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F = forward;  R = reverse

Tables S2-S3
Available for download at http://www.g3journal.org/lookup/suppl/doi:10.1534/g3.113.008763/-/DC1

Table S2  Complete list of SGA interactions with ess1^{H164R}. 429 genetic interactions were identified after setting a calibrated P-value of between +0.2 to +1.0 (aggravating interactions), or ~0.2 to -1.0 (alleviating interactions).

Table S3  Results of SlimMapper Gene Ontology (GO) analysis. List shows functional categories of genes that interact with ESS1 derived from the SGA using a calibrated P-value cutoff value of 0.25. Analysis was done using SlimMapper http://www.yeastgenome.org/cgi-bin/GO/goSlimMapper.pl and yeast GO-Slim:Process was used to sort genes into indicated functional categories.