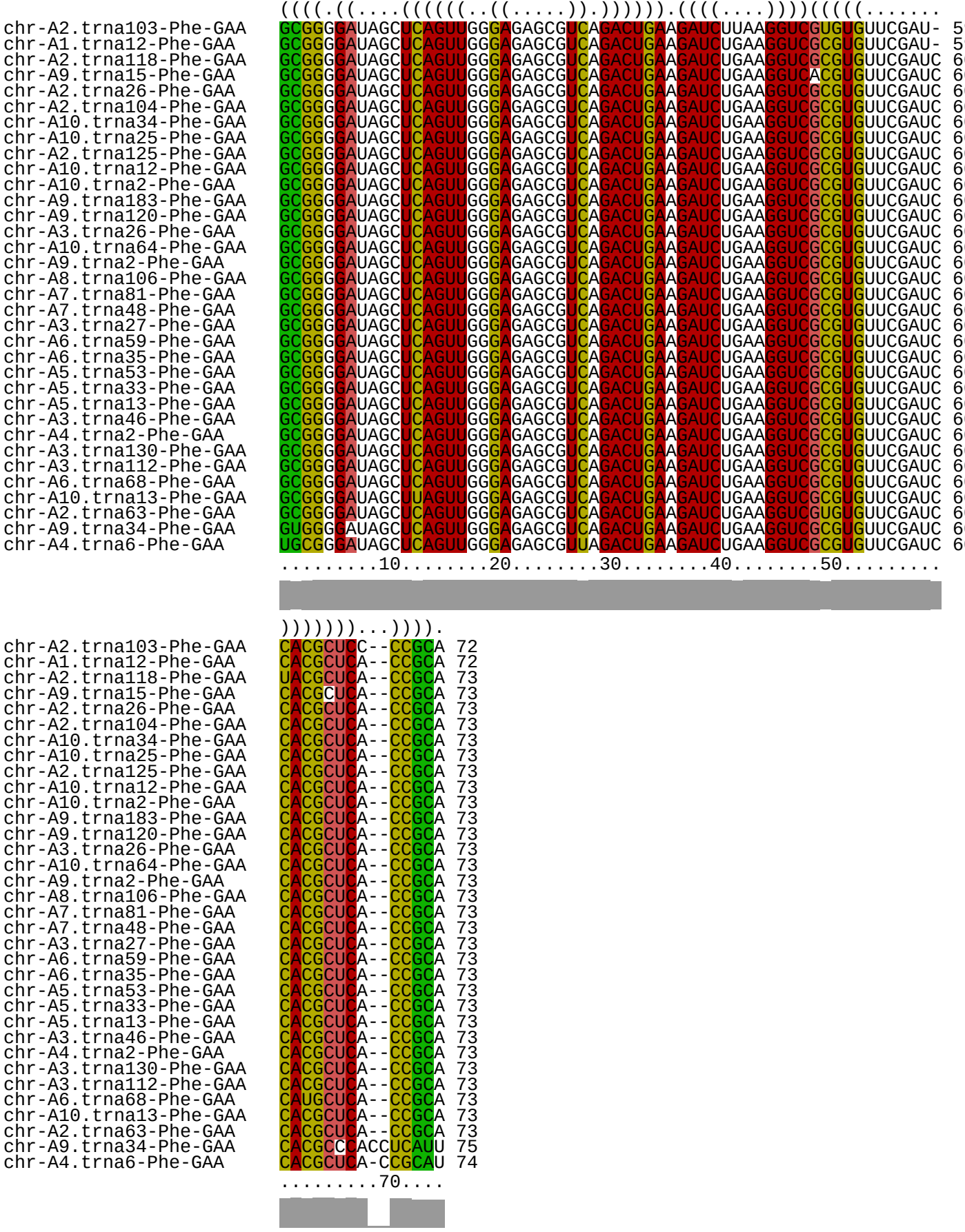
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chr-A1.trna12-Phe-GAA GCGGGGAUAGCUCAGUUGGGAGAGCGUCAGACUGAAGAUCUGAAGGUCGCGUGUUCGAU- 59
chr-A2.trna118-Phe-GAA GCGGGGAUAGCUCAGUUGGGAGAGCGUCAGACUGAAGAUCUGAAGGUCGCGUGUUCGAUC 60
chr-A9.trna15-Phe-GAA GCGGGGAUAGCUCAGUUGGGAGAGCGUCAGACUGAAGAUCUGAAGGUCACGUGUUCGAUC 60
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chr-A2.trna118-Phe-GAA UACGCUCA--CCGCA 73
chr-A9.trna15-Phe-GAA CACGCUCA--CCGCA 73
chr-A2.trna26-Phe-GAA CACGCUCA--CCGCA 73
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