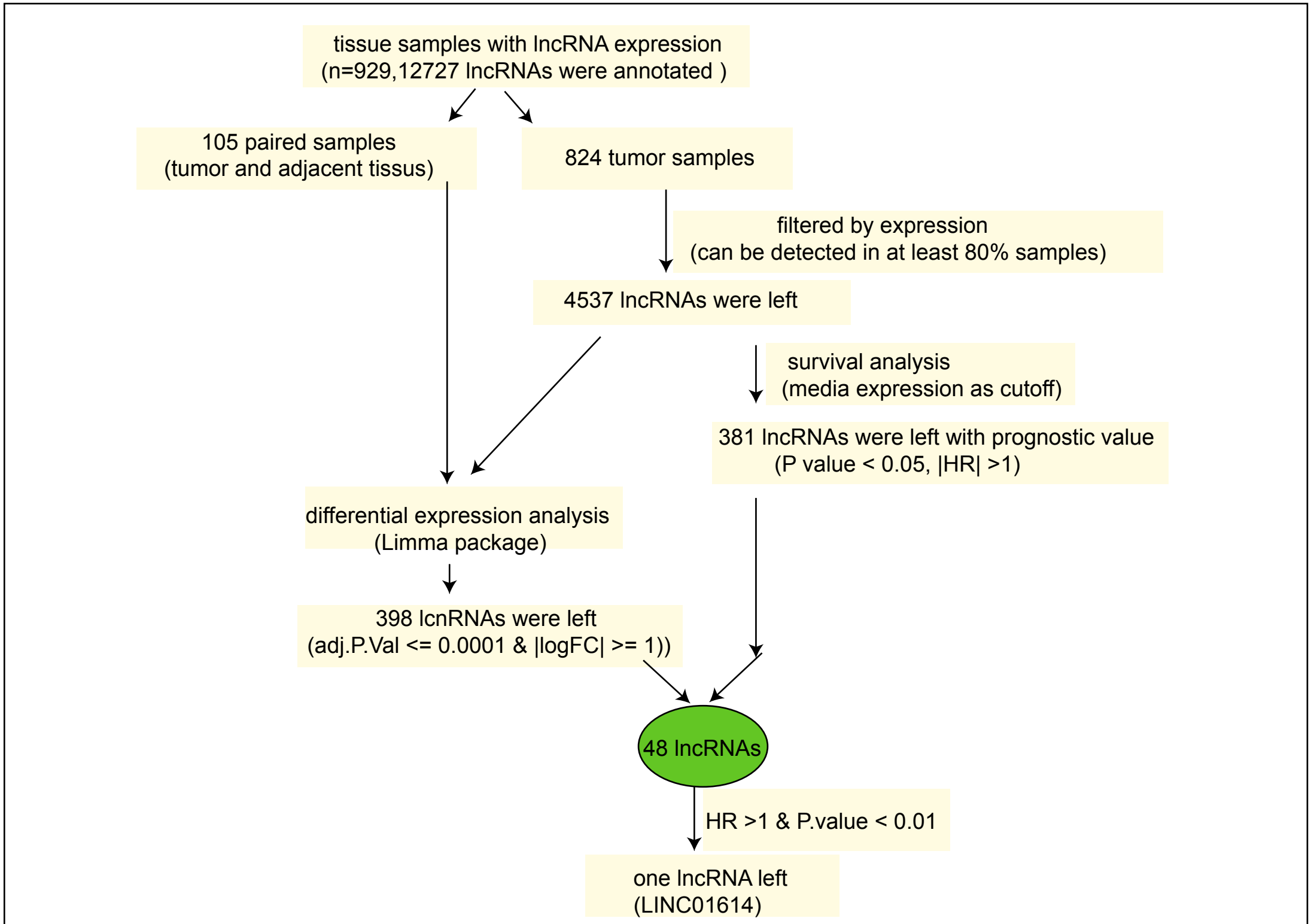
tissue samples with lncRNA expression
(n=929,12727 lncRNAs were annotated)
105 paired samples
(tumor and adjacent tissus)
824 tumor samples
filtered by expression
(can be detected in at least 80% samples)
4537 lncRNAs were left
survival analysis
(media expression as cutoff)
381 lncRNAs were left with prognostic value
(P value < 0.05, |HR| >1)
differential expression analysis
(Limma package)
398 lcnRNAs were left
(adj.P.Val <= 0.0001 & |logFC| >= 1))
48 lncRNAs
HR >1 & P.value < 0.01
one lncRNA left
(LINC01614)