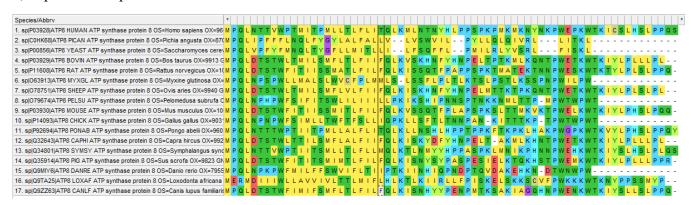
Кондратюк Александр Николаевич, группа 3

1)Скриншот выравнивания для метода ClustalW



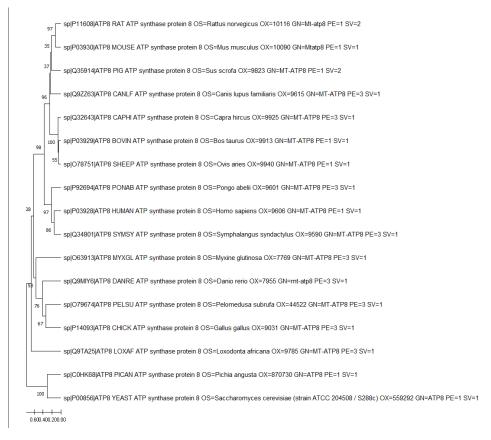
2) Скриншот выравнивания для метода Muscle



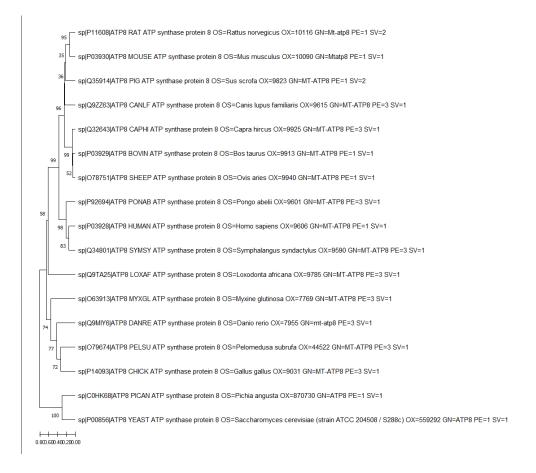
Вывод: Разница в выравнивании есть. Лучше сработал алгоритм выравнивания: Muscle, т.к в нем меньше гэпов.

3) Деревья UPGMA

Muscle



ClustalW



4) Деревья NJ

Muscle

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sp|P03929|ATP8 BOVIN ATP synthase protein 8 OS=Bos taurus OX=9913 GN=MT-ATP8 PE=1 SV=1
sp|Q32643|ATP8 CAPHI ATP synthase protein 8 OS=Capra hircus OX=9925 GN=MT-ATP8 PE=3 SV=1
sp|O78751|ATP8 SHEEP ATP synthase protein 8 OS=Ovis aries OX=9940 GN=MT-ATP8 PE=1 SV=1
splQ9ZZ63IATP8 CANLF ATP synthase protein 8 OS=Canis lupus familiaris OX=9615 GN=MT-ATP8 PE=3 SV=1
sp|Q35914|ATP8 PIG ATP synthase protein 8 OS=Sus scrofa OX=9823 GN=MT-ATP8 PE=1 SV=2
sp|P11608|ATP8 RAT ATP synthase protein 8 OS=Rattus norvegicus OX=10116 GN=Mt-atp8 PE=1 SV=2
sp|P03930|ATP8 MOUSE ATP synthase protein 8 OS=Mus musculus OX=10090 GN=Mtatp8 PE=1 SV=1
     splQ9TA25|ATP8 LOXAF ATP synthase protein 8 OS=Loxodonta africana OX=9785 GN=MT-ATP8 PE=3 SV=1
 sp|P92694|ATP8 PONAB ATP synthase protein 8 OS=Pongo abelii OX=9601 GN=MT-ATP8 PE=3 SV=1
 sp|P03928|ATP8 HUMAN ATP synthase protein 8 OS=Homo sapiens OX=9606 GN=MT-ATP8 PE=1 SV=1
sp|Q34801|ATP8 SYMSY ATP synthase protein 8 OS=Symphalangus syndactylus OX=9590 GN=MT-ATP8 PE=3 SV=1
   splC0HK68IATP8 PICAN ATP synthase protein 8 OS=Pichia angusta OX=870730 GN=ATP8 PE=1 SV=1
       plP00856|ATP8 YEAST ATP synthase protein 8 OS=Saccharomyces cerevisiae (strain ATCC 204508 / S288c) OX=559292 GN=ATP8 PE=1 SV=1
sp|P14093|ATP8 CHICK ATP synthase protein 8 OS=Gallus gallus OX=9031 GN=MT-ATP8 PE=3 SV=1
splO79674IATP8 PELSU ATP synthase protein 8 OS=Pelomedusa subrufa OX=44522 GN=MT-ATP8 PE=3 SV=1
  splQ9MIY6IATP8 DANRE ATP synthase protein 8 OS=Danio rerio OX=7955 GN=mt-atp8 PE=3 SV=1
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ClustalW

