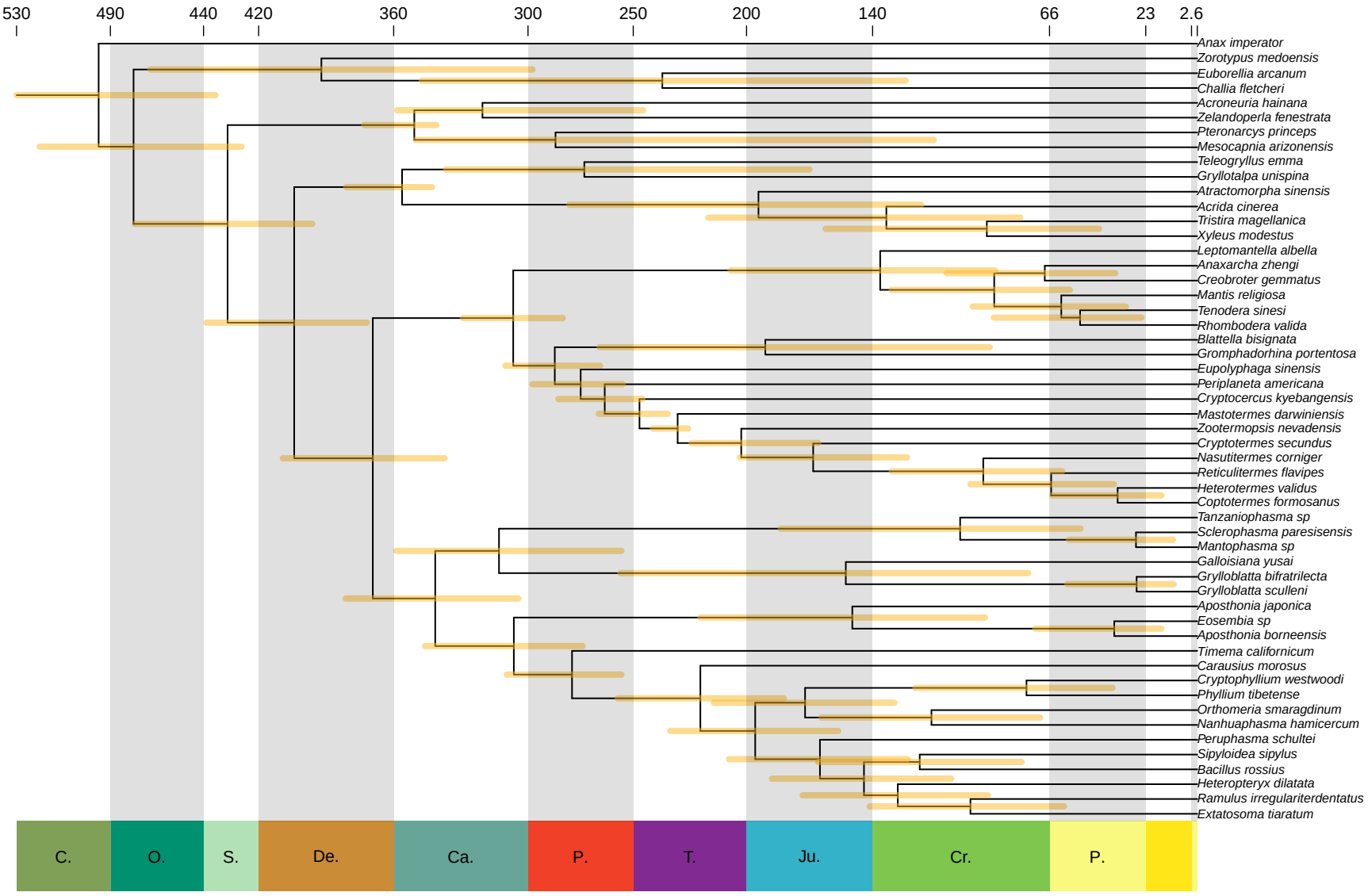
Mitochondrial phylogeny generated using MCMCtree with an indipendent rates clock model and the nucleotide sequences without Embioptera and Zoraptera.



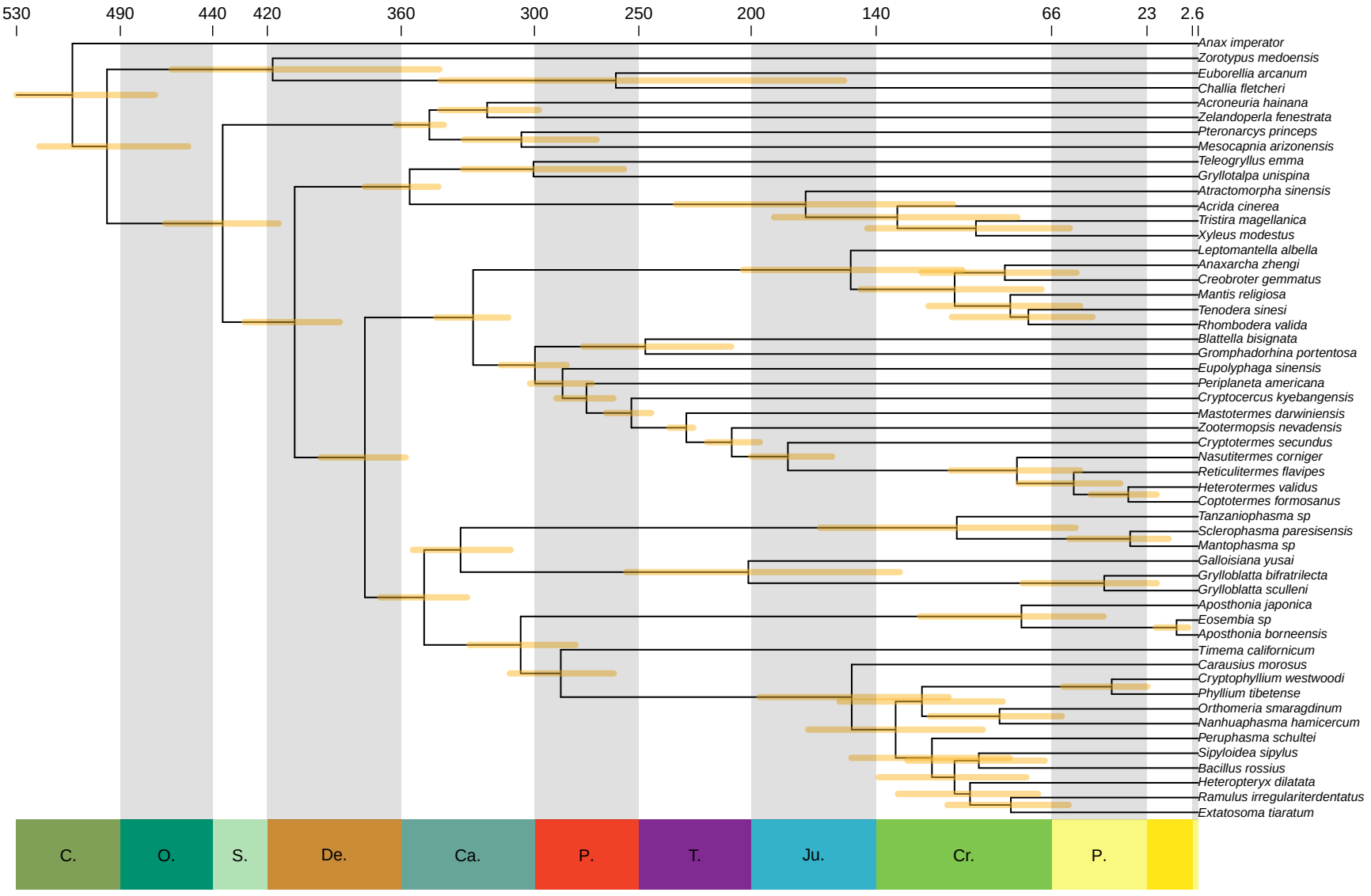
Mitochondrial phylogeny generated using MCMCtree with a correlated rates clock model and the nucleotide sequences without Embioptera and Zoraptera.



