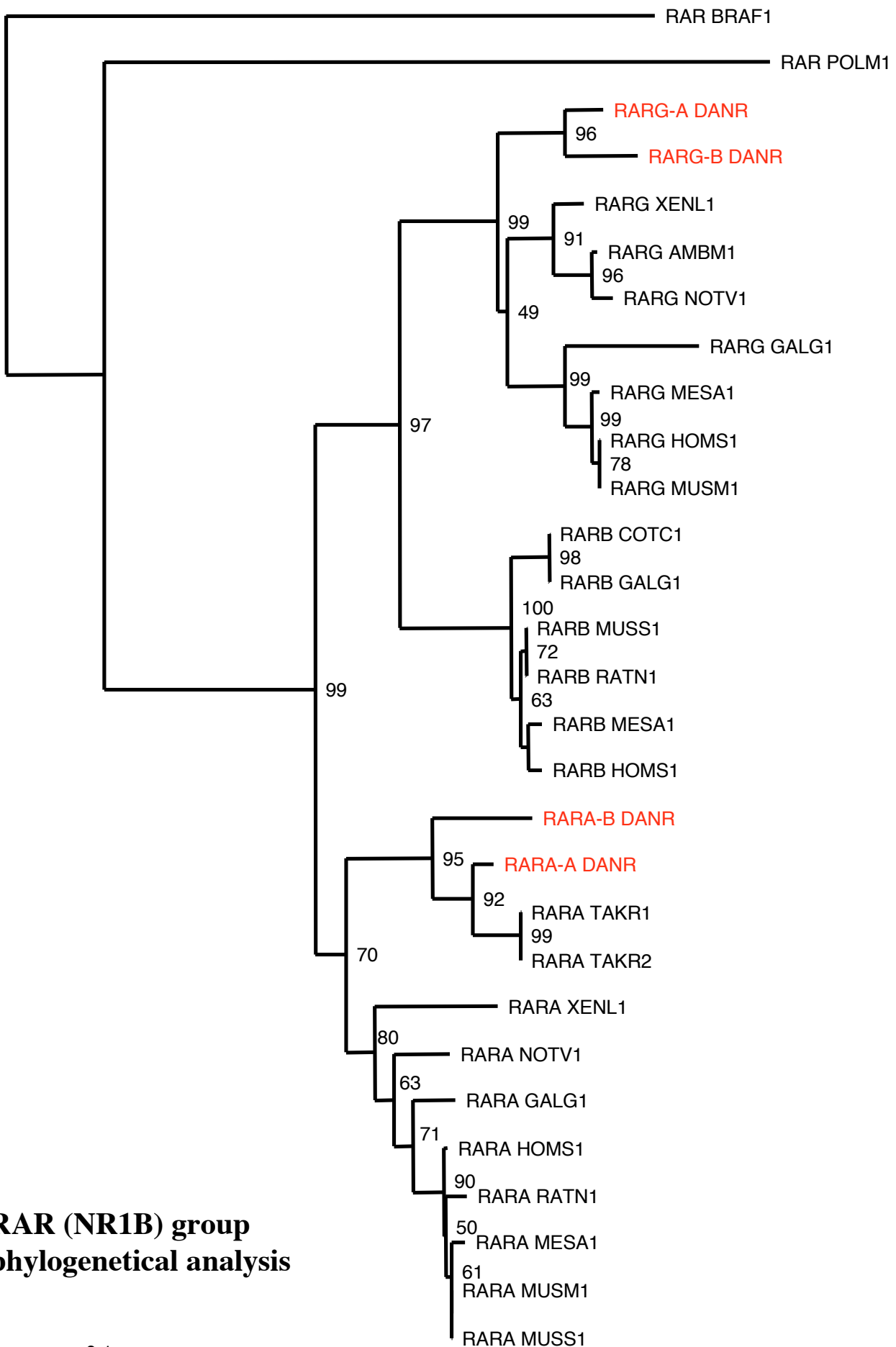


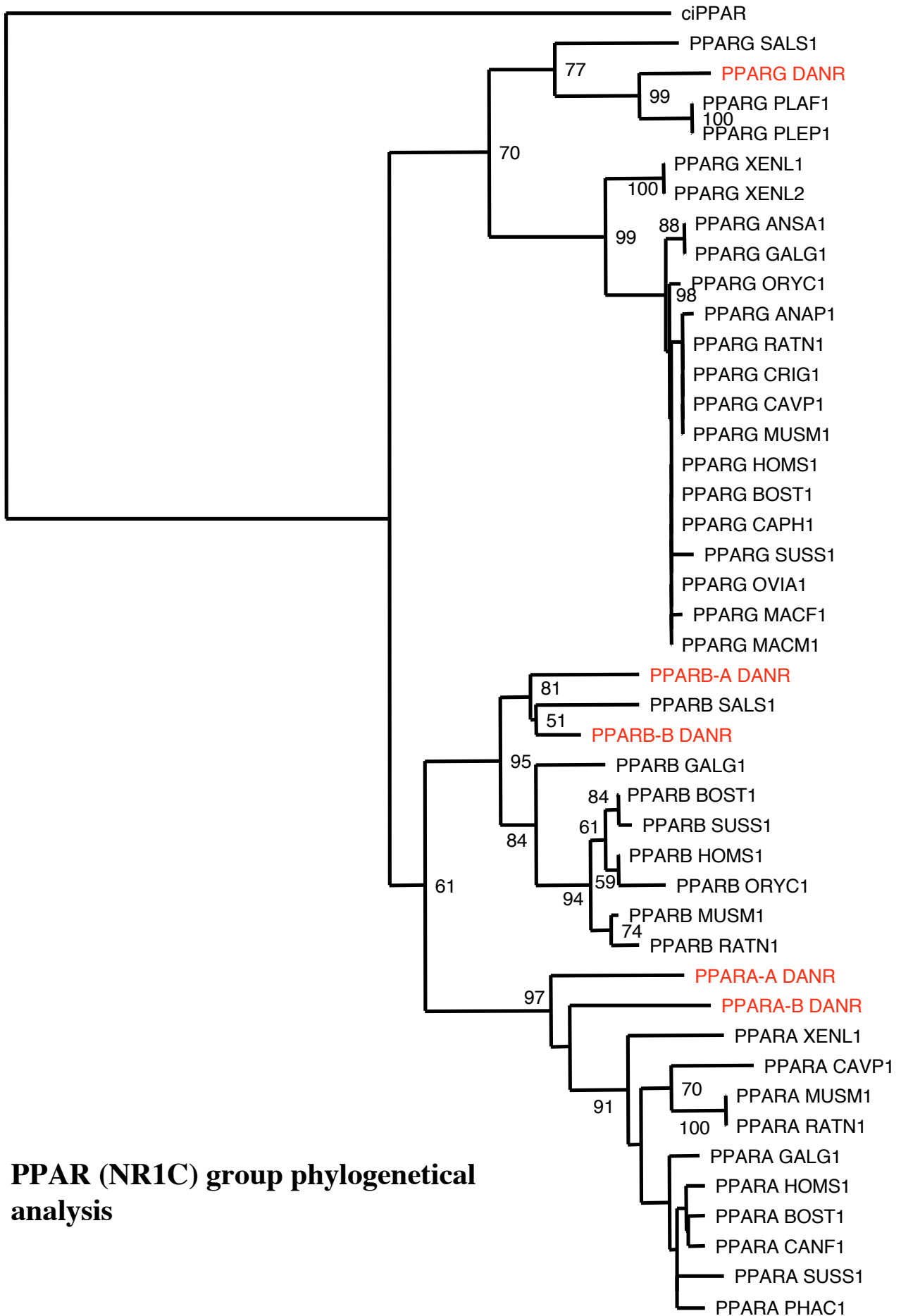
TR (NR1A) group phylogenetical analysis

0.1



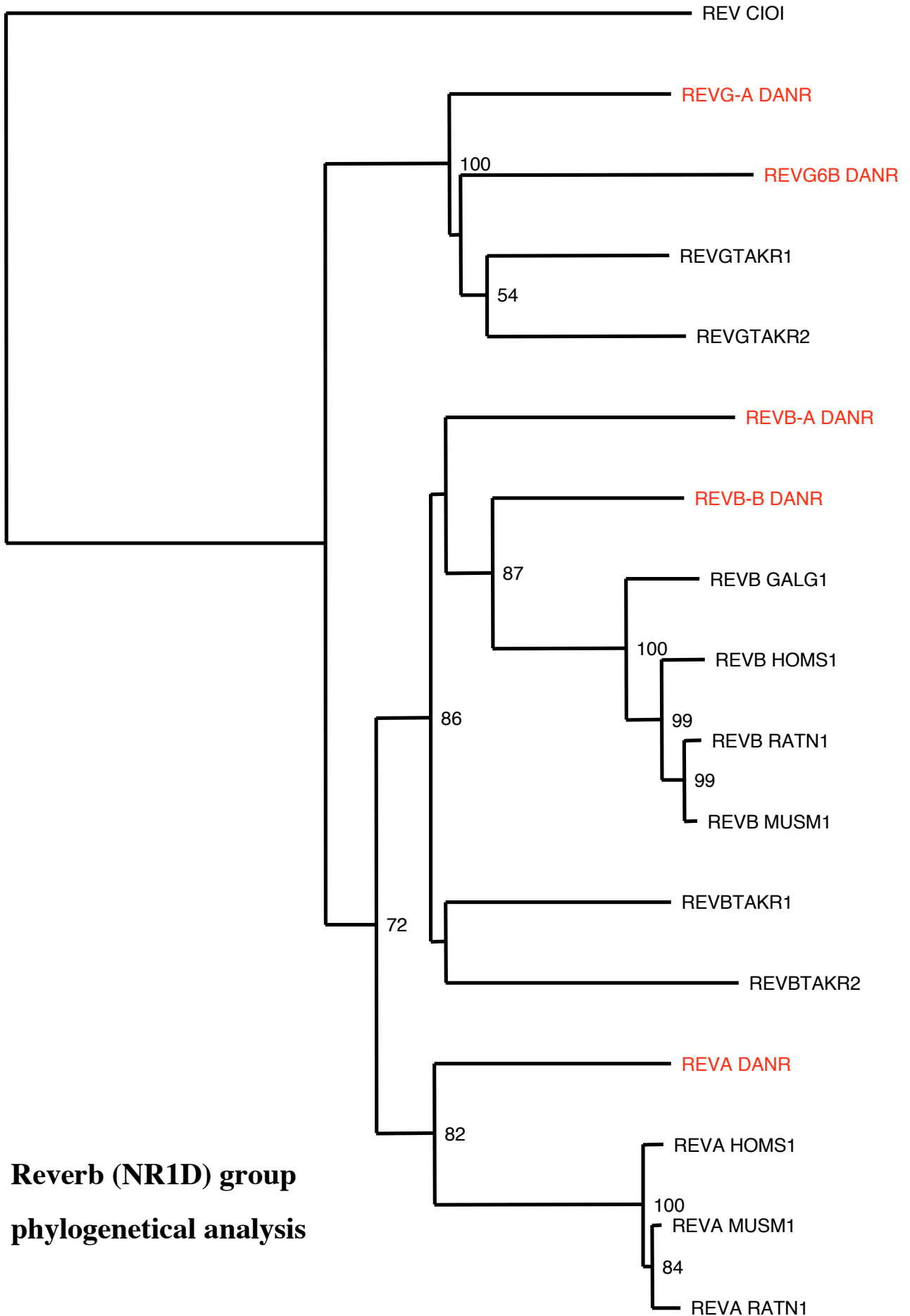
**RAR (NR1B) group
phylogenetical analysis**

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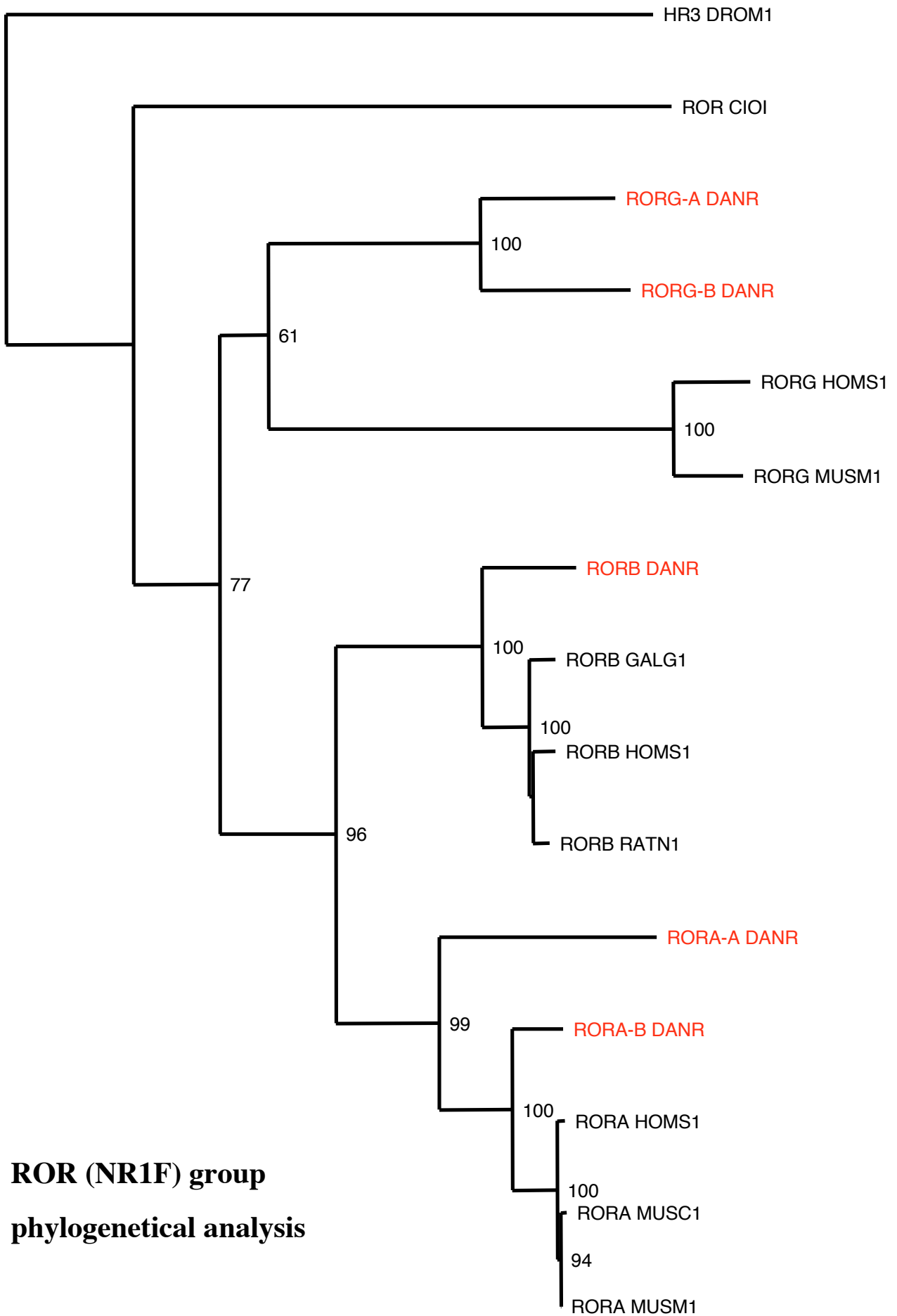
PPAR (NR1C) group phylogenetical analysis

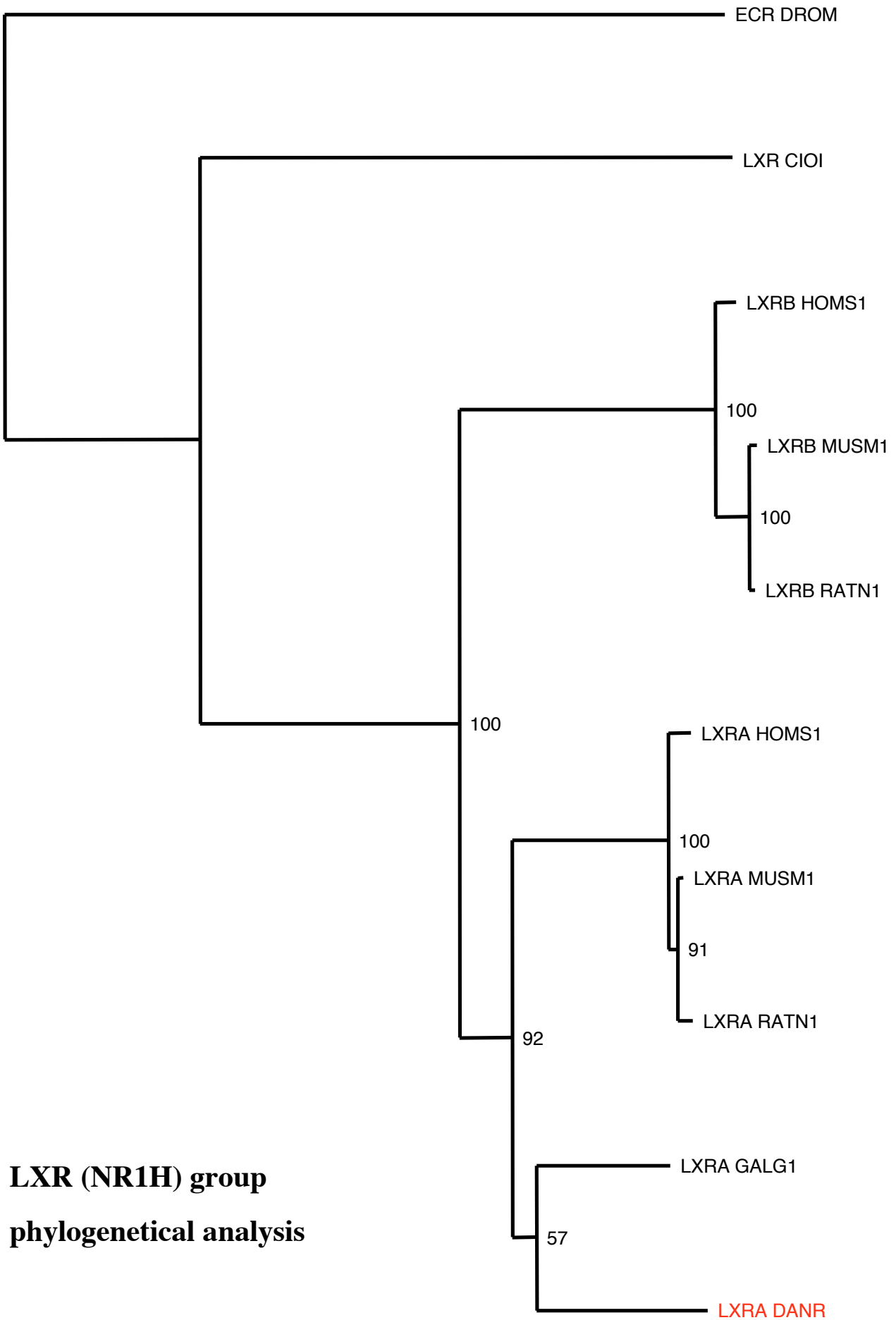
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**Reverb (NR1D) group
phylogenetical analysis**

