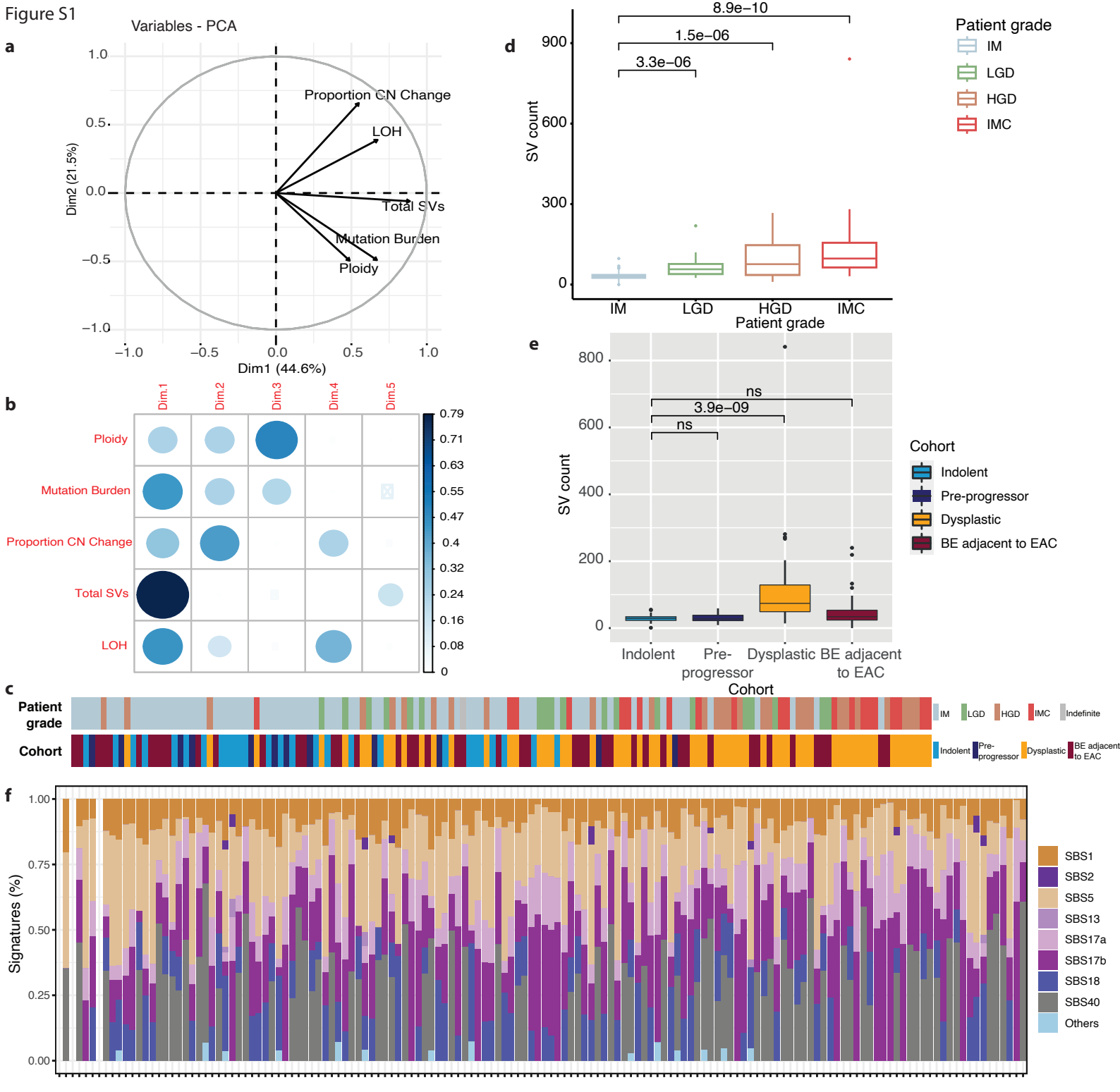
Figure S1


a
Variables - PCA
Proportion CN Change
LOH
Total SVs
Mutation Burden
Ploidy
Dim1 (44.6%)
Dim2 (21.5%)
b
Dim.1
Dim.2
Dim.3
Dim.4
Dim.5
Ploidy
Mutation Burden
Proportion CN Change
Total SVs
LOH
c
Patient grade
Cohort
IM
LGD
HGD
IMC
Indefinite
Indolent
Pre-progressor
Dysplastic
BE adjacent to EAC
d
8.9e−10
1.5e−06
3.3e−06
SV count
Patient grade
IM
LGD
HGD
IMC
e
ns
3.9e−09
ns
SV count
Cohort
Indolent
Pre-progressor
Dysplastic
BE adjacent to EAC
f
Signatures (%)
SBS1
SBS2
SBS5
SBS13
SBS17a
SBS17b
SBS18
SBS40
Others