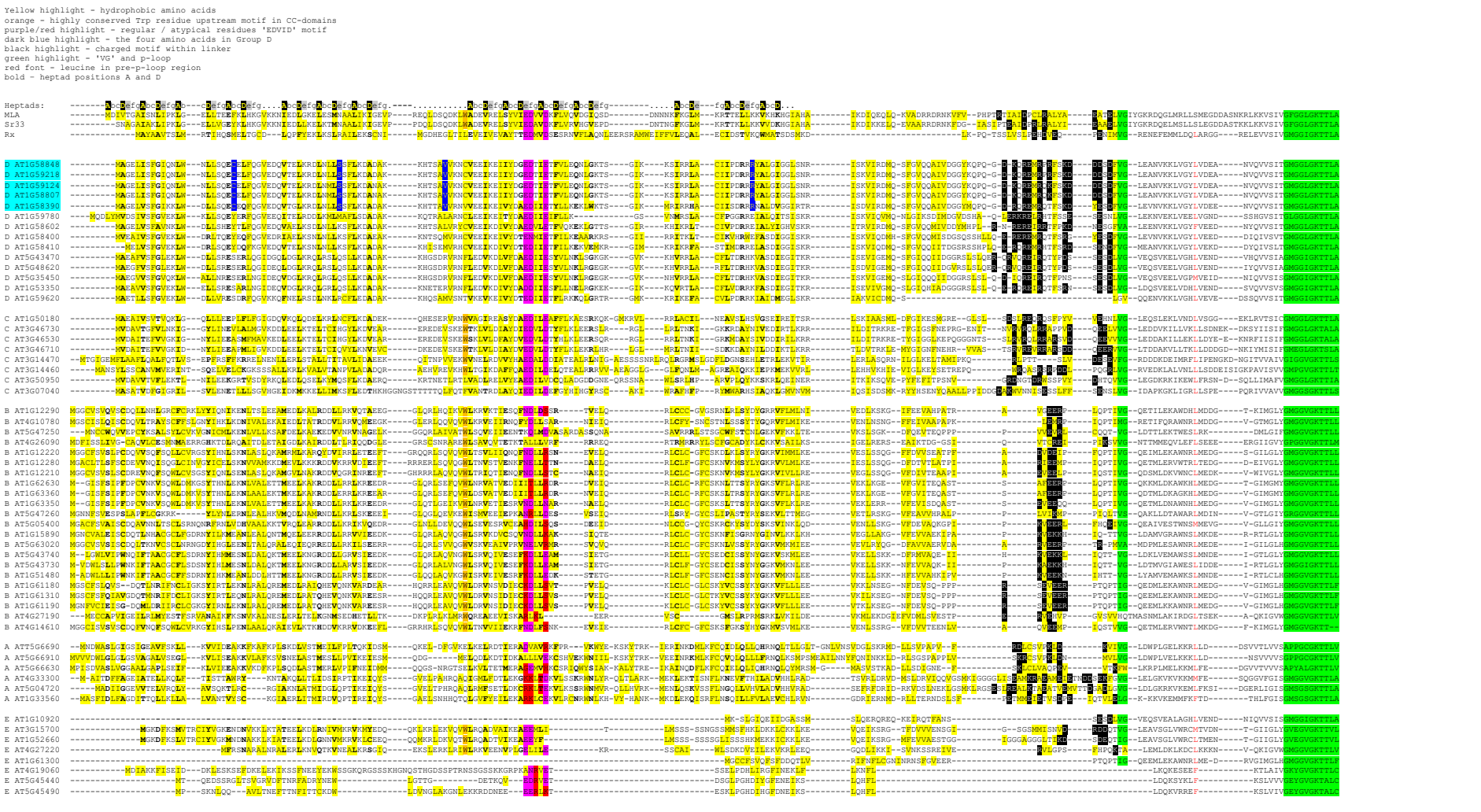
Yellow highlight - hydrophobic amino acids
orange - highly conserved Trp residue upstream motif in CC-domains
purple/red highlight - regular / atypical residues 'EDVID' motif
dark blue highlight - the four amino acids in Group D
black highlight - charged motif within linker
green highlight - 'VG' and p-loop
red font - leucine in pre-p-loop region
bold - heptad positions A and D
Heptads:
MLA
Sr33
Rx
D AT1G58848
D AT1G59218
D AT1G59124
D AT1G58807
D AT1G58390
D AT1G59780
D AT1G58602
D AT1G58400
D AT1G58410
D AT5G43470
D AT5G48620
D AT5G35450
D AT1G53350
D AT1G59620
C AT1G50180
C AT3G46730
C AT3G46530
C AT3G46710
C AT3G14470
C AT3G14460
C AT3G50950
C AT3G07040
B AT1G12290
B AT4G10780
B AT5G47250
B AT4G26090
B AT1G12220
B AT1G12280
B AT1G12210
B AT1G62630
B AT1G60330
B AT1G62350
B AT5G47260
B AT5G05400
B AT1G15890
B AT5G63020
B AT5G43740
B AT5G43730
B AT1G51480
B AT1G61180
B AT1G61310
B AT1G61190
B AT4G27190
B AT4G14610
A ATT5G66690
A AT5G66910
A AT5G66630
A AT4G33300
A AT5G04720
A AT1G33560
E AT1G10920
E AT3G15700
E AT1G52660
E AT4G27220
E AT1G61300
E AT4G19060
E AT5G45440
E AT5G45490