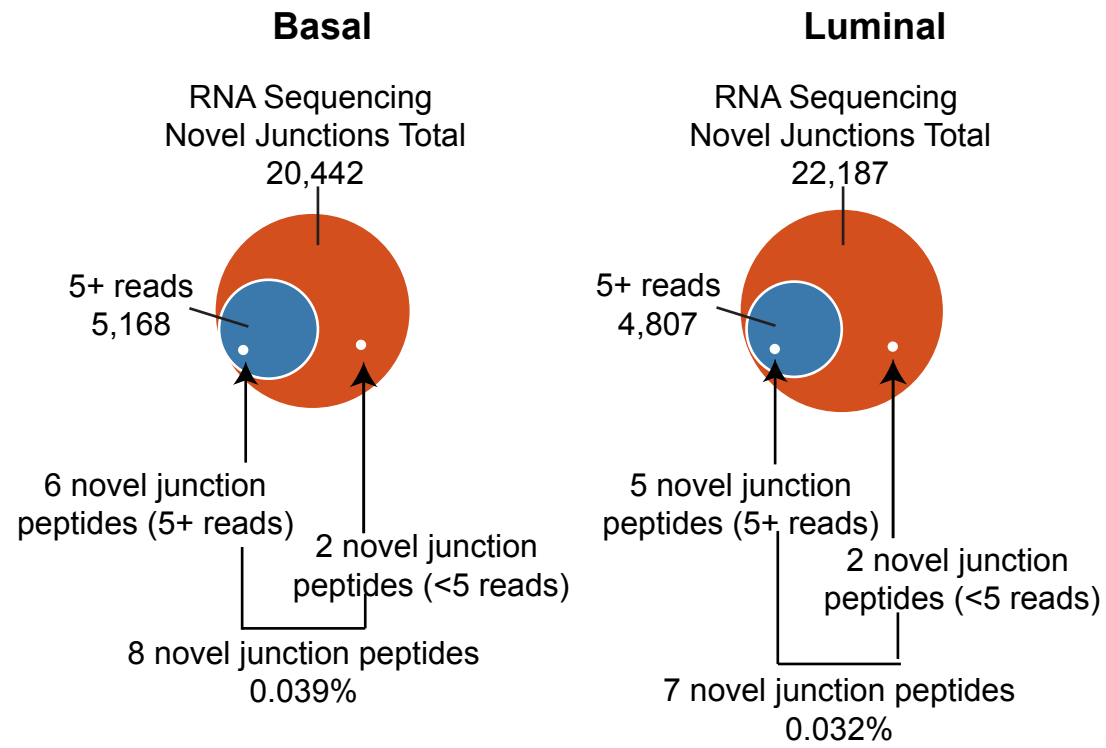
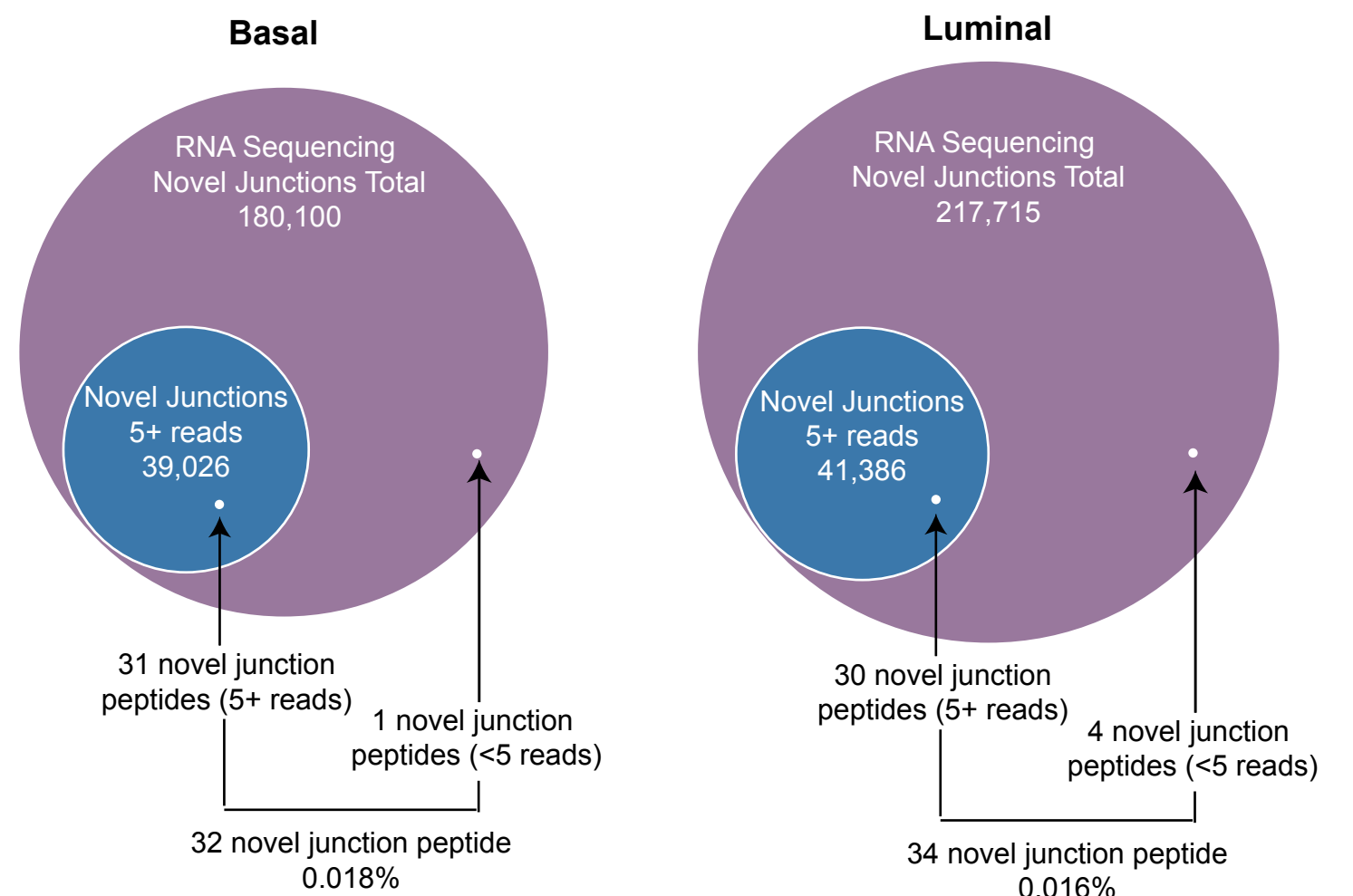