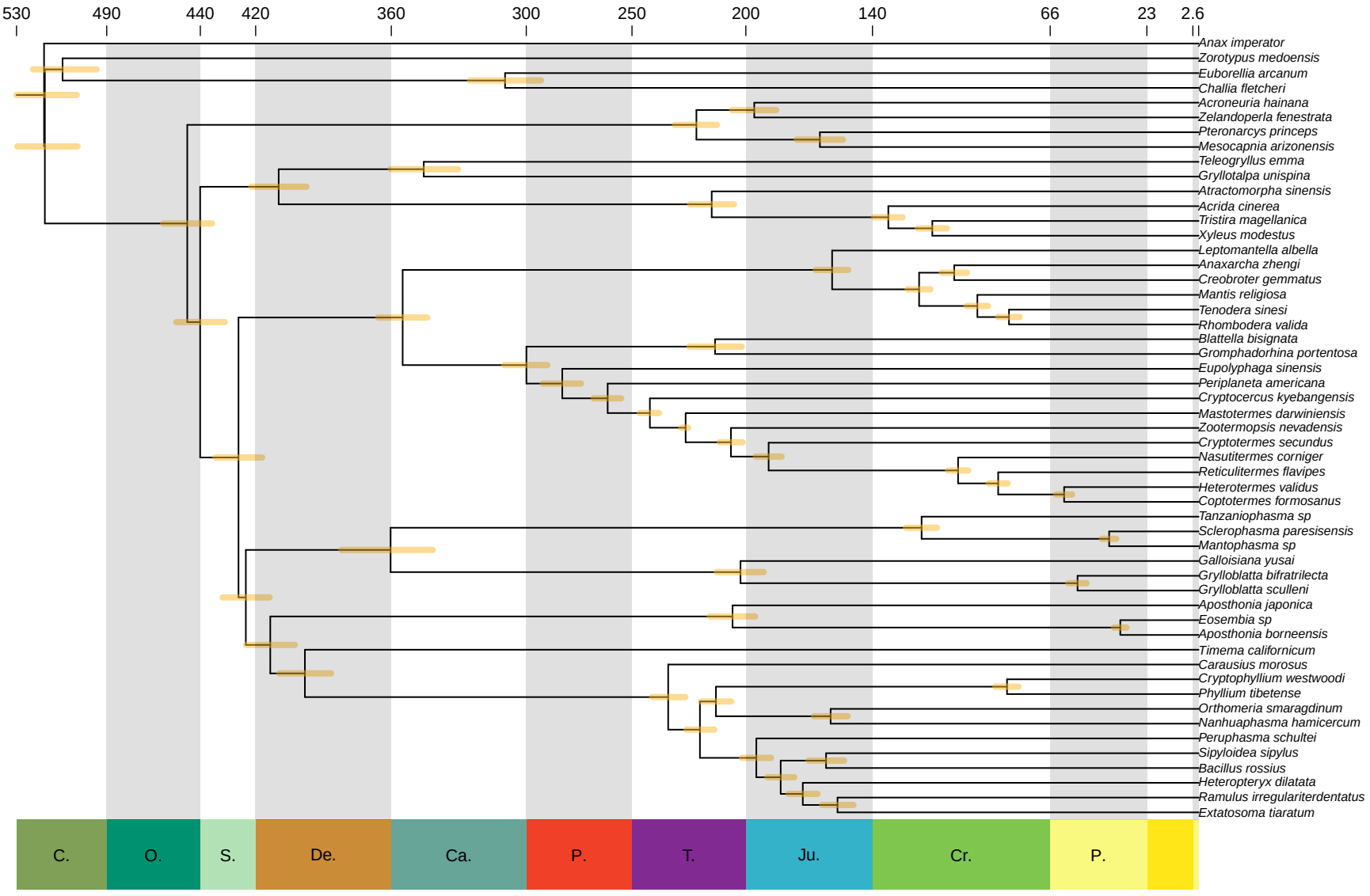
Mitochondrial phylogeny generated using MCMCtree with a strict clock model and the aminoacids sequences including Embioptera and Zoraptera.



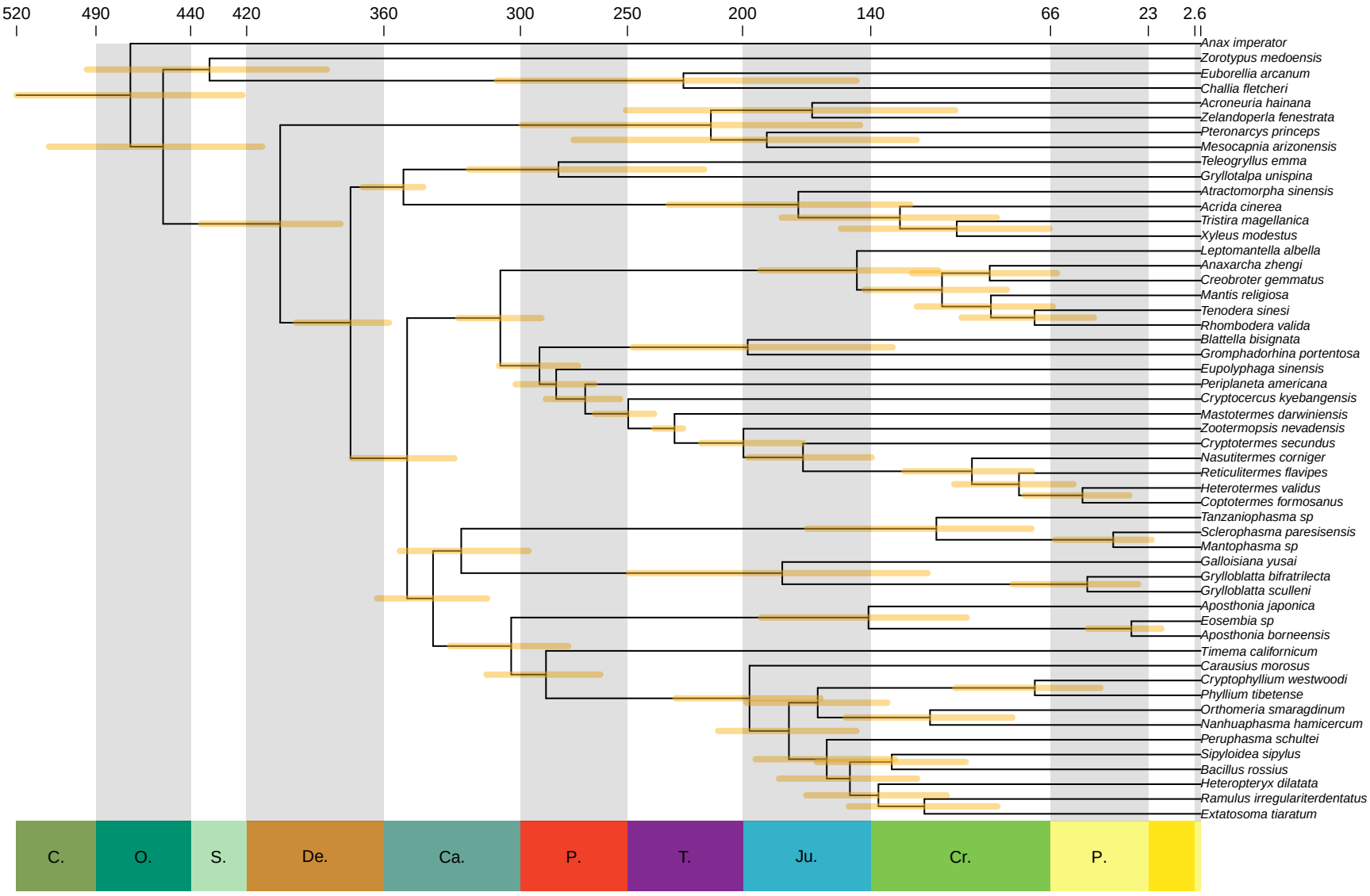
Mitochondrial phylogeny generated using MCMCtree with an independent rates clock model and the aminoacids sequences including Embioptera and Zoraptera.