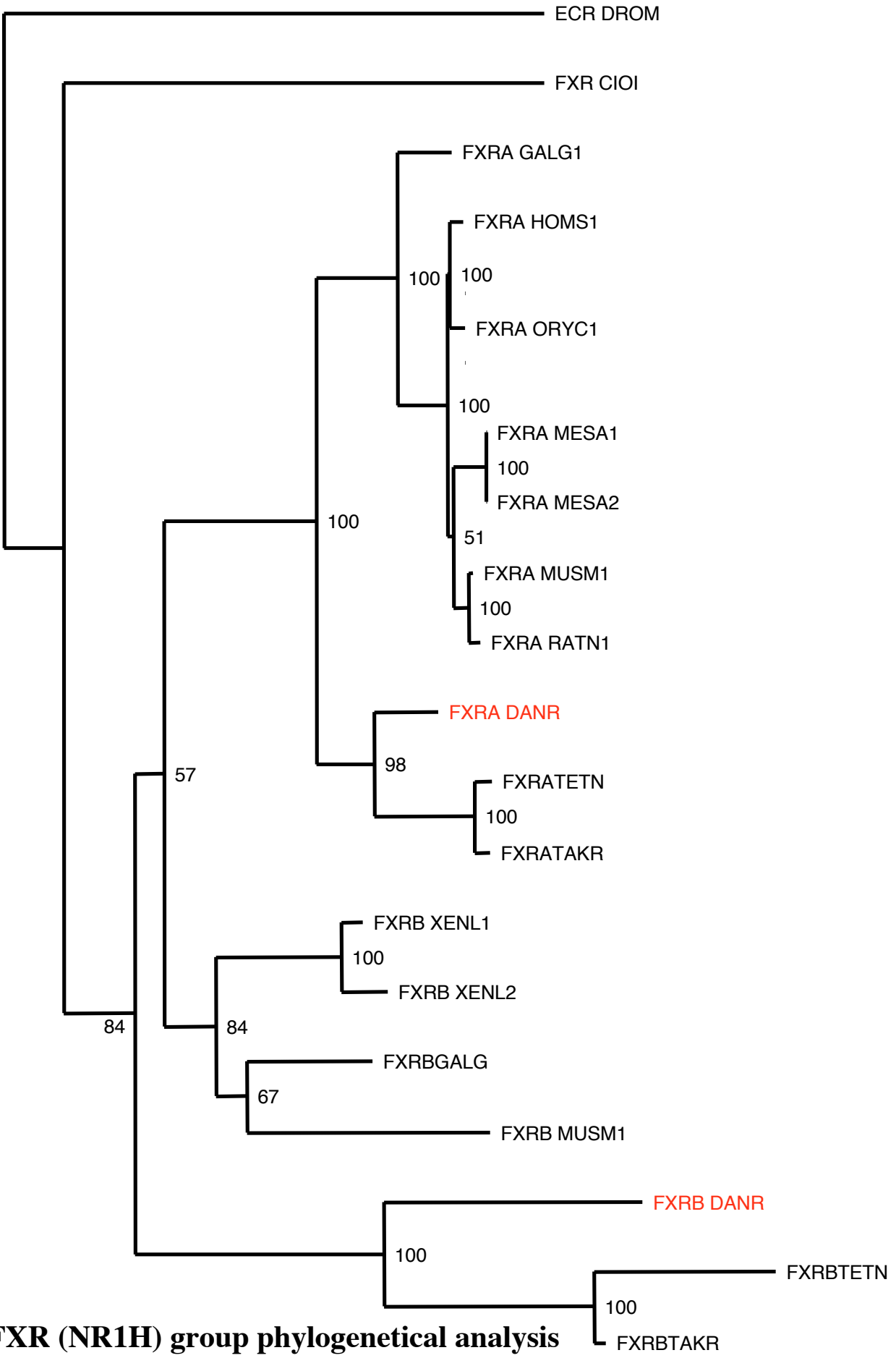
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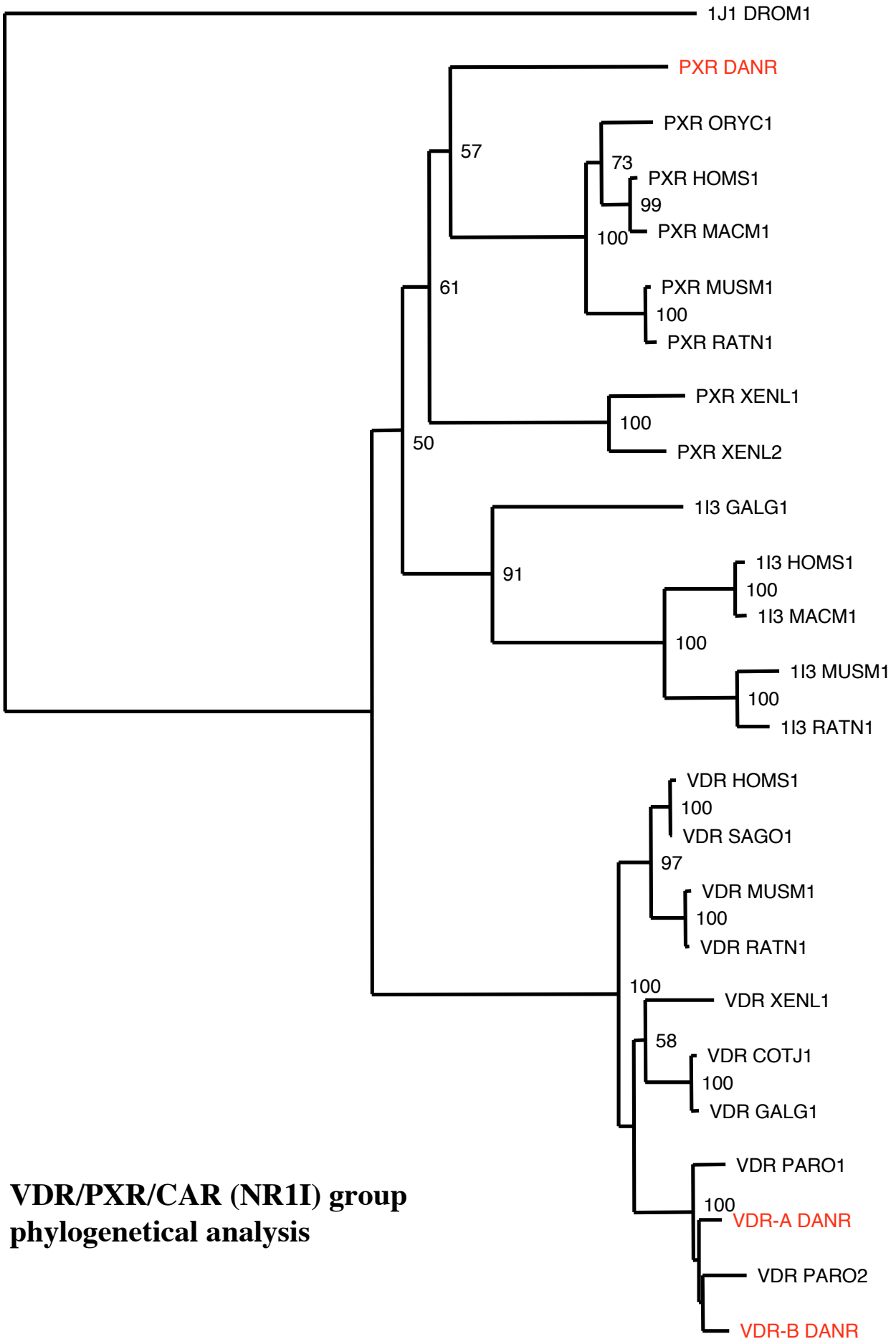
**LXR (NR1H) group
phylogenetical analysis**

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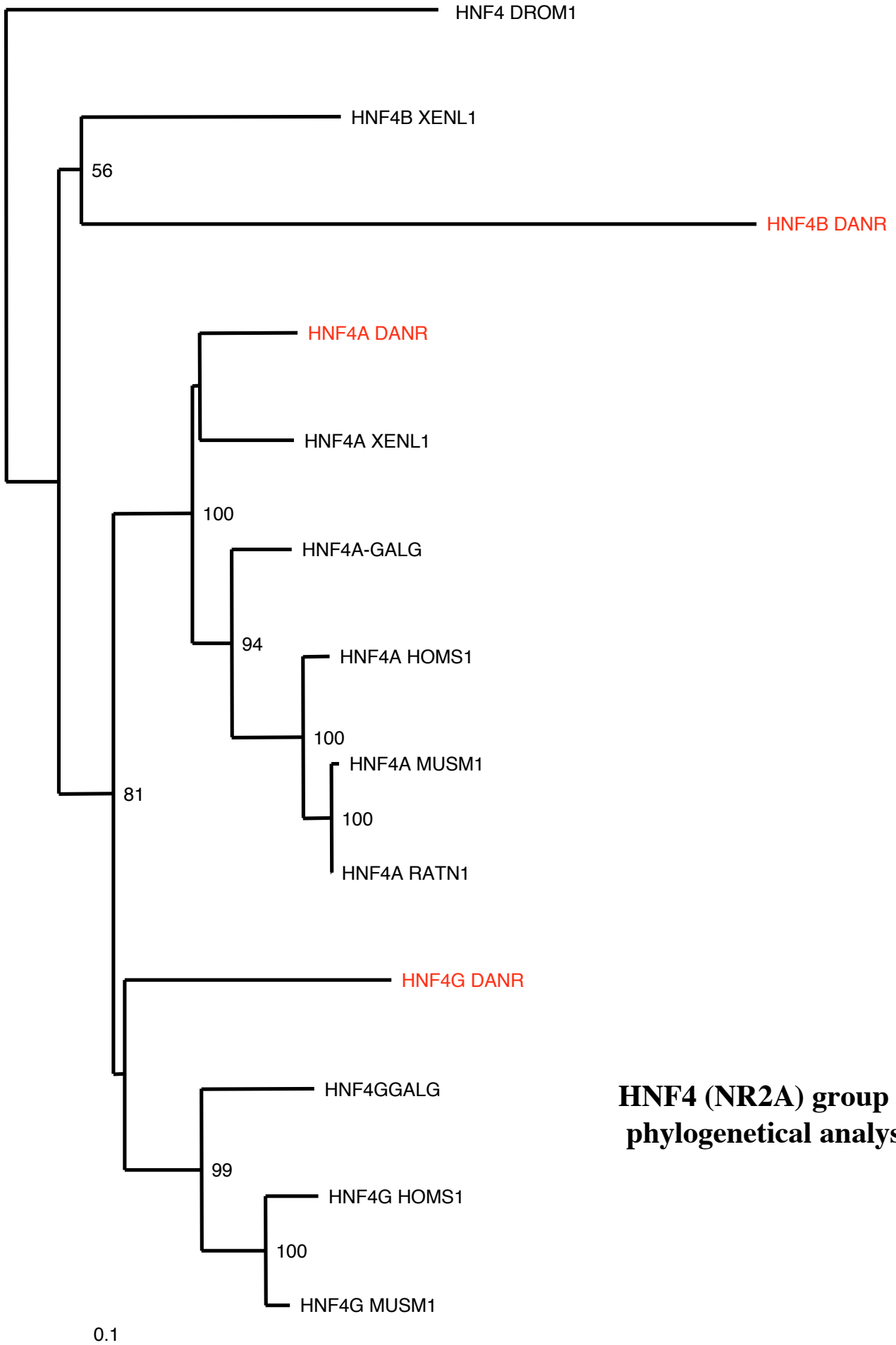


FXR (NR1H) group phylogenetical analysis

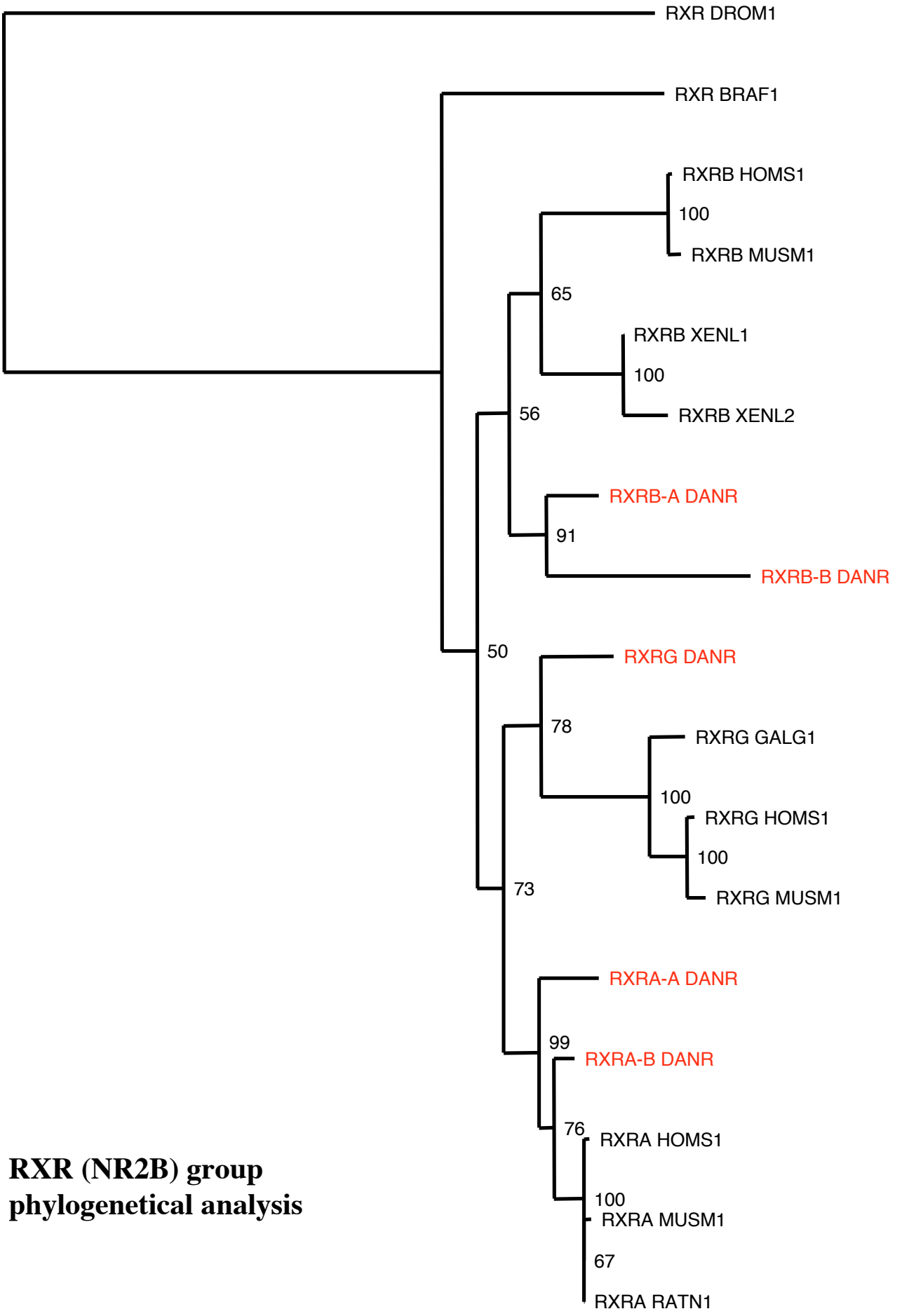
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**VDR/PXR/CAR (NR1) group
phylogenetical analysis**

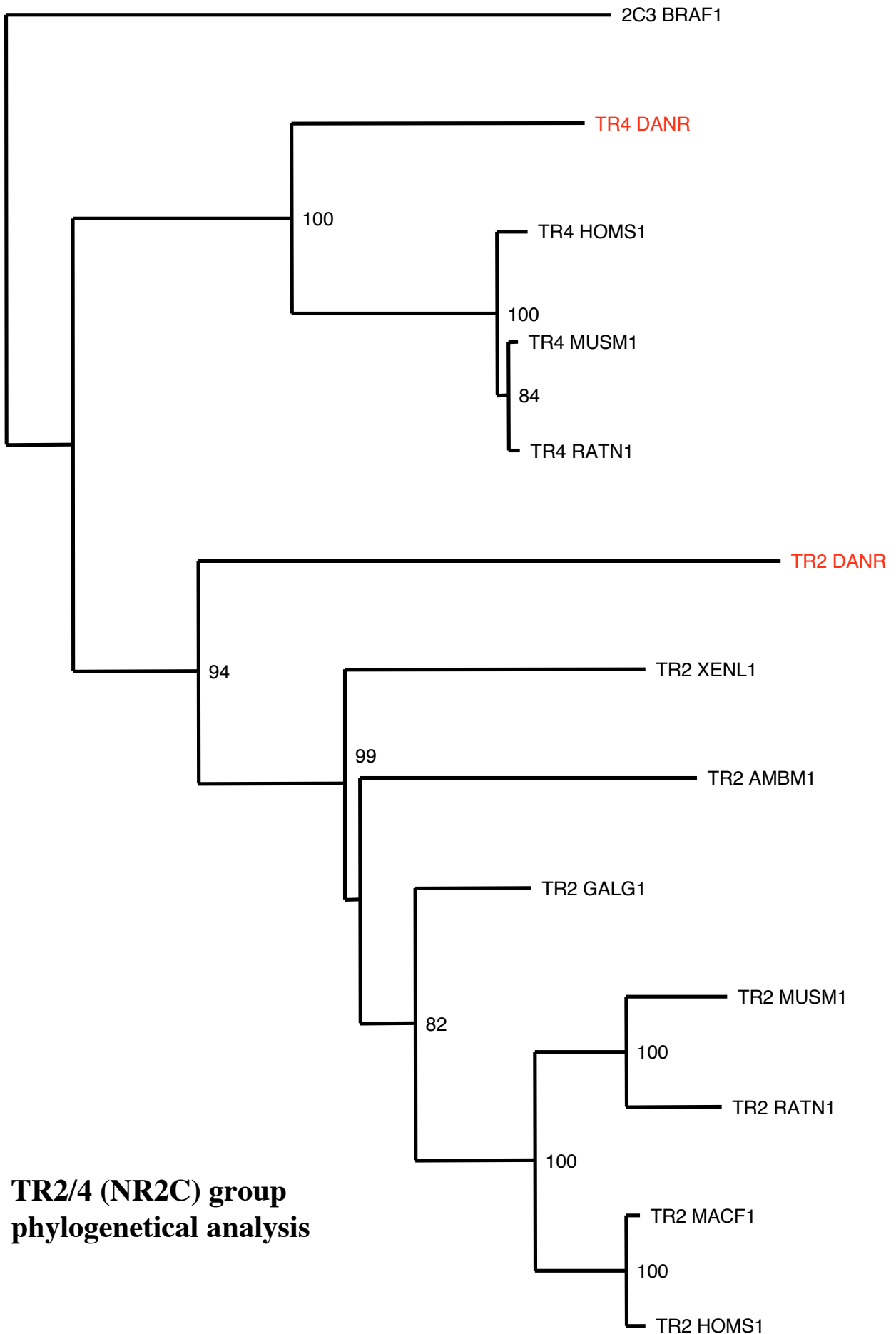


**HNF4 (NR2A) group
phylogenetical analysis**



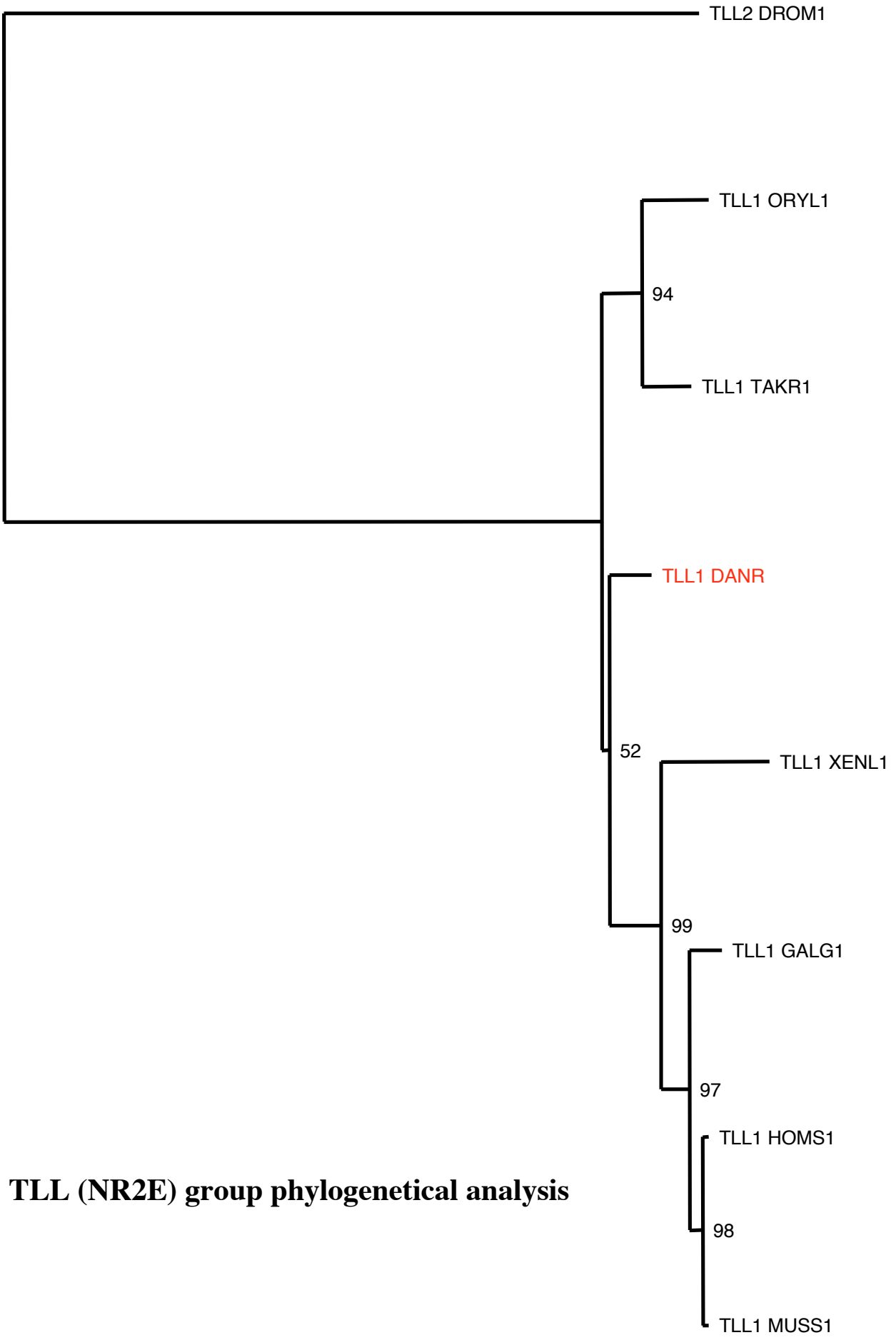
**RXR (NR2B) group
phylogenetical analysis**

