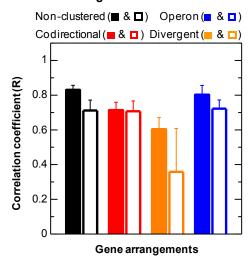
## Gene arrangements with lacl deleted



**Figure S9** Correlation coefficient (R) for different gene arrangements calculated without outliers (cells more than 2 S.D. from the mean). Data symbols indicate the gene arrangements shown in **Figure 1A**. Error bars indicate the s.e.m. of 5-7 replicate measurements.