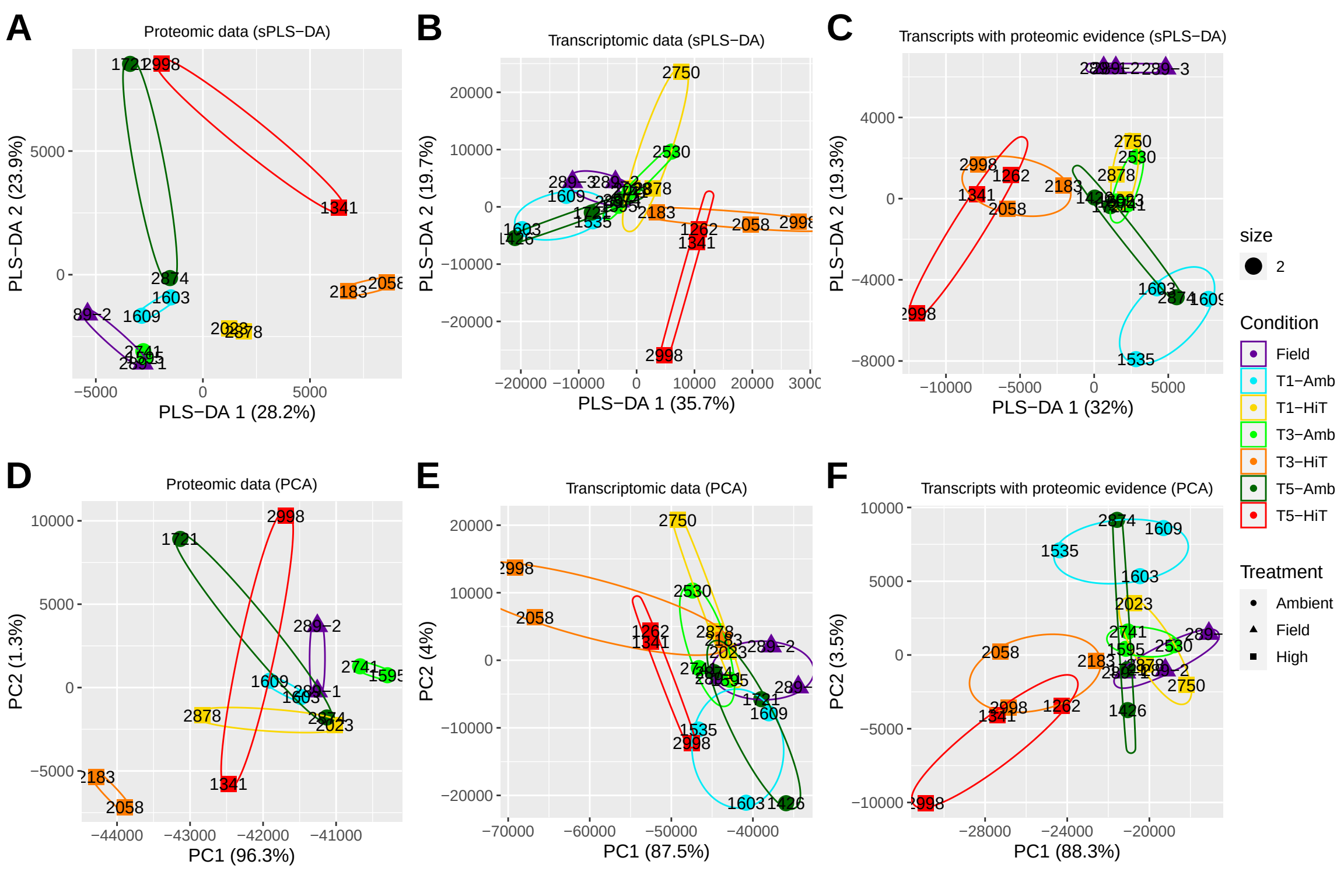
A
Proteomic data (sPLS−DA)
PLS−DA 1 (28.2%)
PLS−DA 2 (23.9%)
B
Transcriptomic data (sPLS−DA)
PLS−DA 1 (35.7%)
PLS−DA 2 (19.7%)
C
Transcripts with proteomic evidence (sPLS−DA)
PLS−DA 1 (32%)
PLS−DA 2 (19.3%)
D
Proteomic data (PCA)
PC1 (96.3%)
PC2 (1.3%)
E
Transcriptomic data (PCA)
PC1 (87.5%)
PC2 (4%)
F
Transcripts with proteomic evidence (PCA)
PC1 (88.3%)
PC2 (3.5%)
size
2
Condition
Field
T1−Amb
T1−HiT
T3−Amb
T3−HiT
T5−Amb
T5−HiT
Treatment
Ambient
Field
High