0.1



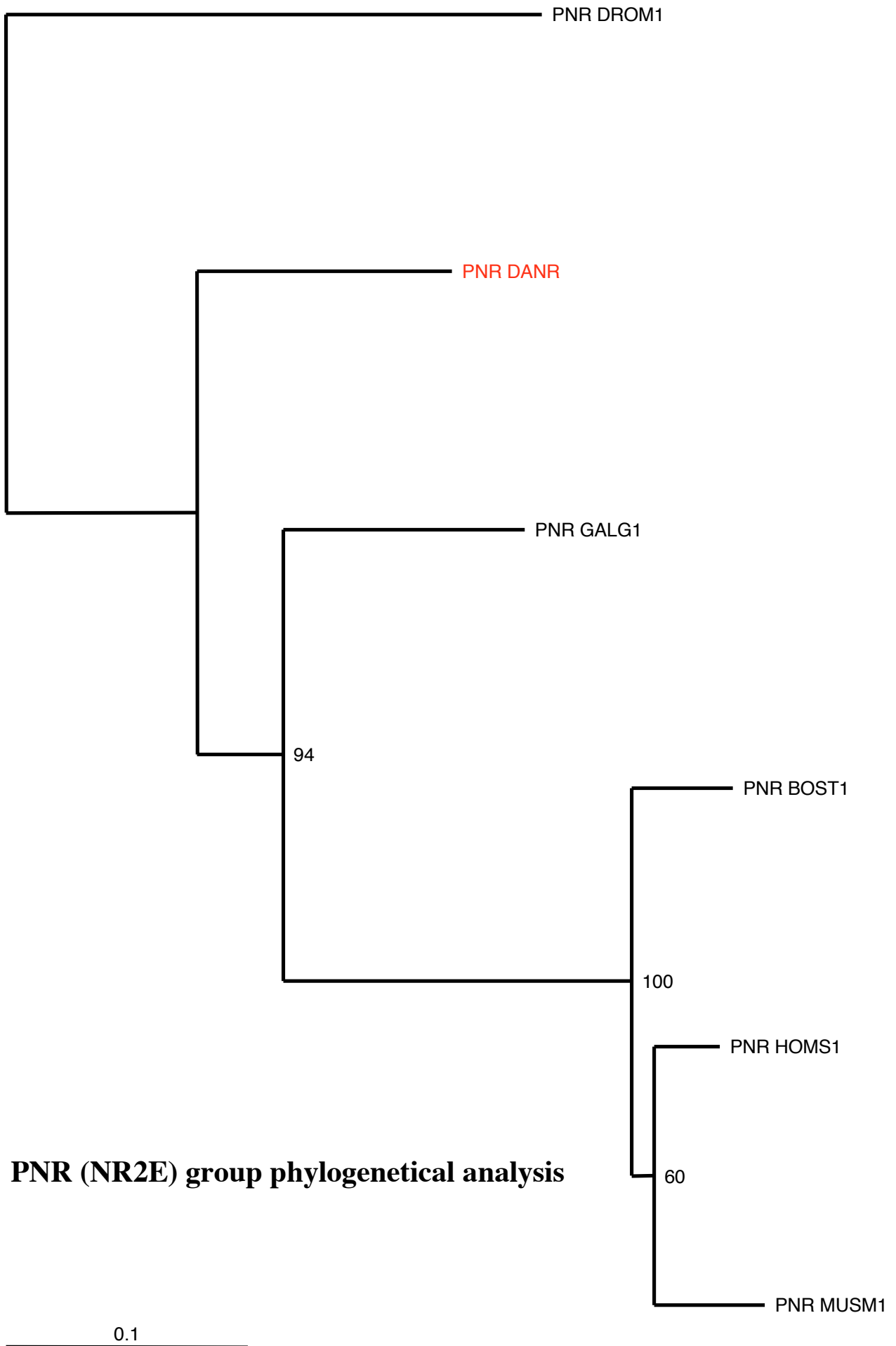
**TR2/4 (NR2C) group
phylogenetical analysis**

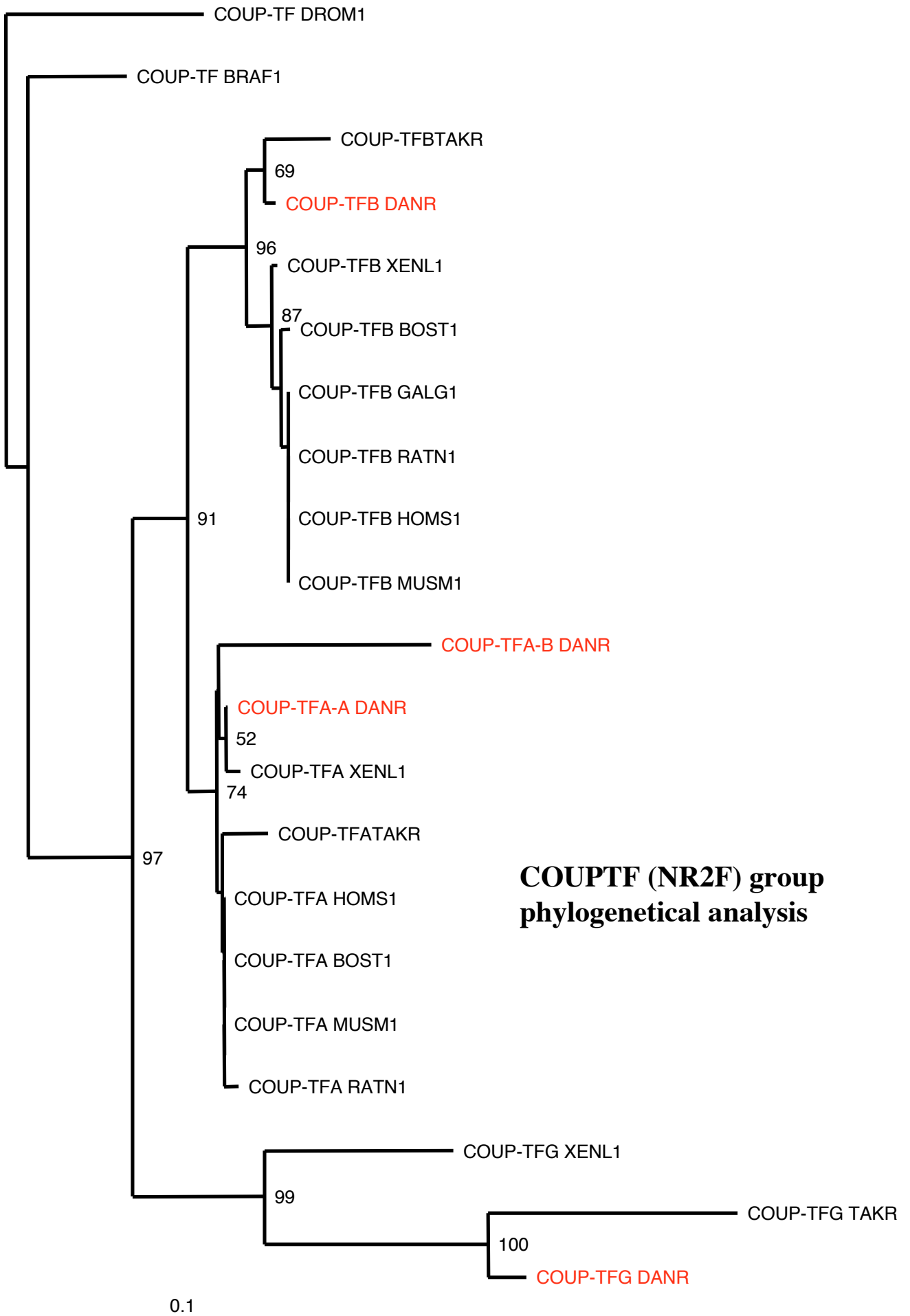
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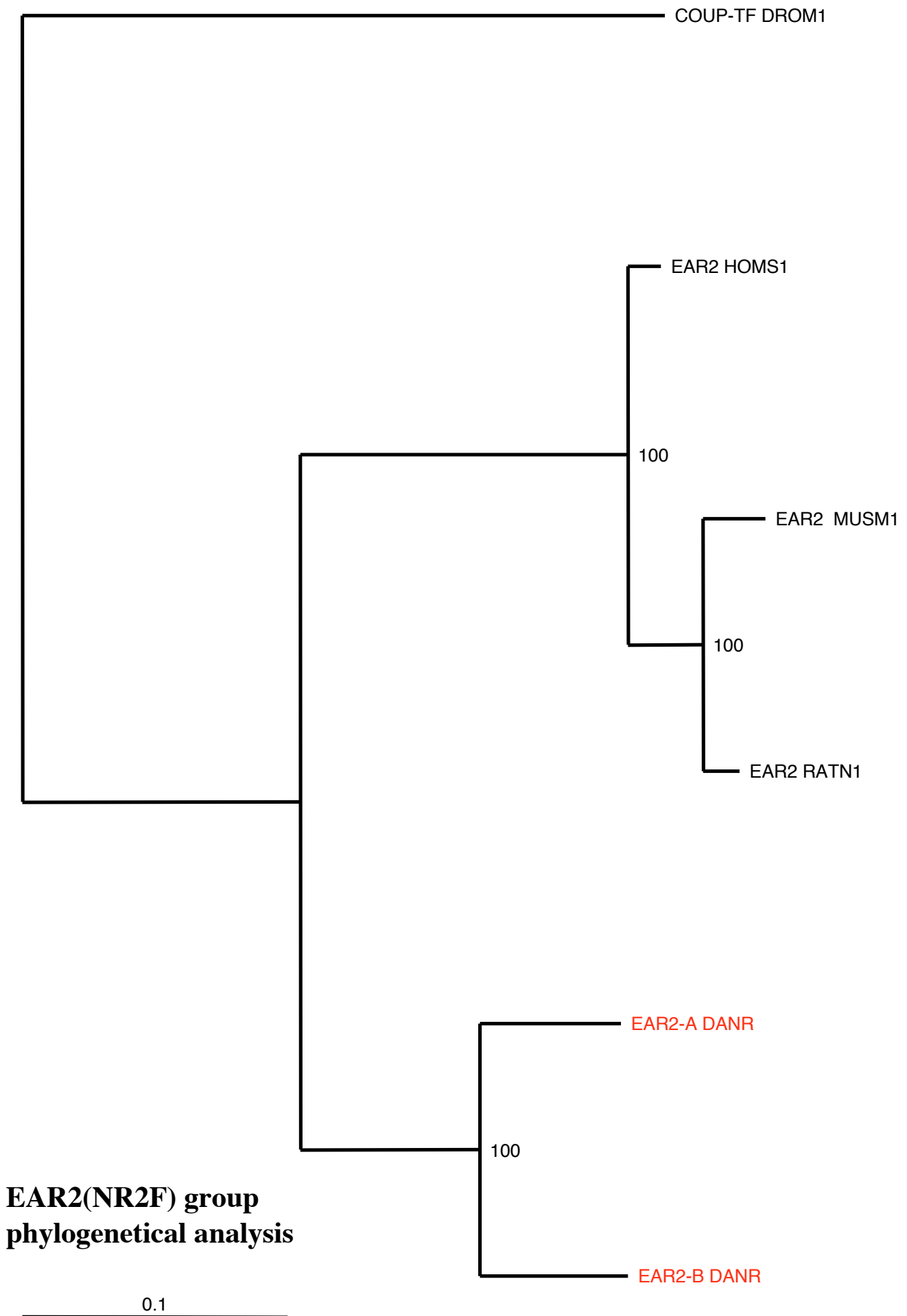


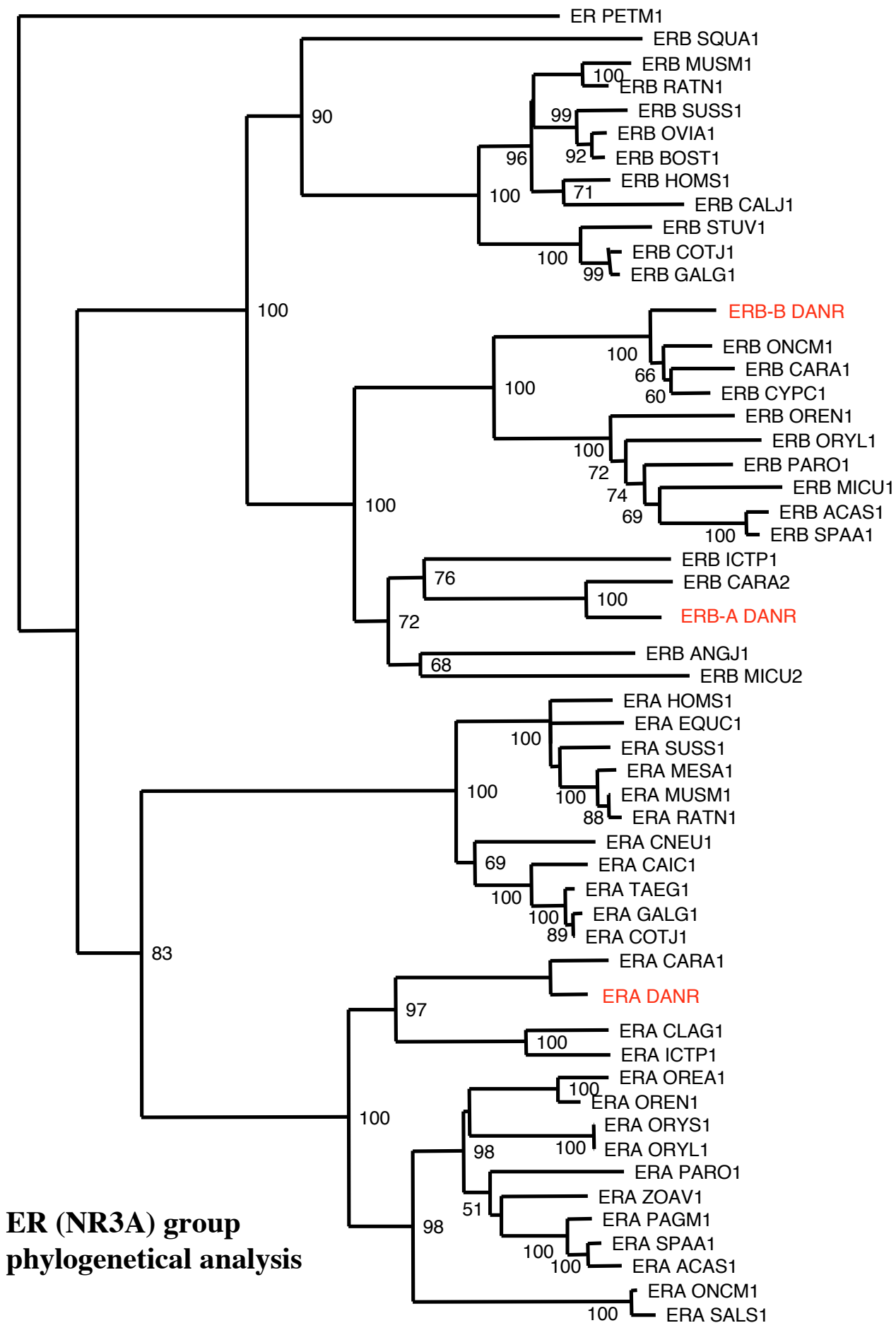
TLL (NR2E) group phylogenetical analysis

0.1









ER PETM1

ERB SQUA1

ERB MUSM1

100

ERB RATN1

99

ERB SUSS1

ERB OVIA1

96

ERB BOST1

ERB HOMS1

100

71

ERB CALJ1

ERB STUV1

100

ERB COTJ1

99

ERB GALG1

90

100

ERB-B DANR

100

ERB ONCM1

66

ERB CARA1

60

ERB CYPC1

ERB OREN1

100

ERB ORYL1

72

ERB PARO1

74

ERB MICU1

69

ERB ACAS1

100

ERB SPAA1

ERB ICTP1

76

ERB CARA2

100

ERB-A DANR

ERB ANGJ1

68

ERB MICU2

ERA HOMS1

ERA EQUC1

100

ERA SUSS1

ERA MESA1

100

ERA MUSM1

88

ERA RATN1

ERA CNEU1

69

ERA CAIC1

100

ERA TAEG1

100

ERA GALG1

89

ERA COTJ1

ERA CARA1

ERA DANR

97

ERA CLAG1

100

ERA ICTP1

100

ERA OREA1

100

ERA OREN1

98

ERA ORYS1

100

ERA ORYL1

98

ERA PARO1

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ERA ACAS1

100

ERA ONCM1

100

ERA SALS1

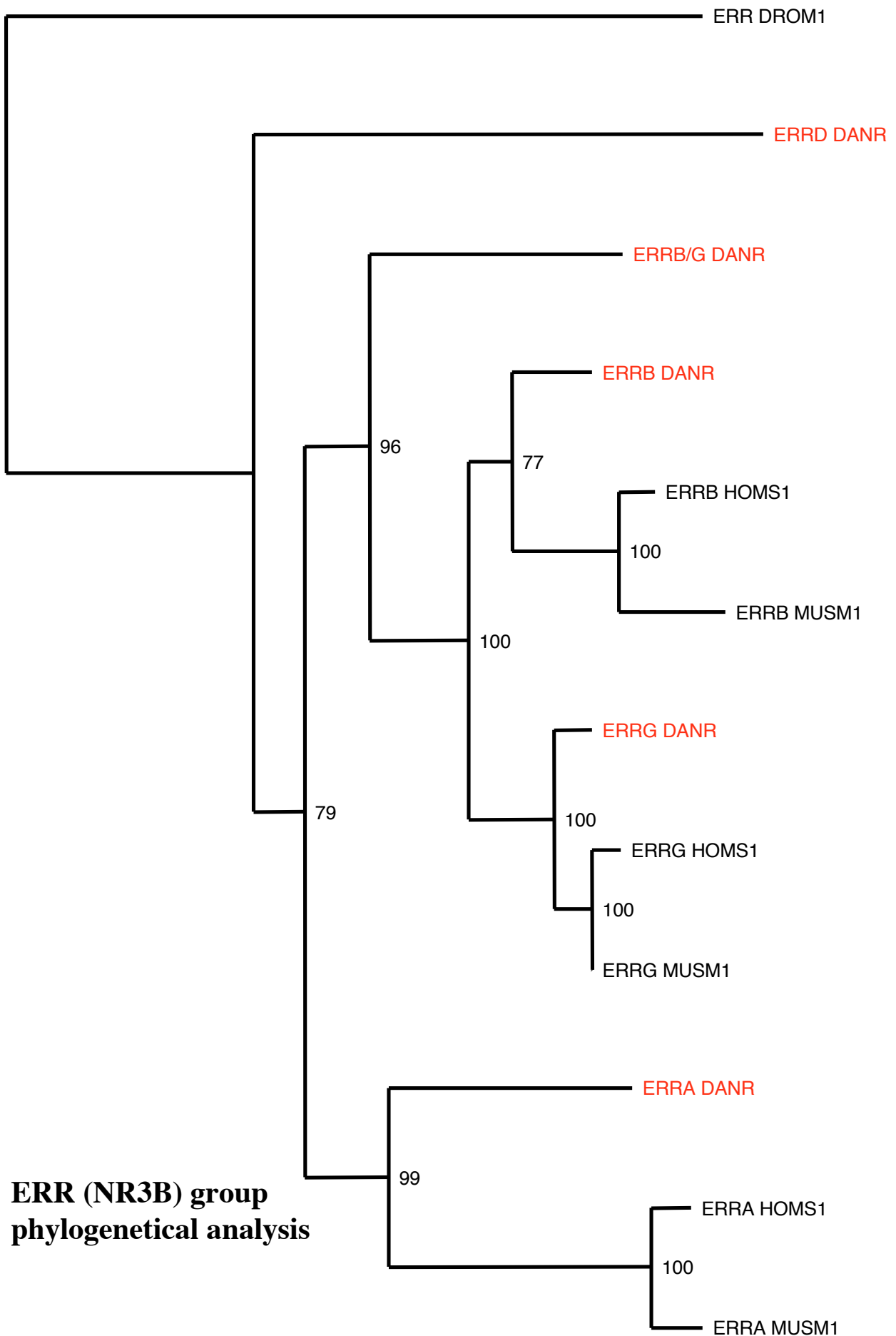
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100

