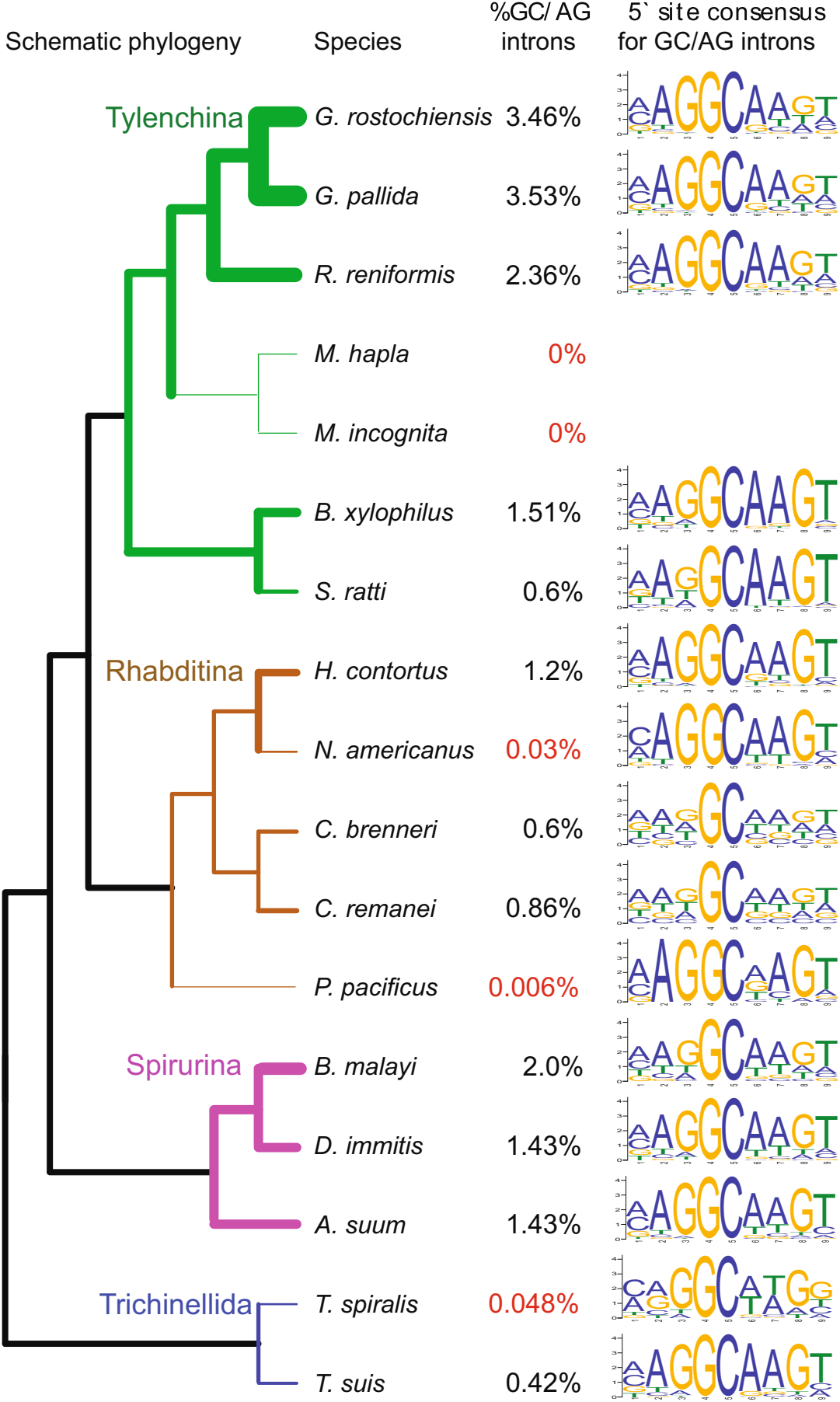
Schematic phylogeny
Species
%GC/ AG introns
5` site consensus for GC/AG introns
Tylenchina
G. rostochiensis 3.46%
G. pallida 3.53%
R. reniformis 2.36%
M. hapla 0%
M. incognita 0%
B. xylophilus 1.51%
S. ratti 0.6%
Rhabditina
H. contortus 1.2%
N. americanus 0.03%
C. brenneri 0.6%
C. remanei 0.86%
P. pacificus 0.006%
Spirurina
B. malayi 2.0%
D. immitis 1.43%
A. suum 1.43%
Trichinellida
T. spiralis 0.048%
T. suis 0.42%