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seqfile = seqfile.txt          * sequence data filename
outfile = results_0.001.txt  * main result file name [CHANGE THIS]

    noisy = 9          * 0,1,2,3,9: how much rubbish on the screen
verbose = 1          * 1:detailed output
runmode = -2         * -2:pairwise

seqtype = 1         * 1:codons
CodonFreq = 3       * 0:equal, 1:F1X4, 2:F3X4, 3:F61
    model = 0        *
NSsites = 0         *
    icode = 0        * 0:universal code

fix_kappa = 0       * 1:kappa fixed, 0:kappa to be estimated
    kappa = 2        * initial or fixed kappa

fix_omega = 1      * 1:omega fixed, 0:omega to be estimated
    omega = 0.001  * 1st fixed omega value [CHANGE THIS]

*NOTES: alternate fixed omega values
*omega = 0.005    * 2nd fixed value
*omega = 0.01     * 3rd fixed value
*omega = 0.05     * 4th fixed value
*omega = 0.10     * 5th fixed value
*omega = 0.20     * 6th fixed value
*omega = 0.40     * 7th fixed value
*omega = 0.80     * 8th fixed value
*omega = 1.60     * 9th fixed value
*omega = 2.00     * 10th fixed value

```