98

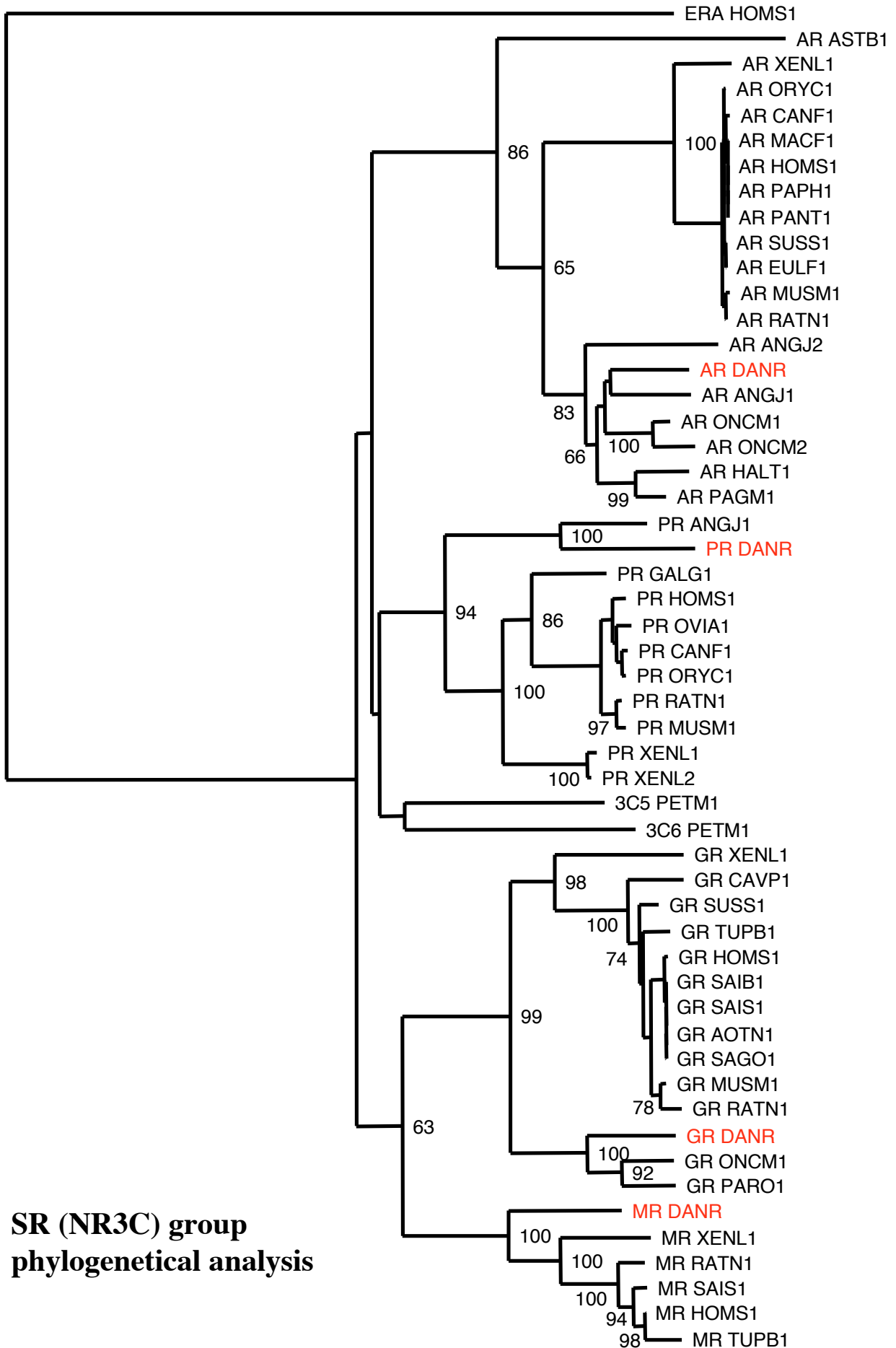
ER (NR3A) group
phylogenetical analysis

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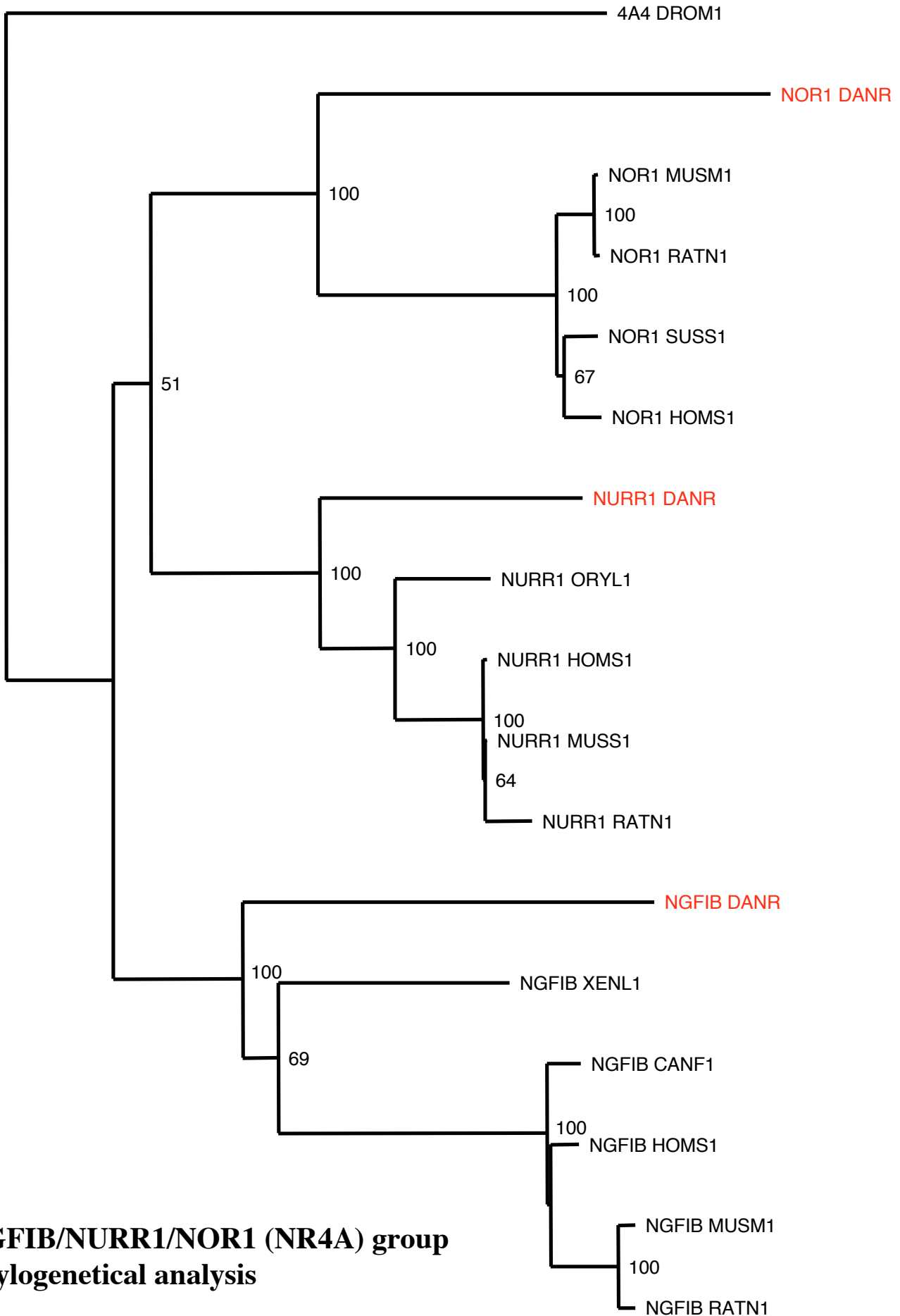


**ERR (NR3B) group
phylogenetical analysis**

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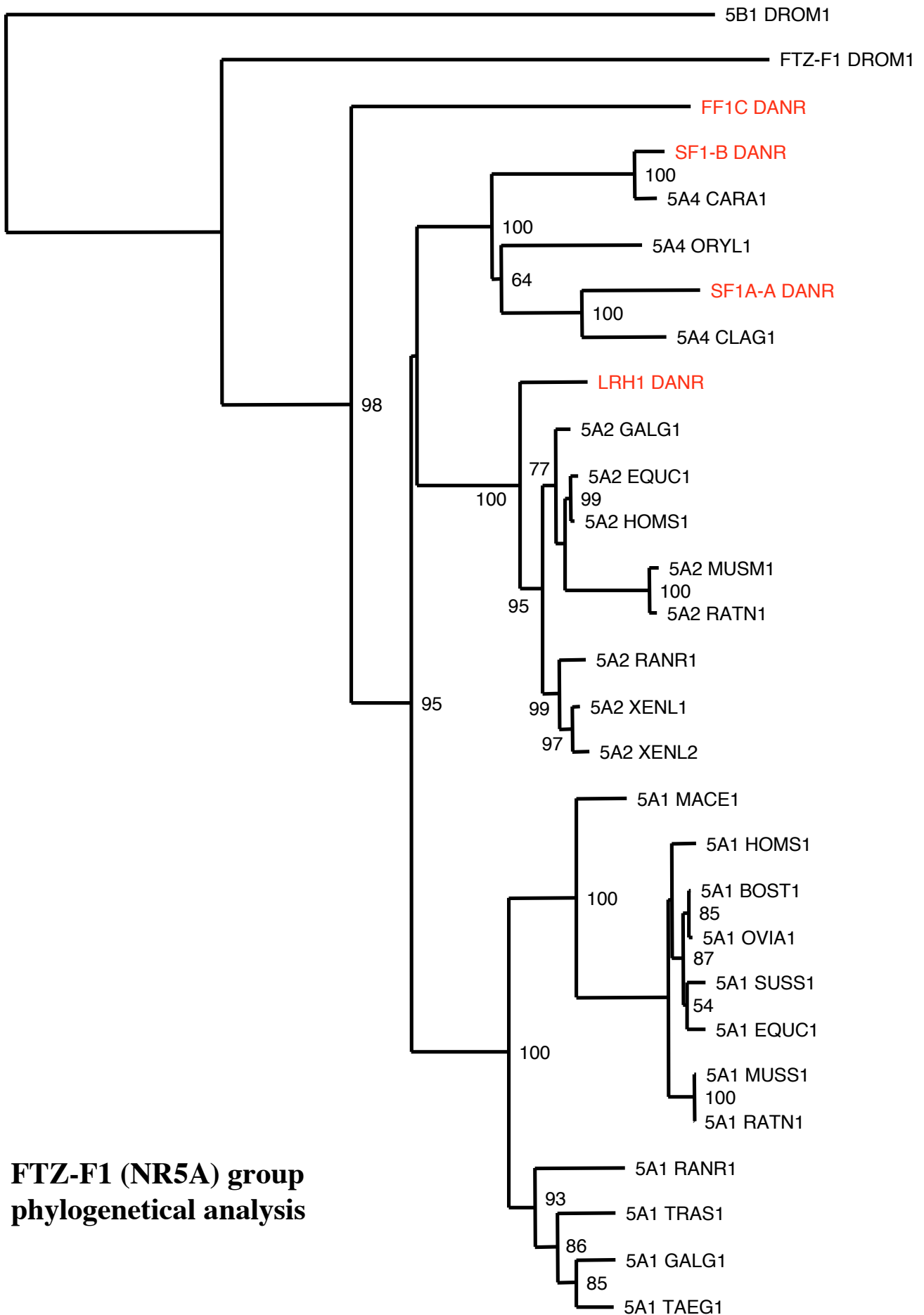


**SR (NR3C) group
phylogenetical analysis**



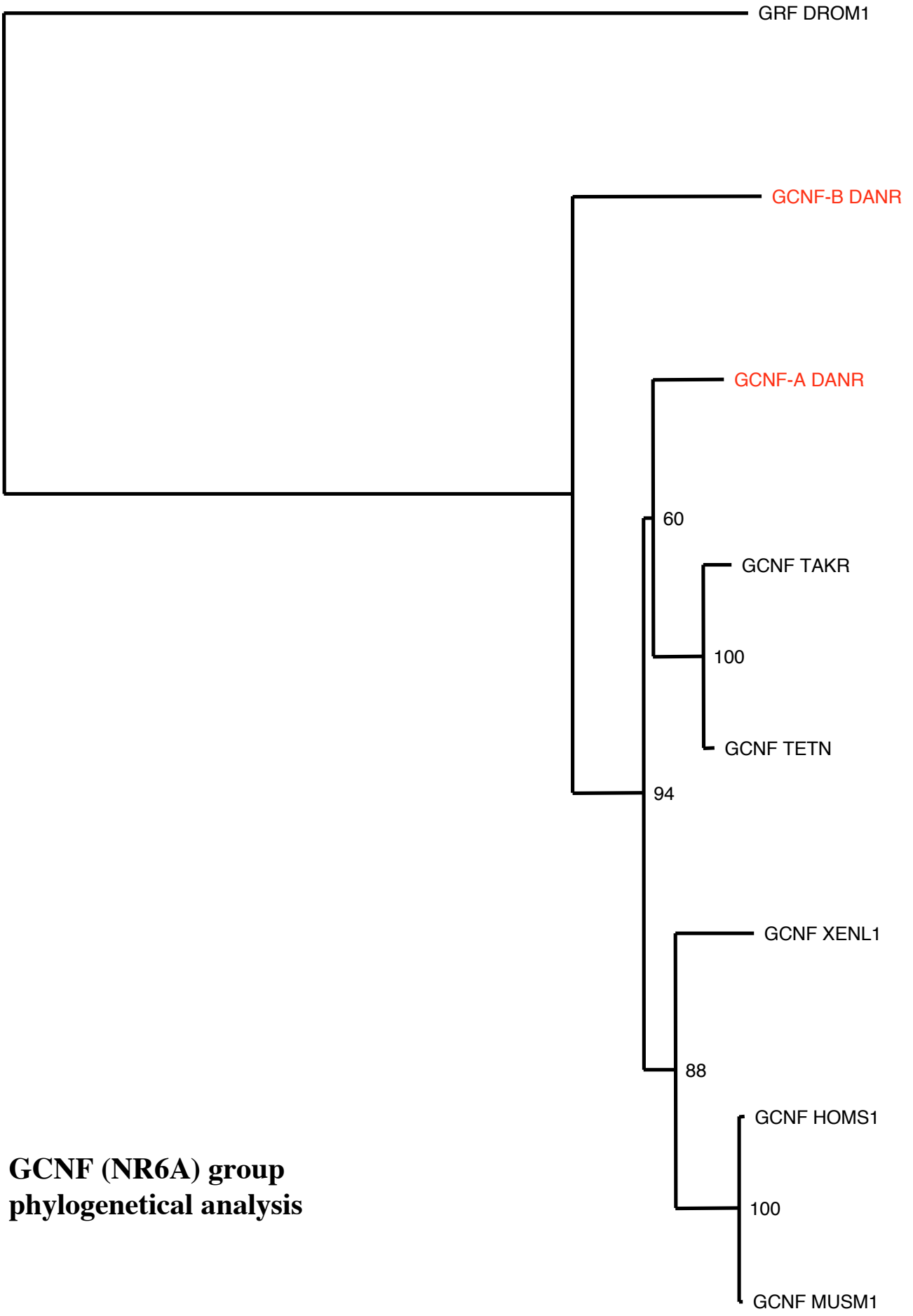
**NGFIB/NURR1/NOR1 (NR4A) group
phylogenetical analysis**

0.1



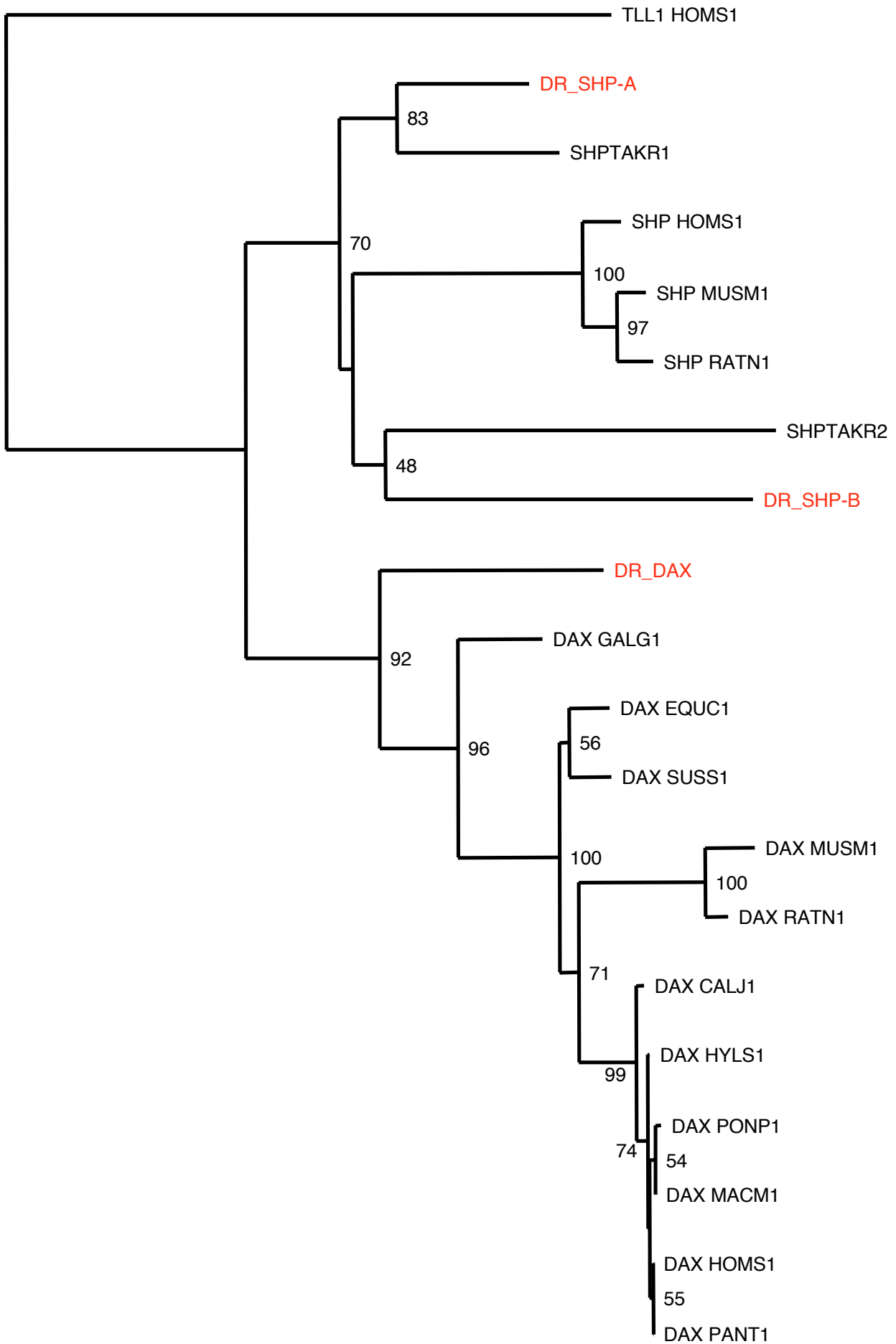
**FTZ-F1 (NR5A) group
phylogenetical analysis**

