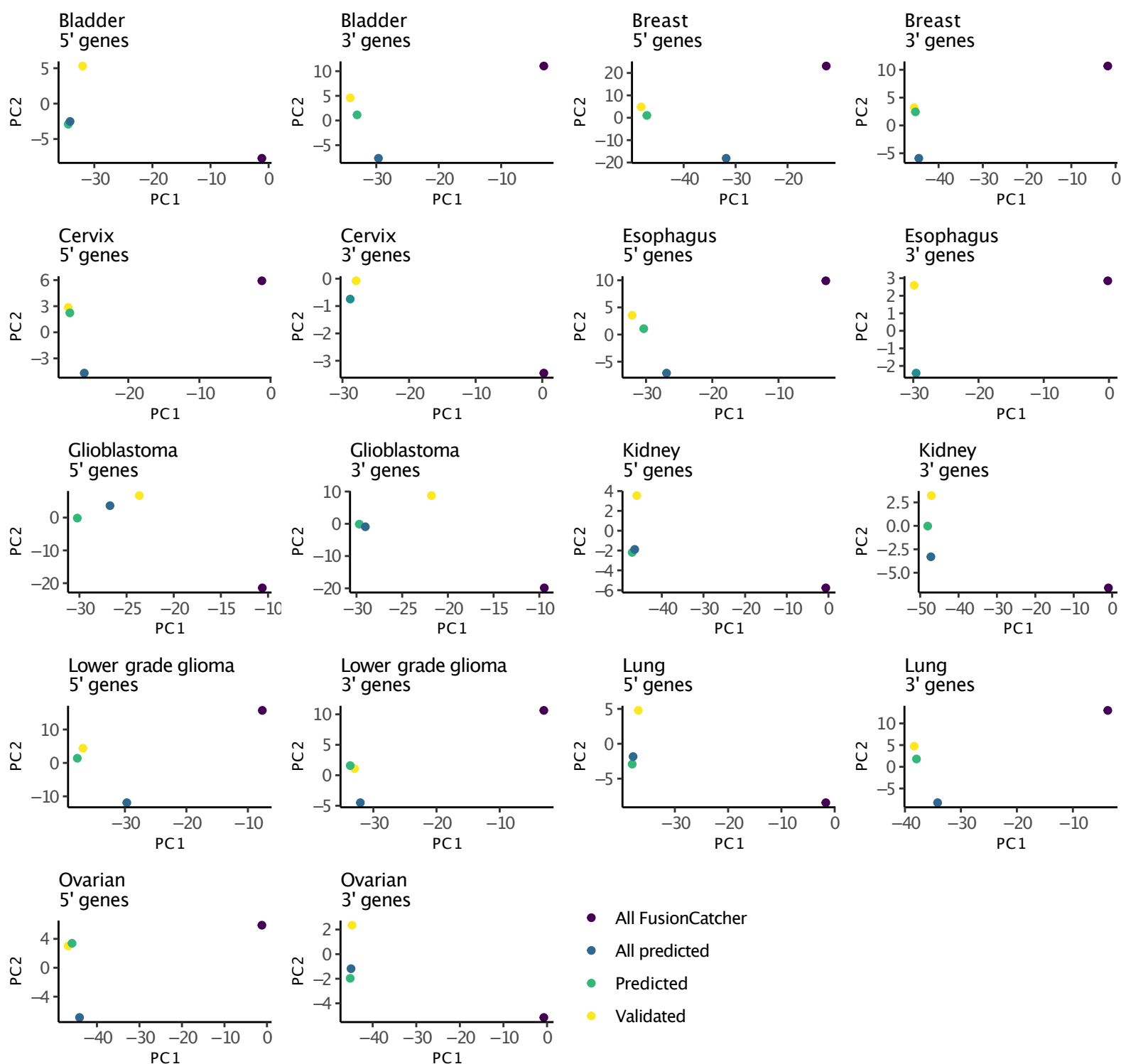




Additional file 5. PCA of gene set over-representation analysis for all fusion genes detected by FusionCatcher, fusion genes validated by our pipeline, and fusion genes predicted by the machine learning classifier in samples with WGS data (predicted) and in all RNA-seq samples (all predicted).