predictors

- log(scholarRefs) + NumberOfSubspecies + log(distrSize) + rand
- log(scholarRefs) + NumberOfSubspecies + log(mass) + log(distrSize) + rand
- log(scholarRefs) + NumberOfSubspecies + log(mass) + rand
- log(scholarRefs) + NumberOfSubspecies + rand
- NumberOfSubspecies + log(distrSize) + rand
- NumberOfSubspecies + log(mass) + log(distrSize) + rand
- NumberOfSubspecies + log(mass) + rand
- NumberOfSubspecies + rand

0.02

mbda