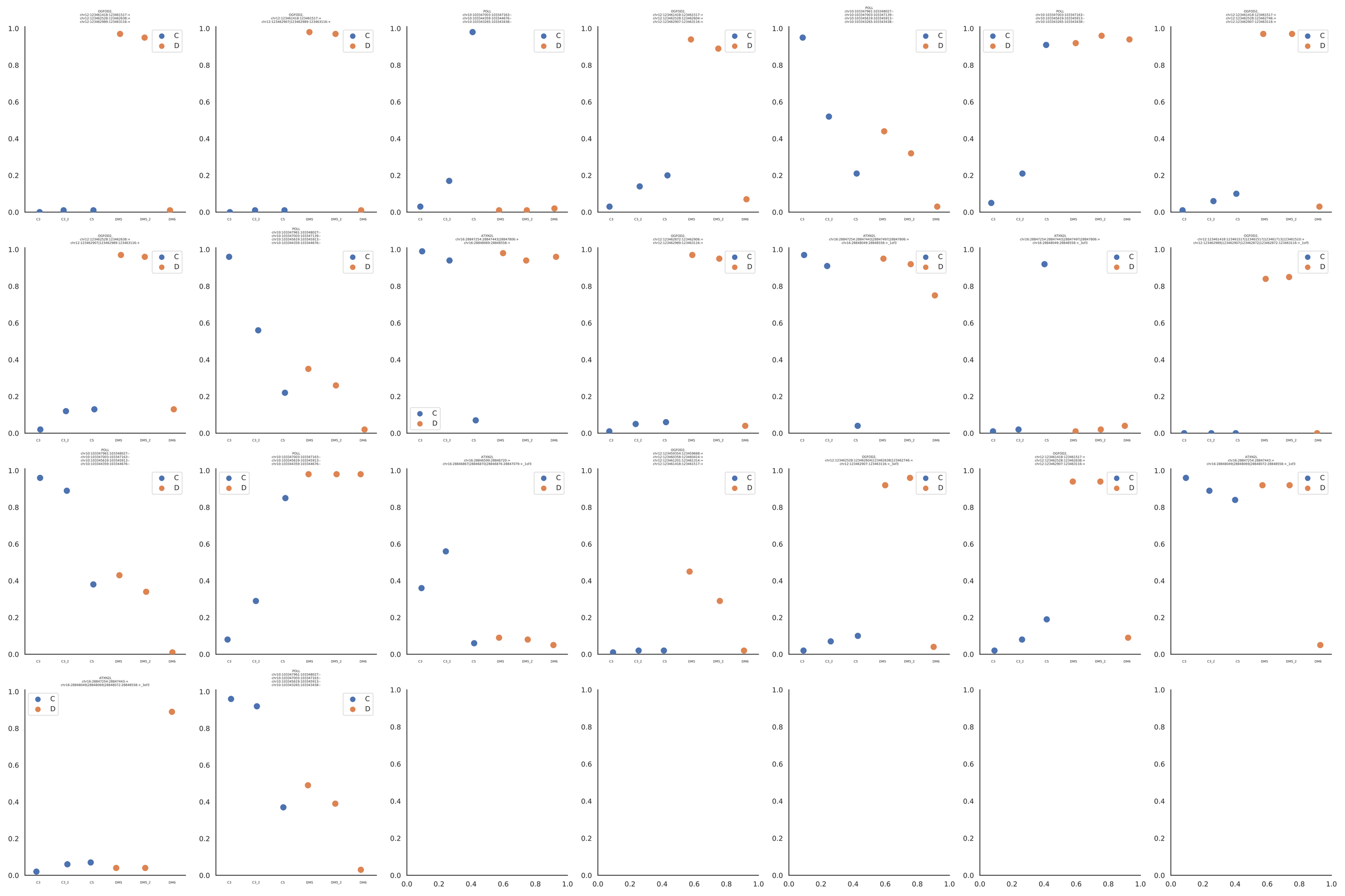
OGFOD2, chr12:123461418:123461517:+ chr12:123462528:123462638:+ chr12:123462989:123463116:+
OGFOD2, chr12:123461418:123461517:+ chr12:123462907|123462989:123463116:+
POLL chr10:103347003:103347163:- chr10:103344359:103344676:-
OGFOD2, chr12:123461418:123461517:+ chr12:123462528:123462604:+ chr12:123462907:123463116:+
POLL chr10:103347961:103348027:- chr10:103347003:103347139:- chr10:103345619:103345913:- chr10:103343265:103343438:-
POLL chr10:103347023:103347163:- chr10:103345619:103345913:- chr10:103343265:103343438:-
OGFOD2, chr12:123461418:123461517:+ chr12:123462528:123462746:+ chr12:123462907:123463116:+
OGFOD2, chr12:123462528:123462638:+ chr12:123462907|123462989:123463116:+
POLL chr10:103347961:103348027:- chr10:103347003:103347139:- chr10:103345619:103345913:- chr10:103344359:103344676:-
ATXN2L chr16:28847254:28847443|28847806:+ chr16:28848069:28848558:+
OGFOD2, chr12:123462872:123462906:+ chr12:123462989:123463116:+
ATXN2L chr16:28847254:28847443|28847497|28847806:+ chr16:28848049:28848558:+_1of3
ATXN2L chr16:28847254:28847443|28847497|28847806:+ chr16:28848049:28848558:+_3of3
OGFOD2, chr12:123461418:123461517|123461713|123461510:+ chr12:123462989|123462907|123462872|123462872:123463116:+_1of5
POLL chr10:103347961:103348027:- chr10:103347003:103347163:- chr10:103345619:103345913:- chr10:103344359:103344676:-
POLL chr10:103347003:103347163:- chr10:103345619:103345913:- chr10:103344359:103344676:-
ATXN2L chr16:28846599:28846720:+ chr16:28846867|28846870|28846876:28847079:+_1of3
OGFOD2, chr12:123459354:123459668:+ chr12:123460358:123460414:+ chr12:123461201:123461314:+ chr12:123461418:123461517:+
OGFOD2, chr12:123462528:123462604|123462638|123462746:+ chr12:123462907:123463116:+_3of3
OGFOD2, chr12:123461418:123461517:+ chr12:123462528:123462638:+ chr12:123462907:123463116:+
ATXN2L chr16:28847254:28847443:+ chr16:28848049|28848069|28848072:28848558:+_1of3
ATXN2L chr16:28847254:28847443:+ chr16:28848049|28848069|28848072:28848558:+_3of3
POLL chr10:103347961:103348027:- chr10:103347003:103347163:- chr10:103345619:103345913:- chr10:103343265:103343438:-
C
D
C3
C3_2
C5
DM5
DM5_2
DM6
0.0
0.2
0.4
0.6
0.8
1.0