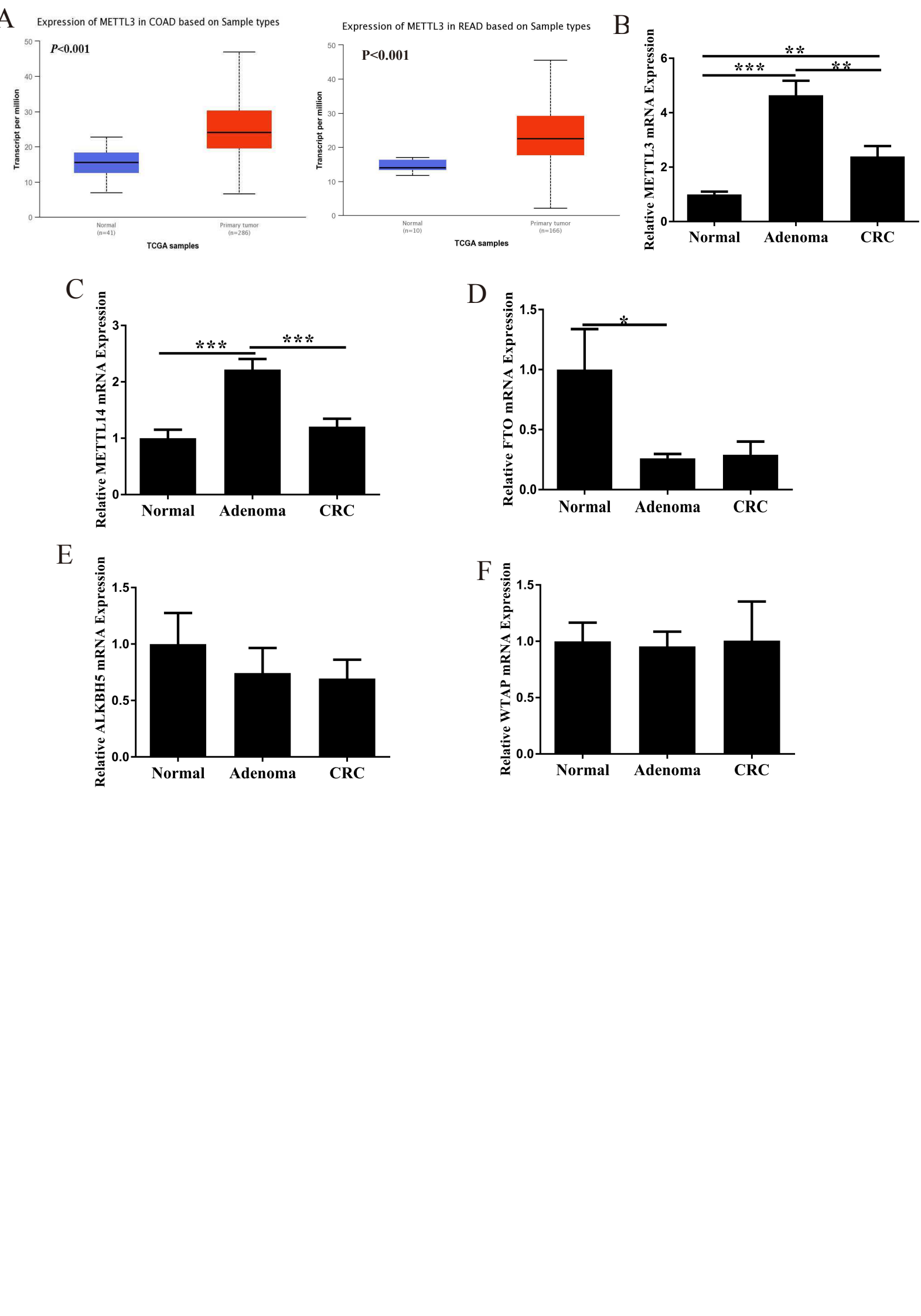
A
Expression of METTL3 in COAD based on Sample types
P<0.001
Transcript per million
Normal (n=41)
Primary tumor (n=286)
TCGA samples
Expression of METTL3 in READ based on Sample types
P<0.001
Transcript per million
Normal (n=10)
Primary tumor (n=166)
TCGA samples
B
Relative METTL3 mRNA Expression
**
***
**
Normal
Adenoma
CRC
C
Relative METTL14 mRNA Expression
***
***
Normal
Adenoma
CRC
D
Relative FTO mRNA Expression
*
Normal
Adenoma
CRC
E
Relative ALKBH5 mRNA Expression
Normal
Adenoma
CRC
F
Relative WTAP mRNA Expression
Normal
Adenoma
CRC