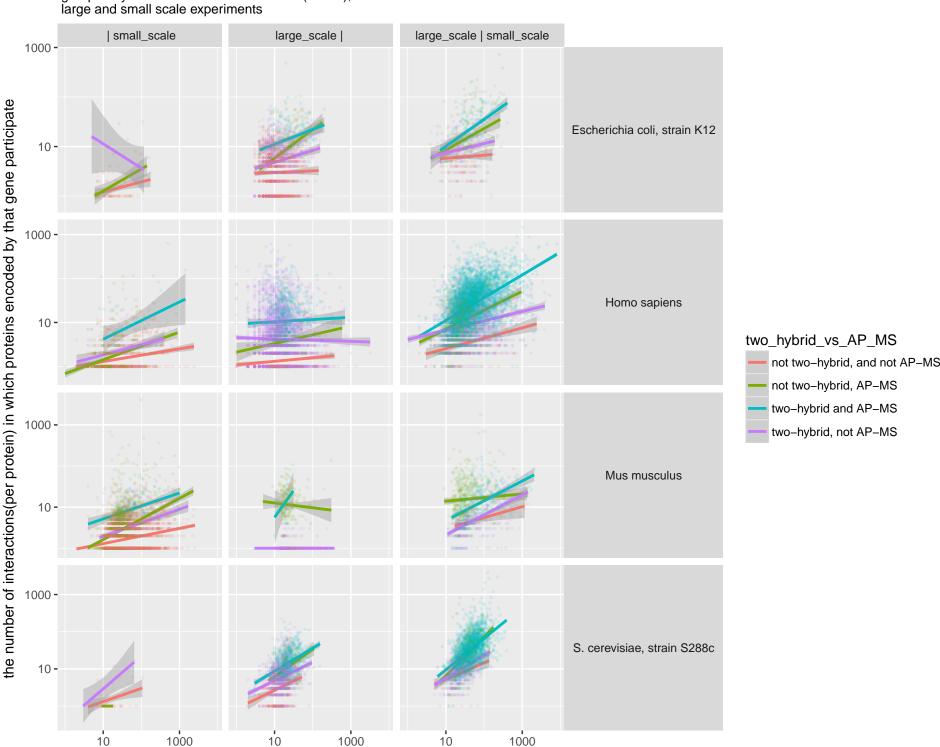
The relationship between the number of publications per gene and the number of interactions in which proteins encoded by that gene participate

line was fitted using robust linear regression, all UniprotAC were coerced to generic UniprotAC, grouped by interaction detection method (colour), large and small scale experiments



the number of publications per gene