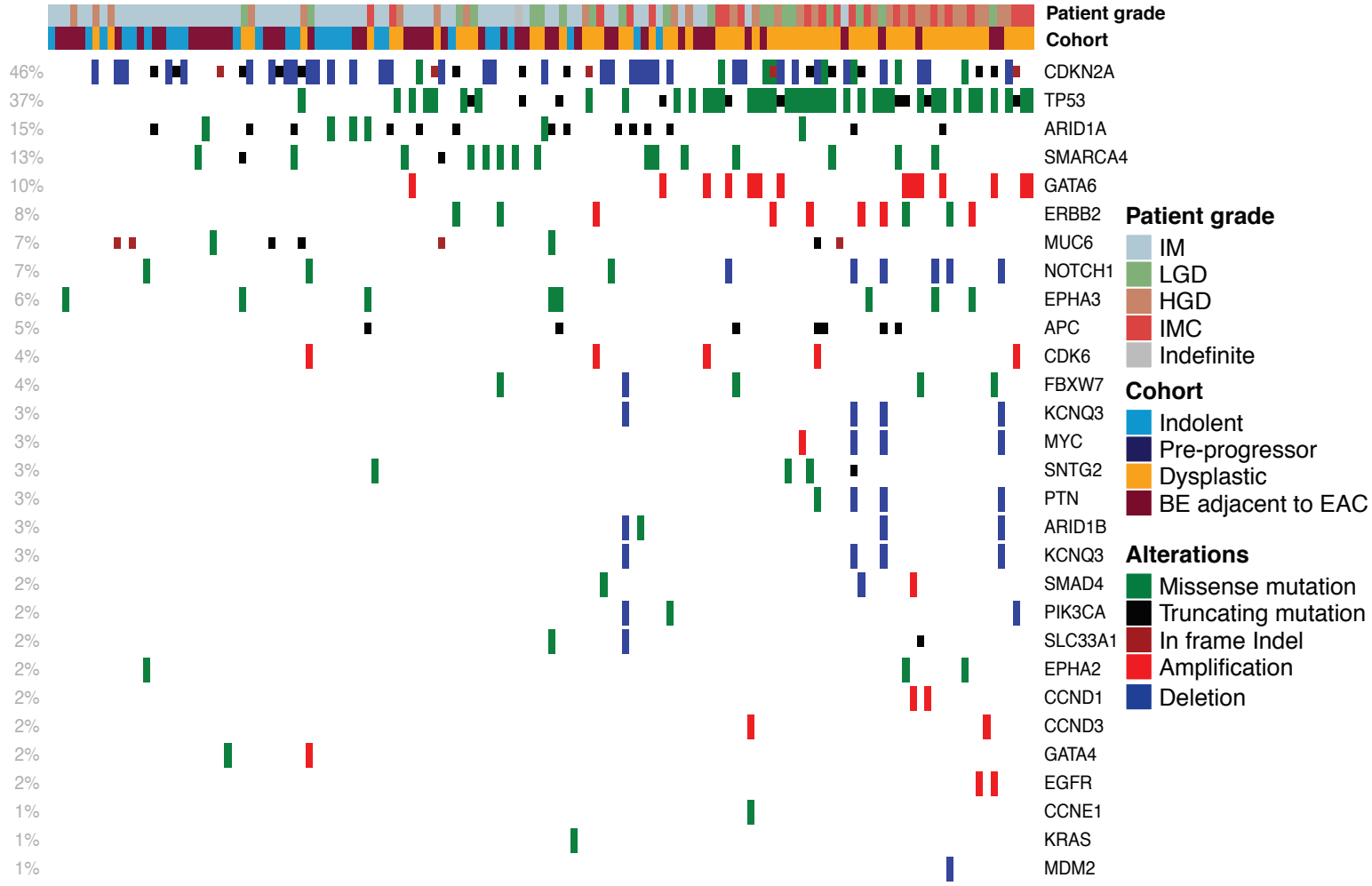
Figure S2


Patient grade
Cohort
46% CDKN2A
37% TP53
15% ARID1A
13% SMARCA4
10% GATA6
8% ERBB2
7% MUC6
7% NOTCH1
6% EPHA3
5% APC
4% CDK6
4% FBXW7
3% KCNQ3
3% MYC
3% SNTG2
3% PTN
3% ARID1B
3% KCNQ3
2% SMAD4
2% PIK3CA
2% SLC33A1
2% EPHA2
2% CCND1
2% CCND3
2% GATA4
2% EGFR
1% CCNE1
1% KRAS
1% MDM2
Patient grade
IM
LGD
HGD
IMC
Indefinite
Cohort
Indolent
Pre-progressor
Dysplastic
BE adjacent to EAC
Alterations
Missense mutation
Truncating mutation
In frame Indel
Amplification
Deletion