```
# STOCKHOLM 1.0
cabr-RF00052-366    AUG-----CUUUCGGCCU-----GUUCCCUGAGACCUCAAGU---GUGAGCGUUCUGA---AC-AUGCUUCACGCCUGGGCUCUCCGGGUACCAGGACG---GUCUGAg--
cael-RF00052-373    AUG-----CUUCCGGCCU-----GUUCCCUGAGACCUCAAGU---GUGAGUGUACUAUUG----AUGCUUCACACCUGGGCUCUCCGGGUACCAGGACG---GUUUGAG--
care-RF00052-367    ca--GCUCa---CGGCCU-----GCUCCCUGAGACCUCAAGU---GUGAGCGUACUGUAG----AUGCUUCACACCUGGGCUCUCUGGGUACCAGGACG---GCguauu--
cavu-RF00052-376    AUG-----CUUUCGGCCU-----GCUCCCUGAGACCUCAAGU---GUGAGCGCACUGUUG----AUGCUUCACGCCUGGGCUCUCUGGGUACCAGGACG---GCUUGAG--
drme-RF00052-372    G--------UUU--GUAUGGCUGAUUCCCUGAGACCCUAACU--UGUGAC----UUUUAAUAC-CAGUUUCACAAGUUUUGAUCUCCGGUAUUGGACGC---AAACUU---
drme-RF00052-375    G--------UUU--GUAUGGCUGAUUCCCUGAGACCCUAACU--UGUGAC----UUUUAAUAC-CAGUUUCACAAGUUUUGAUCUCCGGUAUUGGACGC---AAACUU---
gogo-RF00052-369    ---CCAGACUUU--UCCU-----AGUCCCUGAGACCCUAACU--UGUGAG-GUAUUUUAG----UAACAUCACAAGUCAGGCUCUUGGGACCUAGGCGG---AGGGGAA--
hosa-RF00052-368    UUGCGCUCCU---------CUCAGUCCCUGAGACCCUAACU--UGUGAU---GUUUACCGUUUAAAUCCACGGGUUAGGCUCUUGGGAGCUGCGAGUCGUGCUU-----
hosa-RF00052-370    UUGCGCUCCU---------CUCAGUCCCUGAGACCCUAACU--UGUGAU---GUUUACCGUUUAAAUCCACGGGUUAGGCUCUUGGGAGCUGCGAGUCGUGCUU-----
hosa-RF00052-371    UUGCCAGUCU---------CUAGGUCCCUGAGACCCUuuaccUGUGAG-Gac---A---UC-CAGGGUCACAGGUGAGGUUCUUGGGAGCCUGGCG-----UCUGgccc-
hosa-RF00052-374    ---CCAGACUUU--UCCU-----AGUCCCUGAGACCCUAACU--UGUGAG-GUAUUUUAG----UAACAUCACAAGUCAGGCUCUUGGGACCUAGGCGG---AGGGGAA--
mumu-RF00052-377    cg---AGACUUU--GCCU-----AGUCCCUGAGACCCUAACU--UGUGAG-GUAUUUUAG----UAACAUCACAAGUCAGGUUCUUGGGACCUAGGCGG---AGGGGaua-
#=GC SS_cons        .................(((.....(((.(((((((.(((.((...(((((((..........))))).)).))).))))).)))))).))))))..........
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