

Box plot showing the distribution of the number of genes in the germinant-specific regulon for six genes: *cotB*, *cotH*, *ywrJ*, *gerE*, *gerR*, and *spoIIID*. The y-axis represents the number of genes (0 to 100). The x-axis lists the genes. Each box plot shows the median (thick horizontal line), the interquartile range (box), and the range (whiskers). Outliers are shown as open circles.

Gene	Median	Q1	Q3	Min	Max	Outliers
<i>cotB</i>	~45	~35	~55	~25	~65	1
<i>cotH</i>	~35	~25	~45	~15	~55	2
<i>ywrJ</i>	~45	~35	~55	~25	~65	2
<i>gerE</i>	~55	~45	~75	~35	~95	0
<i>gerR</i>	~55	~45	~75	~35	~95	15
<i>spoIIID</i>	~35	~25	~45	~15	~55	0

spoIIID