

Materials List for:

Using Phylogenetic Analysis to Investigate Eukaryotic Gene Origin

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Materials

| Name | Company | Catalog Number | Comments |
|--------------------|---------|----------------|--|
| Adobe Illustration | | | a graphical tool developed by Adobe Systems Software Ireland Ltd. Copyright © 2017 |
| BioEdit | | | a biological sequence alignment editor written for Windows 95/98/NT/2000/XP/7. Copyright © Tom Hall |
| Clustal Omega | | | a package for making multiple sequence alignments of amino acid or nucleotide sequences. http://www.clustal.org/ |
| CorelDRAW | | | a graphic design software. Copyright © 2017 Corel Corporation |
| FigTree | | | a graphical viewer of phylogenetic trees designed by the University of Edinburgh |
| MEGA | | | MolecularEvolutionary Genetics Analysis version6.0 http://www.megasoftware.net/home |
| MrBayes | | | an Bayesian phylogenetic inference tool |
| NVIDIA | | | a company designs graphics processing units (GPUs) for the gaming and professional markets. Corporation Copyright © 2017 |
| PAUP | | | Phylogenetic Analysis Using Parsimony. David Swofford's program implements the maximum likelihood method under a number of nucleotide models. |
| Photoshop | | | a raster graphics editor developed and published by Adobe Systems Software Ireland Ltd. Copyright © 2017 |
| RHYTHM | | | a knowledge based prediction of heix contacts. Charité Berlin – Protein Formatics Group - Copyright 2007-2009 |
| TMHMM | | | a tool for prediction of transmembrane helices in proteins. |

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| | | | http://www.cbs.dtu.dk/services/TMHMM/ |
| Compter | | | 4 GB memory, Core 2 or above CPU. Windows 7, Windows 10 |