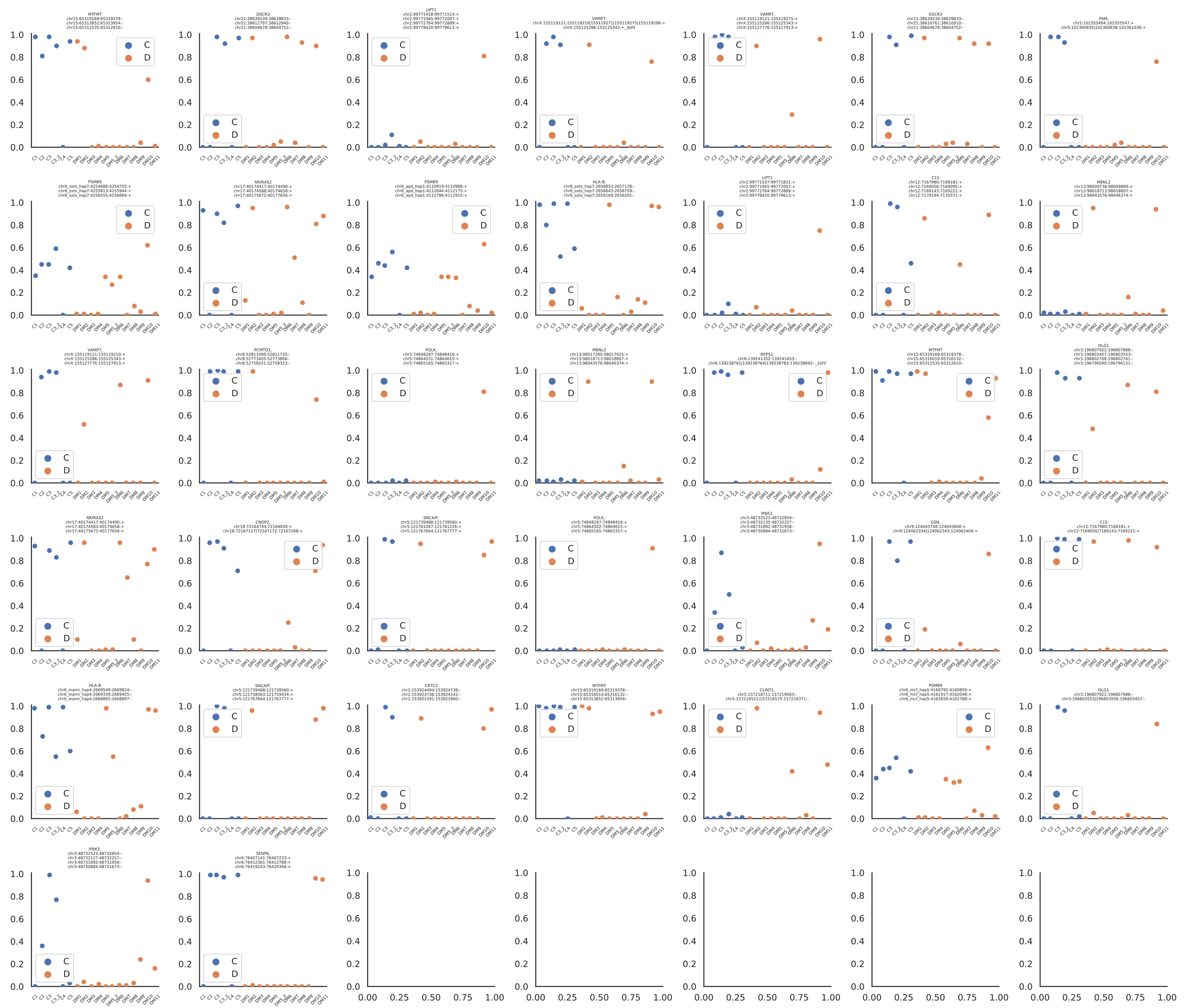
MTFMT
chr15:65319169:65319378:-
chr15:65313852:65313954:-
chr15:65312535:65312610:-
DSCR3
chr21:38639539:38639833:-
chr21:38612797:38612940:-
chr21:38604678:38604752:-
LIPT1
chr2:99771418:99771514:+
chr2:99771945:99772007:+
chr2:99772764:99772889:+
chr2:99778420:99779613:+
VAMP7,
chrX:155119121:155119210|155119271|155119275|155119296:+
chrX:155125286:155125343:+_3of4
VAMP7,
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chrX:155125286:155125343:+
chrX:155127776:155127913:+
DSCR3
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chr21:38610761:38610910:-
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PAM,
chr5:102355494:102355547:+
chr5:102360835|102360838:102361038:+
PSMB9
chr6_ssto_hap7:4254688:4254755:+
chr6_ssto_hap7:4255813:4255944:+
chr6_ssto_hap7:4256555:4256684:+
NKIRAS2
chr17:40174417:40174490:+
chr17:40174588:40174658:+
chr17:40175672:40177656:+
PSMB9
chr6_apd_hap1:4110919:4110986:+
chr6_apd_hap1:4112044:4112175:+
chr6_apd_hap1:4112786:4112915:+
HLA-B,
chr6_ssto_hap7:2656853:2657128:-
chr6_ssto_hap7:2656643:2656759:-
chr6_ssto_hap7:2656169:2656201:-
LIPT1
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chr2:99771945:99772007:+
chr2:99772764:99772889:+
chr2:99778420:99779613:+
C1S
chr12:7167980:7168181:+
chr12:7169056:7169095:+
chr12:7169143:7169221:+
chr12:7170194:7170371:+
MBNL2
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chr13:98018713:98018807:+
chr13:98043576:98046374:+
VAMP7,
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chrX:155127776:155127913:+
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chr8:52758221:52758323:-
POLK,
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chr15:65312535:65312610:-
DLG1