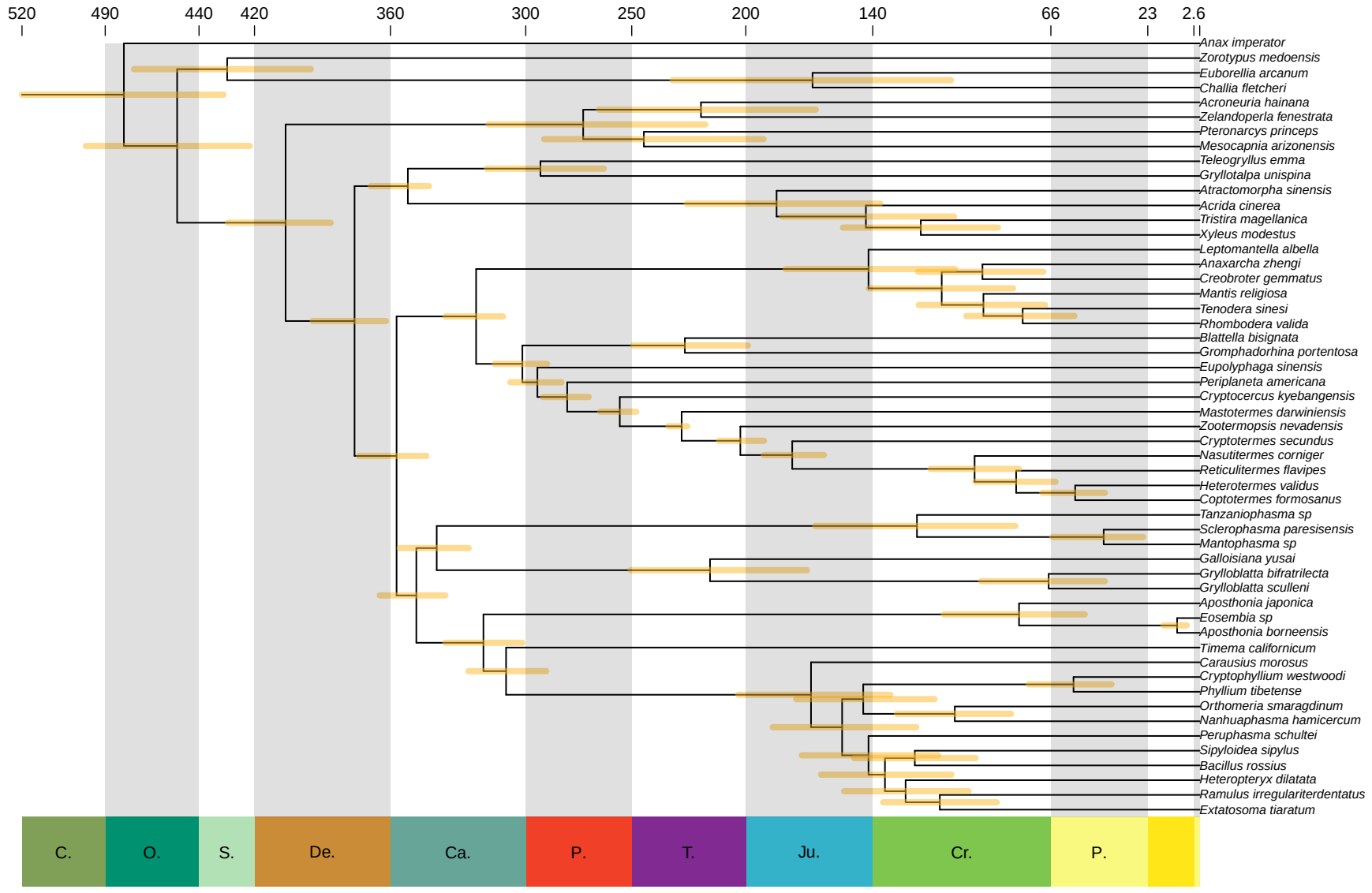
Mitochondrial phylogeny generated using MCMCtree with a correlated rates clock model and the aminoacids sequences including Embioptera and Zoraptera.



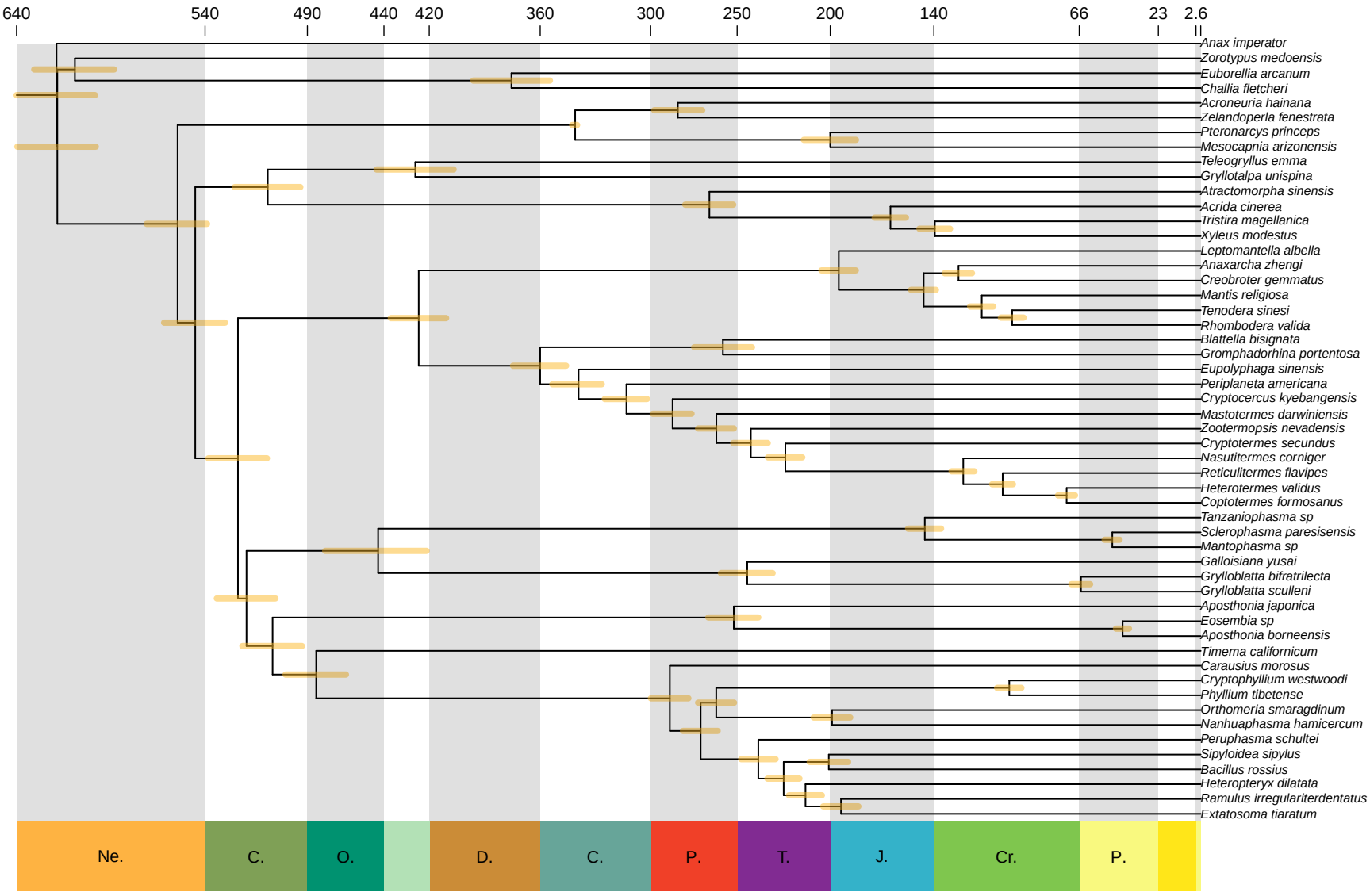
Mitochondrial phylogeny generated using MCMCtree with a strict clock model and the nucletide sequences including Embioptera and Zoraptera and removing crown Plecoptera calibration.



