[Skills] How to perform gene set enrichment analysis with python?

gseapy



	Gene	Rank
0	PPP4C	-5.175958
1	PRMT5	-4.945346
2	HDGFRP2	-3.997596
3	SMNDC1	-3.855049
4	PRMT1	-3.723365
1113	UBE3D	3.370299
1114	METTL3	3.446301
1115	ATR	3.623120
1116	METTL14	4.369509
1117	CCT7	4.561326

