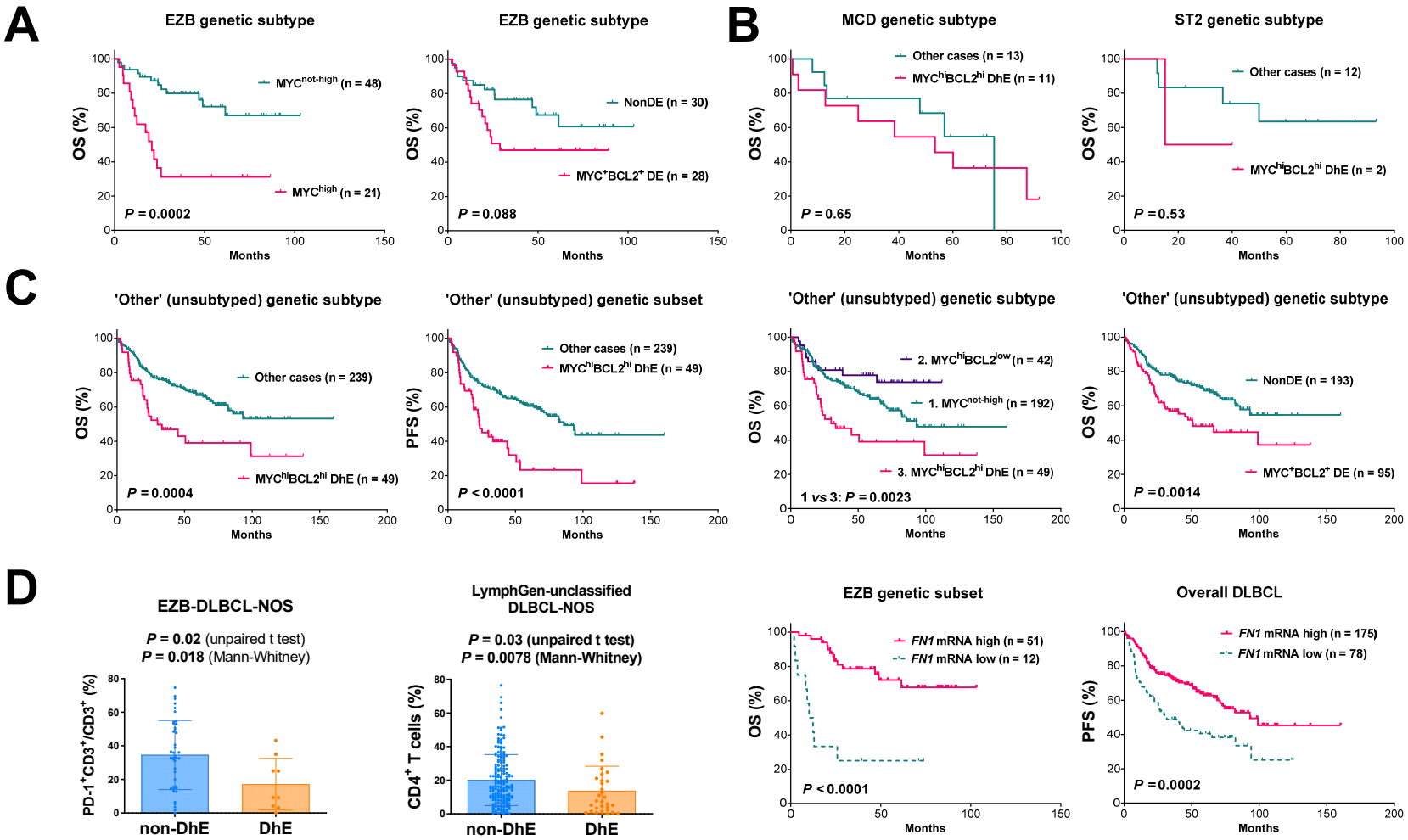
Figure S3


A
EZB genetic subtype
MYC^not-high (n = 48)
MYC^high (n = 21)
P = 0.0002
OS (%)
Months
EZB genetic subtype
NonDE (n = 30)
MYC+BCL2+ DE (n = 28)
P = 0.088
B
MCD genetic subtype
Other cases (n = 13)
MYC^hiBCL2^hi DhE (n = 11)
P = 0.65
ST2 genetic subtype
Other cases (n = 12)
MYC^hiBCL2^hi DhE (n = 2)
P = 0.53
C
'Other' (unsubtyped) genetic subtype
Other cases (n = 239)
MYC^hiBCL2^hi DhE (n = 49)
P = 0.0004
'Other' (unsubtyped) genetic subset
PFS (%)
P < 0.0001
'Other' (unsubtyped) genetic subtype
2. MYC^hiBCL2^low (n = 42)
1. MYC^not-high (n = 192)
3. MYC^hiBCL2^hi DhE (n = 49)
1 vs 3: P = 0.0023
'Other' (unsubtyped) genetic subtype
NonDE (n = 193)
MYC+BCL2+ DE (n = 95)
P = 0.0014
D
EZB-DLBCL-NOS
P = 0.02 (unpaired t test)
P = 0.018 (Mann-Whitney)
PD-1+CD3+/CD3+ (%)
non-DhE
DhE
LymphGen-unclassified DLBCL-NOS
P = 0.03 (unpaired t test)
P = 0.0078 (Mann-Whitney)
CD4+ T cells (%)
EZB genetic subset
FN1 mRNA high (n = 51)
FN1 mRNA low (n = 12)
P < 0.0001
Overall DLBCL
FN1 mRNA high (n = 175)
FN1 mRNA low (n = 78)
P = 0.0002