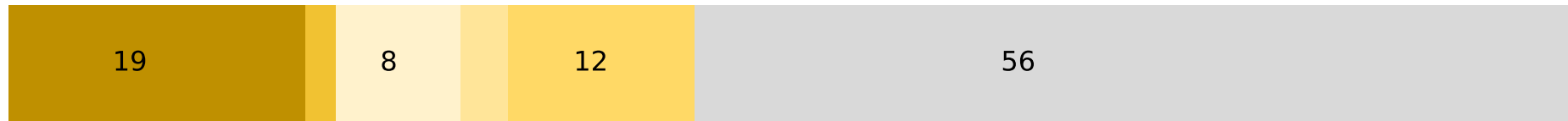
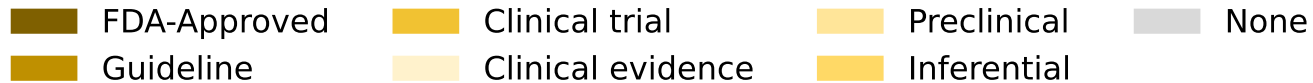


Counts of profiles (n=100) with at least one observed molecular feature
related to disease prognosis, by evidence



0%

20%

40%

60%

80%

100%