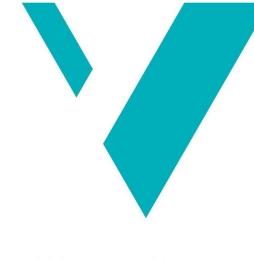
This is only a draft!

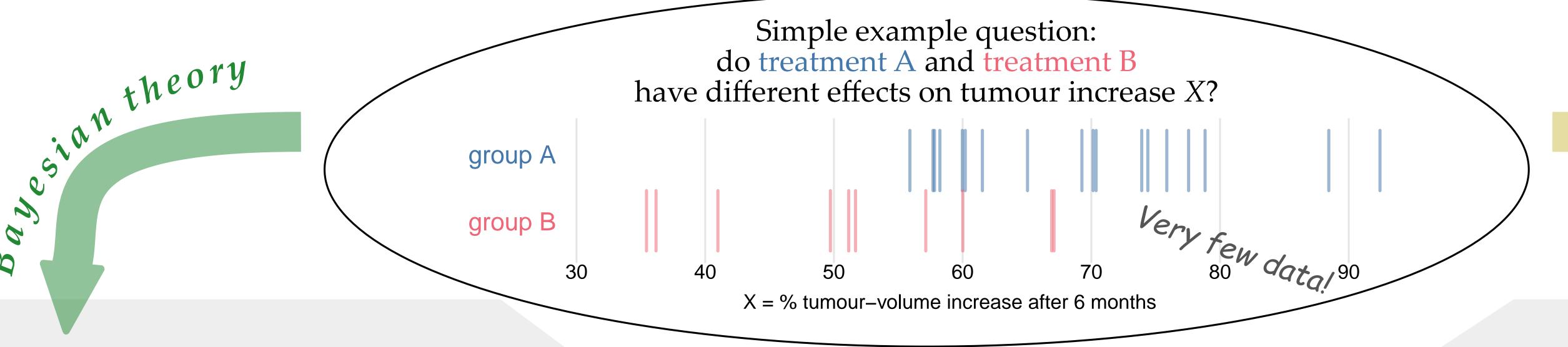


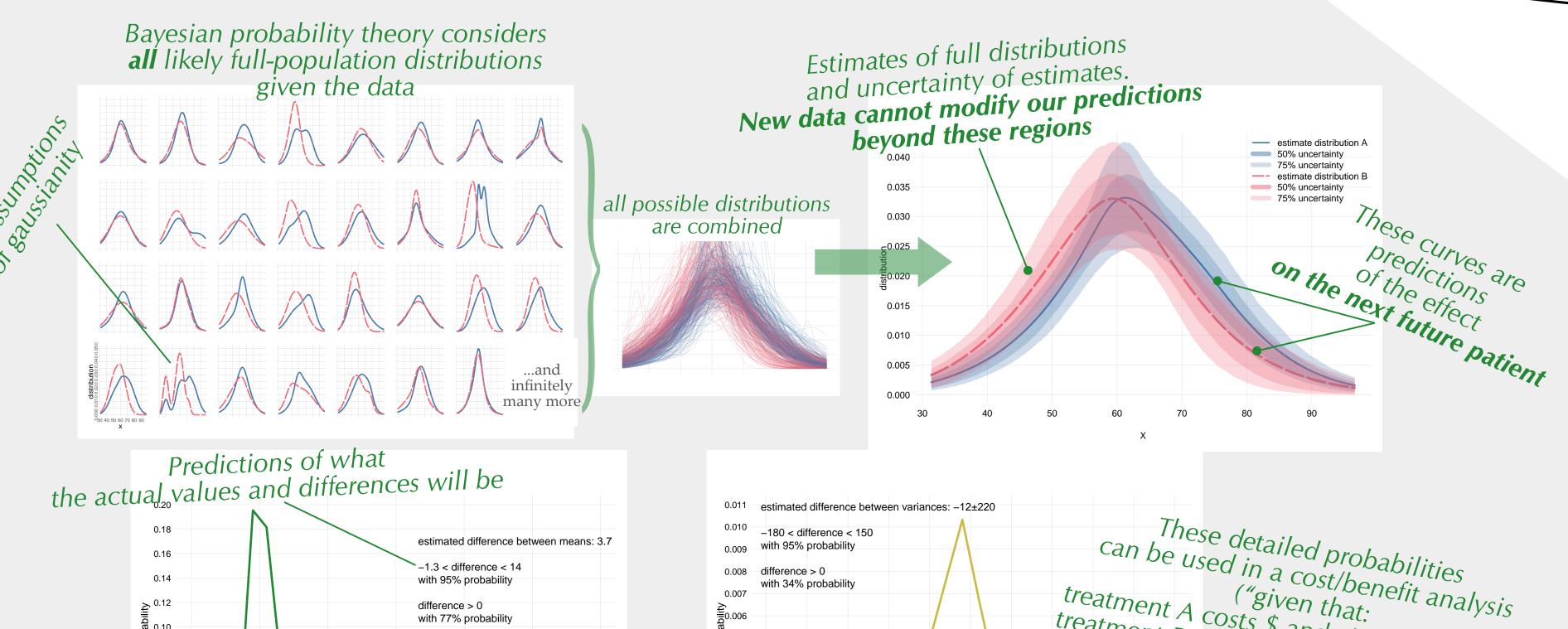
User-friendly software for Bayesian analysis of medical data