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chr3:196803457:196803553:-
chr3:196802708:196802741:-
chr3:196796090:196796131:-
NKIRAS2
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chr17:40174583:40174658:+
chr17:40175672:40177656:+
CNDP2,
chr18:72164744:72164830:+
chr18:72167117|72167172:72167268:+
SNCAIP,
chr5:121739488:121739560:+
chr5:121761047:121761226:+
chr5:121767664:121767777:+
POLK,
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chr5:74864502:74864615:+
chr5:74865165:74865317:+
IP6K2,
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chr3:48732135:48732257:-
chr3:48731892:48731958:-
chr3:48730884:48731673:-
GSN,
chr9:124043748:124043840:+
chr9:124062334|124062343:124062404:+
C1S
chr12:7167980:7168181:+
chr12:7169056|7169143:7169221:+
HLA-B
chr6_mann_hap4:2669549:2669824:-
chr6_mann_hap4:2669339:2669455:-
chr6_mann_hap4:2668865:2668897:-
SNCAIP,
chr5:121739488:121739560:+
chr5:121758563:121759434:+
chr5:121767664:121767777:+
CRTC2
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chr1:153923736:153924142:-
chr1:153921591:153921860:-
MTFMT
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chr15:65316010:65316132:-
chr15:65313852:65313954:-
CLINT1,
chr5:157218711:157219003:-
chr5:157216521|157216575:157216371:-
PSMB9
chr6_mcf_hap5:4160792:4160859:+
chr6_mcf_hap5:4161917:4162048:+
chr6_mcf_hap5:4162659:4162788:+
DLG1
chr3:196807922:196807988:-
chr3:196803553|196803556:196803457:-
IP6K2,
chr3:48732523:48732854:-
chr3:48732127:48732257:-
chr3:48731892:48731958:-
chr3:48730884:48731673:-
SENP6,
chr6:76407141:76407233:+
chr6:76412361:76412788:+
chr6:76419243:76419346:+
C
D
C1 C2 C3 C3_2 C4 C5 DM1 DM2 DM3 DM4 DM5 DM5_2 DM6 DM7 DM8 DM9 DM10 DM11
0.0 0.2 0.4 0.6 0.8 1.0
0.00 0.25 0.50 0.75 1.00