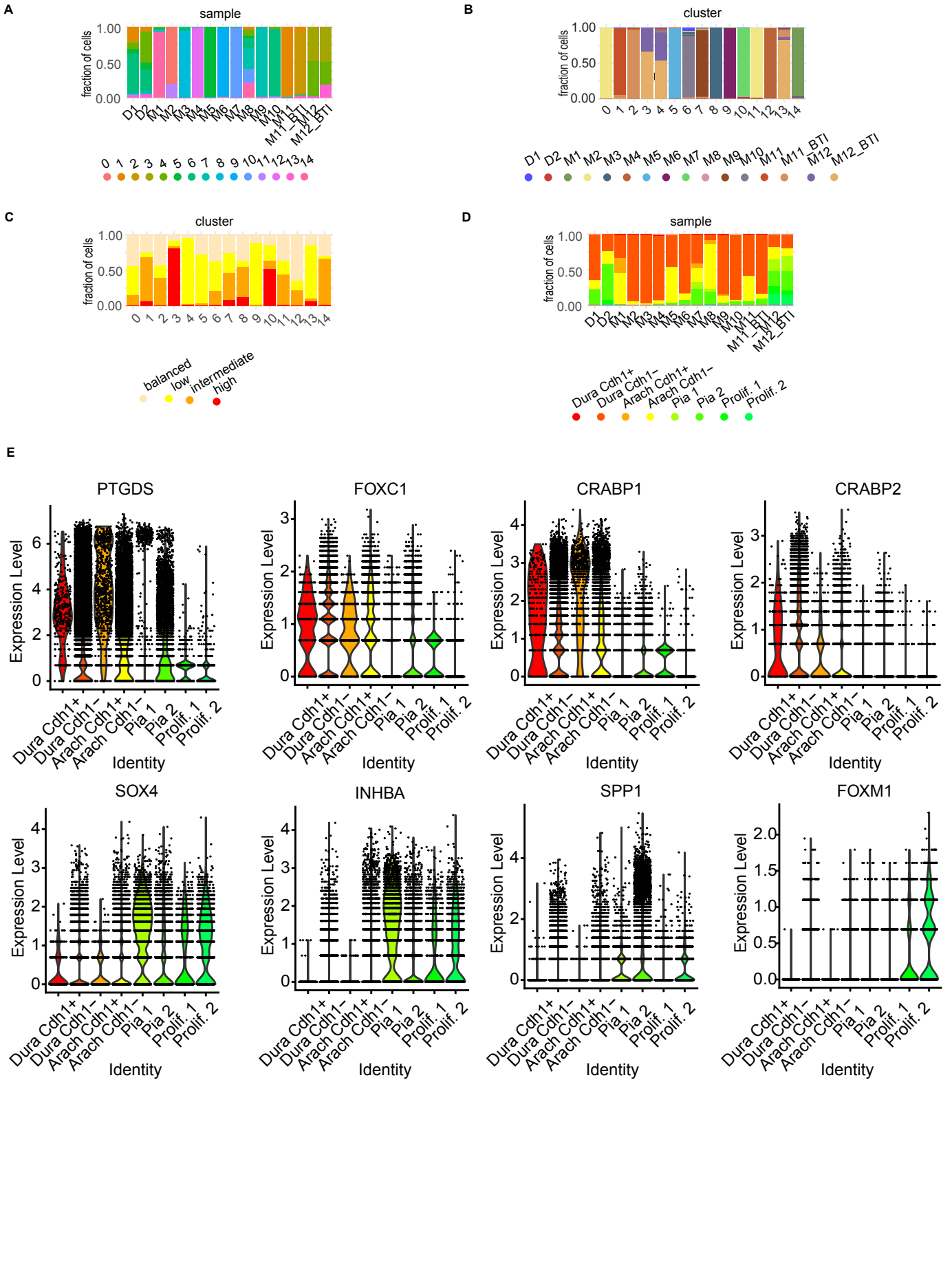
A
sample
fraction of cells
1.00
0.50
0.00
D1 D2 M1 M2 M3 M4 M5 M6 M7 M8 M9 M10 M11 M11_BTI M12 M12_BTI
0 1 2 3 4 5 6 7 8 9 10 11 12 13 14
B
cluster
fraction of cells
1.00
0.50
0.00
0 1 2 3 4 5 6 7 8 9 10 11 12 13 14
D1 D2 M1 M2 M3 M4 M5 M6 M7 M8 M9 M10 M11 M11_BTI M12 M12_BTI
C
cluster
fraction of cells
1.00
0.50
0.00
0 1 2 3 4 5 6 7 8 9 10 11 12 13 14
balanced
low
intermediate
high
D
sample
fraction of cells
1.00
0.50
0.00
D1 D2 M1 M2 M3 M4 M5 M6 M7 M8 M9 M10 M11 M11_BTI M12 M12_BTI
Dura Cdh1+
Dura Cdh1−
Arach Cdh1+
Arach Cdh1−
Pia 1
Pia 2
Prolif. 1
Prolif. 2
E
PTGDS
FOXC1
CRABP1
CRABP2
SOX4
INHBA
SPP1
FOXM1
Expression Level
Identity
Dura Cdh1+
Dura Cdh1−
Arach Cdh1+
Arach Cdh1−
Pia 1
Pia 2
Prolif. 1
Prolif. 2