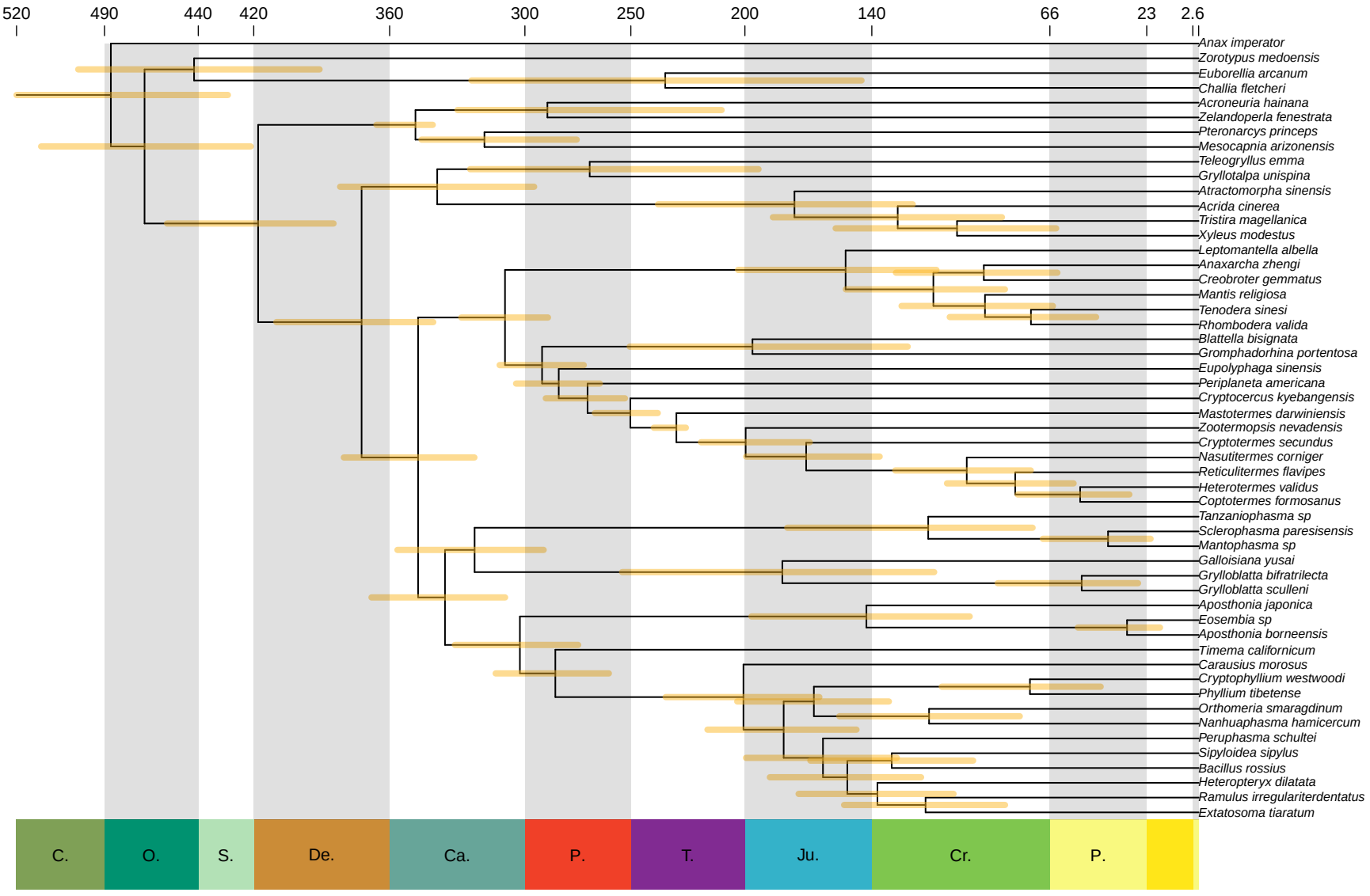
Mitochondrial phylogeny generated using MCMCtree with a indipendent rates clock model and the nucletide sequences including Embioptera and Zoraptera and removing crown Plecoptera calibration.



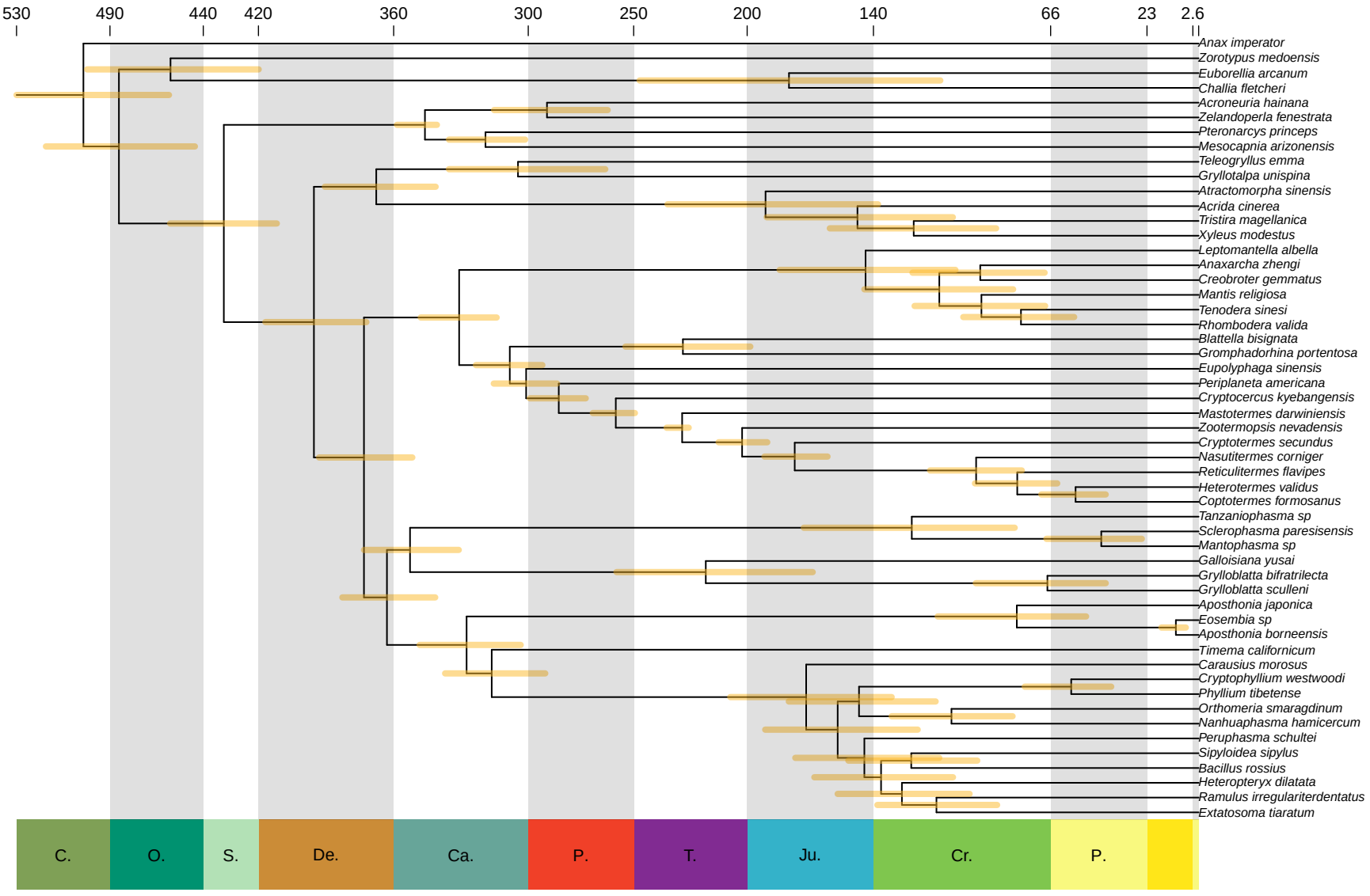
Mitochondrial phylogeny generated using MCMCtree with a correlated rates clock model and the nucletide sequences including Embioptera and Zoraptera and removing crown Plecoptera calibration.



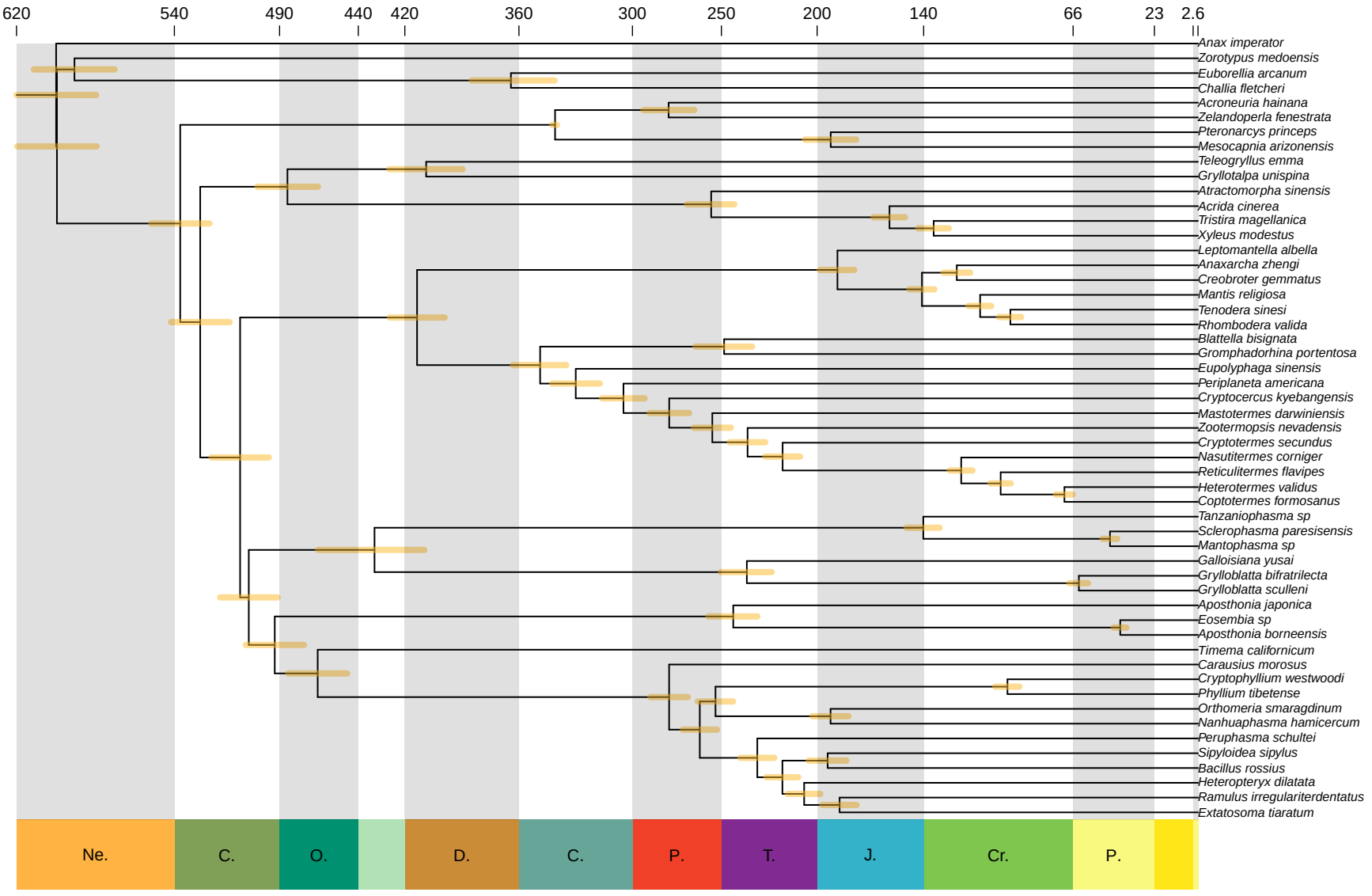
Mitochondrial phylogeny generated using MCMCtree with a strict clock model and the nucletide sequences including Embioptera and Zoraptera and removing crown Orthoptera calibration.