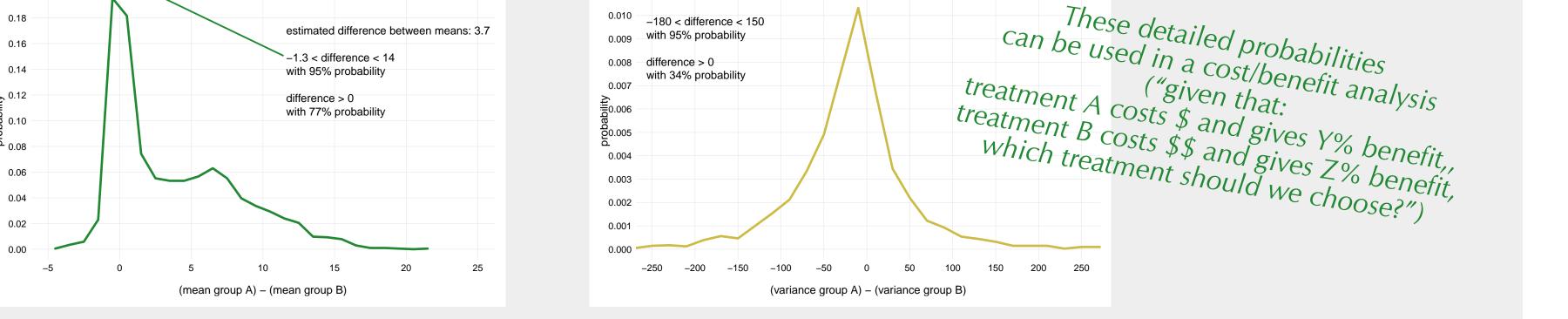
PierGianLuca Porta Mana, Kjetil Dyrland, Alexander Selvikvåg Lundervold [add addresses]



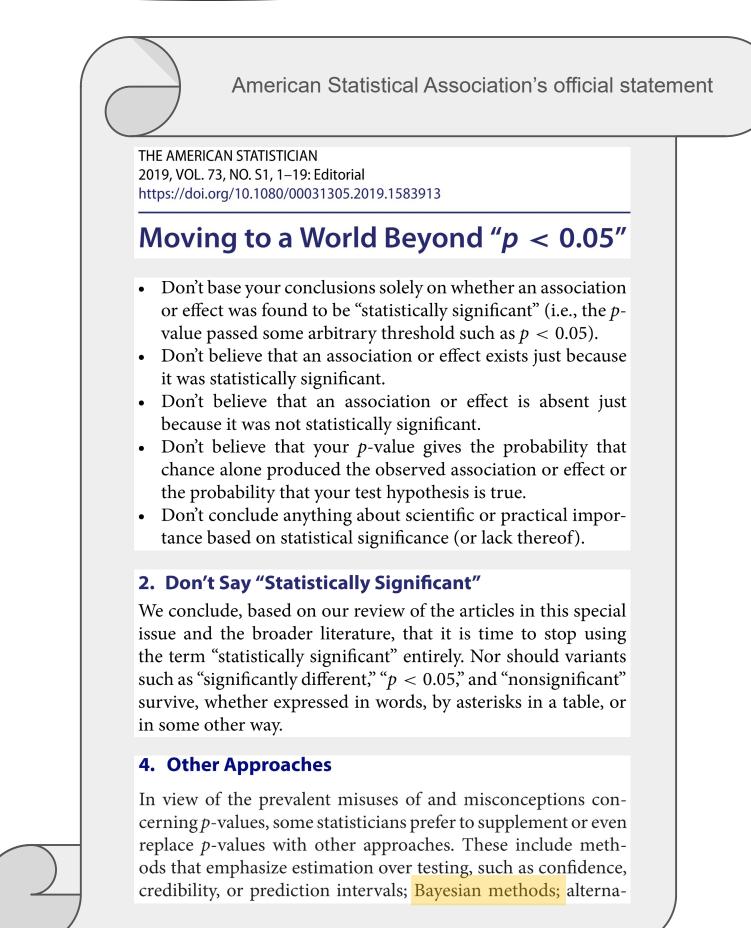
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- "Assuming that the two full-population distributions are smooth, we predict:
- The distributions of **future treatment outcomes** will be as in the plots, within the uncertainties shown
- Future patients under treatment A will have on average X=63, and 59 < X < 69 with 95% probability
- Future patients under treatment B will have on average X=59, and 52 < X < 66 with 95% probability
- The difference between the mean X of the treatments will be within -1.3 and 14 with 95% probability
- Average *X* under treatment A will be **larger** than under treatment B with 77% probability
- Variance of *X* under treatment A will be **smaller** than under treatment B with 66% probability "



- "Assuming that the two full populations:
- are gaussian and independent
- have identical variances (F-test, p = 0.44)

The hypothesis that their means are equal has a *p*-value 0.00043 (two-tail *t*-test: +4.0).

Does the value "0.00043" have a meaning easy to grasp?

The sample mean difference of X is 17.6 The 95% confidence interval is [8.6, 26.7]"

> This does **not** mean that 8.6 < X < 26.7 with 95% probability! It means that this technique to construct the interval contains the true value in 95% of all imaginary datasets. But we care about the dataset actually observed!

Compare the two analyses

The predictions of Bayesian analysis are:

- detailed
- quantitative
- easy to understand

(eg, "fraction x% of population will have effect y")

The statements of sampling theory are:

- vague
- obscure or misleading
- heavily dependent on tacit assumptions

This was just a simple example. Bayesian theory deals in the same way with multiple covariates and correlation questions



there's very little friendly software for doing this!"

True! That's why we are preparing a user-friendly software to do Bayesian analysis on (non-imaging) medical data

The maths will be taken care of under the hood Works with continuous and categorical variables - allows predictions about their correlations and relevance

No assumptions of gaussianity or other assumptions typical of sampling theory No need for corrections of any kind

The software will suggest meaningful questions to be asked (in line with ASA's statement)

We're already using a prototype version for drug-discovery research

Please get in touch if you want to test it and help us making a great sofware!