A

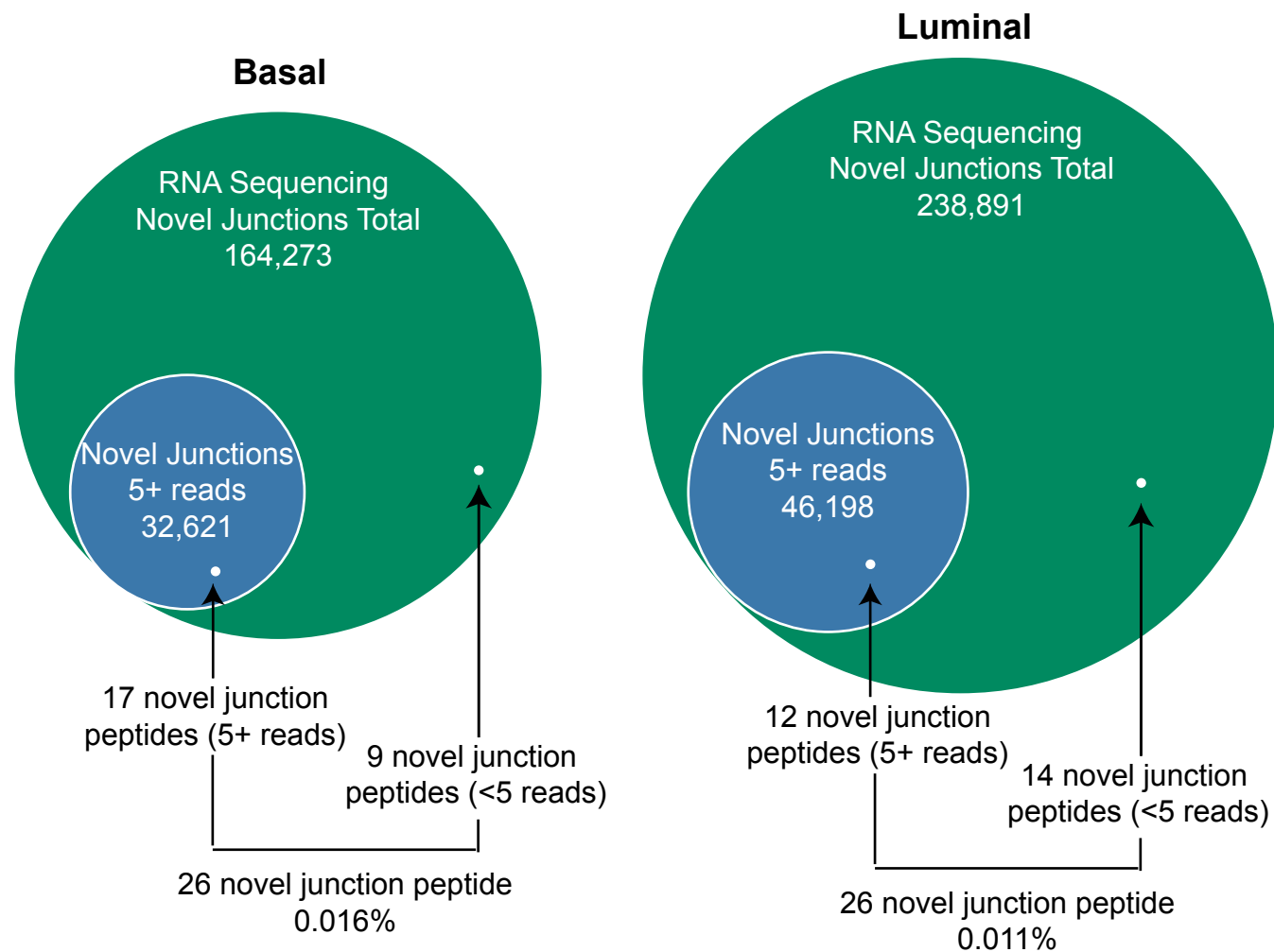
Novel Junction Peptides with 2 known exons
Predicted by RNA Sequencing, Identified by MS/MS

**B**

Novel Junction Peptides with 2 known exons
Predicted by RNA Sequencing, Identified by MS/MS

**C**

Novel Junction Peptides with 0 known exons
Predicted by RNA Sequencing, Identified by MS/MS



Supplementary Figure 8. Novel junction peptide detection in each splice junction class. Proportion of predicted junctions based on RNA sequencing (all junctions, red/purple/green; junctions 5+ reads blue) which were identified by MS/MS proteomics for luminal and basal breast tumors from splicing between (A) two known exons, (B) one known exon and a novel region and (C) no known exons (2 novel coding regions).