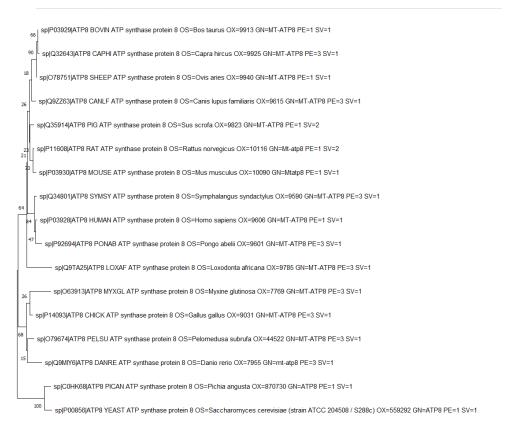


5) Деревья ML

Muscle

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spi(03298/ATP8 BOVIN ATP synthase protein 8 OS=Bos taurus OX=9913 GN=MT-ATP8 PE=1 SV=1
spi(032643)ATP8 CAPHI ATP synthase protein 8 OS=Capra hircus OX=9925 GN=MT-ATP8 PE=3 SV=1
spi(078751)ATP8 SHEEP ATP synthase protein 8 OS=Capra hircus OX=9926 GN=MT-ATP8 PE=3 SV=1
spi(078751)ATP8 SHEEP ATP synthase protein 8 OS=Caris lupus familiaris OX=9615 GN=MT-ATP8 PE=3 SV=1
spi(035914)ATP8 PG ATP synthase protein 8 OS=Caris lupus familiaris OX=9615 GN=MT-ATP8 PE=3 SV=1
spi(035914)ATP8 PG ATP synthase protein 8 OS=Sus scrofa OX=9823 GN=MT-ATP8 PE=1 SV=2
spi(035914)ATP8 PONAB ATP synthase protein 8 OS=Loxodonta africana OX=9785 GN=MT-ATP8 PE=3 SV=1
spi(03628)ATP8 HUMAN ATP synthase protein 8 OS=Pongo abelii OX=9601 GN=MT-ATP8 PE=3 SV=1
spi(0364801)ATP8 SYMSY ATP synthase protein 8 OS=Symphalangus syndactylus OX=9590 GN=MT-ATP8 PE=3 SV=1
spi(0364801)ATP8 SYMSY ATP synthase protein 8 OS=Symphalangus syndactylus OX=9590 GN=MT-ATP8 PE=3 SV=1
spi(03930)ATP8 MOUSE ATP synthase protein 8 OS=Rattus norvegicus OX=10116 GN=Mt-atp8 PE=1 SV=1
spi(03930)ATP8 MOUSE ATP synthase protein 8 OS=Pongo and since and substance of the synthase protein 8 OS=Saccharomyces cerevisiae (strain ATCC 204508 / S288c) OX=559292 GN=ATP8 PE=1 SV=1
spi(03930)ATP8 MYXGL ATP synthase protein 8 OS=Saccharomyces cerevisiae (strain ATCC 204508 / S288c) OX=559292 GN=ATP8 PE=1 SV=1
spi(079674)ATP8 PELSU ATP synthase protein 8 OS=Gallus gallus OX=9031 GN=MT-ATP8 PE=3 SV=1
spi(079674)ATP8 PELSU ATP synthase protein 8 OS=Danio rerio OX=7955 GN=mt-atp8 PE=3 SV=1
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ClustalW



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При построении обоими методами топология деревьев получилась разная, бустрэпзначения тоже различные.

Таким образом, деревья не совпадают.