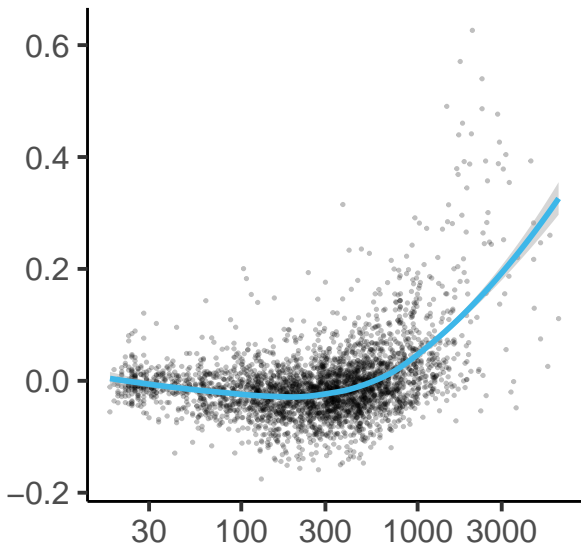


GAM residual



Distance from first TM/signal  
to end of polypeptide (AAs)