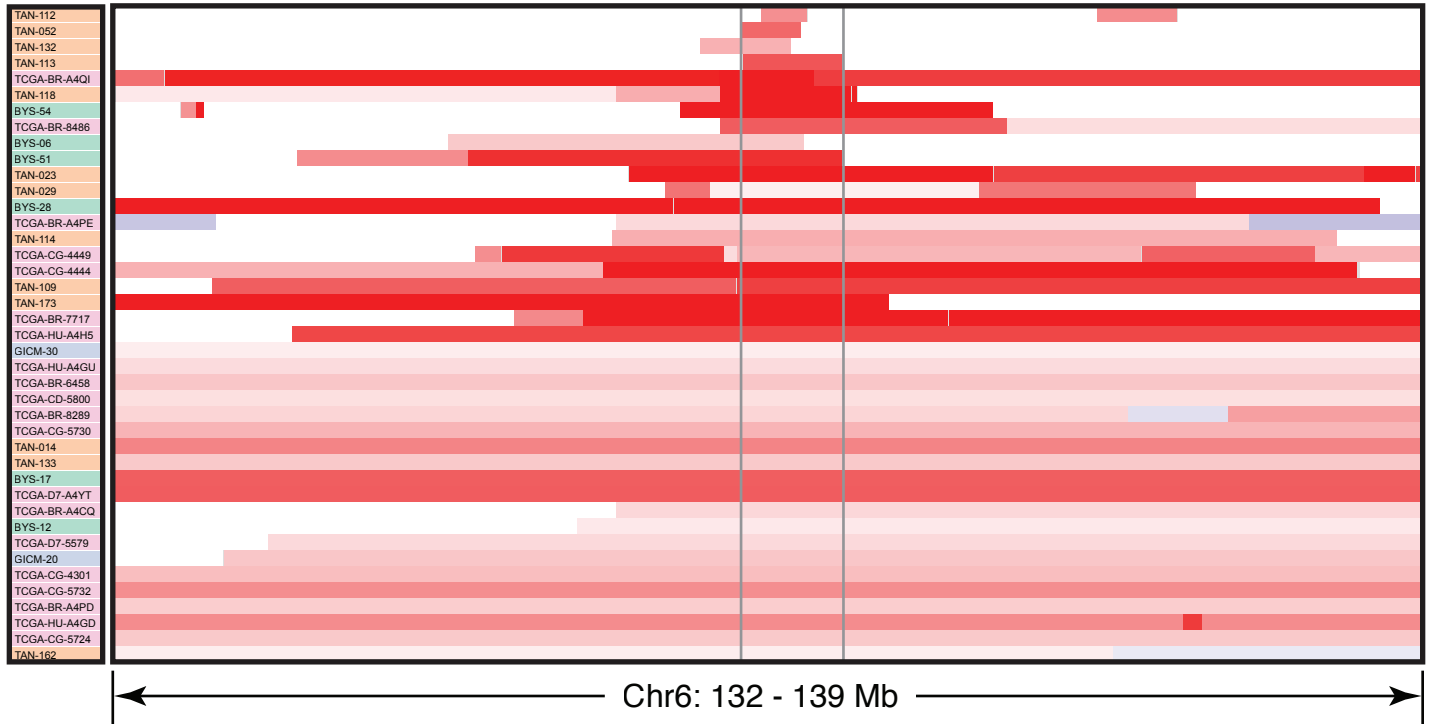


**S1 Fig. Copy number alterations contributing to novel gastric cancer peaks.** Amplified segments are marked red and deleted segments blue across the genome (X-axis) and selected patients (Y-axis). Peak regions are delimited by vertical gray lines. Genes in or near the peak are placed above the heat map with the presumed driver highlighted in green. Shown are (A) the significantly amplified region containing *MYB* on chromosome 6 and (B) the significantly deleted region containing *SMARCA4* on chromosome 19.

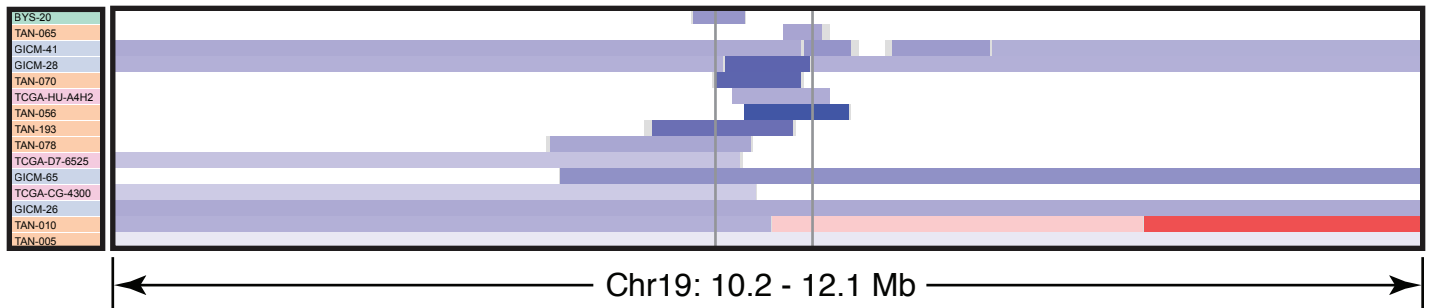
**A** *MYB* amplification region

*HBS1L* *MYB* *AHI1* *LINC00271*



**B** *SMARCA4* deletion region

*C19orf52* *SMARCA4* *LDLR*



**Key**

Source

- TCGA
- Korea
- Singapore
- Italy

Copy Number

