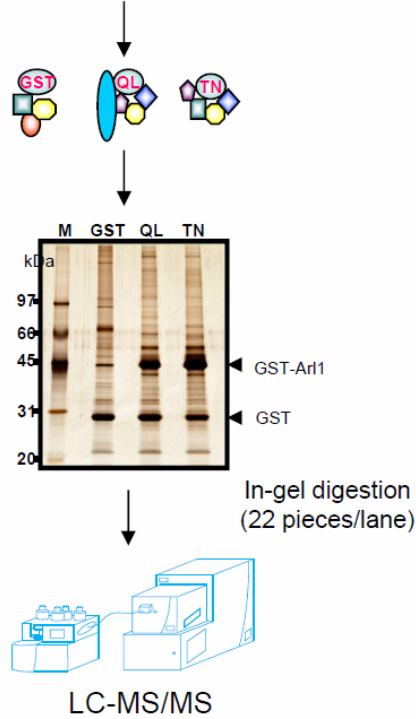
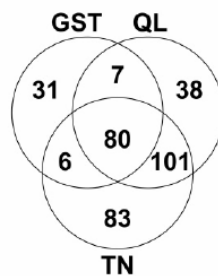


Differential affinity chromatography with immobilized GST-Ar11 (GST-Ar11Q71L and GST-Ar11T31N)



Database search using Mascot algorithm

Identification probability analysis by Scaffold 2 software (peptide thresholds >95.0%; protein thresholds >95.0%, and 2 peptides minimum)



Ar11 specific interaction proteins

S1 Fig.