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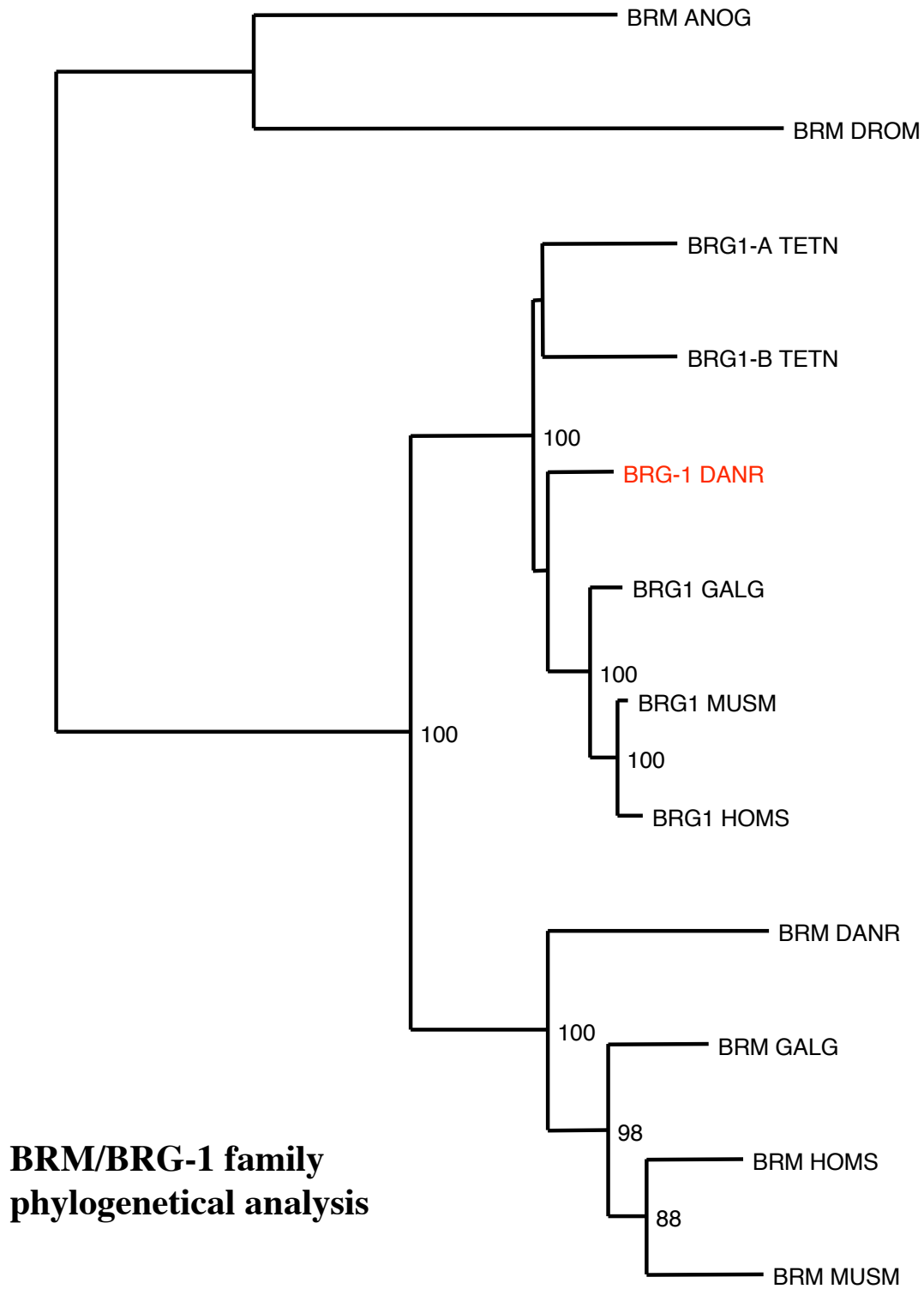


**GCNF (NR6A) group
phylogenetical analysis**

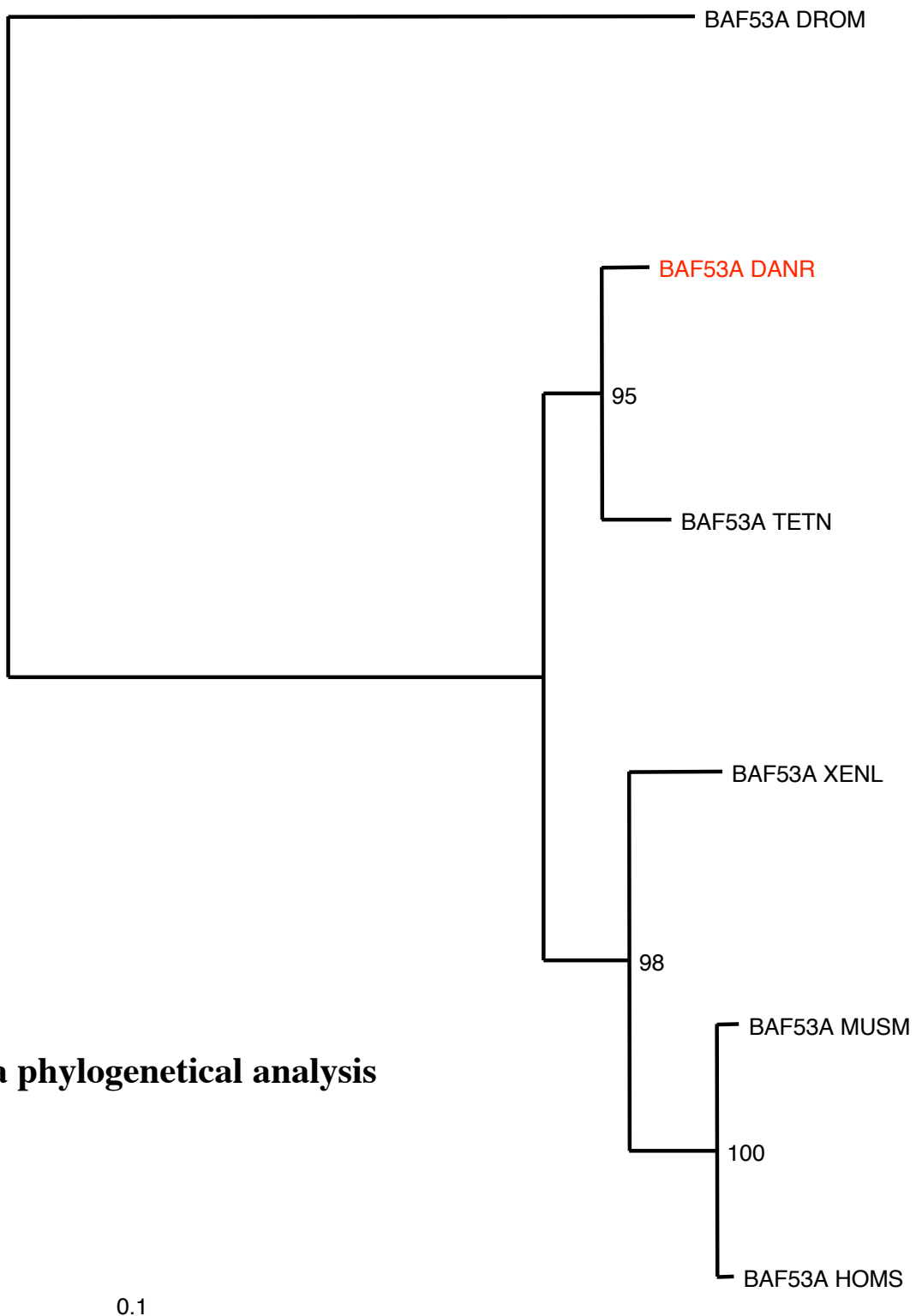
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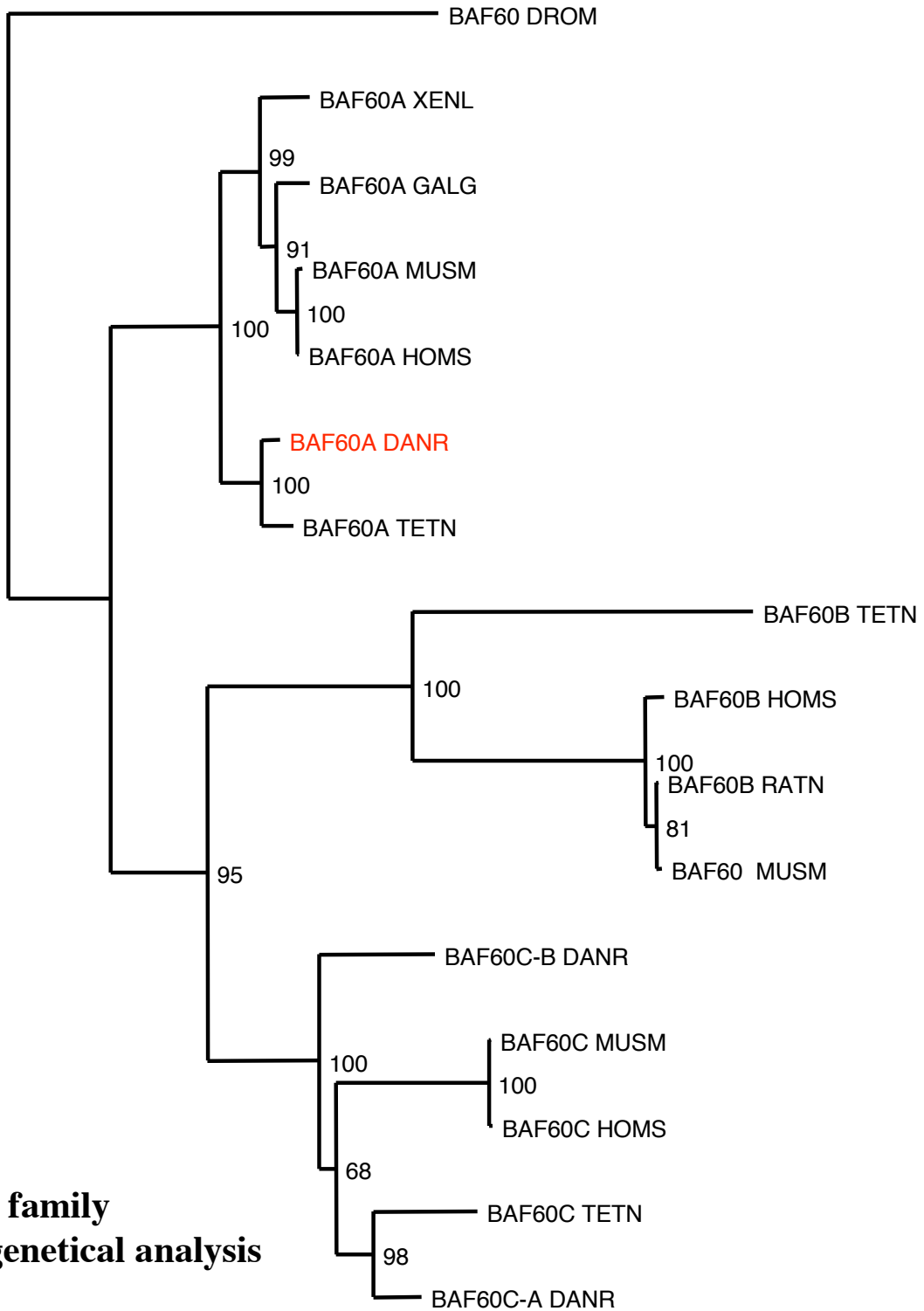


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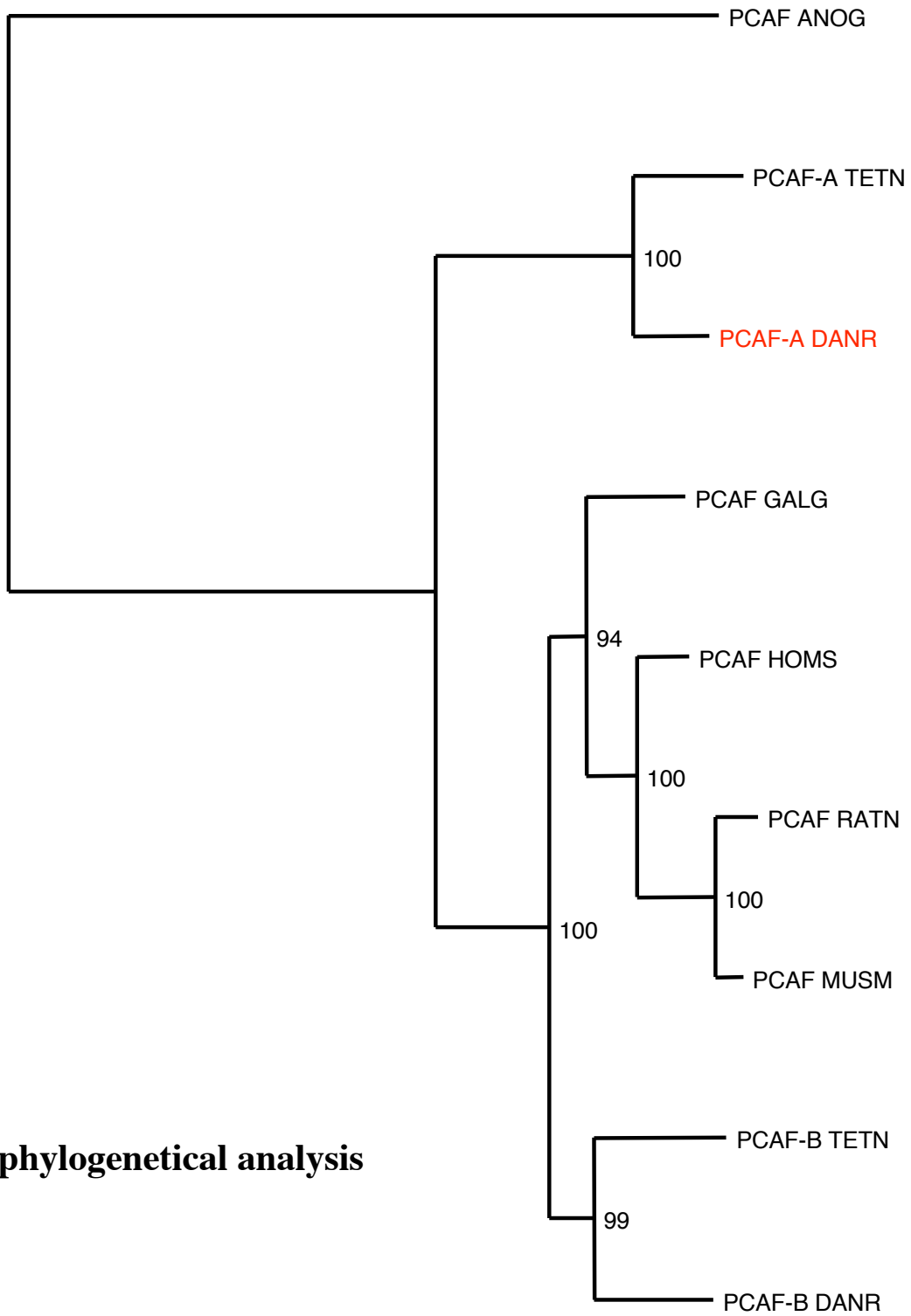


