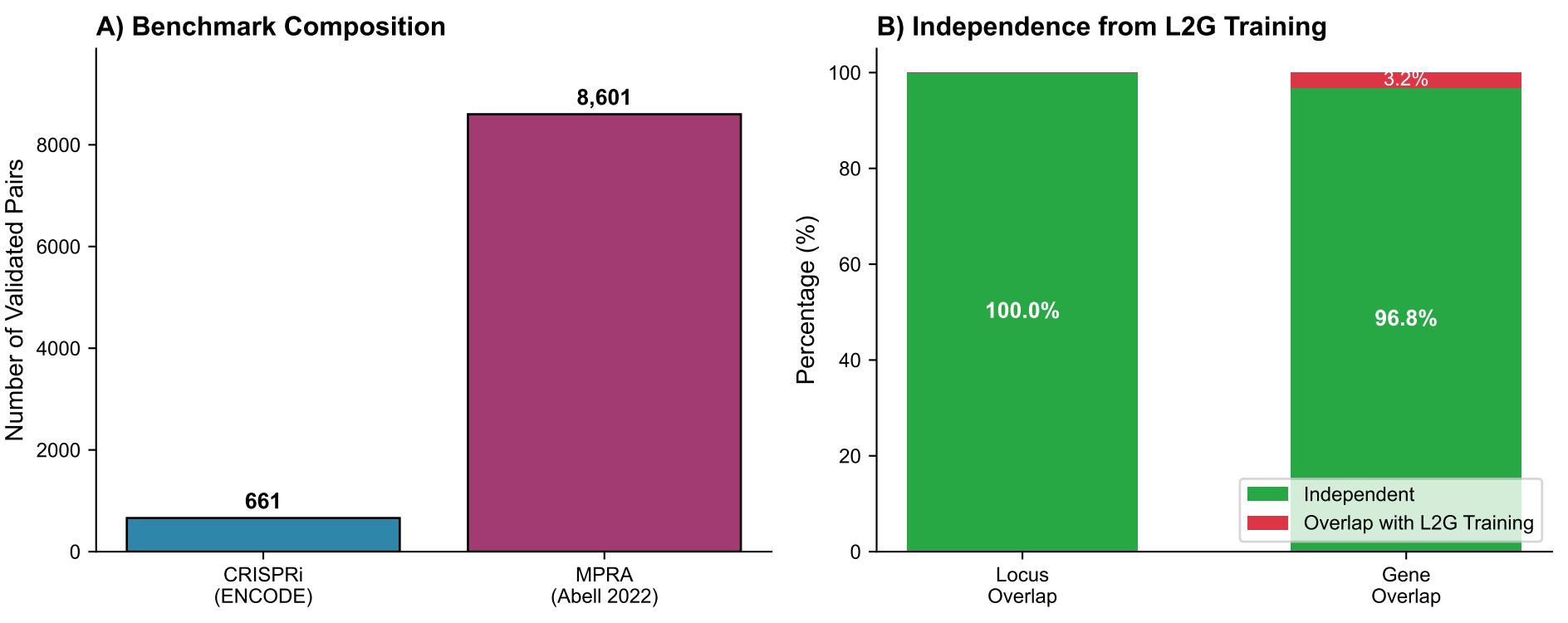
A) Benchmark Composition
Number of Validated Pairs
0
2000
4000
6000
8000
661
8,601
CRISPRi
(ENCODE)
MPRA
(Abell 2022)
B) Independence from L2G Training
Percentage (%)
0
20
40
60
80
100
100.0%
3.2%
96.8%
Independent
Overlap with L2G Training
Locus
Overlap
Gene
Overlap