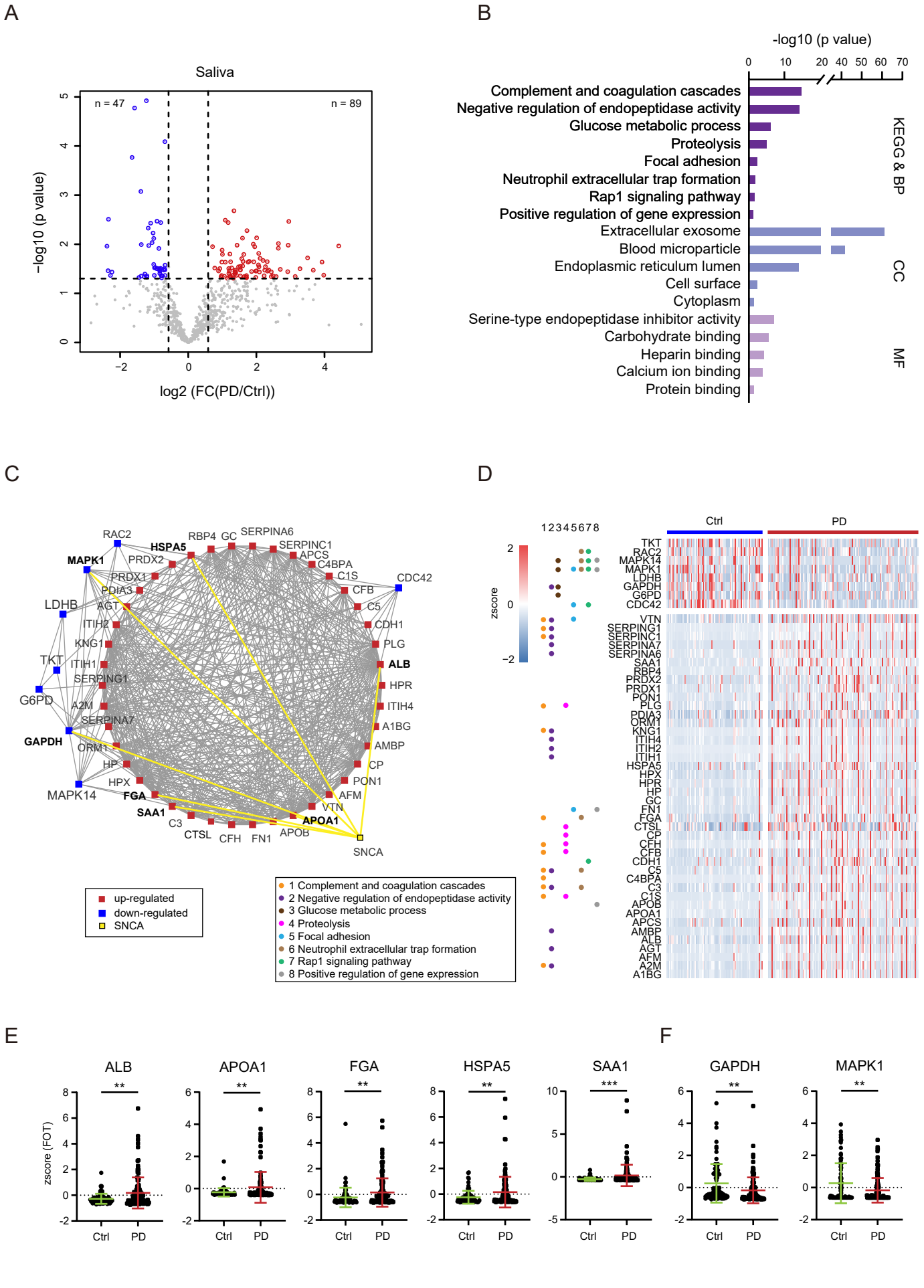
Figure S2


A
Saliva
n = 47
n = 89
−log10 (p value)
log2 (FC(PD/Ctrl))
B
-log10 (p value)
Complement and coagulation cascades
Negative regulation of endopeptidase activity
Glucose metabolic process
Proteolysis
Focal adhesion
Neutrophil extracellular trap formation
Rap1 signaling pathway
Positive regulation of gene expression
Extracellular exosome
Blood microparticle
Endoplasmic reticulum lumen
Cell surface
Cytoplasm
Serine-type endopeptidase inhibitor activity
Carbohydrate binding
Heparin binding
Calcium ion binding
Protein binding
KEGG & BP
CC
MF
C
up-regulated
down-regulated
SNCA
1 Complement and coagulation cascades
2 Negative regulation of endopeptidase activity
3 Glucose metabolic process
4 Proteolysis
5 Focal adhesion
6 Neutrophil extracellular trap formation
7 Rap1 signaling pathway
8 Positive regulation of gene expression
D
Ctrl
PD
zscore
E
ALB
APOA1
FGA
HSPA5
SAA1
zscore (FOT)
Ctrl
PD
F
GAPDH
MAPK1