Baf53a phylogenetical analysis



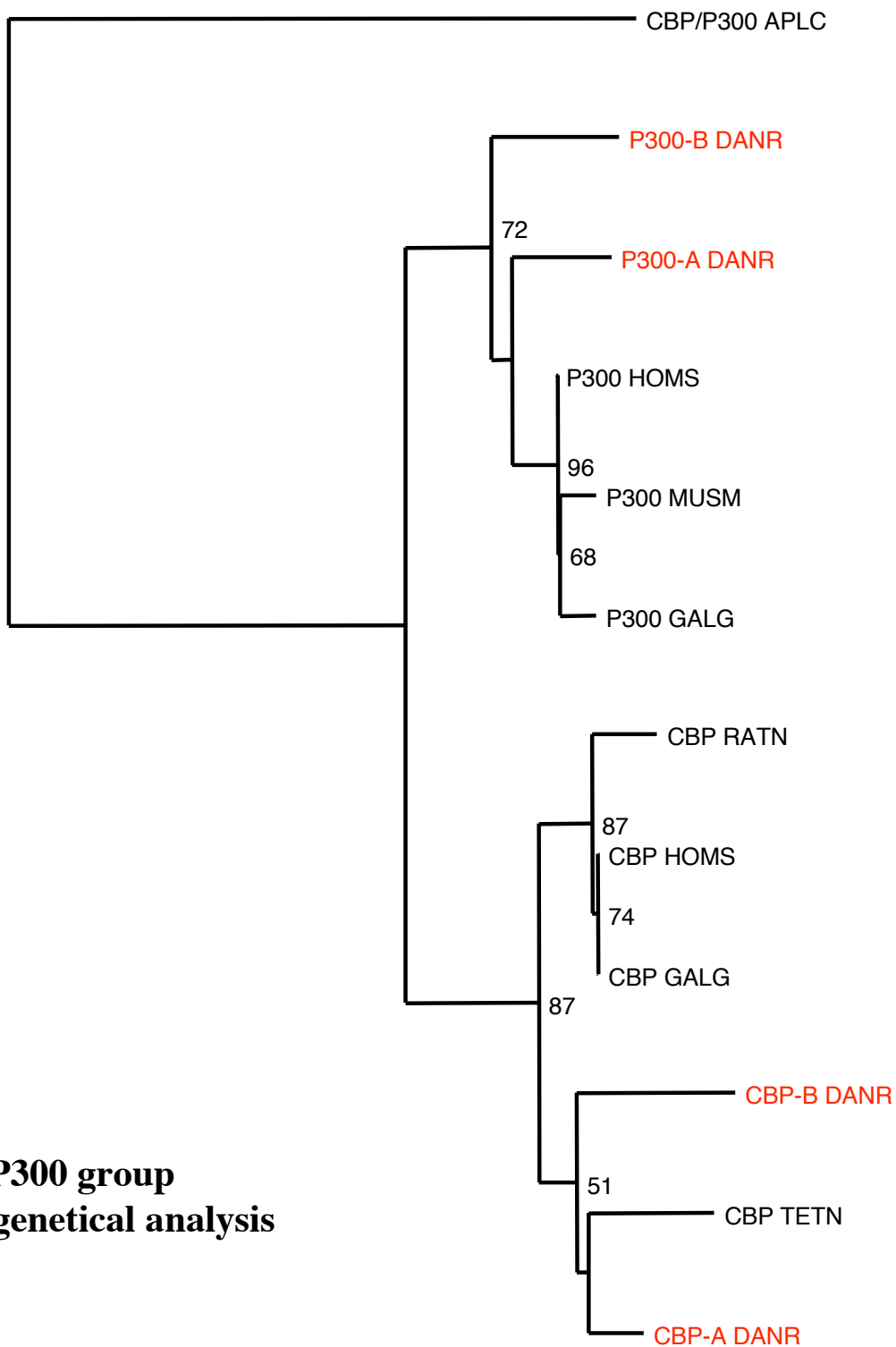


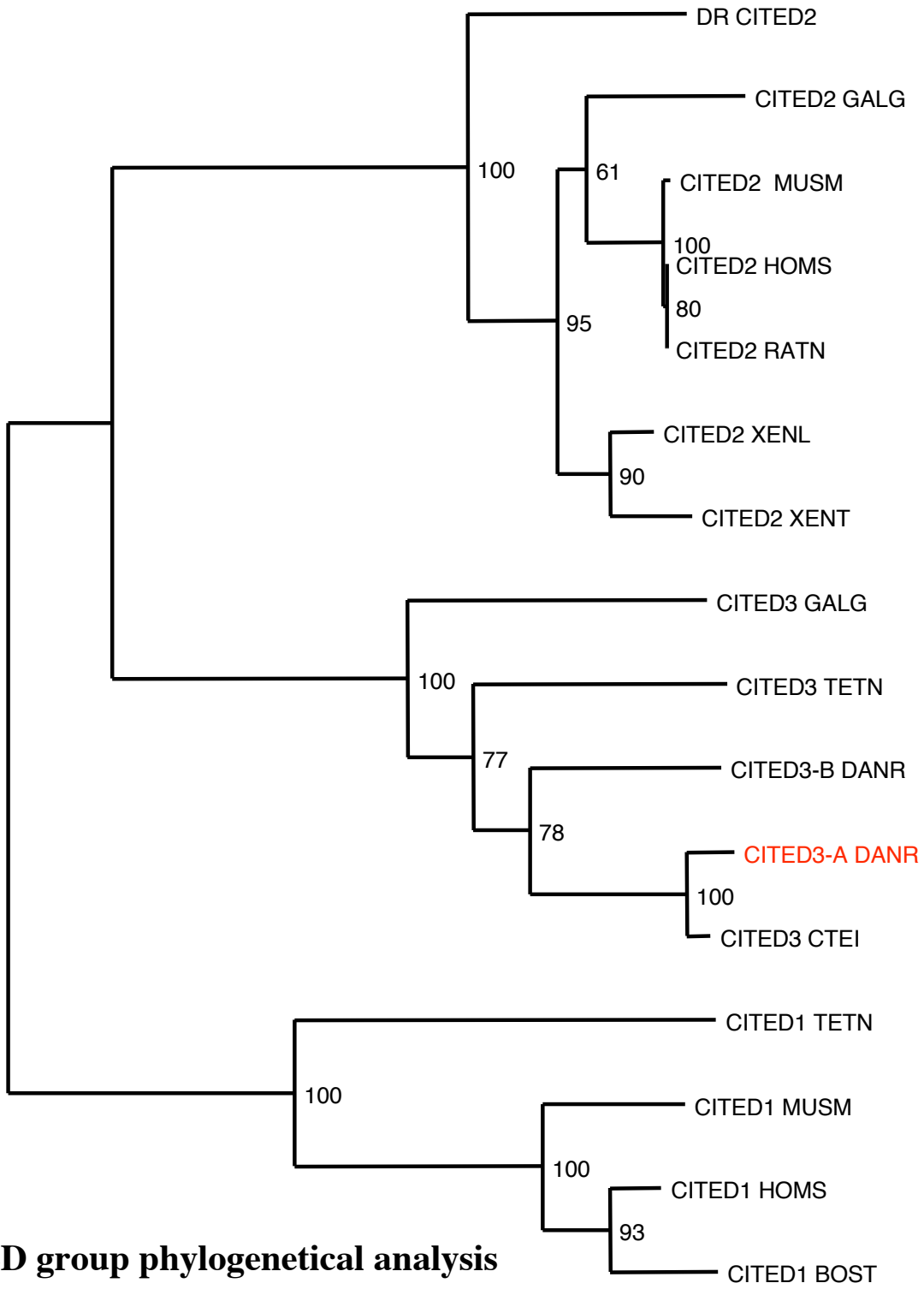
PCAF phylogenetical analysis



0.1

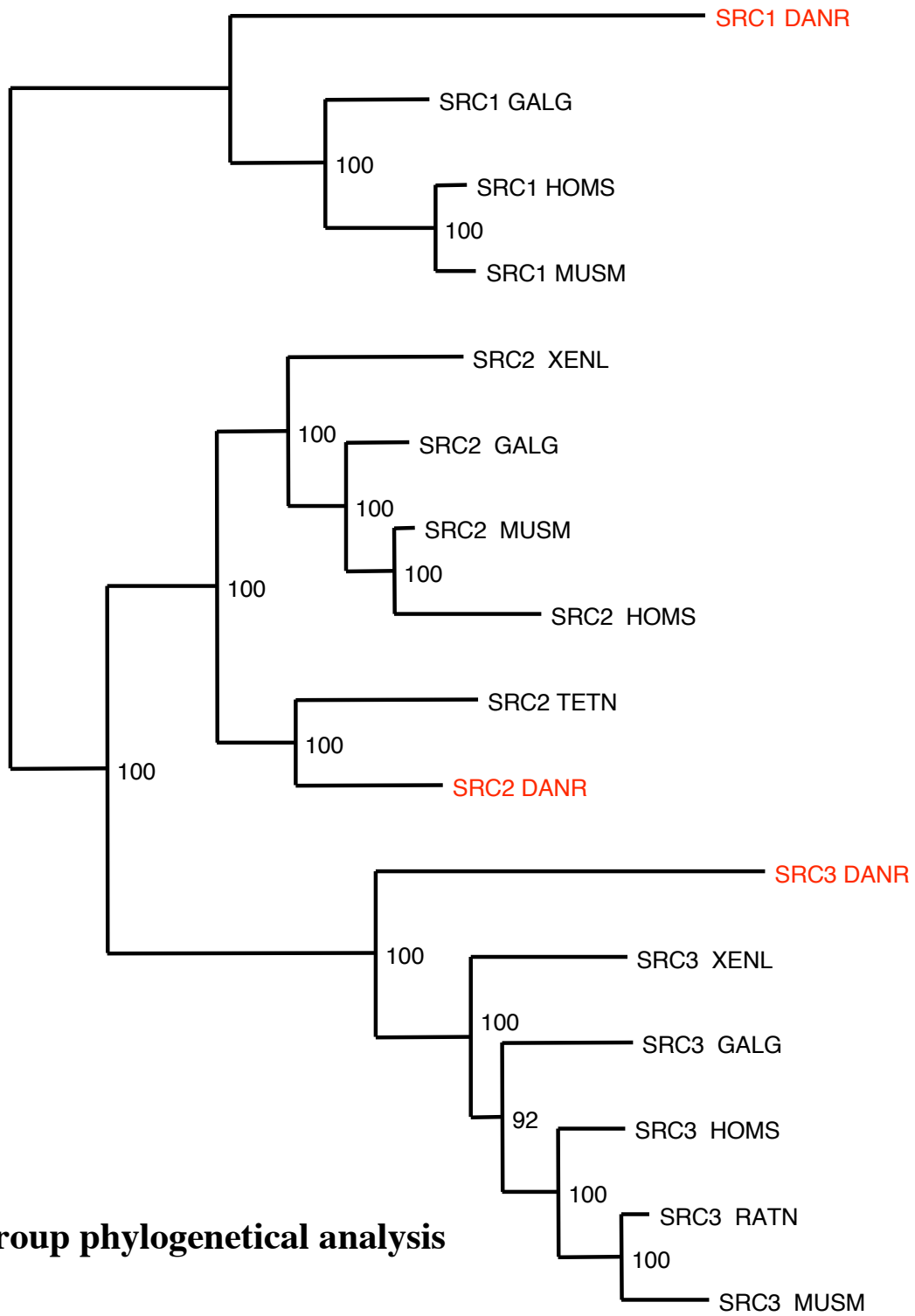
**CBP/P300 group
phylogenetical analysis**





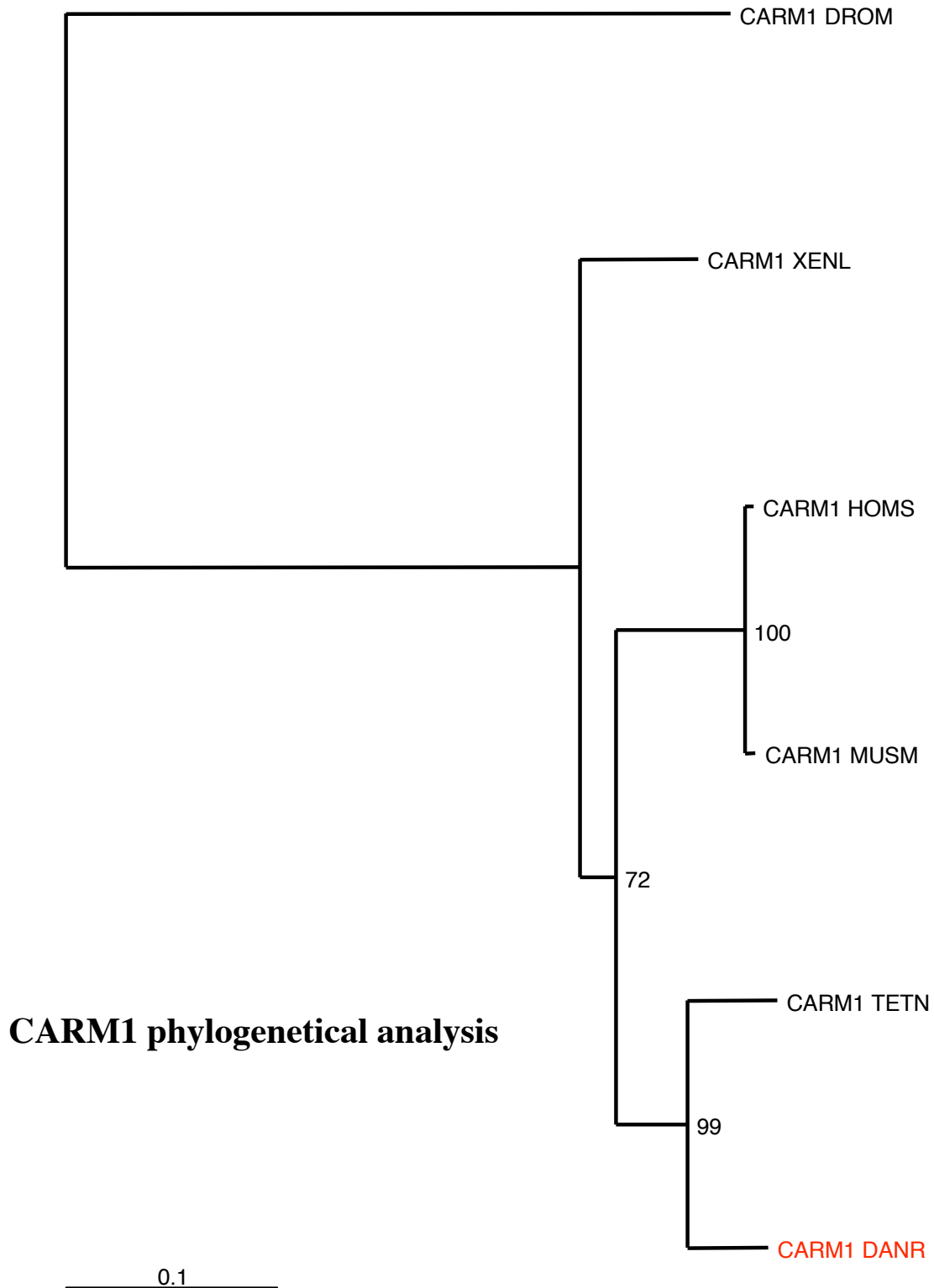
CITED group phylogenetical analysis

0.1

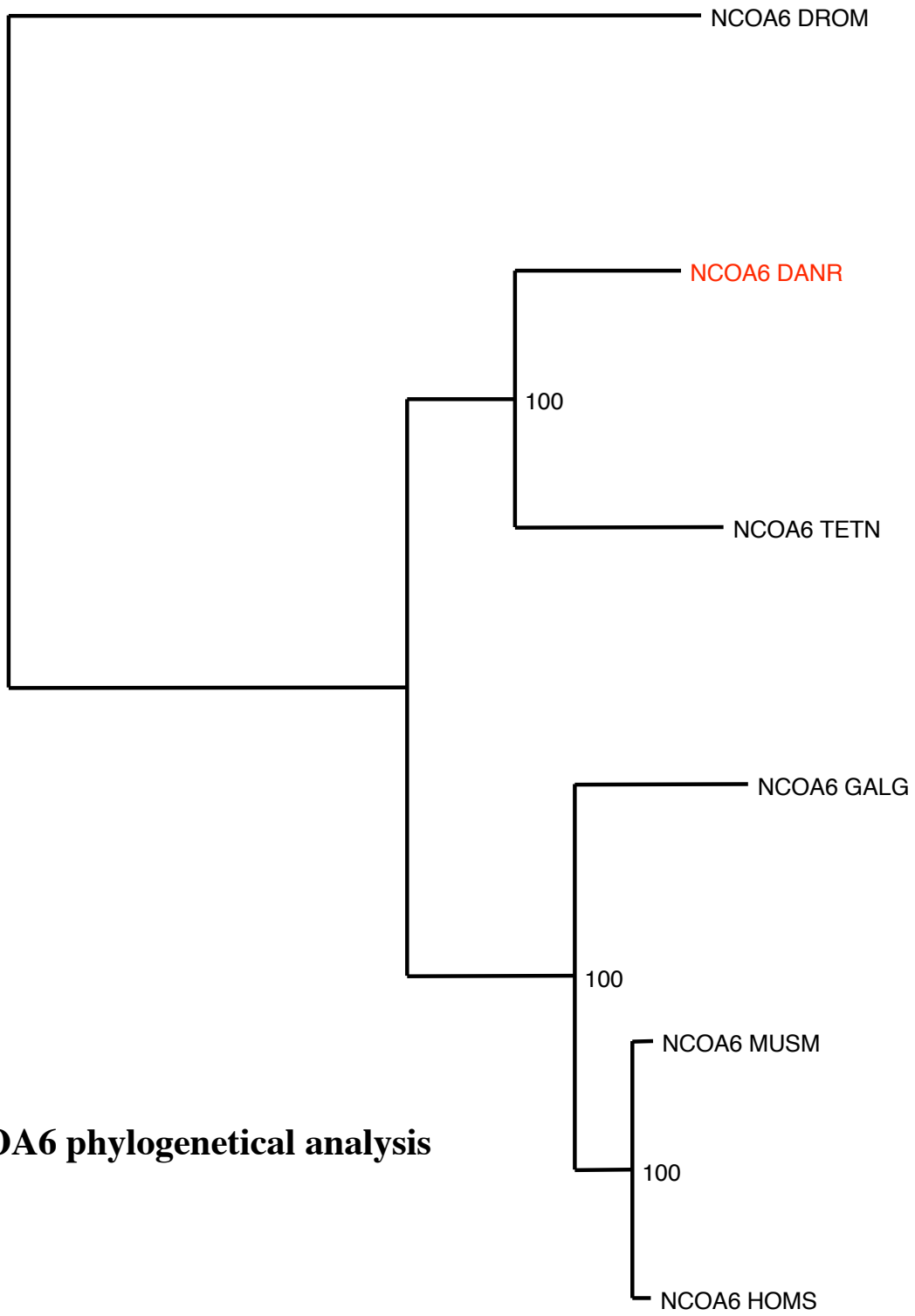


SRC group phylogenetical analysis

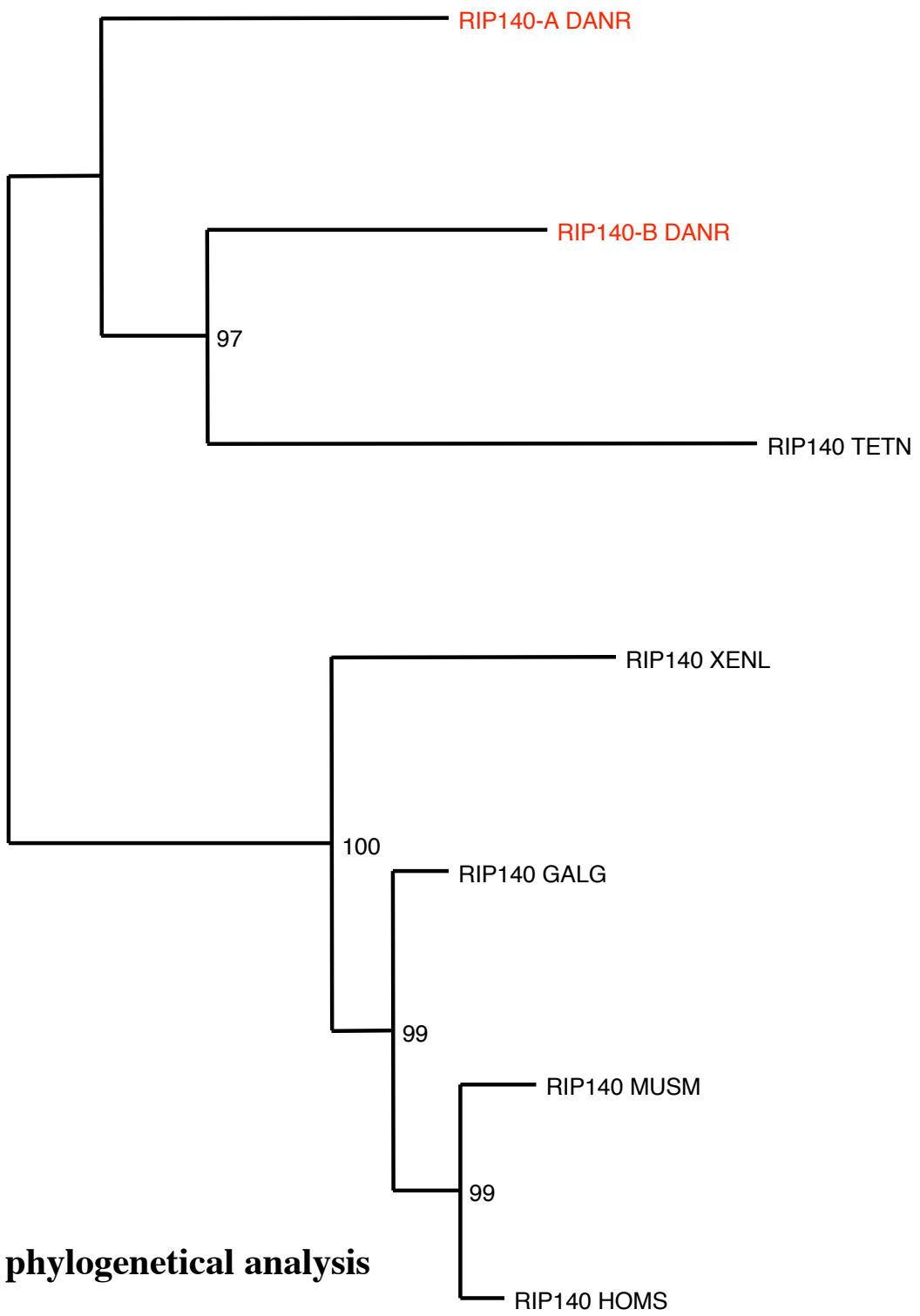
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NCOA6 phylogenetical analysis

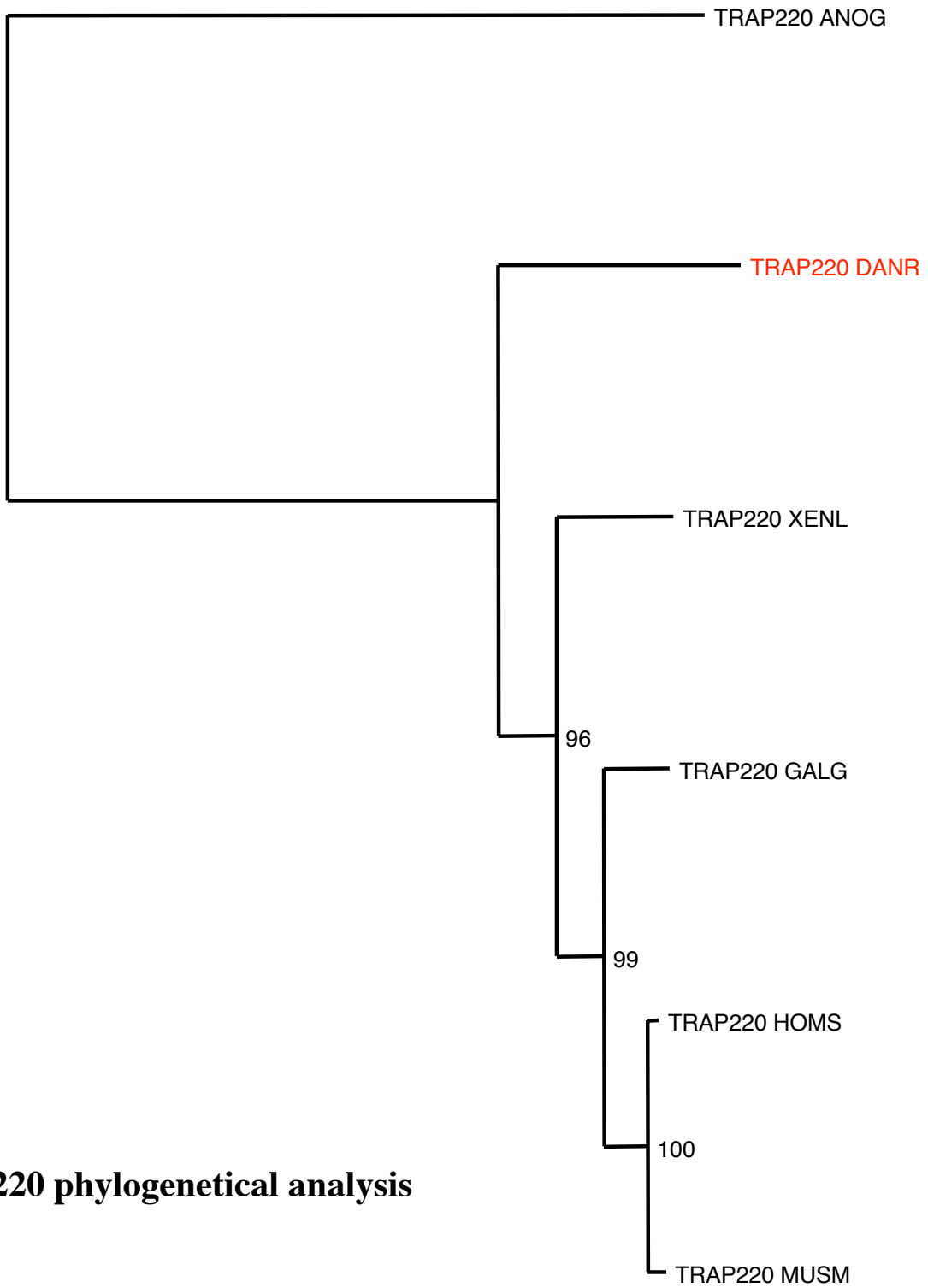


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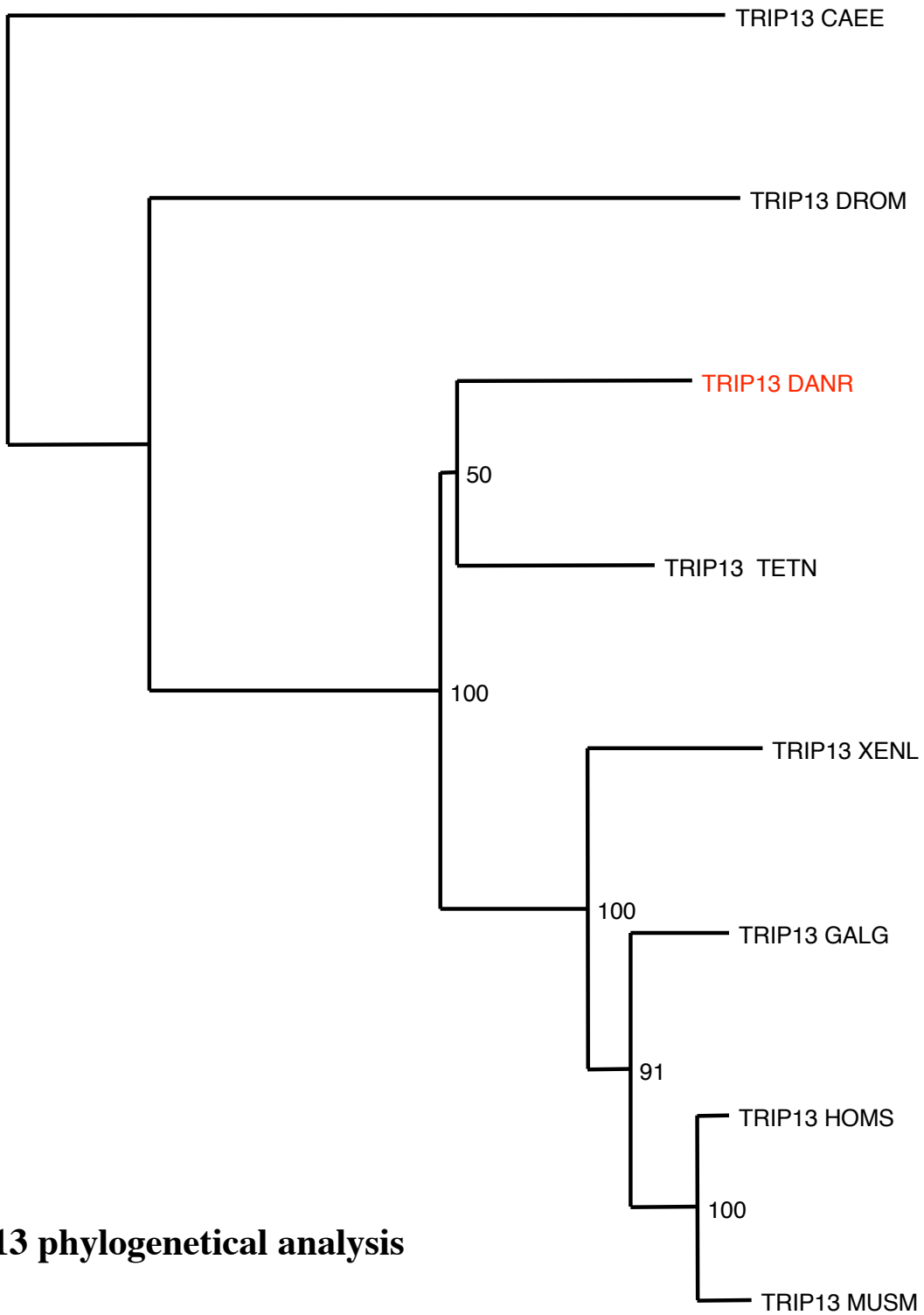
RIP140 phylogenetical analysis

