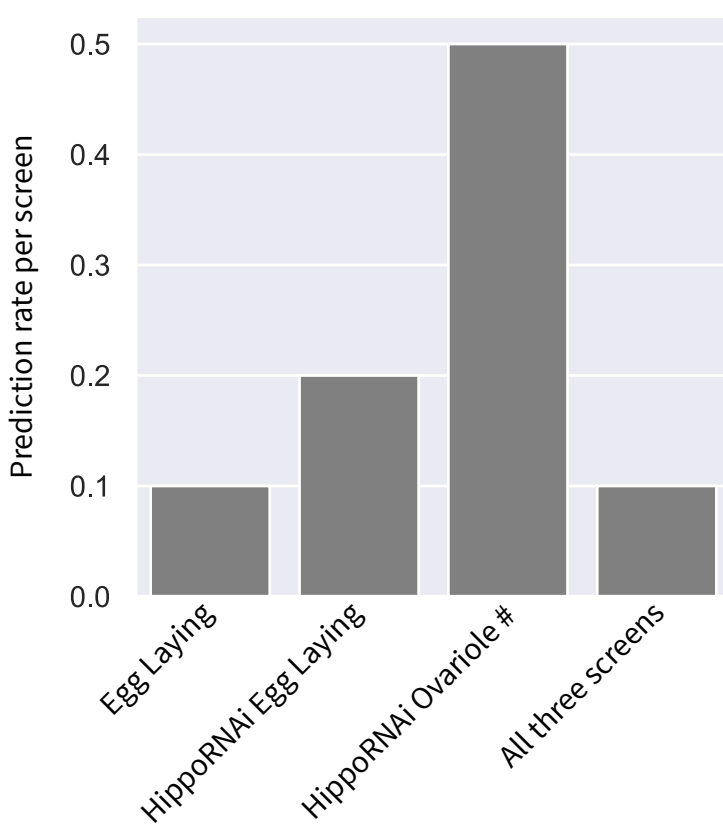
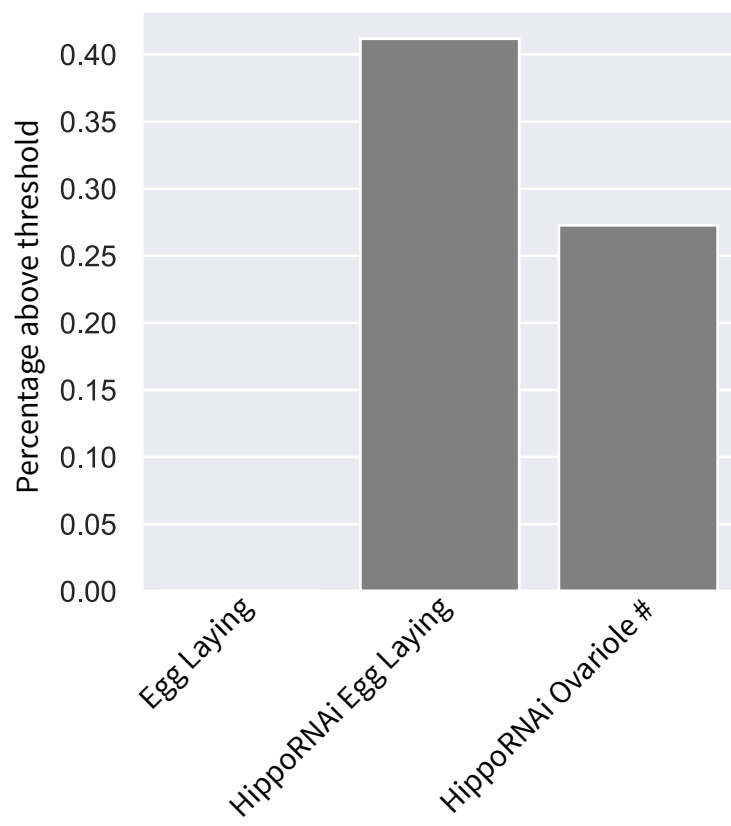


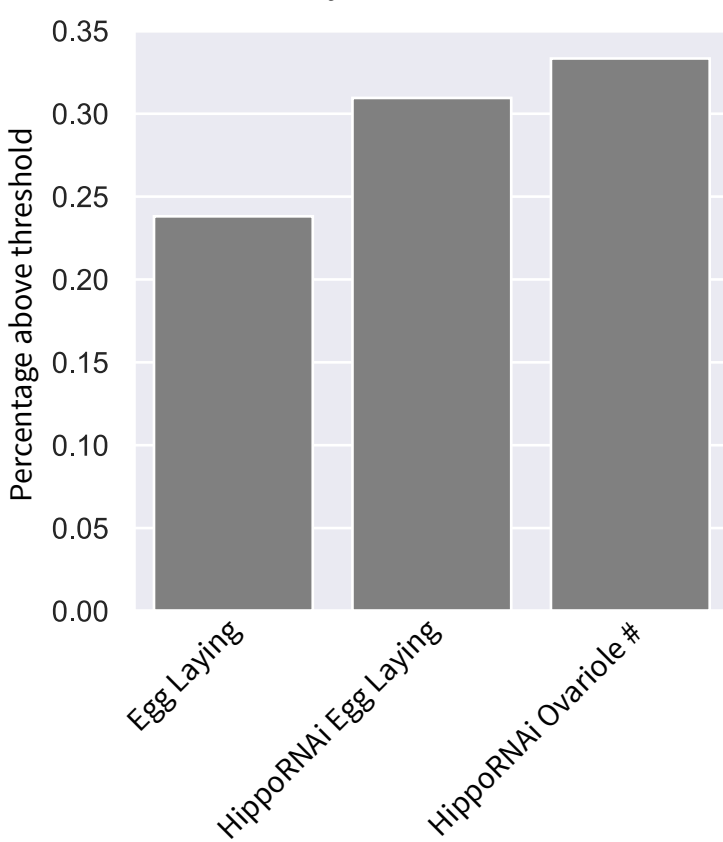
**a** Prediction rates of the Core module



**b** Prediction rates of the three modules



**c** Positive candidates per screen divided by all tested connectors



**d** Positive candidates per screen divided by all tested signaling genes