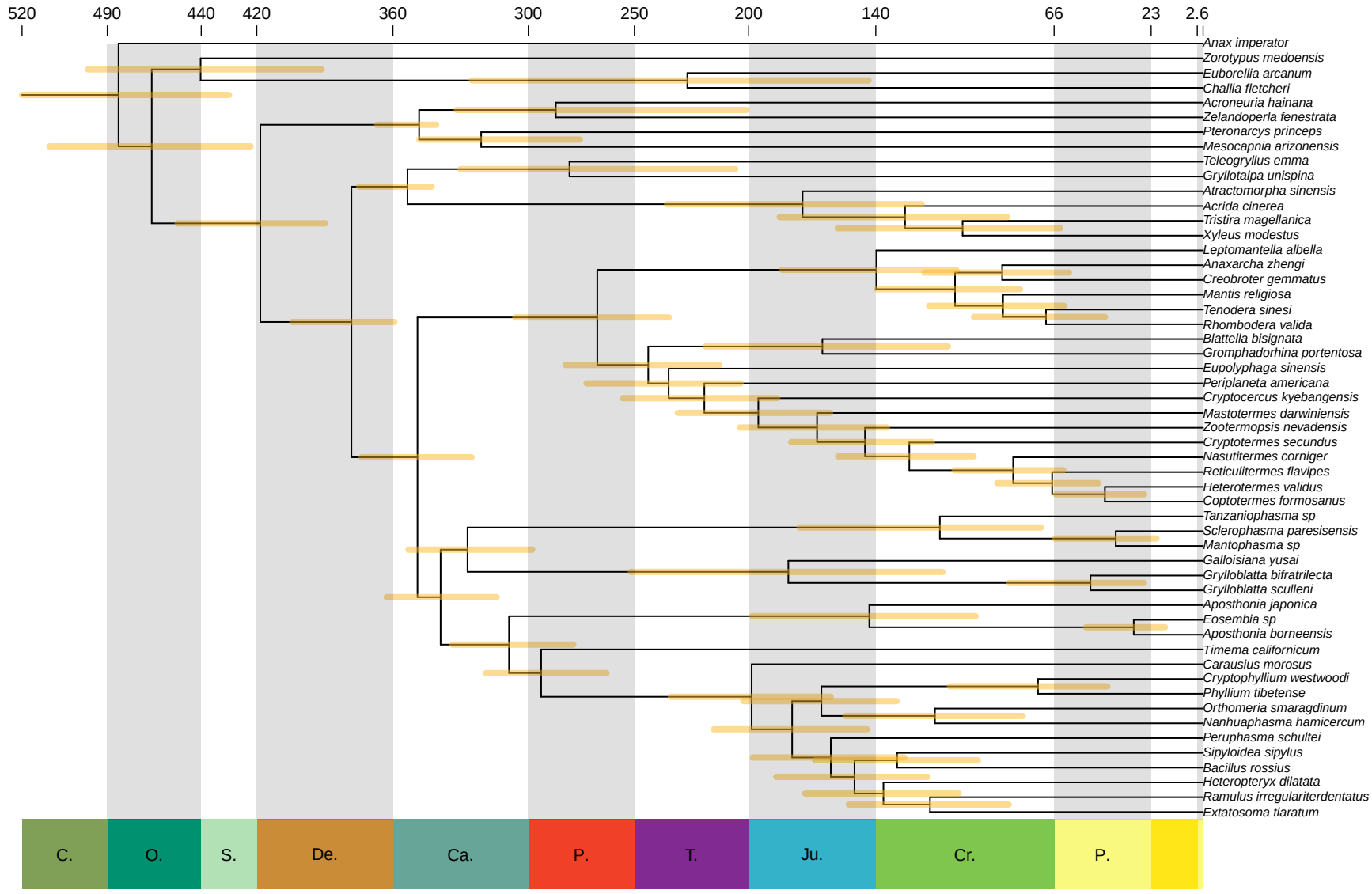
Mitochondrial phylogeny generated using MCMCtree with an independent rates clock model and the nucletide sequences including Embioptera and Zoraptera and removing crown Orthoptera calibration.



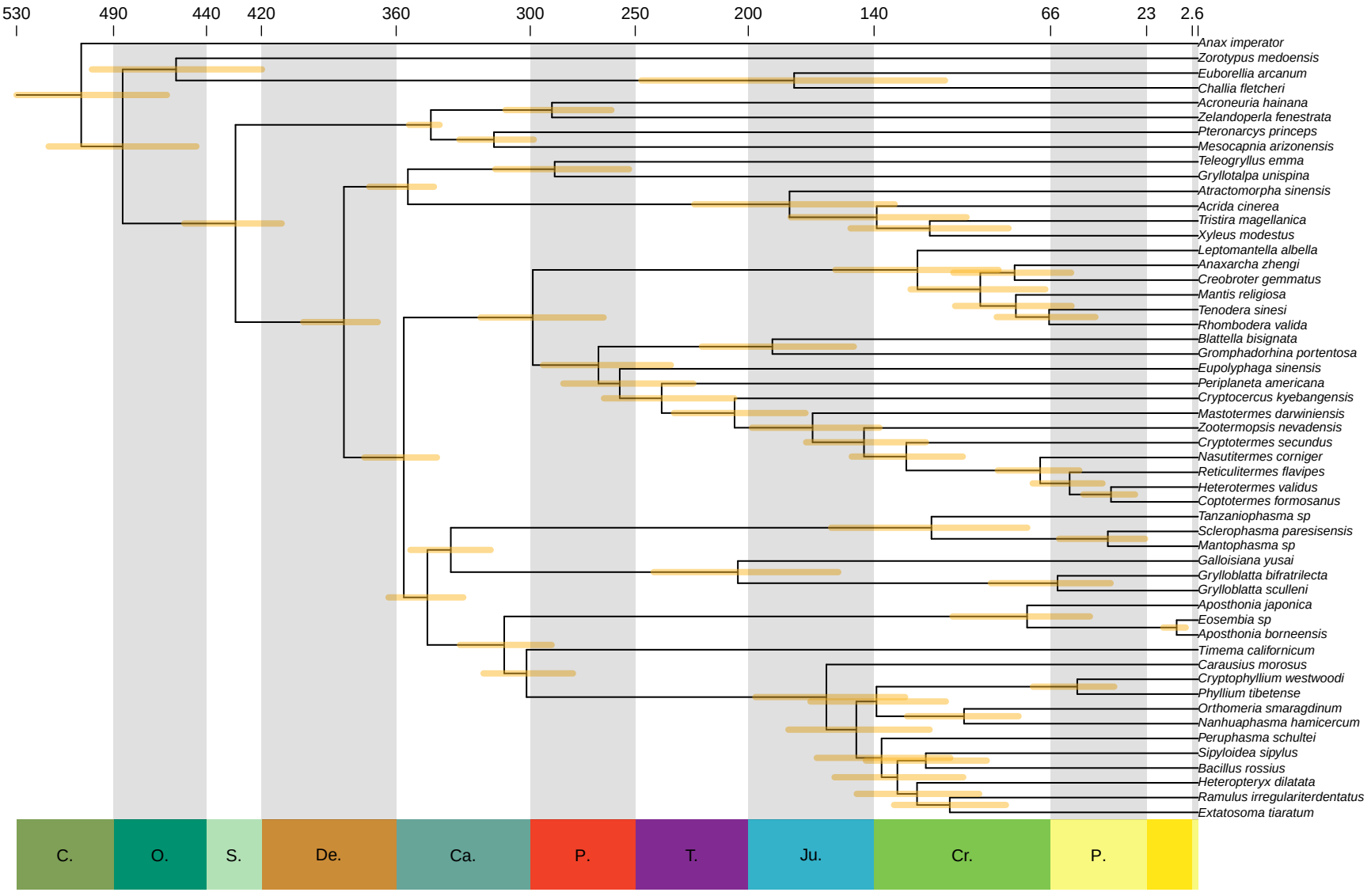
Mitochondrial phylogeny generated using MCMCtree with a correlated rates clock model and the nucletide sequences including Embioptera and Zoraptera and removing crown Orthoptera calibration.