0.1



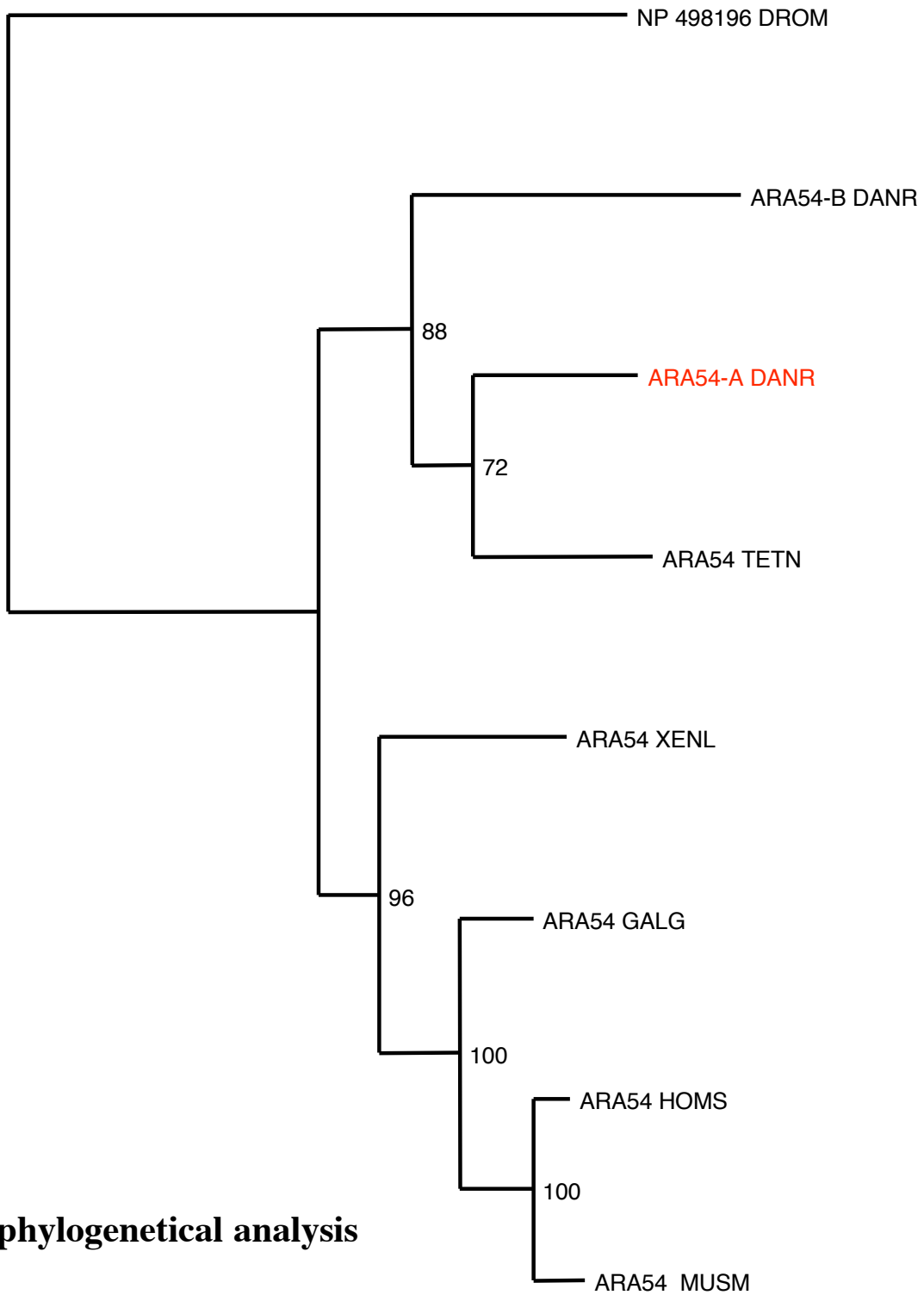
TRAP220 phylogenetical analysis

0.1



TRIP13 phylogenetical analysis

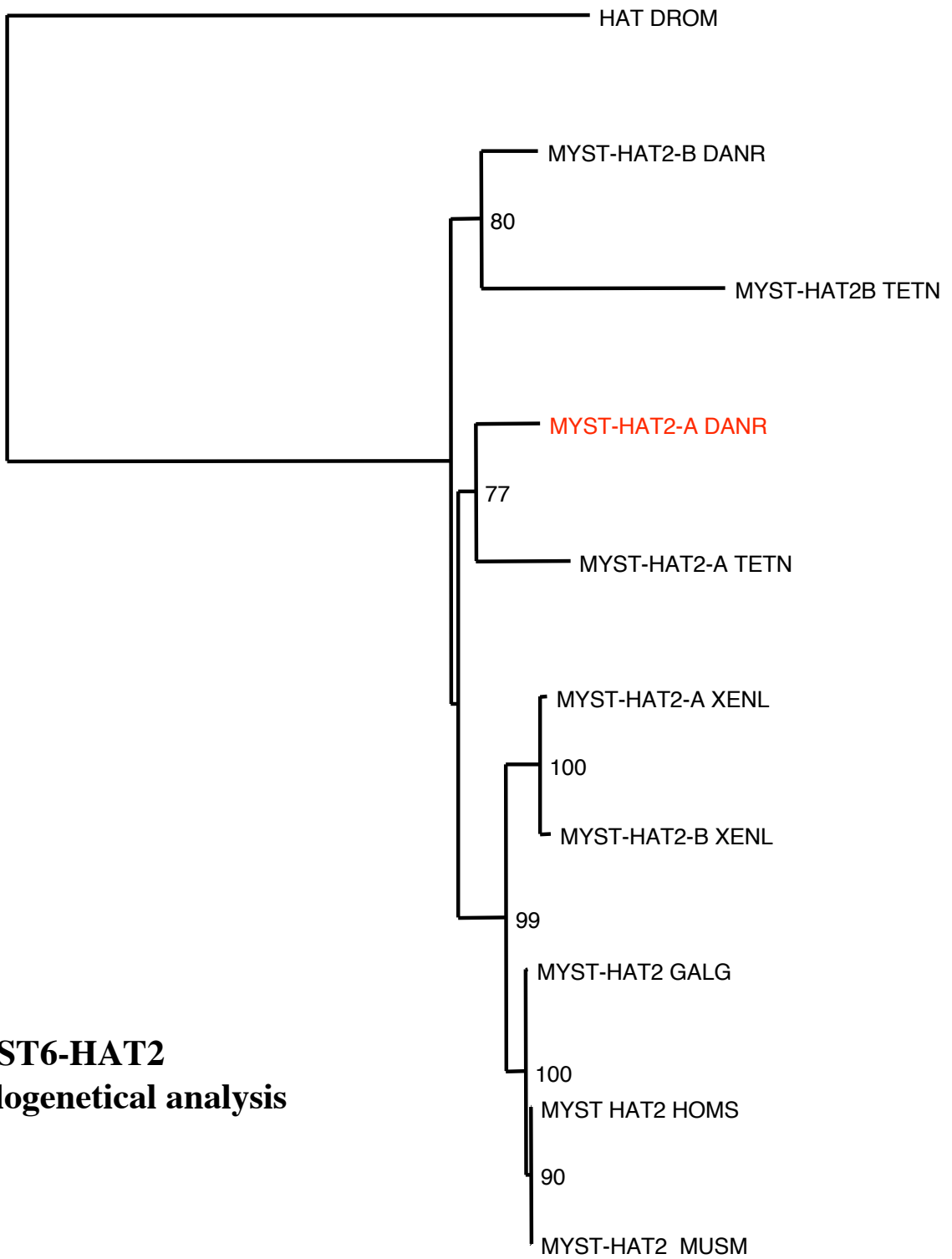
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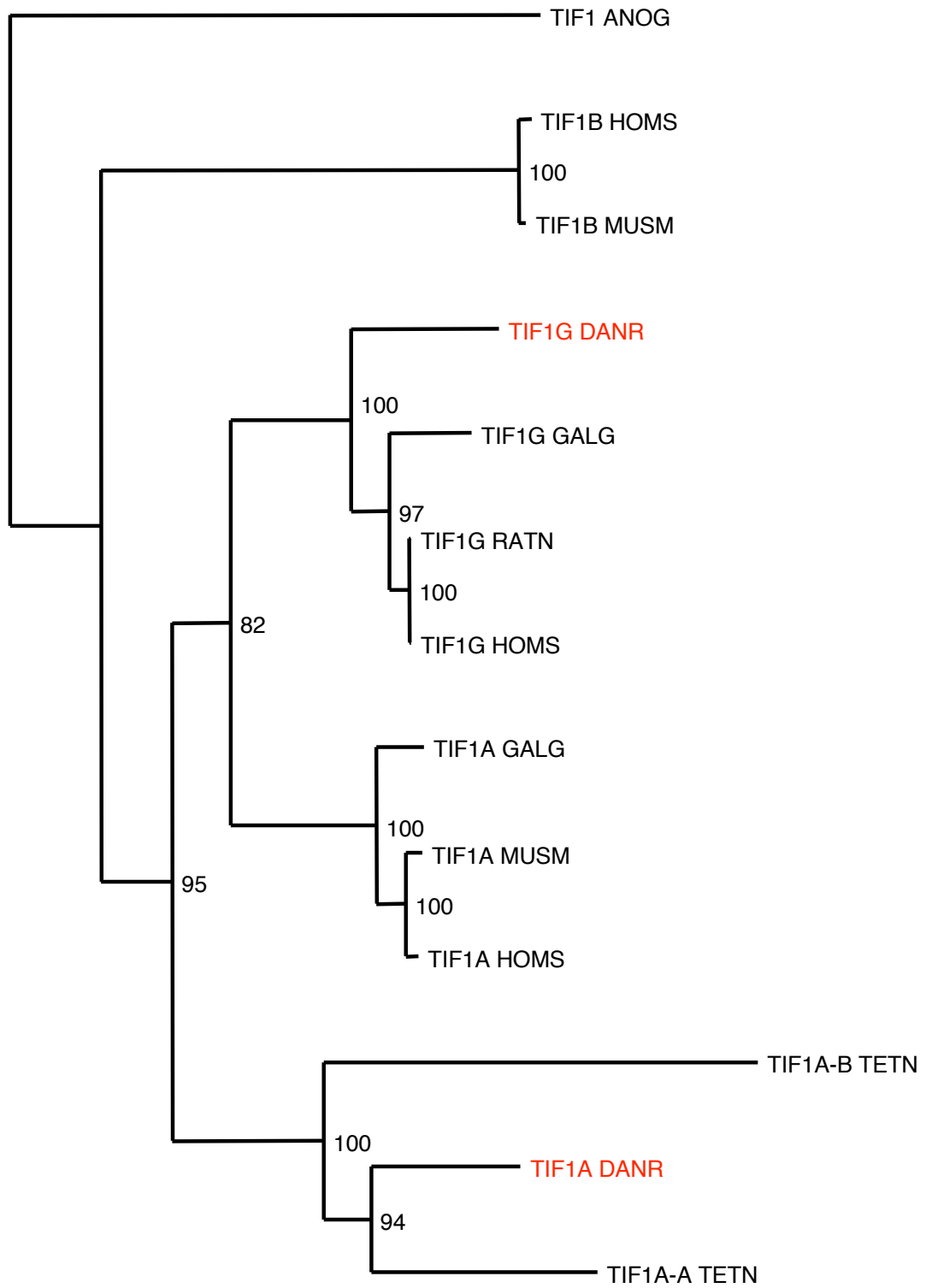


ARA54 phylogenetical analysis

0.1

**MYST6-HAT2
phylogenetical analysis**

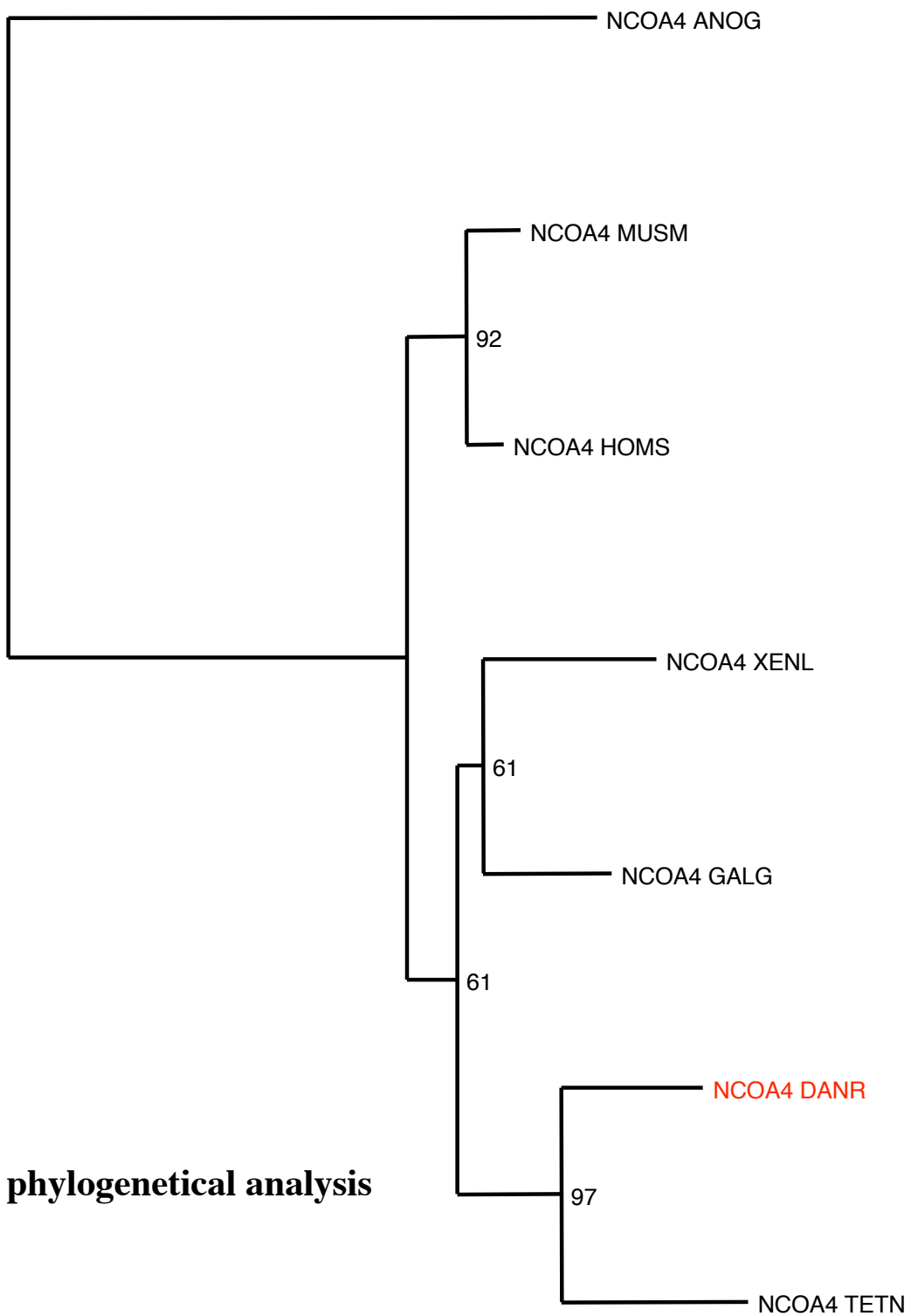




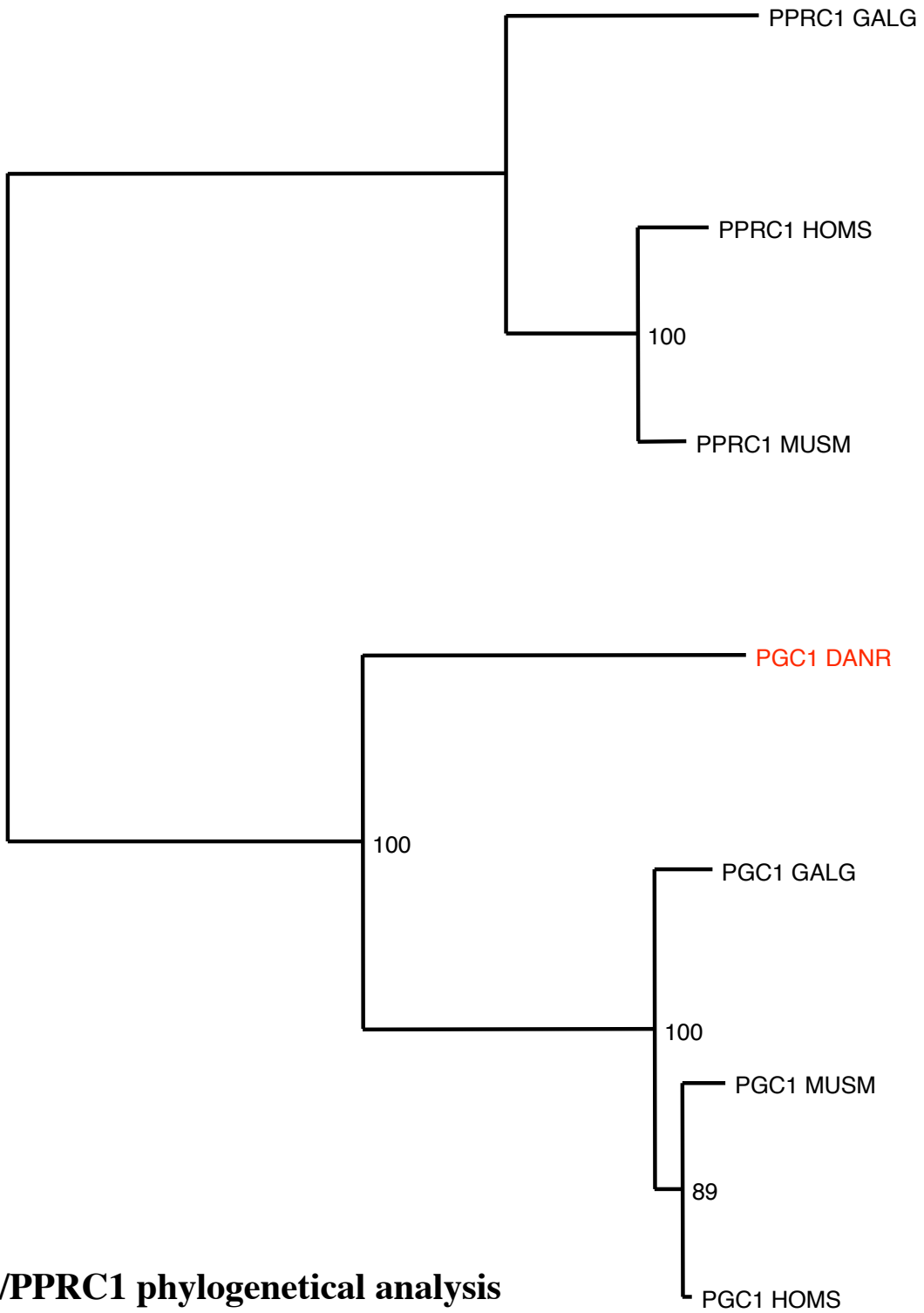
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TIF1 group phylogenetical analysis

