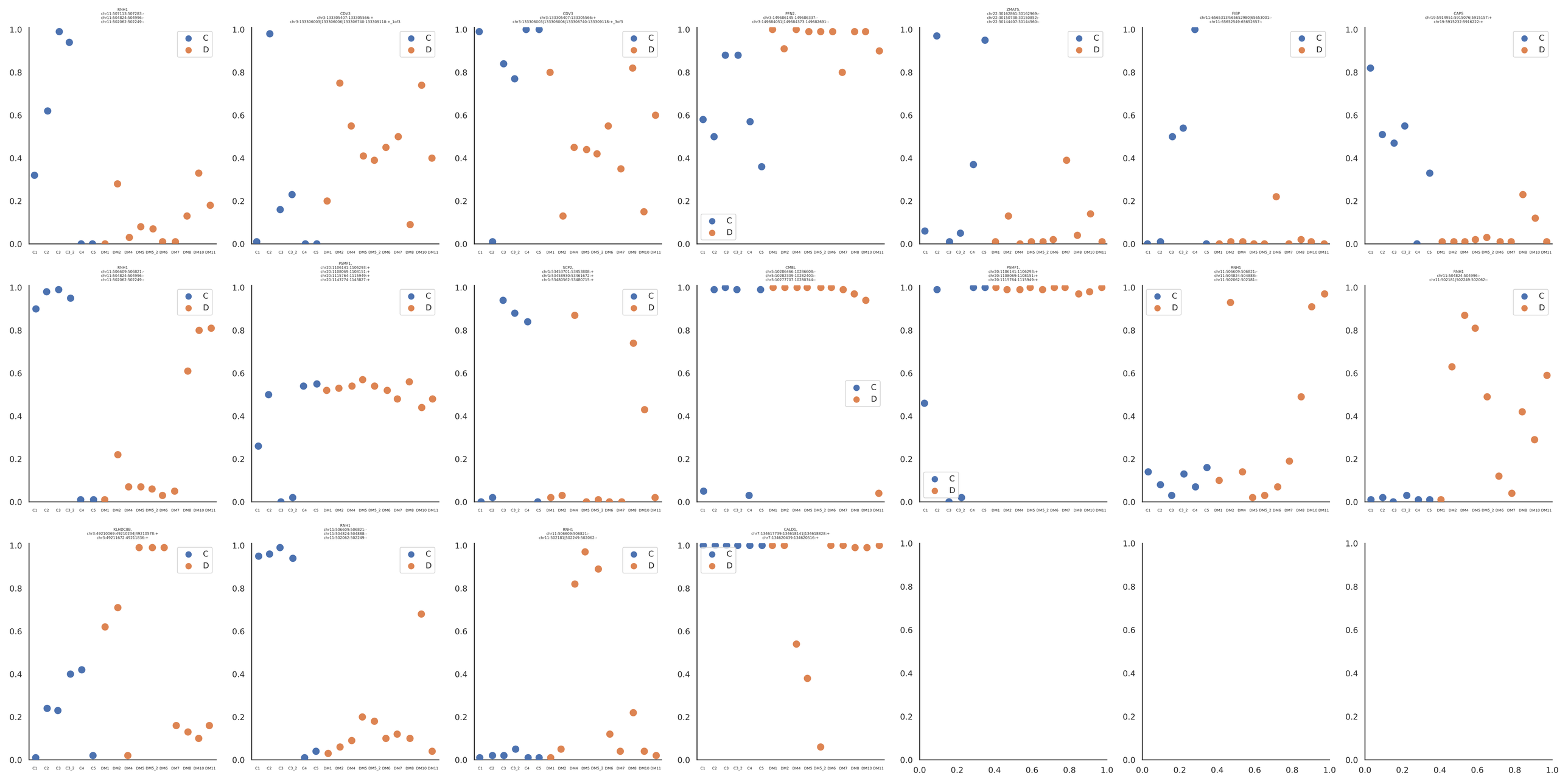
RNH1
chr11:507113:507283:-
chr11:504824:504996:-
chr11:502062:502249:-
CDV3
chr3:133305407:133305566:+
chr3:133306003|133306006|133306740:133309118:+_1of3
CDV3
chr3:133305407:133305566:+
chr3:133306003|133306006|133306740:133309118:+_3of3
PFN2,
chr3:149686145:149686337:-
chr3:149684053|149684373:149682691:-
ZMATS,
chr22:30162861:30162969:-
chr22:30150738:30150852:-
chr22:30144407:30144660:-
FIBP
chr11:65653134:65652980|65653001:-
chr11:65652549:65652657:-
CAPS
chr19:5914951:5915076|5915157:+
chr19:5915232:5916222:+
RNH1
chr11:506609:506821:-
chr11:504824:504996:-
chr11:502062:502249:-
PSMF1,
chr20:1106141:1106293:+
chr20:1108069:1108151:+
chr20:1115764:1115949:+
chr20:1143774:1143827:+
SCP2,
chr1:53453701:53453808:+
chr1:53458930:53461672:+
chr1:53480562:53480715:+
CMBL
chr5:10286466:10286608:-
chr5:10282309:10282400:-
chr5:10277707:10280744:-
PSMF1,
chr20:1106141:1106293:+
chr20:1108069:1108151:+
chr20:1115764:1115949:+
RNH1
chr11:506609:506821:-
chr11:504824:504888:-
chr11:502062:502181:-
RNH1
chr11:504824:504996:-
chr11:502181|502249:502062:-
KLHDC8B,
chr3:49210069:49210234|49210578:+
chr3:49211672:49211836:+
RNH1
chr11:506609:506821:-
chr11:504824:504888:-
chr11:502062:502249:-
RNH1
chr11:506609:506821:-
chr11:502181|502249:502062:-
CALD1,
chr7:134617739:134618141|134618828:+
chr7:134620439:134620516:+
C
D
0.0
0.2
0.4
0.6
0.8
1.0
C1
C2
C3
C3_2
C4
C5
DM1
DM2
DM4
DM5
DM5_2
DM6
DM7
DM8
DM10
DM11