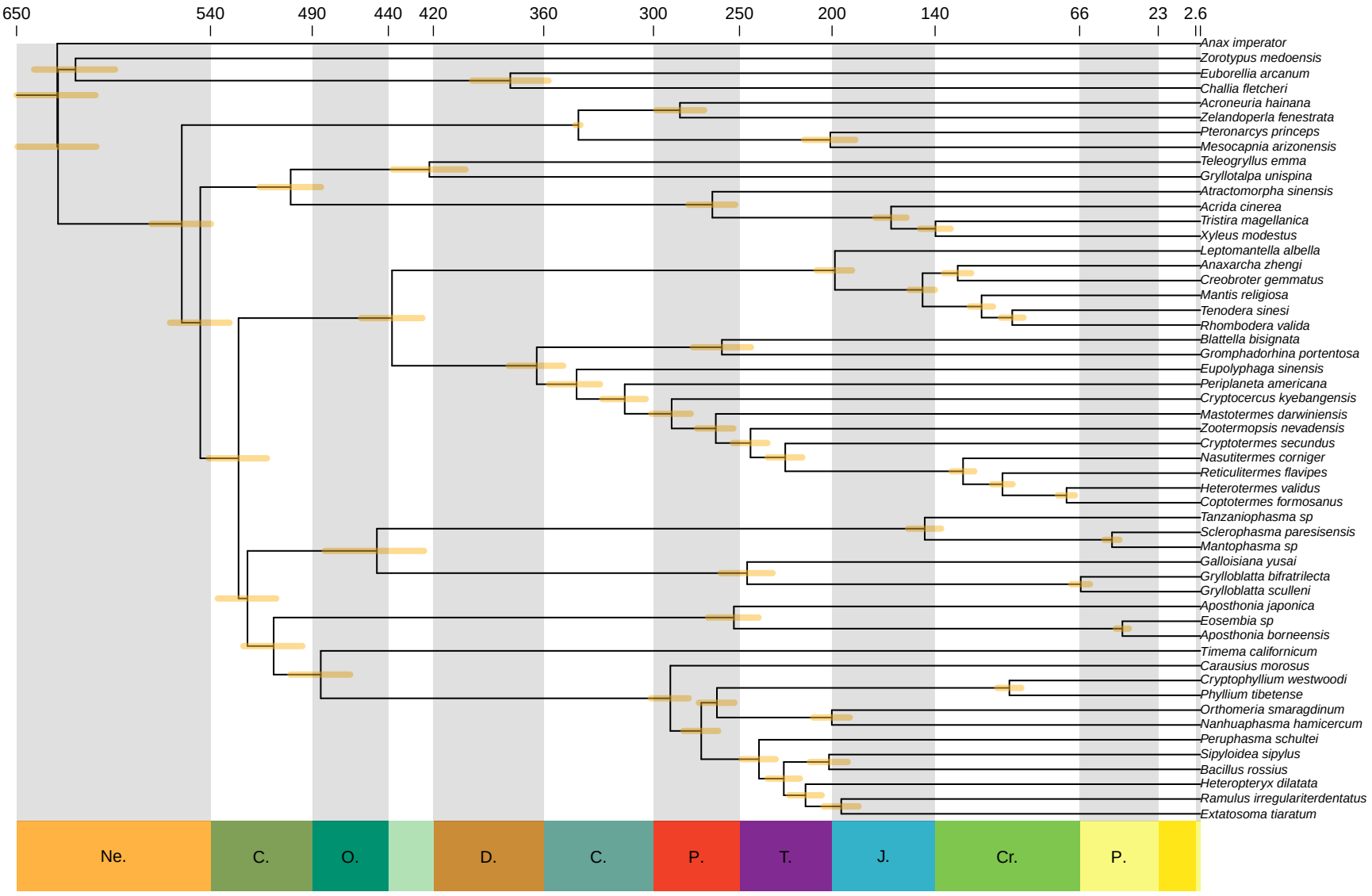
Mitochondrial phylogeny generated using MCMCtree with a strict clock model and the nucletide sequences including Embioptera and Zoraptera and removing crown Isoptera calibration.



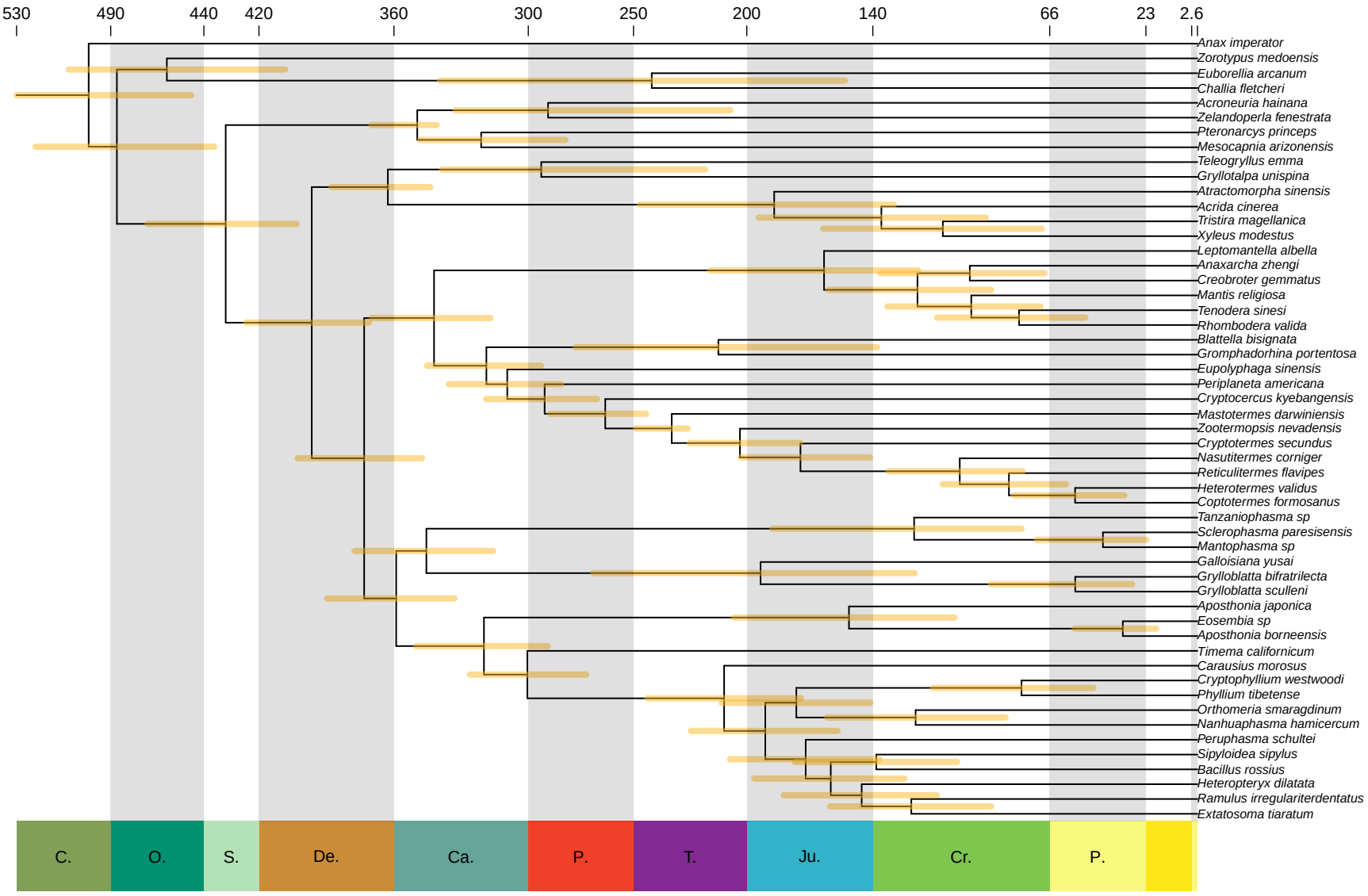
Mitochondrial phylogeny generated using MCMCtree with an independent rates clock model and the nucletide sequences including Embioptera and Zoraptera and removing crown Isoptera calibration.