NCOA4 phylogenetical analysis

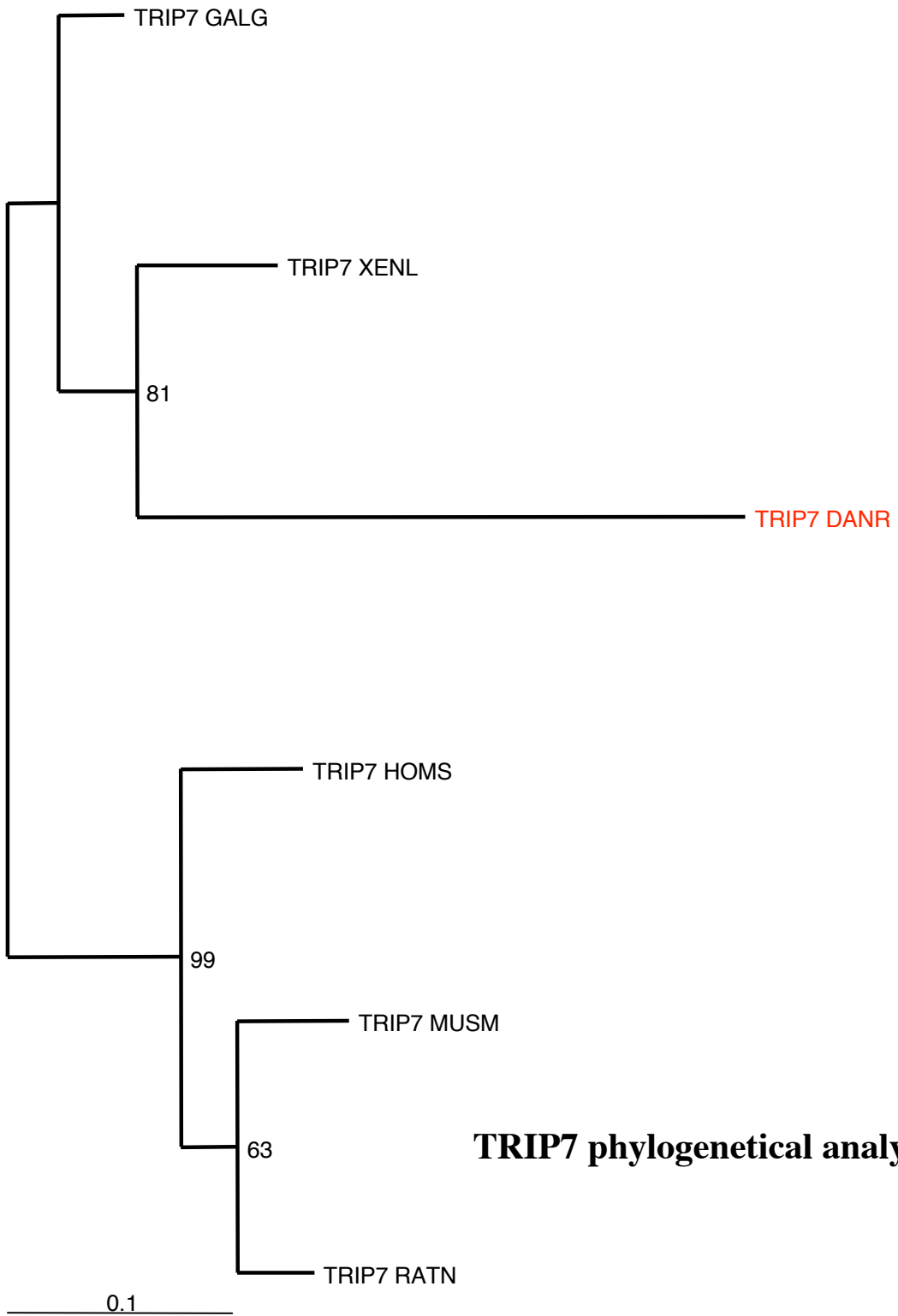


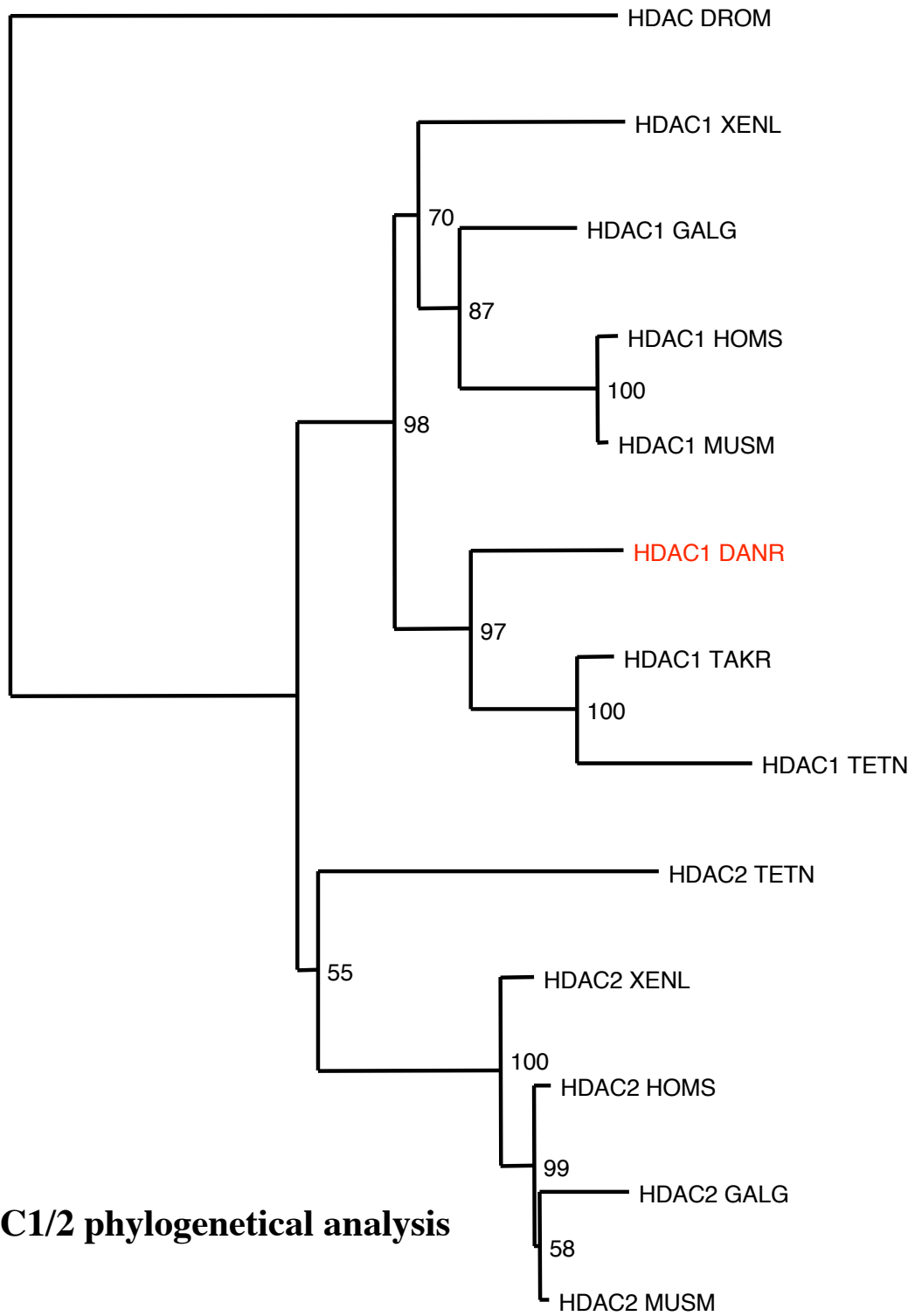
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PGC1/PPRC1 phylogenetical analysis

0.1

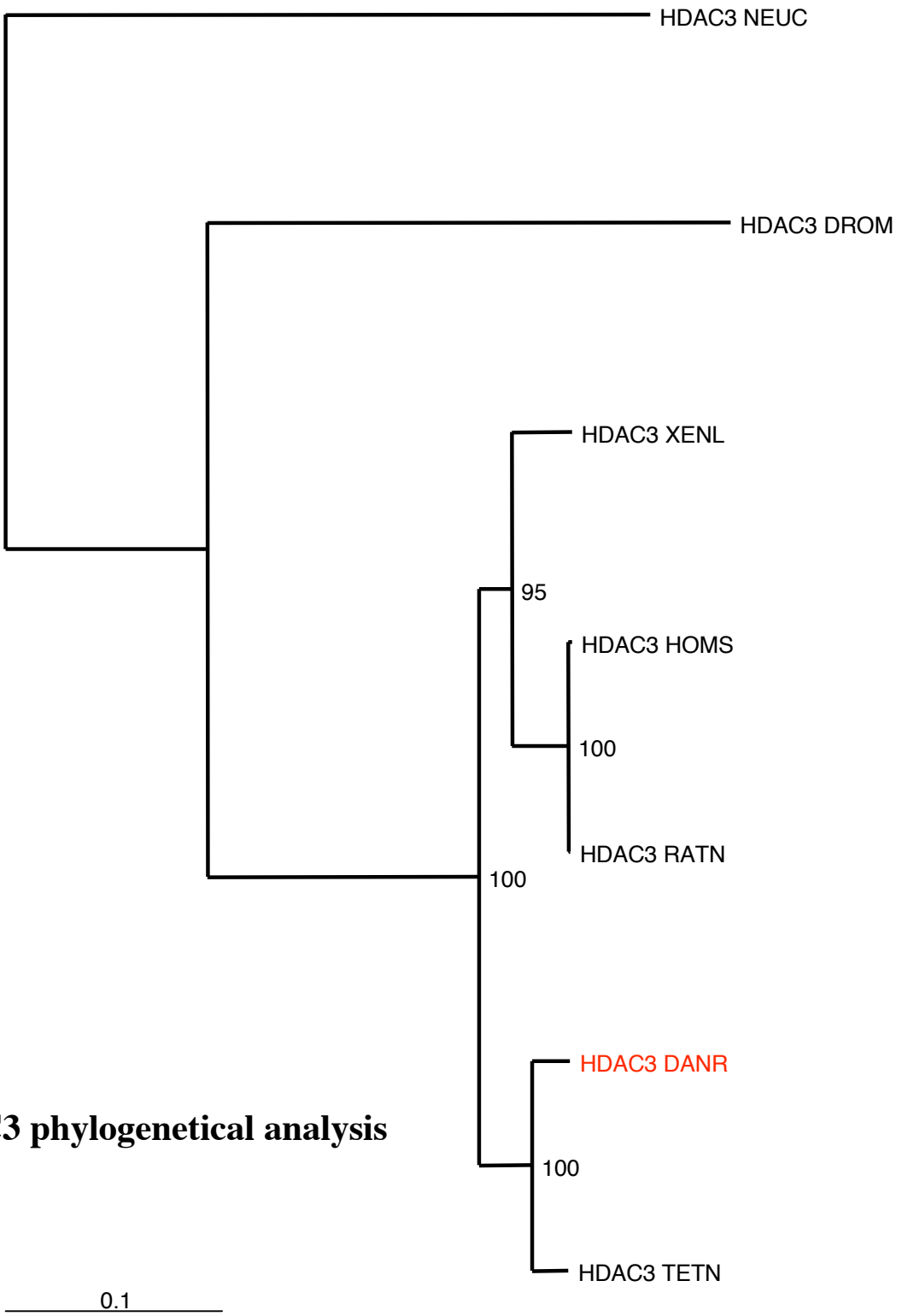


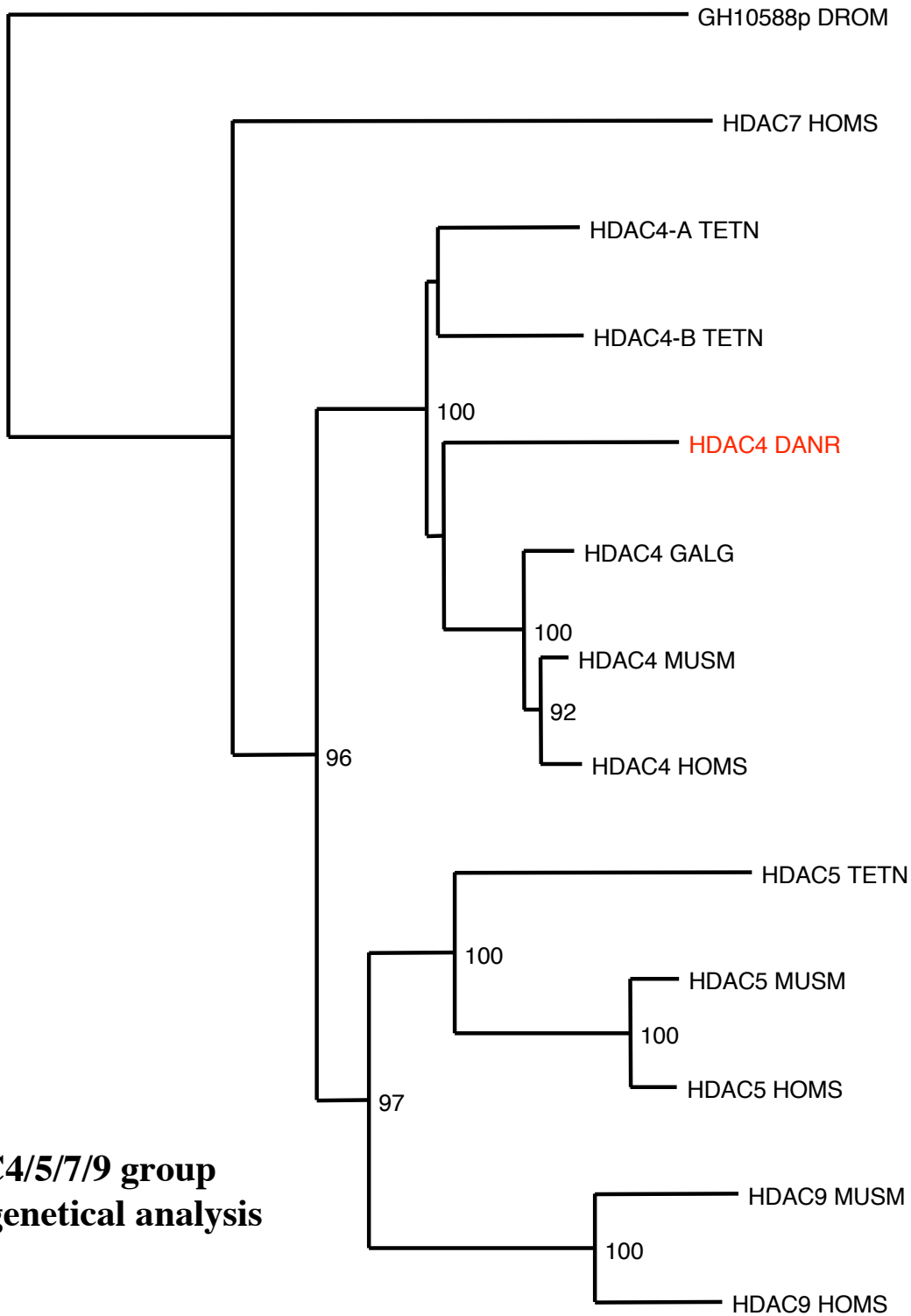


HDAC1/2 phylogenetical analysis

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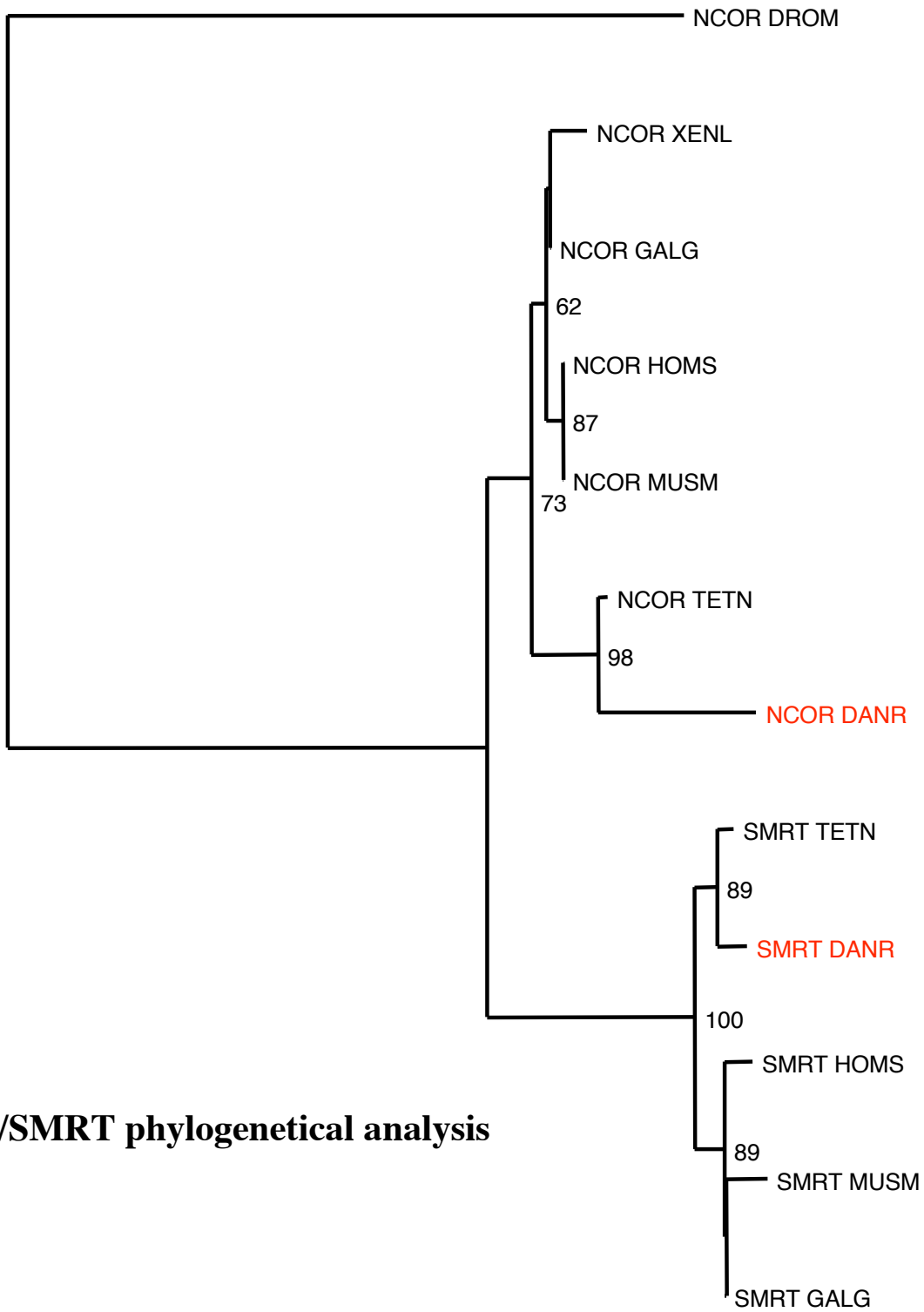
HDAC3 phylogenetical analysis





**HDAC4/5/7/9 group
phylogenetical analysis**

0.1



NCOR/SMRT phylogenetical analysis

0.1

