

CAA AA A AAGT T G G A C TT T ATAT T TTA T TTT T G G A G C T T G A G C T G G A A T A G T T G G A A C A T C T T T A A G A A T T T T A A T
 C K K AA G AAGT T G G A C TT T ATAT T TTA T TTT T G G A G C T T G A G C T G G A A T A G T T G G A A C A T C T T T A A G A A T T T T A A T



T C G AGC T G AAT T A GG AC AT CC T G G AGC AT TA AT T G G A GAT G AT C AAA T T TA T AA T G TA A T T G T AA C T G C A CA T GC T T T T A

77

T A T A A T T T T T T T A T A G T T A T A C C T A T T A T A A T T G G T G G A T T T G G A A A T T G A T A G T G C C T T T A A T A T T A G G T G C T C C T
T A T A A T T T T T T T A T A G T T A T A C C T A T T A T A A T T G G T G G A T T T G G A A A T T G A T A G T G C C T T T A A T A T T A G G T G C T C C T

58

ATATAGCATTTCCACGAATAAATAATATAAGATTTTGACTTCTACCTCC TGTCTTTCTTTACATATAGTAAGTAGAATA
 ATATAGCATTTCCACGAATAAATAATATAA AATTTTGACTTCTACCTCCGGTCTTTCTTTACAAATAGTAAGGTAA AAAA

238

T T G A A A T G G A G C T G G G A C A G G A T G A A C T G T T T A T C C A C C T C T A T C C G C T G G A A T T G C T C A T G G T G G A G C T T C A G T T G A T
T T G A A A T G G A G C T G G G A C A G G A T G A A C T G T T T A T C C A C C T C T A T C C G C G G G A A T T G C T C A G G G G G G A G C T T C A G T T G A T

319

TT AGC TAT TTT TT CT CT AC AT TTA G AGG AA TTTT TT CAA TTT T AGG AGCT GT AAA TTT TATT AC AAC TG TAA TT A AT AT
TT A CC TAT TTT TT CT CT AC AT TTA G AGG AA TTTT TT CAA TTT T AGG AGCT GT AAA TTT TATT AC AAC TG TAA TT A AT AT

669

C G A T C A A C A G G A A T T T C A T T A G A T C G T A T A C C T T T A T T T G T T T G A T C A G T A G T T A T T A C T G C T T T A T T A T T A T T A T T A T C
 C G A T C A A C A G G A A T T T C A T T A G A T C G T A T A C C T T T A T T T G T T T G A T C A G T A G T T A T T A C T G C T T T A T T A T T A T T A T T A T C

180

A C T T C G A G T A C T A G C A C G A G C T A T G A C T A T A T G A T T A A C A G A T C G A A G T T G A A A T A C A T C A A T T T T C G A C C T A G C G G C A G
 A C T T C T A G T A C T A G C A C G A G C T A T G A C T A T A T G A T T A A C A G A T C G A A G T T G A A A T A C A T C A S T T T T A G A C C T A G C G G C A G

560

A GG AG A TC G T A T T T A AT AC G AG CA T T G AT T T C G AG T T T T AG G TC A C C G T GTA G TTC A GT C ATA G C TGA TT ACG T G A
 A GG AG A TC G T A T T T A AT AC T AG CA T T C AT T T C G AG T T T T AG G TC A C C G T GCA G TTC A AT C ATA G C RGA TT GCA A G A

541