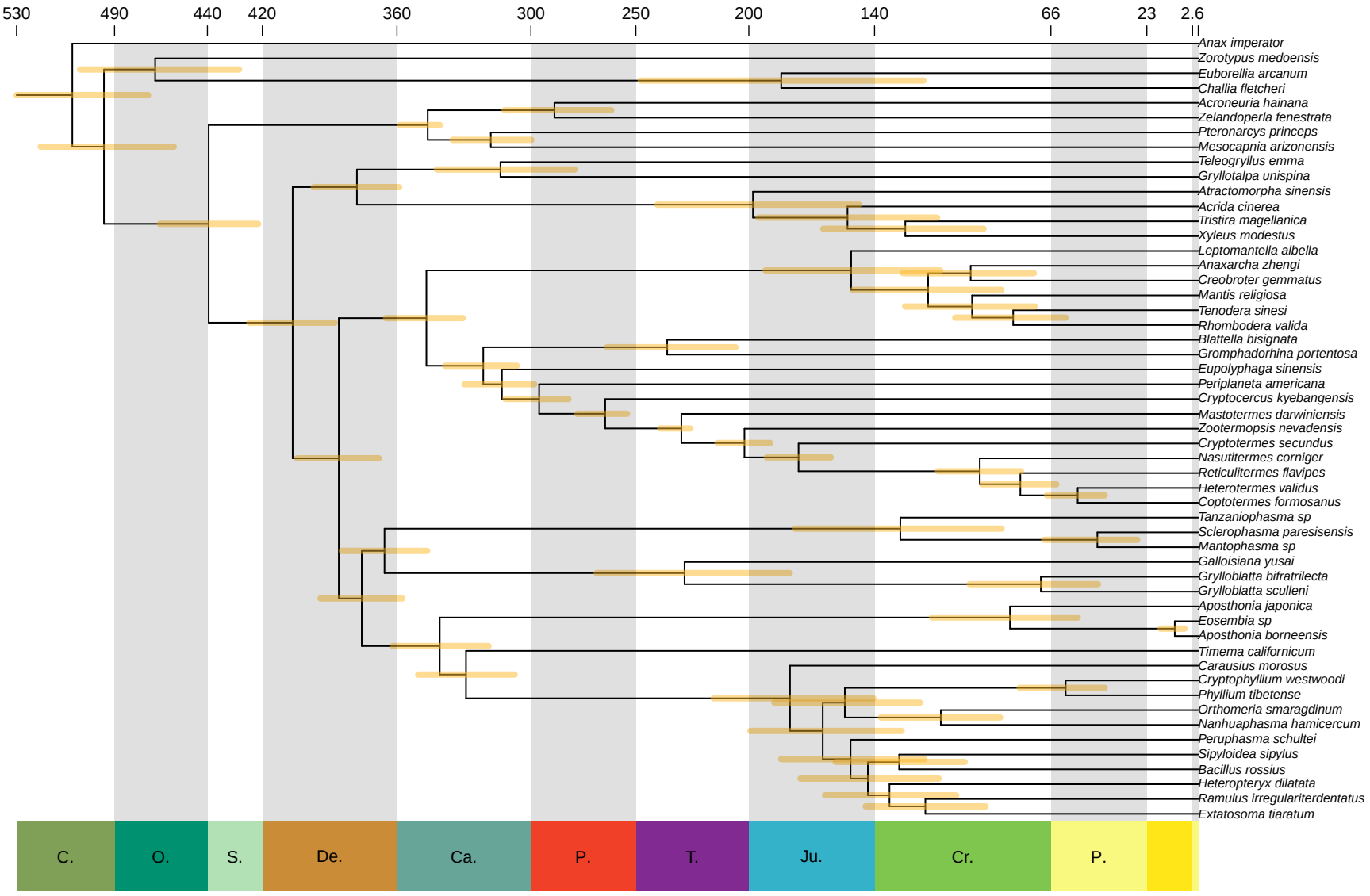
Mitochondrial phylogeny generated using MCMCtree with a correlated rates clock model and the nucletide sequences including Embioptera and Zoraptera and removing crown Isoptera calibration.



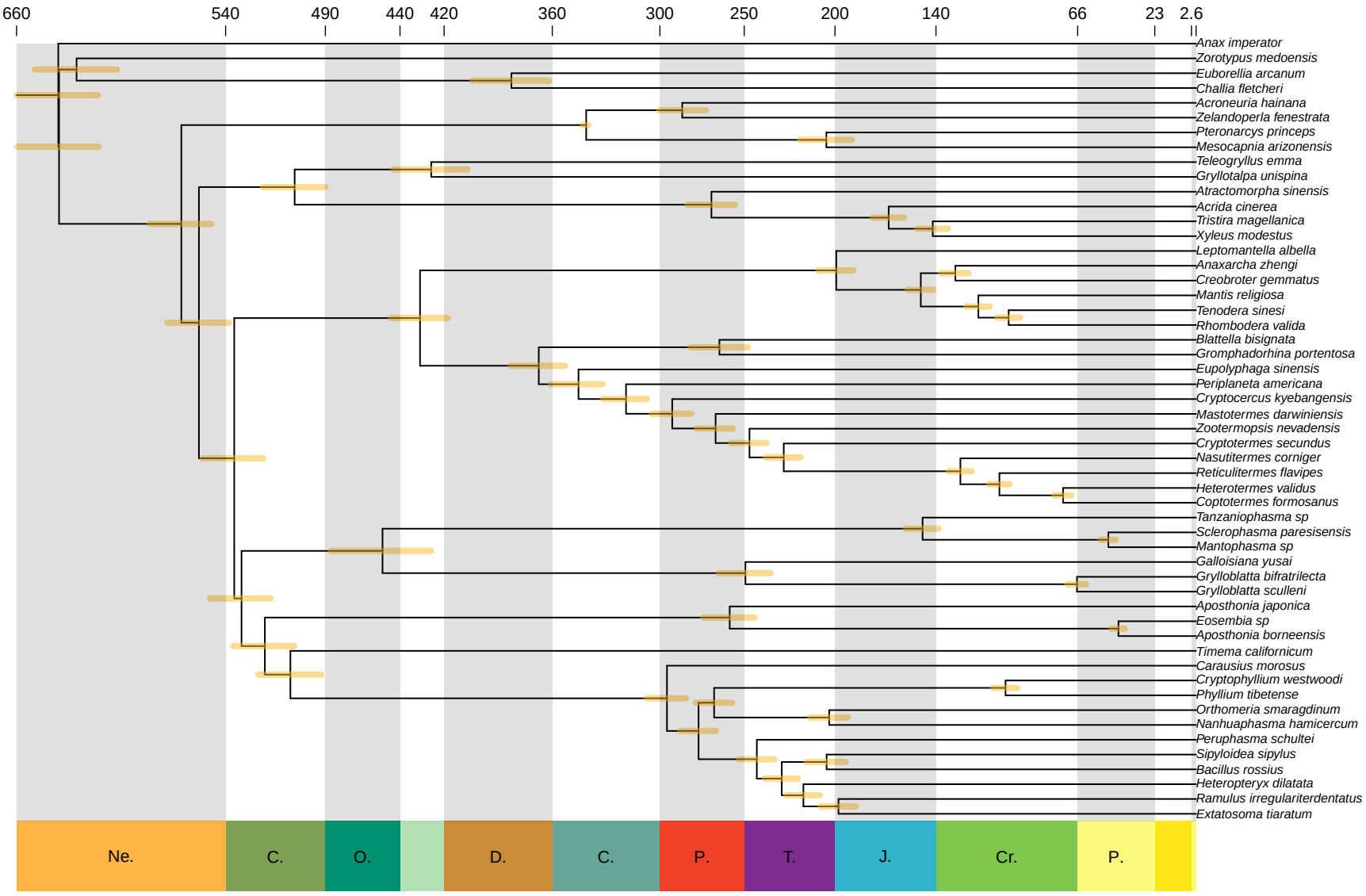
Mitochondrial phylogeny generated using MCMCtree with a strict clock model and the nucletide sequences including Embioptera and Zoraptera and removing crown Dictyoptera calibration.



