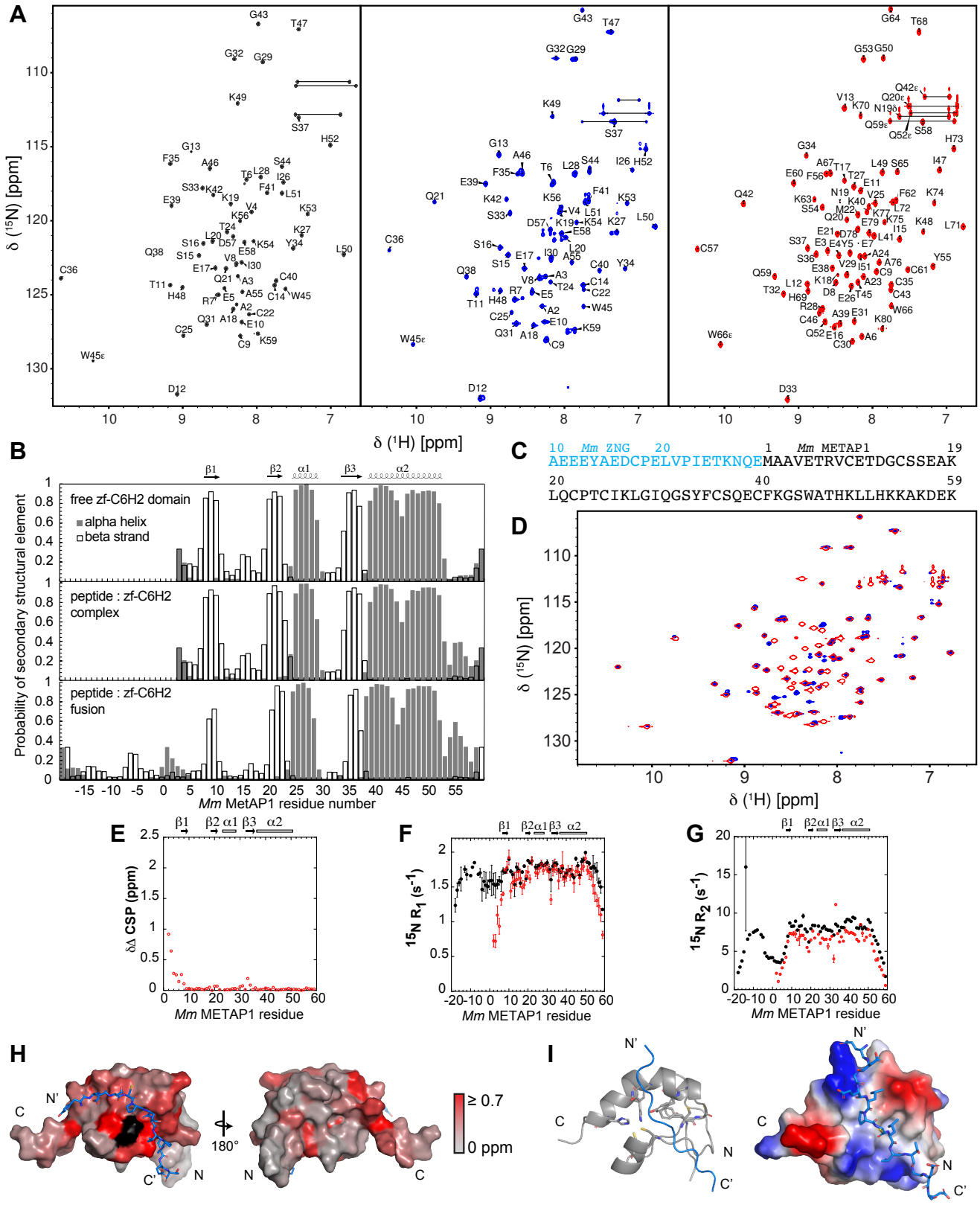
A
δ (15N) [ppm]
δ (1H) [ppm]
B
free zf-C6H2 domain
alpha helix
beta strand
peptide : zf-C6H2 complex
peptide : zf-C6H2 fusion
Probability of secondary structural element
Mm MetAP1 residue number
C
10 Mm ZNG 20 1 Mm METAP1 19
AEEEYAEDCPELVPIETKNQEMAAVETRVCETDGCSSEAK
20 40 59
LQCPTCIKLGIQGSYFCSQECFKGSWATHKLLHKKAKDEK
D
δ (15N) [ppm]
δ (1H) [ppm]
E
δΔ CSP (ppm)
Mm METAP1 residue
F
15N R1 (s-1)
Mm METAP1 residue
G
15N R2 (s-1)
Mm METAP1 residue
H
180°
≥ 0.7
0 ppm
I