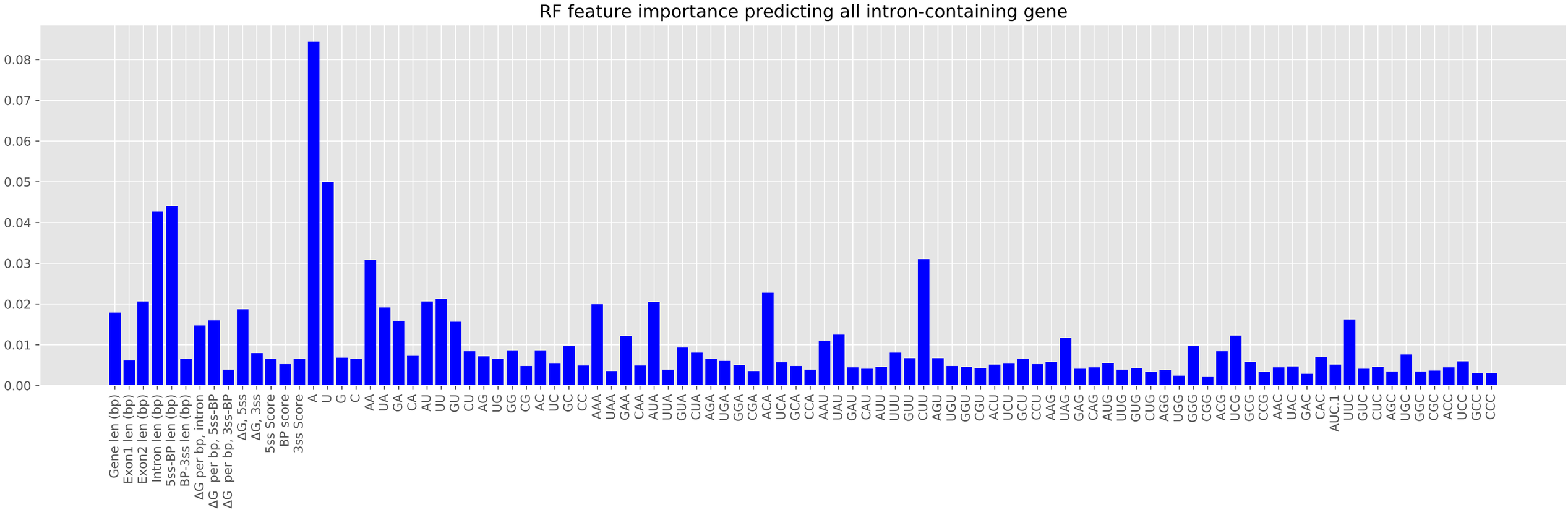
RF feature importance predicting all intron-containing gene
0.08
0.07
0.06
0.05
0.04
0.03
0.02
0.01
0.00
Gene len (bp)
Exon1 len (bp)
Exon2 len (bp)
Intron len (bp)
5ss-BP len (bp)
BP-3ss len (bp)
ΔG per bp, intron
ΔG per bp, 5ss-BP
ΔG per bp, 3ss-BP
ΔG, 5ss
ΔG, 3ss
5ss Score
BP score
3ss Score
A
U
G
C
AA
UA
GA
CA
AU
UU
GU
CU
AG
UG
GG
CG
AC
UC
GC
CC
AAA
UAA
GAA
CAA
AUA
UUA
GUA
CUA
AGA
UGA
GGA
CGA
ACA
UCA
GCA
CCA
AAU
UAU
GAU
CAU
AUU
UUU
GUU
CUU
AGU
UGU
GGU
CGU
ACU
UCU
GCU
CCU
AAG
UAG
GAG
CAG
AUG
UUG
GUG
CUG
AGG
UGG
GGG
CGG
ACG
UCG
GCG
CCG
AAC
UAC
GAC
CAC
AUC.1
UUC
GUC
CUC
AGC
UGC
GGC
CGC
ACC
UCC
GCC
CCC