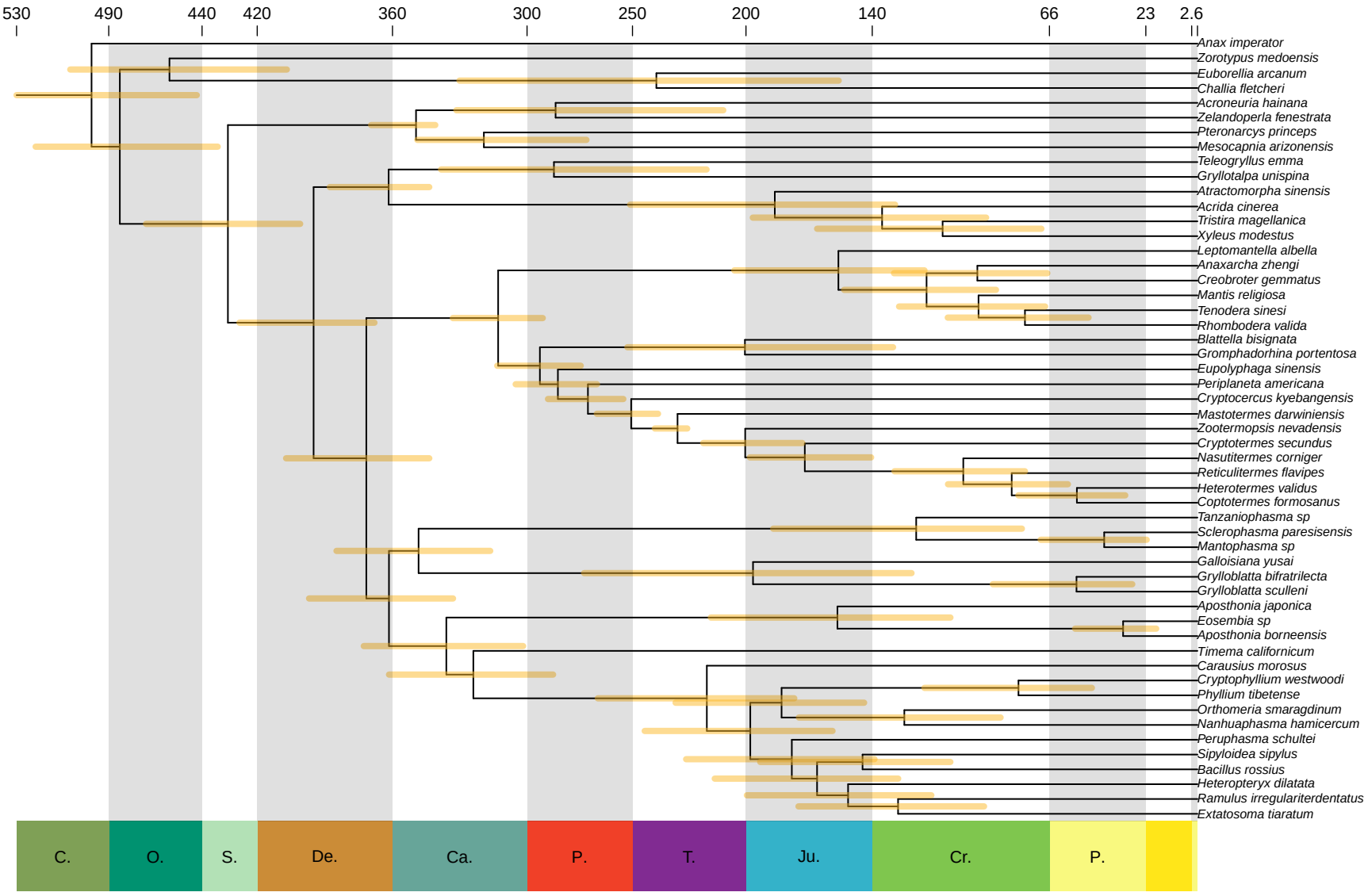
Mitochondrial phylogeny generated using MCMCtree with an independent rates clock model and the nucleotide sequences including Embioptera and Zoraptera and removing crown Dictyoptera calibration.



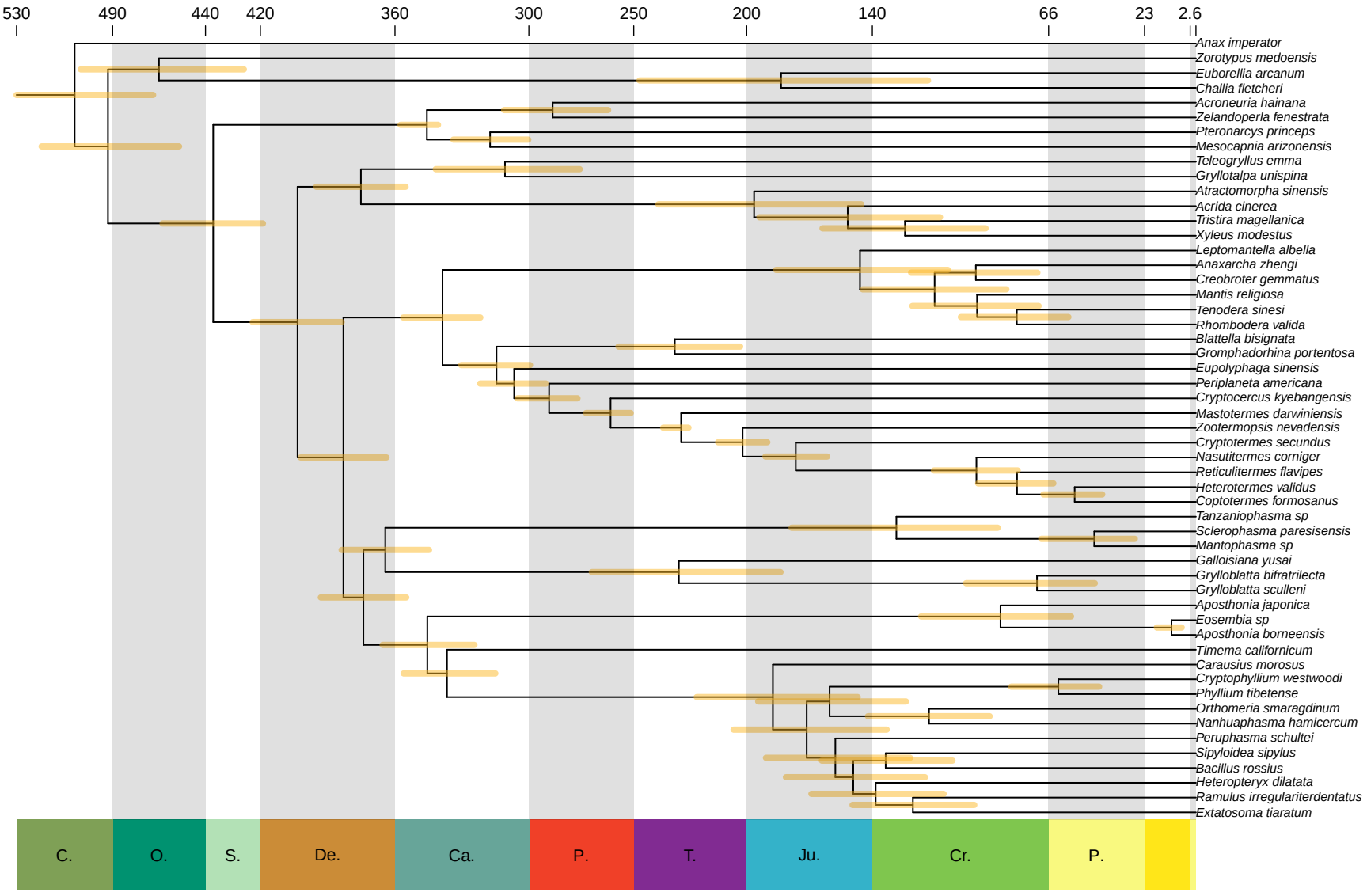
Mitochondrial phylogeny generated using MCMCtree with a correlated rates clock model and the nucleotide sequences including Embioptera and Zoraptera and removing crown Dictyoptera calibration.



Mitochondrial phylogeny generated using MCMCtree with a strict clock model and the nucletide sequences including Embioptera and Zoraptera and removing crown Phasmatodea calibration.



Mitochondrial phylogeny generated using MCMCtree with an independent rates model and the nucletide sequences including Embioptera and Zoraptera and removing crown Phasmatodea calibration.



Mitochondrial phylogeny generated using MCMCtree with a correlated rates clock model and the nucletide sequences including Embioptera and Zoraptera and removing crown Phasmatodea calibration.